



Codebook for Incident Events 2008-2021 April 2025

**Prepared by the Collaborative Studies Coordinating Center
Version 1.0**

HCHS/SOL Incident Events 2008-2021 Codebook

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Chapter 1: Endpoint Ascertainment

Section A: Events Eligibility Ascertainment

Events Eligibility Form: FORM CODE=EEF, VERSION=B,C

<i>ID</i>		<i>HCHS Participant ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9011	Present	Text suppressed

<i>VISIT</i>		<i>Visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1035	1	Visit 1
994	2	Visit 2
930	3	Visit 3
813	4	Visit 4
836	5	Visit 5
592	6	Visit 6
603	7	Visit 7
584	8	Visit 8
548	9	Visit 9
503	10	Visit 10
592	11	Visit 11
382	12	Visit 12
366	13	Visit 13
183	14	Visit 14
42	15	Visit 15
8	16	Visit 16

Chapter 1: Endpoint Ascertainment

Section A: Events Eligibility Ascertainment

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6718	1	Occurrence 1
1355	2	Occurrence 2
477	3	Occurrence 3
189	4	Occurrence 4
105	5	Occurrence 5
59	6	Occurrence 6
36	7	Occurrence 7
25	8	Occurrence 8
13	9	Occurrence 9
11	10	Occurrence 10
7	11	Occurrence 11
6	12	Occurrence 12
2	13	Occurrence 13
1	14	Occurrence 14
1	15	Occurrence 15
1	16	Occurrence 16
2	17	Occurrence 17
2	18	Occurrence 18
1	19	Occurrence 19

<i>EEF0A_FUTIME</i>		<i>Follow-up time from baseline for EEF0A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
9010	Range	-116 - 6214 (median=2736 mean=2971.3 std=1217.0)

<i>EEF0AY</i>		<i>Year of EEF0A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
9010	Range	2009 - 2025 (median=2017 mean=2017.6 std=3.4)

<i>EEF0C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9011	Present	Text suppressed

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<i>EEF0D_FUTIME</i>		<i>Follow-up time from baseline for EEF0D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
9008	Range	-297 - 5877 (median=1825 mean=2077.6 std=1406.0)

<i>EEF0DY</i>		<i>Year of EEF0D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
9008	Range	2008 - 2025 (median=2015 mean=2015.2 std=3.9)

<i>EEF1</i>		<i>Event location</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7		Missing
1766	1	1= In hospital
899	2	2= Emergency Dept. (ED)
6243	3	3= Both ED and in hospital
96	4	4= other

<i>EEF1A</i>		<i>Other event location</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
8914	~	Legitimately skipped
96	Present	Text suppressed

<i>EEF2_FUTIME</i>		<i>Follow-up time from baseline for EEF2 (Admission date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
677		Missing
8334	Range	-196 - 5853 (median=1902.5 mean=2123.50 std=1389.24)

<i>EEF2Y</i>		<i>Year of EEF2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
677		Missing
8334	Range	2008 - 2024 (median=2015 mean=2015.4 std=3.8)

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<i>EEF3_FUTIME</i>		<i>Follow-up time from baseline for EEF3 (Discharge date/date of death)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
232		Missing
8779	Range	-297 - 5877 (median=1861 mean=2106.7 std=1401.1)

<i>EEF3Y</i>		<i>Year of EEF3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
232		Missing
8779	Range	2008 - 2025 (median=2015 mean=2015.3 std=3.9)

<i>EEF4</i>		<i>Vital status at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24		Missing
254	0	0= Dead
8733	1	1= Alive

<i>EEF5A</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
989		Missing
8022	Present	Text suppressed

<i>EEF5B</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1532		Missing
7479	Present	Text suppressed

<i>EEF5C</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2063		Missing
6948	Present	Text suppressed

<i>EEF5D</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2656		Missing
6355	Present	Text suppressed

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Section A: Events Eligibility Ascertainment

<i>EEF5E</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3278		Missing
5733	Present	Text suppressed

<i>EEF5F</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3864		Missing
5147	Present	Text suppressed

<i>EEF5G</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4405		Missing
4606	Present	Text suppressed

<i>EEF5H</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4961		Missing
4050	Present	Text suppressed

<i>EEF5I</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5424		Missing
3587	Present	Text suppressed

<i>EEF5J</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5843		Missing
3168	Present	Text suppressed

<i>EEF5K</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6280		Missing
2731	Present	Text suppressed

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Section A: Events Eligibility Ascertainment

<i>EEF5L</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6622		Missing
2389	Present	Text suppressed

<i>EEF5M</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6929		Missing
2082	Present	Text suppressed

<i>EEF5N</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7165		Missing
1846	Present	Text suppressed

<i>EEF5O</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7404		Missing
1607	Present	Text suppressed

<i>EEF5P</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7615		Missing
1396	Present	Text suppressed

<i>EEF5Q</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7818		Missing
1193	Present	Text suppressed

<i>EEF5R</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7930		Missing
1081	Present	Text suppressed

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Section A: Events Eligibility Ascertainment

<i>EEF5S</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8087		Missing
924	Present	Text suppressed

<i>EEF5T</i>		<i>ICD discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8207		Missing
804	Present	Text suppressed

<i>EEF6A</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5070		Missing
3941	Present	Text suppressed

<i>EEF6B</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6451		Missing
2560	Present	Text suppressed

<i>EEF6C</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7344		Missing
1667	Present	Text suppressed

<i>EEF6D</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7922		Missing
1089	Present	Text suppressed

<i>EEF6E</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8328		Missing
683	Present	Text suppressed

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Section A: Events Eligibility Ascertainment

<i>EEF6F</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8531		Missing
480	Present	Text suppressed

<i>EEF6G</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8666		Missing
345	Present	Text suppressed

<i>EEF6H</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8763		Missing
248	Present	Text suppressed

<i>EEF6I</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8834		Missing
177	Present	Text suppressed

<i>EEF6J</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8879		Missing
132	Present	Text suppressed

<i>EEF6K</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8916		Missing
95	Present	Text suppressed

<i>EEF6L</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8940		Missing
71	Present	Text suppressed

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<i>EEF6M</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8956		Missing
55	Present	Text suppressed

<i>EEF6N</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8972		Missing
39	Present	Text suppressed

<i>EEF6O</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8982		Missing
29	Present	Text suppressed

<i>EEF6P</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8988		Missing
23	Present	Text suppressed

<i>EEF6Q</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8992		Missing
19	Present	Text suppressed

<i>EEF6R</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8997		Missing
14	Present	Text suppressed

<i>EEF6S</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9000		Missing
11	Present	Text suppressed

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Section A: Events Eligibility Ascertainment

<i>EEF6T</i>		<i>ICD procedure code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9003		Missing
8	Present	Text suppressed

<i>EEF7</i>		<i>discharge summary mentions cardiac</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7281	0	0=No
1727	1	1=Yes

<i>EEF7A</i>		<i>MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1323	0	0=No
400	1	1=Yes

<i>EEF7AA</i>		<i>pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1653	0	0=No
71	1	1=Yes

<i>EEF7B</i>		<i>angina</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1604	0	0=No
120	1	1=Yes

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Section A: Events Eligibility Ascertainment

<i>EEF7C</i>		<i>ischemic heart disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1638	0	0=No
85	1	1=Yes

<i>EEF7D</i>		<i>CHD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1025	0	0=No
699	1	1=Yes

<i>EEF7E</i>		<i>unstable angina</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1634	0	0=No
89	1	1=Yes

<i>EEF7F</i>		<i>coronary insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1718	0	0=No
5	1	1=Yes

<i>EEF7G</i>		<i>cardiac arrest</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1584	0	0=No
139	1	1=Yes

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Section A: Events Eligibility Ascertainment

<i>EEF7H</i>		<i>AHD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1456	0	0=No
268	1	1=Yes

<i>EEF7I</i>		<i>heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1433	0	0=No
290	1	1=Yes

<i>EEF7J</i>		<i>CHF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1092	0	0=No
632	1	1=Yes

<i>EEF7K</i>		<i>cardiomyopathy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1433	0	0=No
291	1	1=Yes

<i>EEF7L</i>		<i>atherosclerosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1705	0	0=No
18	1	1=Yes

Chapter 1: Endpoint Ascertainment

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<i>EEF7M</i>		<i>PVD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1546	0	0=No
178	1	1=Yes

<i>EEF7N</i>		<i>claudication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7284	.S	Legitimately skipped
1696	0	0=No
28	1	1=Yes

<i>EEF7O</i>		<i>acute PE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1589	0	0=No
134	1	1=Yes

<i>EEF7P</i>		<i>aortic aneurysm</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
7284	.S	Legitimately skipped
1719	0	0=No
3	1	1=Yes

<i>EEF7Q</i>		<i>CCU</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1618	0	0=No
105	1	1=Yes

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<i>EEF7R</i>		<i>cardiac cath</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1290	0	0=No
433	1	1=Yes

<i>EEF7S</i>		<i>CABG</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1503	0	0=No
220	1	1=Yes

<i>EEF7T</i>		<i>coronary stent</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1413	0	0=No
310	1	1=Yes

<i>EEF7U</i>		<i>elevated CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1324	0	0=No
399	1	1=Yes

<i>EEF7V</i>		<i>coronary angiography</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1504	0	0=No
219	1	1=Yes

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<i>EEF7W</i>		<i>cardiac angioplasty</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1663	0	0=No
60	1	1=Yes

<i>EEF7X</i>		<i>leg amputation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1711	0	0=No
12	1	1=Yes

<i>EEF7Y</i>		<i>leg angioplasty</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1710	0	0=No
13	1	1=Yes

<i>EEF7Z</i>		<i>leg revascularization</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7284	.S	Legitimately skipped
1698	0	0=No
25	1	1=Yes

<i>EEF8</i>		<i>discharge summary mentions stroke</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
8590	0	0=No
419	1	1=Yes

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<i>EEF8A</i>		<i>TIA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
303	0	0=No
116	1	1=Yes

<i>EEF8B</i>		<i>stroke</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
203	0	0=No
216	1	1=Yes

<i>EEF8C</i>		<i>cerebral infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
292	0	0=No
127	1	1=Yes

<i>EEF8D</i>		<i>lacunar (syndrome or infarction)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
403	0	0=No
16	1	1=Yes

<i>EEF8E</i>		<i>cerebral embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
412	0	0=No
7	1	1=Yes

<i>EEF8F</i>		<i>cerebral thrombosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
415	0	0=No
4	1	1=Yes

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<i>EEF8G</i>		<i>central or cerebral venous thrombosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
419	0	0=No
0	1	1=Yes

<i>EEF8H</i>		<i>intracranial stenting</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
419	0	0=No
0	1	1=Yes

<i>EEF8I</i>		<i>carotid endarterectomy or carotid ste</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
405	0	0=No
14	1	1=Yes

<i>EEF8J</i>		<i>arteriovenous malformation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
414	0	0=No
5	1	1=Yes

<i>EEF8K</i>		<i>cerebral aneurysm</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
401	0	0=No
18	1	1=Yes

<i>EEF8L</i>		<i>cerebral, intracerebral, brain hemorr</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
343	0	0=No
76	1	1=Yes

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<i>EEF8M</i>		<i>subarachnoid hemorrhage</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
374	0	0=No
45	1	1=Yes

<i>EEF8N</i>		<i>cerebrovascular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
403	0	0=No
16	1	1=Yes

<i>EEF8O</i>		<i>CT/MRI scan showing new cerebrovascul</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8592	.S	Legitimately skipped
232	0	0=No
187	1	1=Yes

<i>EEF9</i>		<i>discharge summary mentions HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
7893	0	0=No
1116	1	1=Yes

<i>EEF9A</i>		<i>heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
811	0	0=No
304	1	1=Yes

<i>EEF9B</i>		<i>CHF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
453	0	0=No
662	1	1=Yes

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<i>EEF9C</i>		<i>pump failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
1115	0	0=No
0	1	1=Yes

<i>EEF9D</i>		<i>cardiomyopathy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
831	0	0=No
284	1	1=Yes

<i>EEF9E</i>		<i>ventricular failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
1111	0	0=No
4	1	1=Yes

<i>EEF9F</i>		<i>LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
976	0	0=No
139	1	1=Yes

<i>EEF9G</i>		<i>jugular venous distension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
1063	0	0=No
52	1	1=Yes

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<i>EEF9H</i>		<i>orthopnea</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7895	.S	Legitimately skipped
967	0	0=No
149	1	1=Yes

<i>EEF9I</i>		<i>paroxysmal nocturnal dyspnea</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
7895	.S	Legitimately skipped
1063	0	0=No
51	1	1=Yes

<i>EEF9J</i>		<i>low ejection fraction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1899		Missing
6110	.S	Legitimately skipped
781	0	0=No
221	1	1=Yes

<i>EEF9K</i>		<i>ejection fraction less than 40</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1899		Missing
6110	.S	Legitimately skipped
787	0	0=No
215	1	1=Yes

<i>EEF9L</i>		<i>cardiomegaly</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
779	0	0=No
336	1	1=Yes

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<i>EEF9M</i>		<i>impaired systolic function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
891	0	0=No
224	1	1=Yes

<i>EEF9N</i>		<i>impaired diastolic function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
772	0	0=No
343	1	1=Yes

<i>EEF9O</i>		<i>pulmonary edema device</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
1096	0	0=No
19	1	1=Yes

<i>EEF9P</i>		<i>heart biopsy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
1115	0	0=No
0	1	1=Yes

<i>EEF9Q</i>		<i>AICD check</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
1097	0	0=No
18	1	1=Yes

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<i>EEF9R</i>		<i>implantation of cardiac resync therapy device</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7895	.S	Legitimately skipped
1103	0	0=No
12	1	1=Yes

<i>EEF10</i>		<i>discharge summary mentions pulmonary</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
7231	0	0=No
1778	1	1=Yes

<i>EEF10A</i>		<i>acute bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1487	0	0=No
288	1	1=Yes

<i>EEF10B</i>		<i>acute bronchitis with exacerbation of asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1743	0	0=No
32	1	1=Yes

<i>EEF10C</i>		<i>obstructive chronic bronchitis with acute bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1754	0	0=No
21	1	1=Yes

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<i>EEF10D</i>		<i>acute exacerbation of asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
7233	.S	Legitimately skipped
1208	0	0=No
569	1	1=Yes

<i>EEF10E</i>		<i>chronic obstruction asthma with acute exacerbation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1450	0	0=No
325	1	1=Yes

<i>EEF10F</i>		<i>chronic obstructive asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1306	0	0=No
469	1	1=Yes

<i>EEF10G</i>		<i>status asthmaticus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1761	0	0=No
14	1	1=Yes

<i>EEF10H</i>		<i>chronic airway obstruction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1758	0	0=No
17	1	1=Yes

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<i>EEF10I</i>		<i>chronic bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1746	0	0=No
29	1	1=Yes

<i>EEF10J</i>		<i>pneumonia with underlying COPD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
7233	.S	Legitimately skipped
1720	0	0=No
56	1	1=Yes

<i>EEF10K</i>		<i>emphysema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1694	0	0=No
81	1	1=Yes

<i>EEF10L</i>		<i>bronchiectasis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1733	0	0=No
42	1	1=Yes

<i>EEF10M</i>		<i>extrinsic allergic alveolitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1775	0	0=No
0	1	1=Yes

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<i>EEF10N</i>		<i>acute respiratory failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1143	0	0=No
632	1	1=Yes

<i>EEF10O</i>		<i>acute and chronic respiratory failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
7233	.S	Legitimately skipped
1684	0	0=No
91	1	1=Yes

<i>EEF10P</i>		<i>positive spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
7233	.S	Legitimately skipped
1773	0	0=No
1	1	1=Yes

<i>EEF10Q</i>		<i>pulmonary insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
7233	.S	Legitimately skipped
1767	0	0=No
6	1	1=Yes

<i>EEF11</i>		<i>discharge summary mentions pregnancy-related complications</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2330		Missing
6536	0	0=No
145	1	1=Yes

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<i>EEF11A</i>		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1836		Missing
7030	.S	Legitimately skipped
99	0	0=No
46	1	1=Yes

<i>EEF11B</i>		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
140	0	0=No
4	1	1=Yes

<i>EEF11C</i>		<i>PEC</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
142	0	0=No
2	1	1=Yes

<i>EEF11D</i>		<i>PreX</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
144	0	0=No
0	1	1=Yes

<i>EEF11E</i>		<i>PET</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
144	0	0=No
0	1	1=Yes

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<i>EEF11F</i>		<i>Toxemia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
144	0	0=No
0	1	1=Yes

<i>EEF11G</i>		<i>pregnancy-induced hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
135	0	0=No
9	1	1=Yes

<i>EEF11H</i>		<i>pregestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
140	0	0=No
4	1	1=Yes

<i>EEF11I</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
111	0	0=No
33	1	1=Yes

<i>EEF11J</i>		<i>chronic hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
124	0	0=No
20	1	1=Yes

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<i>EEF11K</i>		<i>hyperglycemia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
143	0	0=No
1	1	1=Yes

<i>EEF11L</i>		<i>gestational diabetes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
66	0	0=No
78	1	1=Yes

<i>EEF11M</i>		<i>insulin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
125	0	0=No
19	1	1=Yes

<i>EEF11N</i>		<i>HELLP syndrome</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
143	0	0=No
1	1	1=Yes

<i>EEF11O</i>		<i>seizure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
144	0	0=No
0	1	1=Yes

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<i>EEF11P</i>		<i>renal failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
144	0	0=No
0	1	1=Yes

<i>EEF11Q</i>		<i>proteinuria</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
135	0	0=No
9	1	1=Yes

<i>EEF11R</i>		<i>oliguria</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1837		Missing
7030	.S	Legitimately skipped
144	0	0=No
0	1	1=Yes

<i>EEF12A</i>		<i>no additional form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
408	~	Legitimately skipped
4755	N	
3848	Y	

<i>EEF12B</i>		<i>MI form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
486	~	Legitimately skipped
5583	N	
2941	Y	

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<i>EEF12C</i>		<i>Stroke form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
589	~	Legitimately skipped
7852	N	
569	Y	

<i>EEF12D</i>		<i>HF form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
582	~	Legitimately skipped
6823	N	
1591	Y	

<i>EEF12E</i>		<i>Pulmonary form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
494	~	Legitimately skipped
5488	N	
3029	Y	

<i>EEF12F</i>		<i>Pregnancy-Related Complications form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1835		Missing
633	~	Legitimately skipped
2777	Hn	
45	Hy	
3595	N	
126	Y	

<i>EEF12G</i>		<i>Death form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1835		Missing
644	~	Legitimately skipped
6278	N	
254	Y	

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Section B: Myocardial Infarction Ascertainment

Myocardial Infarction Abstraction Form: FORM CODE=MIF, VERSION=B

ID		HCHS Participant ID
N	Value	Description
2921	Present	Text suppressed

VISIT		Visit
N	Value	Description
207	1	Visit 1
222	2	Visit 2
261	3	Visit 3
255	4	Visit 4
254	5	Visit 5
201	6	Visit 6
211	7	Visit 7
266	8	Visit 8
217	9	Visit 9
192	10	Visit 10
261	11	Visit 11
153	12	Visit 12
148	13	Visit 13
67	14	Visit 14
6	15	Visit 15

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<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1968	1	Occurrence 1
530	2	Occurrence 2
194	3	Occurrence 3
83	4	Occurrence 4
52	5	Occurrence 5
30	6	Occurrence 6
15	7	Occurrence 7
10	8	Occurrence 8
8	9	Occurrence 9
9	10	Occurrence 10
4	11	Occurrence 11
4	12	Occurrence 12
1	13	Occurrence 13
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19
7	31	Occurrence 31
1	32	Occurrence 32
2	33	Occurrence 33

<i>MIF0A_FUTIME</i>		<i>Follow-up time from baseline for MIF0A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
2917	Range	594 - 6135 (median=3222 mean=3381.9 std=1122.0)

<i>MIF0AY</i>		<i>Year of MIF0A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
2917	Range	2010 - 2025 (median=2018 mean=2018.8 std=3.1)

<i>MIF0C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2921	Present	Text suppressed

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<i>MIF0D_FUTIME</i>		<i>Follow-up time from baseline for MIF0D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2921	Range	5 - 5416 (median=2350 mean=2375.3 std=1368.4)

<i>MIF0DY</i>		<i>Year of MIF0D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2921	Range	2008 - 2024 (median=2016 mean=2016.0 std=3.8)

<i>MIF1</i>		<i>Type of Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
400	1	1= In hospital only
3	2	2= Emergency Dept. visit only (ED)
2516	3	3= Both ED and in hospital

<i>MIF2_FUTIME</i>		<i>Follow-up time from baseline for MIF2 (Date of arrival)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
66		Missing
2855	Range	5 - 5416 (median=2402 mean=2396.0 std=1372.1)

<i>MIF2A</i>		<i>Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
237		Missing
2684	Range	0:00:00 - 23:59:00

<i>MIF2A1</i>		<i>AM/PM of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42		Missing
2679	.S	Legitimately skipped
92	1	1=AM
108	2	2=PM

<i>MIF2B_FUTIME</i>		<i>Follow-up time from baseline for MIF2B (Date of admission)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2919	Range	5 - 5416 (median=2350 mean=2375.1 std=1367.4)

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Section B: Myocardial Infarction Ascertainment

<i>MIF2BY</i>		<i>Year of MIF2B</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2919	Range	2008 - 2024 (median=2016 mean=2016.0 std=3.8)

<i>MIF2Y</i>		<i>Year of MIF2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
66		Missing
2855	Range	2008 - 2024 (median=2016 mean=2016.1 std=3.8)

<i>MIF3_FUTIME</i>		<i>Follow-up time from baseline for MIF3 (Date of discharge)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
2907	Range	9 - 5418 (median=2353 mean=2382.2 std=1367.1)

<i>MIF3A</i>		<i>Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1245		Missing
1676	Range	0:03:00 - 23:59:00

<i>MIF3A1</i>		<i>AM/PM of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
130		Missing
2679	.S	Legitimately skipped
15	1	1=AM
97	2	2=PM

<i>MIF3Y</i>		<i>Year of MIF3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
2907	Range	2008 - 2024 (median=2016 mean=2016.1 std=3.8)

<i>MIF4</i>		<i>Primary admitting diagnosis code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
365		Missing
2556	Present	Text suppressed

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<i>MIF5</i>		<i>Primary discharge diagnosis code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
290		Missing
2631	Present	Text suppressed

<i>MIF6</i>		<i>EMS unit transport patient to hospital</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2131	0	0=No/NR
790	1	1=Yes

<i>MIF7</i>		<i>Patient transferred to this hospital?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2811	0	0=No/NR
110	1	1=Yes

<i>MIF8</i>		<i>Patient's code ever 'no-code' or 'DNR'</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2800	0	0=No/NR
121	1	1=Yes

<i>MIF9</i>		<i>Patient alive at discharge?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
159	0	0=No/NR
2762	1	1=Yes

<i>MIF9A</i>		<i>Patient dead on arrival?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2762	.S	Legitimately skipped
159	0	0=No
0	1	1=Yes

<i>MIF9B</i>		<i>Patient die in emergency department?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2762	.S	Legitimately skipped
158	0	0=No
1	1	1=Yes

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<i>MIF9C</i>		<i>Was an autopsy performed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
35		Missing
2762	.S	Legitimately skipped
116	0	0=No
8	1	1=Yes

<i>MIF10</i>		<i>Acute episode occur prior to admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
51	0	0=No
2822	1	1=Yes
47	9	9=NR

<i>MIF10A</i>		<i>Time estimation of acute episode</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
99	.S	Legitimately skipped
56	1	1= (< 1 hr)
129	2	2= (>= 1 - < 3 hrs)
88	3	3= (>= 3 - < 6 hrs)
72	4	4= (>= 6 - < 12 hrs)
75	5	5= (>= 12 - < 24 hrs)
1642	6	6= (>= 24 hrs)
759	7	7= Unsure

<i>MIF11</i>		<i>Mention of acute CHD at hospital</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2610	0	0=No
113	1	1=Yes
197	9	9=NR

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<i>MIF12</i>		<i>Acute episode within 72 hours to arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1326	0	0=No
1058	1	1=Yes
537	9	9=NR

<i>MIF12A</i>		<i>Pain/discomfort involve the chest?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1863	.S	Legitimately skipped
13	0	0=No
1038	1	1=Yes
7	9	9=NR

<i>MIF12B</i>		<i>Pain crescendo over time?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1863	.S	Legitimately skipped
388	0	0=No
122	1	1=Yes
548	9	9=NR

<i>MIF12C</i>		<i>Pain/discomfort have non-cardiac origin?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1863	.S	Legitimately skipped
490	0	0=No
130	1	1=Yes
438	9	9=NR

<i>MIF13</i>		<i>Nausea/vomiting related?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1417	0	0=No
696	1	1=Yes
808	9	9=NR

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<i>MIF14</i>		<i>Diaphoresis associated?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1130	0	0=No
211	1	1=Yes
1580	9	9=NR

<i>MIF15</i>		<i>Fatigue/malaise associated ?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
458	0	0=No
563	1	1=Yes
1900	9	9=NR

<i>MIF16A1</i>		<i>Sys. blood pressure at arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
347		Missing
2574	Range	0 - 271 (median=137 mean=139.7 std=30.2)

<i>MIF16A2</i>		<i>Dia. blood pressure at arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
356		Missing
2565	Range	0 - 191 (median=75 mean=76.2 std=16.8)

<i>MIF16B</i>		<i>Heart rate (bpm)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
394		Missing
2527	Range	0 - 234 (median=84 mean=86.7 std=22.5)

<i>MIF17A</i>		<i>Myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2468	0	0=No/NR
452	1	1=Yes

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<i>MIF17A1</i>		<i>If history of MI, MI within 4 weeks?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2469	.S	Legitimately skipped
424	0	0=No/NR
27	1	1=Yes

<i>MIF17B</i>		<i>Angina</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2590	0	0=No/NR
331	1	1=Yes

<i>MIF17C</i>		<i>Percutaneous coronary intervention</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2494	0	0=No/NR
427	1	1=Yes

<i>MIF17D</i>		<i>CABG</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2567	0	0=No/NR
354	1	1=Yes

<i>MIF17E</i>		<i>Coronary artery disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1807	0	0=No/NR
1114	1	1=Yes

<i>MIF17F</i>		<i>Heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2209	0	0=No/NR
712	1	1=Yes

<i>MIF17G</i>		<i>Arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2428	0	0=No/NR
492	1	1=Yes

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<i>MIF17G1</i>		<i>Atrial Fibrillation/Flutter</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2428	.S	Legitimately skipped
89	0	0=No/NR
401	1	1=Yes

<i>MIF17G2</i>		<i>Ventricular Fibrillation/Tachycardia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2428	.S	Legitimately skipped
451	0	0=No/NR
39	1	1=Yes

<i>MIF17G3</i>		<i>Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2428	.S	Legitimately skipped
417	0	0=No/NR
73	1	1=Yes

<i>MIF18A</i>		<i>Indicated Angina</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2470	0	0=No/NR
451	1	1=Yes

<i>MIF18B</i>		<i>Indicated Acute myocardial Infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2623	0	0=No/NR
298	1	1=Yes

<i>MIF18C</i>		<i>Indicated ST elevation not on ECG</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2893	0	0=No/NR
28	1	1=Yes

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<i>MIF18D</i>		<i>Indicated congestive heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2382	0	0=No/NR
539	1	1=Yes

<i>MIF18D1</i>		<i>If yes, heart failure/pulmonary edema?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2382	.S	Legitimately skipped
456	0	0=No/NR
83	1	1=Yes

<i>MIF18E</i>		<i>Indicated shock or cardiogenic shock</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2860	0	0=No/NR
61	1	1=Yes

<i>MIF18E1</i>		<i>If yes, shock occur within 24 hours</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2860	.S	Legitimately skipped
44	0	0=No/NR
17	1	1=Yes

<i>MIF18F</i>		<i>Indicated ventricular fibrillation/asystole</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2782	0	0=No/NR
139	1	1=Yes

<i>MIF18F1</i>		<i>If yes, arrest occur within 24 hours</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2782	.S	Legitimately skipped
109	0	0=No/NR
30	1	1=Yes

<i>MIF18G</i>		<i>Indicated Ventricular Tachycardia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2872	0	0=No/NR
49	1	1=Yes

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<i>MIF18H</i>		<i>Indicated Atrial fibrillation or atrial flut</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2625	0	0=No/NR
296	1	1=Yes

<i>MIF19</i>		<i>Cardiac enzymes reported within days</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
1075	0	0=No/NR
1844	1	1=Yes

<i>MIF19A</i>		<i>Cardiac enzymes reported arrival day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
969	.S	Legitimately skipped
210	0	0=No/NR
1741	1	1=Yes

<i>MIF19B</i>		<i>Cardiac enzymes reported day after arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
969	.S	Legitimately skipped
1237	0	0=No/NR
714	1	1=Yes

<i>MIF19C</i>		<i>Cardiac enzymes reported third day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
969	.S	Legitimately skipped
1762	0	0=No/NR
189	1	1=Yes

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<i>MIF19D</i>		<i>Cardiac enzymes reported fourth day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
969	.S	Legitimately skipped
1863	0	0=No/NR
88	1	1=Yes

<i>MIF20A1</i>		<i>Set 1 CK Upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1211		Missing
1076	.S	Legitimately skipped
634	Range	20 - 499 (median=200 mean=201.1 std=56.2)

<i>MIF20A2</i>		<i>Set 1 CK units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1143		Missing
1076	.S	Legitimately skipped
3	1	1=ng/mL
699	2	2=Units/L
0	3	3=Åµg/L

<i>MIF20B1</i>		<i>Set 1 CK-MB upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1396		Missing
1076	.S	Legitimately skipped
449	Range	0.5 - 10 (median=5 mean=5.4 std=1.7)

<i>MIF20B2</i>		<i>Set 1 CK-MB units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1388		Missing
1076	.S	Legitimately skipped
456	1	1=ng/mL
0	2	2=Units/L
1	3	3=Åµg/L

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<i>MIF20C1</i>		<i>Set 1 Total LDH upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1766		Missing
1076	.S	Legitimately skipped
79	Range	179 - 618 (median=214 mean=248.6 std=108.6)

<i>MIF20C2</i>		<i>Set 1 Total LDH units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1762		Missing
1076	.S	Legitimately skipped
0	1	1=ng/mL
83	2	2=Units/L
0	3	3=Åµg/L

<i>MIF20D1</i>		<i>Set 1 LDH-1 upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1844		Missing
1076	.S	Legitimately skipped
1	Range	100 - 100 (median=100 mean=100.0 std=.)

<i>MIF20D2</i>		<i>Set 1 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1843		Missing
1076	.S	Legitimately skipped
0	1	1=ng/mL
2	2	2=Units/L
0	3	3=Åµg/L

<i>MIF20E1</i>		<i>Set 1 LDH-2 upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1844		Missing
1076	.S	Legitimately skipped
1	Range	100 - 100 (median=100 mean=100.0 std=.)

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<i>MIF20E2</i>		<i>Set 1 LDH-2 units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1844		Missing
1076	.S	Legitimately skipped
1	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF20F1</i>		<i>Set 1 Troponin upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
656		Missing
1076	.S	Legitimately skipped
1189	Range	0.01 - 79 (median=0.056 mean=2.5514 std=8.5977)

<i>MIF20F2</i>		<i>Set 1 Troponin units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
637		Missing
1076	.S	Legitimately skipped
1188	1	1=ng/mL
1	2	2=Units/L
19	3	3=Åµg/L

<i>MIF20F3</i>		<i>Set 1 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
200		Missing
1076	.S	Legitimately skipped
167	1	1= Troponin, type not specified
834	2	2= Troponin I
506	3	3= Troponin T
136	4	4= High Sensitivity Troponin (HS)
2	5	5= Unsure

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<i>MIF20G</i>		<i>second set of normal values</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
22		Missing
969	.S	Legitimately skipped
1886	0	0=No/NR
44	1	1=Yes

<i>MIF21A1</i>		<i>Set 2 Total CK upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
169		Missing
2744	.S	Legitimately skipped
8	Range	41 - 374 (median=250 mean=238.1 std=102.1)

<i>MIF21A2</i>		<i>Set 2 Total CK units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
170		Missing
2744	.S	Legitimately skipped
0	1	1=ng/mL
7	2	2=Units/L
0	3	3=Åµg/L

<i>MIF21B1</i>		<i>Set 2 CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
171		Missing
2744	.S	Legitimately skipped
6	Range	4.2 - 10 (median=5.85 mean=6.800 std=2.612)

<i>MIF21B2</i>		<i>Set 2 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172		Missing
2744	.S	Legitimately skipped
5	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF21C1</i>		<i>Set 2 Total LDH Upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
177		Missing
2744	.S	Legitimately skipped

<i>MIF21C2</i>		<i>Set 2 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
177		Missing
2744	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF21D1</i>		<i>Set 2 LDH-1 upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
177		Missing
2744	.S	Legitimately skipped

<i>MIF21D2</i>		<i>Set 2 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
177		Missing
2744	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF21E1</i>		<i>Set 2 LDH-2 upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
177		Missing
2744	.S	Legitimately skipped

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<i>MIF21E2</i>		<i>Set 2 LDH-2 units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
177		Missing
2744	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF21F1</i>		<i>Set 2 Troponin upper limit of normal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
138		Missing
2744	.S	Legitimately skipped
39	Range	0.01 - 79 (median=0.07 mean=2.230 std=12.641)

<i>MIF21F2</i>		<i>Set 2 Troponin units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
141		Missing
2744	.S	Legitimately skipped
36	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF21F3</i>		<i>Set 2 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
135		Missing
2744	.S	Legitimately skipped
4	1	1= Troponin, type not specified
30	2	2= Troponin I
6	3	3= Troponin T
2	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure

<i>MIF22_FUTIME</i>		<i>Follow-up time from baseline for MIF22 (Date of Day 1/Set 1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1183		Missing
1738	Range	-78 - 5302 (median=2483 mean=2448.5 std=1392.2)

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MIF22A		Day 1/Set 1 Total CK (CPK)
N	Value	Description
1029		Missing
1173	.S	Legitimately skipped
719	Range	11 - 28480 (median=102 mean=371.6 std=1776.8)

MIF22A1		Day 1/Set 1 Total CK (CPK) Units
N	Value	Description
1148		Missing
1173	.S	Legitimately skipped
4	1	1=ng/mL
596	2	2=Units/L
0	3	3=Åµg/L

MIF22A2		Day 1/Set 1 Total CK Range Set
N	Value	Description
1095		Missing
1173	.S	Legitimately skipped
653	1	1=Range Set 1
0	2	2=Range Set 2

MIF22A3		Day 1/Set 1 Total CK Words Code
N	Value	Description
1590		Missing
1173	.S	Legitimately skipped
114	1	1=absent/negative/normal
2	2	2=trace/weak positive
42	3	3=present/positive/abnormal

MIF22B		Day 1/Set 1 CK-MB
N	Value	Description
1293		Missing
1173	.S	Legitimately skipped
455	Range	0.22 - 300 (median=2.6 mean=6.73 std=23.72)

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<i>MIF22B1</i>		<i>Day 1/Set 1 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1375		Missing
1173	.S	Legitimately skipped
372	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L

<i>MIF22B2</i>		<i>Day 1/Set 1 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1360		Missing
1173	.S	Legitimately skipped
383	1	1=Range Set 1
5	2	2=Range Set 2

<i>MIF22B3</i>		<i>Day 1/Set 1 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1670		Missing
1173	.S	Legitimately skipped
69	1	1=absent/negative/normal
1	2	2=trace/weak positive
8	3	3=present/positive/abnormal

<i>MIF22C</i>		<i>Day 1/Set 1 Total LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1657		Missing
1173	.S	Legitimately skipped
91	Range	19 - 3840 (median=273 mean=420.5 std=494.4)

<i>MIF22C1</i>		<i>Day 1/Set 1 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1688		Missing
1173	.S	Legitimately skipped
1	1	1=ng/mL
59	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF22C2</i>		<i>Day 1/Set 1 Total LDH Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1665		Missing
1173	.S	Legitimately skipped
82	1	1=Range Set 1
1	2	2=Range Set 2

<i>MIF22C3</i>		<i>Day 1/Set 1 Total LDH Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1730		Missing
1173	.S	Legitimately skipped
8	1	1=absent/negative/normal
0	2	2=trace/weak positive
10	3	3=present/positive/abnormal

<i>MIF22D</i>		<i>Day 1/Set 1 LDH-1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1584		Missing
1173	.S	Legitimately skipped
164	Range	1 - 192 (median=16 mean=17.3 std=20.2)

<i>MIF22D1</i>		<i>Day 1/Set 1 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1746		Missing
1173	.S	Legitimately skipped
0	1	1=ng/mL
2	2	2=Units/L
0	3	3=Åµg/L

<i>MIF22D2</i>		<i>Day 1/Set 1 LDH-1 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1746		Missing
1173	.S	Legitimately skipped
2	1	1=Range Set 1
0	2	2=Range Set 2

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<i>MIF22D3</i>		<i>Day 1/Set 1 LDH-1 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1748		Missing
1173	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF22E</i>		<i>Day 1/Set 1 LDH-2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1746		Missing
1173	.S	Legitimately skipped
2	Range	26 - 32.7 (median=29.35 mean=29.350 std=4.738)

<i>MIF22E1</i>		<i>Day 1/Set 1 LDH-2 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1748		Missing
1173	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF22E2</i>		<i>Day 1/Set 1 LDH-2 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1748		Missing
1173	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF22E3</i>		<i>Day 1/Set 1 LDH-2 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1747		Missing
1173	.S	Legitimately skipped
1	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

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<i>MIF22F</i>		<i>Day 1/Set 1 Troponin <</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
159		Missing
1173	.S	Legitimately skipped
896	0	0=n/a
693	1	1=<

<i>MIF22F0</i>		<i>Day 1/Set 1 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134		Missing
1173	.S	Legitimately skipped
1614	Range	0 - 11551 (median=0.03 mean=24.960 std=343.233)

<i>MIF22F1</i>		<i>Day 1/Set 1 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
575		Missing
1173	.S	Legitimately skipped
1157	1	1=ng/mL
2	2	2=Units/L
14	3	3=Åµg/L

<i>MIF22F2</i>		<i>Day 1/Set 1 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
245		Missing
1173	.S	Legitimately skipped
1497	1	1=Range Set 1
6	2	2=Range Set 2

<i>MIF22F3</i>		<i>Day 1/Set 1 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
946		Missing
1173	.S	Legitimately skipped
496	1	1=absent/negative/normal
28	2	2=trace/weak positive
278	3	3=present/positive/abnormal

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<i>MIF22F4</i>		<i>Day 1/Set 1 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79		Missing
1173	.S	Legitimately skipped
206	1	1= Troponin, type not specified
828	2	2= Troponin I
492	3	3= Troponin T
134	4	4= High Sensitivity Troponin (HS)
9	5	5= Unsure

<i>MIF22G</i>		<i>second set of enzymes reported the first day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
1066	.S	Legitimately skipped
1188	0	0=No/NR
665	1	1=Yes

<i>MIF22Y</i>		<i>Year of MIF22</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1181		Missing
1740	Range	2008 - 2024 (median=2016 mean=2016.2 std=3.8)

<i>MIF23_FUTIME</i>		<i>Follow-up time from baseline for MIF23 (Date of Day 1/Set 2)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2260		Missing
661	Range	-78 - 5302 (median=2344 mean=2363.1 std=1396.3)

<i>MIF23A</i>		<i>Day 1/Set 2 Total CK (CPK)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
494		Missing
2184	.S	Legitimately skipped
243	Range	20 - 30698 (median=108 mean=510.2 std=2229.4)

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<i>MIF23A1</i>		<i>Day 1/Set 2 Total CK Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
517		Missing
2184	.S	Legitimately skipped
2	1	1=ng/mL
218	2	2=Units/L
0	3	3=Åµg/L

<i>MIF23A2</i>		<i>Day 1/Set 2 Total CK Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
514		Missing
2184	.S	Legitimately skipped
221	1	1=Range Set 1
2	2	2=Range Set 2

<i>MIF23A3</i>		<i>Day 1/Set 2 Total CK Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
692		Missing
2184	.S	Legitimately skipped
39	1	1=absent/negative/normal
0	2	2=trace/weak positive
6	3	3=present/positive/abnormal

<i>MIF23B</i>		<i>Day 1/ Set 2 CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
559		Missing
2184	.S	Legitimately skipped
178	Range	0.28 - 962.4 (median=2.6 mean=25.07 std=92.68)

<i>MIF23B1</i>		<i>Day 1/ Set 2 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
581		Missing
2184	.S	Legitimately skipped
156	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF23B2</i>		<i>Day 1/ Set 2 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
577		Missing
2184	.S	Legitimately skipped
156	1	1=Range Set 1
4	2	2=Range Set 2

<i>MIF23B3</i>		<i>Day 1/ Set 2 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
708		Missing
2184	.S	Legitimately skipped
25	1	1=absent/negative/normal
1	2	2=trace/weak positive
3	3	3=present/positive/abnormal

<i>MIF23C</i>		<i>Day 1/Set 2 Total LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
724		Missing
2184	.S	Legitimately skipped
13	Range	140 - 1280 (median=203 mean=322.9 std=308.2)

<i>MIF23C1</i>		<i>Day 1/Set 2 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
725		Missing
2184	.S	Legitimately skipped
0	1	1=ng/mL
12	2	2=Units/L
0	3	3=Åµg/L

<i>MIF23C2</i>		<i>Day 1/Set 2 Total LDH Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
724		Missing
2184	.S	Legitimately skipped
13	1	1=Range Set 1
0	2	2=Range Set 2

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<i>MIF23C3</i>		<i>Day 1/Set 2 Total LDH Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
735		Missing
2184	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
2	3	3=present/positive/abnormal

<i>MIF23D</i>		<i>Day 1/Set 2 LDH-1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
736		Missing
2184	.S	Legitimately skipped
1	Range	26 - 26 (median=26 mean=26.0 std=.)

<i>MIF23D1</i>		<i>Day 1/Set 2 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
737		Missing
2184	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF23D2</i>		<i>Day 1/Set 2 LDH-1 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
736		Missing
2184	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF23D3</i>		<i>Day 1/Set 2 LDH-1 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
737		Missing
2184	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

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MIF23E		Day 1/Set 2 LDH-2
N	Value	Description
736		Missing
2184	.S	Legitimately skipped
1	Range	30.4 - 30.4 (median=30.4 mean=30.40 std=.)

MIF23E1		Day 1/Set 2 LDH-2 Units
N	Value	Description
737		Missing
2184	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

MIF23E2		Day 1/Set 2 LDH-2 Range Set
N	Value	Description
736		Missing
2184	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2

MIF23E3		Day 1/Set 2 LDH-2 Words Code
N	Value	Description
737		Missing
2184	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

MIF23F		Day 1/Set 2 Troponin <
N	Value	Description
115		Missing
2184	.S	Legitimately skipped
404	0	0=n/a
218	1	1=<

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<i>MIF23F0</i>		<i>Day 1/Set 2 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
109		Missing
2184	.S	Legitimately skipped
628	Range	0 - 4858.5 (median=0.07 mean=48.196 std=365.432)

<i>MIF23F1</i>		<i>Day 1/Set 2 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
270		Missing
2184	.S	Legitimately skipped
462	1	1=ng/mL
0	2	2=Units/L
5	3	3=Åµg/L

<i>MIF23F2</i>		<i>Day 1/Set 2 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
145		Missing
2184	.S	Legitimately skipped
571	1	1=Range Set 1
21	2	2=Range Set 2

<i>MIF23F3</i>		<i>Day 1/Set 2 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
414		Missing
2184	.S	Legitimately skipped
169	1	1=absent/negative/normal
9	2	2=trace/weak positive
145	3	3=present/positive/abnormal

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<i>MIF23F4</i>		<i>Day 1/Set 2 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
91		Missing
2184	.S	Legitimately skipped
71	1	1= Troponin, type not specified
328	2	2= Troponin I
183	3	3= Troponin T
64	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure

<i>MIF23Y</i>		<i>Year of MIF23</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2258		Missing
663	Range	2008 - 2024 (median=2016 mean=2015.9 std=3.8)

<i>MIF24_FUTIME</i>		<i>Follow-up time from baseline for MIF24 (Date of Day 2/Set 1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2209		Missing
712	Range	6 - 5208 (median=2147.5 mean=2206.34 std=1327.57)

<i>MIF24A</i>		<i>Day 2/Set 1 Total CK (CPK)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
451		Missing
2137	.S	Legitimately skipped
333	Range	18 - 25413 (median=95 mean=563.7 std=2265.0)

<i>MIF24A1</i>		<i>Day 2/Set 1 Total CK (CPK) Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
491		Missing
2137	.S	Legitimately skipped
1	1	1=ng/mL
292	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF24A2</i>		<i>Day 2/Set 1 Total CK (CPK) Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
479		Missing
2137	.S	Legitimately skipped
303	1	1=Range Set 1
2	2	2=Range Set 2

<i>MIF24A3</i>		<i>Day 2/Set 1 Total CK (CPK) Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
734		Missing
2137	.S	Legitimately skipped
37	1	1=absent/negative/normal
0	2	2=trace/weak positive
13	3	3=present/positive/abnormal

<i>MIF24B</i>		<i>Day 2/Set 1 CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
545		Missing
2137	.S	Legitimately skipped
239	Range	0 - 613 (median=2.5 mean=16.32 std=56.17)

<i>MIF24B1</i>		<i>Day 2/Set 1 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
572		Missing
2137	.S	Legitimately skipped
212	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF24B2</i>		<i>Day 2/Set 1 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
567		Missing
2137	.S	Legitimately skipped
212	1	1=Range Set 1
5	2	2=Range Set 2

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<i>MIF24B3</i>		<i>Day 2/Set 1 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
746		Missing
2137	.S	Legitimately skipped
33	1	1=absent/negative/normal
0	2	2=trace/weak positive
5	3	3=present/positive/abnormal

<i>MIF24C</i>		<i>Day 2/Set 1 Total LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
763		Missing
2137	.S	Legitimately skipped
21	Range	131 - 1487 (median=291 mean=483.7 std=406.7)

<i>MIF24C1</i>		<i>Day 2/Set 1 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
765		Missing
2137	.S	Legitimately skipped
1	1	1=ng/mL
18	2	2=Units/L
0	3	3=µg/L

<i>MIF24C2</i>		<i>Day 2/Set 1 Total LDH Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
764		Missing
2137	.S	Legitimately skipped
20	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF24C3</i>		<i>Day 2/Set 1 Total LDH Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
781		Missing
2137	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
3	3	3=present/positive/abnormal

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MIF24D		Day 2/Set 1 LDH-1
N	Value	Description
705		Missing
2137	.S	Legitimately skipped
79	Range	3 - 31 (median=14 mean=15.9 std=8.2)

MIF24D1		Day 2/Set 1 LDH-1 Units
N	Value	Description
784		Missing
2137	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

MIF24D2		Day 2/Set 1 LDH-1 Range Set
N	Value	Description
783		Missing
2137	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2

MIF24D3		Day 2/Set 1 LDH-1 Words Code
N	Value	Description
784		Missing
2137	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

MIF24E		Day 2/Set 1 LDH-2
N	Value	Description
782		Missing
2137	.S	Legitimately skipped
2	Range	28.9 - 195 (median=111.95 mean=111.950 std=117.450)

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<i>MIF24E1</i>		<i>Day 2/Set 1 LDH-2 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
783		Missing
2137	.S	Legitimately skipped
0	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L

<i>MIF24E2</i>		<i>Day 2/Set 1 LDH-2 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
782		Missing
2137	.S	Legitimately skipped
2	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF24E3</i>		<i>Day 2/Set 1 LDH-2 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
784		Missing
2137	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF24F</i>		<i>Day 2/Set 1 Troponin <</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
137		Missing
2137	.S	Legitimately skipped
412	0	0=n/a
235	1	1=<

<i>MIF24F0</i>		<i>Day 2/Set 1 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126		Missing
2137	.S	Legitimately skipped
658	Range	0 - 8229 (median=0.05 mean=57.918 std=525.489)

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<i>MIF24F1</i>		<i>Day 2/Set 1 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
246		Missing
2137	.S	Legitimately skipped
530	1	1=ng/mL
0	2	2=Units/L
8	3	3=Åµg/L

<i>MIF24F2</i>		<i>Day 2/Set 1 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
173		Missing
2137	.S	Legitimately skipped
598	1	1=Range Set 1
13	2	2=Range Set 2

<i>MIF24F3</i>		<i>Day 2/Set 1 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
489		Missing
2137	.S	Legitimately skipped
157	1	1=absent/negative/normal
6	2	2=trace/weak positive
132	3	3=present/positive/abnormal

<i>MIF24F4</i>		<i>Day 2/Set 1 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
119		Missing
2137	.S	Legitimately skipped
39	1	1= Troponin, type not specified
345	2	2= Troponin I
242	3	3= Troponin T
35	4	4= High Sensitivity Troponin (HS)
4	5	5= Unsure

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<i>MIF24G</i>		<i>second set of enzymes reported second day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2030	.S	Legitimately skipped
599	0	0=No/NR
291	1	1=Yes

<i>MIF24Y</i>		<i>Year of MIF24</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2208		Missing
713	Range	2008 - 2025 (median=2015 mean=2015.5 std=3.7)

<i>MIF25_FUTIME</i>		<i>Follow-up time from baseline for MIF25 (Date of Day 2/Set 2)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2631		Missing
290	Range	6 - 5446 (median=2134 mean=2172.8 std=1254.2)

<i>MIF25A</i>		<i>Day 2/Set 2 Total CK (CPK)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
258		Missing
2517	.S	Legitimately skipped
146	Range	12 - 14559 (median=103 mean=566.0 std=1772.9)

<i>MIF25A1</i>		<i>Day 2/Set 2 Total CK (CPK) Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
266		Missing
2517	.S	Legitimately skipped
0	1	1=ng/mL
138	2	2=Units/L
0	3	3=Åµg/L

<i>MIF25A2</i>		<i>Day 2/Set 2 Total CK (CPK) Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
262		Missing
2517	.S	Legitimately skipped
139	1	1=Range Set 1
3	2	2=Range Set 2

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<i>MIF25A3</i>		<i>Day 2/Set 2 Total CK (CPK) Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
372		Missing
2517	.S	Legitimately skipped
22	1	1=absent/negative/normal
0	2	2=trace/weak positive
10	3	3=present/positive/abnormal

<i>MIF25B</i>		<i>Day 2/Set 2 CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
295		Missing
2517	.S	Legitimately skipped
109	Range	0.06 - 207.3 (median=2.3 mean=14.79 std=39.06)

<i>MIF25B1</i>		<i>Day 2/Set 2 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
298		Missing
2517	.S	Legitimately skipped
104	1	1=ng/mL
2	2	2=Units/L
0	3	3=Åµg/L

<i>MIF25B2</i>		<i>Day 2/Set 2 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
297		Missing
2517	.S	Legitimately skipped
102	1	1=Range Set 1
5	2	2=Range Set 2

<i>MIF25B3</i>		<i>Day 2/Set 2 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
381		Missing
2517	.S	Legitimately skipped
19	1	1=absent/negative/normal
0	2	2=trace/weak positive
4	3	3=present/positive/abnormal

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MIF25C		Day 2/Set 2 Total LDH
N	Value	Description
397		Missing
2517	.S	Legitimately skipped
7	Range	110 - 493 (median=175 mean=223.1 std=129.3)

MIF25C1		Day 2/Set 2 Total LDH Units
N	Value	Description
397		Missing
2517	.S	Legitimately skipped
0	1	1=ng/mL
7	2	2=Units/L
0	3	3=Åµg/L

MIF25C2		Day 2/Set 2 Total LDH Range Set
N	Value	Description
397		Missing
2517	.S	Legitimately skipped
7	1	1=Range Set 1
0	2	2=Range Set 2

MIF25C3		Day 2/Set 2 Total LDH Words Code
N	Value	Description
404		Missing
2517	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

MIF25D		Day 2/Set 2 LDH-1
N	Value	Description
378		Missing
2517	.S	Legitimately skipped
26	Range	4 - 29 (median=15.5 mean=16.08 std=7.58)

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<i>MIF25D1</i>		<i>Day 2/Set 2 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
404		Missing
2517	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF25D2</i>		<i>Day 2/Set 2 LDH-1 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
404		Missing
2517	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF25D3</i>		<i>Day 2/Set 2 LDH-1 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
404		Missing
2517	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF25E</i>		<i>Day 2/Set 2 LDH-2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
404		Missing
2517	.S	Legitimately skipped

<i>MIF25E1</i>		<i>Day 2/Set 2 LDH-2 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
404		Missing
2517	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF25E2</i>		<i>Day 2/Set 2 LDH-2 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
404		Missing
2517	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF25E3</i>		<i>Day 2/Set 2 LDH-2 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
404		Missing
2517	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF25F</i>		<i>Day 2/Set 2 Troponin <</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
143		Missing
2517	.S	Legitimately skipped
184	0	0=n/a
77	1	1=<

<i>MIF25F0</i>		<i>Day 2/Set 2 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
137		Missing
2517	.S	Legitimately skipped
267	Range	0 - 11000 (median=0.1 mean=108.45 std=841.03)

<i>MIF25F1</i>		<i>Day 2/Set 2 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
177		Missing
2517	.S	Legitimately skipped
221	1	1=ng/mL
0	2	2=Units/L
6	3	3=Åµg/L

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<i>MIF25F2</i>		<i>Day 2/Set 2 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
147		Missing
2517	.S	Legitimately skipped
251	1	1=Range Set 1
6	2	2=Range Set 2

<i>MIF25F3</i>		<i>Day 2/Set 2 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
281		Missing
2517	.S	Legitimately skipped
54	1	1=absent/negative/normal
1	2	2=trace/weak positive
68	3	3=present/positive/abnormal

<i>MIF25F4</i>		<i>Day 2/Set 2 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
135		Missing
2517	.S	Legitimately skipped
13	1	1= Troponin, type not specified
148	2	2= Troponin I
92	3	3= Troponin T
16	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure

<i>MIF25Y</i>		<i>Year of MIF25</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2630		Missing
291	Range	2008 - 2024 (median=2015 mean=2015.5 std=3.6)

<i>MIF26_FUTIME</i>		<i>Follow-up time from baseline for MIF26 (Date of Day 3/Set 1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2732		Missing
189	Range	-108 - 5083 (median=2125 mean=2171.5 std=1294.5)

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MIF26A		Day 3/Set 1 Total CK (CPK)
N	Value	Description
229		Missing
2608	.S	Legitimately skipped
84	Range	24 - 14206 (median=172.5 mean=989.75 std=2516.06)

MIF26A1		Day 3/Set 1 Total CK (CPK) Units
N	Value	Description
238		Missing
2608	.S	Legitimately skipped
0	1	1=ng/mL
75	2	2=Units/L
0	3	3=Åµg/L

MIF26A2		Day 3/Set 1 Total CK (CPK) Range Set
N	Value	Description
235		Missing
2608	.S	Legitimately skipped
77	1	1=Range Set 1
1	2	2=Range Set 2

MIF26A3		Day 3/Set 1 Total CK (CPK) Words Code
N	Value	Description
301		Missing
2608	.S	Legitimately skipped
4	1	1=absent/negative/normal
0	2	2=trace/weak positive
8	3	3=present/positive/abnormal

MIF26B		Day 3/Set 1 CK-MB
N	Value	Description
255		Missing
2608	.S	Legitimately skipped
58	Range	0.22 - 300 (median=2.75 mean=13.261 std=41.165)

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<i>MIF26B1</i>		<i>Day 3/Set 1 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
258		Missing
2608	.S	Legitimately skipped
54	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L

<i>MIF26B2</i>		<i>Day 3/Set 1 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
257		Missing
2608	.S	Legitimately skipped
55	1	1=Range Set 1
1	2	2=Range Set 2

<i>MIF26B3</i>		<i>Day 3/Set 1 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
305		Missing
2608	.S	Legitimately skipped
7	1	1=absent/negative/normal
0	2	2=trace/weak positive
1	3	3=present/positive/abnormal

<i>MIF26C</i>		<i>Day 3/Set 1 Total LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
296		Missing
2608	.S	Legitimately skipped
17	Range	137 - 5984 (median=305 mean=827.4 std=1385.5)

<i>MIF26C1</i>		<i>Day 3/Set 1 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
302		Missing
2608	.S	Legitimately skipped
0	1	1=ng/mL
11	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF26C2</i>		<i>Day 3/Set 1 Total LDH Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
299		Missing
2608	.S	Legitimately skipped
14	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF26C3</i>		<i>Day 3/Set 1 Total LDH Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
308		Missing
2608	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
5	3	3=present/positive/abnormal

<i>MIF26D</i>		<i>Day 3/Set 1 LDH-1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
296		Missing
2608	.S	Legitimately skipped
17	Range	1 - 30 (median=14 mean=14.7 std=8.0)

<i>MIF26D1</i>		<i>Day 3/Set 1 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
313		Missing
2608	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF26D2</i>		<i>Day 3/Set 1 LDH-1 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
312		Missing
2608	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2

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<i>MIF26D3</i>		<i>Day 3/Set 1 LDH-1 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
313		Missing
2608	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF26E</i>		<i>Day 3/Set 1 LDH-2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
312		Missing
2608	.S	Legitimately skipped
1	Range	29.8 - 29.8 (median=29.8 mean=29.80 std=.)

<i>MIF26E1</i>		<i>Day 3/Set 1 LDH-2 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
313		Missing
2608	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF26E2</i>		<i>Day 3/Set 1 LDH-2 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
312		Missing
2608	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF26E3</i>		<i>Day 3/Set 1 LDH-2 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
313		Missing
2608	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

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<i>MIF26F</i>		<i>Day 3/Set 1 Troponin <</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
168		Missing
2608	.S	Legitimately skipped
108	0	0=n/a
37	1	1=<

<i>MIF26F0</i>		<i>Day 3/Set 1 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
165		Missing
2608	.S	Legitimately skipped
148	Range	0 - 12000 (median=0.135 mean=176.0745 std=1373.2054)

<i>MIF26F1</i>		<i>Day 3/Set 1 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
181		Missing
2608	.S	Legitimately skipped
128	1	1=ng/mL
0	2	2=Units/L
4	3	3=Åµg/L

<i>MIF26F2</i>		<i>Day 3/Set 1 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
170		Missing
2608	.S	Legitimately skipped
138	1	1=Range Set 1
5	2	2=Range Set 2

<i>MIF26F3</i>		<i>Day 3/Set 1 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
261		Missing
2608	.S	Legitimately skipped
21	1	1=absent/negative/normal
0	2	2=trace/weak positive
31	3	3=present/positive/abnormal

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<i>MIF26F4</i>		<i>Day 3/Set 1 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
163		Missing
2608	.S	Legitimately skipped
11	1	1= Troponin, type not specified
82	2	2= Troponin I
48	3	3= Troponin T
7	4	4= High Sensitivity Troponin (HS)
2	5	5= Unsure

<i>MIF26G</i>		<i>second set of enzymes reported third day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2501	.S	Legitimately skipped
384	0	0=No/NR
34	1	1=Yes

<i>MIF26Y</i>		<i>Year of MIF26</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2732		Missing
189	Range	2008 - 2023 (median=2016 mean=2015.5 std=3.6)

<i>MIF27_FUTIME</i>		<i>Follow-up time from baseline for MIF27 (Date of Day 3/Set 2)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2888		Missing
33	Range	504 - 5070 (median=1854 mean=2112.2 std=1167.7)

<i>MIF27A</i>		<i>Day 3/Set 2 Total CK (CPK)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2755	.S	Legitimately skipped
14	Range	57 - 10891 (median=194 mean=1252.7 std=2933.0)

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<i>MIF27A1</i>		<i>Day 3/Set 2 Total CK (CPK) Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2755	.S	Legitimately skipped
0	1	1=ng/mL
14	2	2=Units/L
0	3	3=Åµg/L

<i>MIF27A2</i>		<i>Day 3/Set 2 Total CK (CPK) Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2755	.S	Legitimately skipped
14	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF27A3</i>		<i>Day 3/Set 2 Total CK (CPK) Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
161		Missing
2755	.S	Legitimately skipped
2	1	1=absent/negative/normal
0	2	2=trace/weak positive
3	3	3=present/positive/abnormal

<i>MIF27B</i>		<i>Day 3/Set 2 CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
155		Missing
2755	.S	Legitimately skipped
11	Range	0.83 - 7.1 (median=2.7 mean=3.62 std=2.37)

<i>MIF27B1</i>		<i>Day 3/Set 2 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
155		Missing
2755	.S	Legitimately skipped
11	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF27B2</i>		<i>Day 3/Set 2 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
155		Missing
2755	.S	Legitimately skipped
11	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF27B3</i>		<i>Day 3/Set 2 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
164		Missing
2755	.S	Legitimately skipped
2	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF27C</i>		<i>Day 3/Set 2 Total LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped

<i>MIF27C1</i>		<i>Day 3/Set 2 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF27C2</i>		<i>Day 3/Set 2 Total LDH Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

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<i>MIF27C3</i>		<i>Day 3/Set 2 Total LDH Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF27D</i>		<i>Day 3/Set 2 LDH-1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped

<i>MIF27D1</i>		<i>Day 3/Set 2 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF27D2</i>		<i>Day 3/Set 2 LDH-1 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF27D3</i>		<i>Day 3/Set 2 LDH-1 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

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<i>MIF27E</i>		<i>Day 3/Set 2 LDH-2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped

<i>MIF27E1</i>		<i>Day 3/Set 2 LDH-2 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF27E2</i>		<i>Day 3/Set 2 LDH-2 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF27E3</i>		<i>Day 3/Set 2 LDH-2 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
166		Missing
2755	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF27F</i>		<i>Day 3/Set 2 Troponin <</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
135		Missing
2755	.S	Legitimately skipped
24	0	0=n/a
7	1	1=<

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<i>MIF27F0</i>		<i>Day 3/Set 2 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
137		Missing
2755	.S	Legitimately skipped
29	Range	0.01 - 1576.3 (median=0.09 mean=56.249 std=292.399)

<i>MIF27F1</i>		<i>Day 3/Set 2 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
139		Missing
2755	.S	Legitimately skipped
26	1	1=ng/mL
0	2	2=Units/L
1	3	3=Åµg/L

<i>MIF27F2</i>		<i>Day 3/Set 2 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
138		Missing
2755	.S	Legitimately skipped
28	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF27F3</i>		<i>Day 3/Set 2 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
154		Missing
2755	.S	Legitimately skipped
5	1	1=absent/negative/normal
0	2	2=trace/weak positive
7	3	3=present/positive/abnormal

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MIF27F4		Day 3/Set 2 Type of Troponin
N	Value	Description
137		Missing
2755	.S	Legitimately skipped
1	1	1= Troponin, type not specified
16	2	2= Troponin I
10	3	3= Troponin T
2	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure

MIF27Y		Year of MIF27
N	Value	Description
2888		Missing
33	Range	2010 - 2022 (median=2015 mean=2015.5 std=3.3)

MIF28_FUTIME		Follow-up time from baseline for MIF28 (Date of Day 4/Set 1)
N	Value	Description
2833		Missing
88	Range	50 - 5071 (median=1979 mean=2199.9 std=1347.1)

MIF28A		Day 4/Set 1 Total CK (CPK)
N	Value	Description
171		Missing
2707	.S	Legitimately skipped
43	Range	20 - 44034 (median=98 mean=1541.8 std=6831.8)

MIF28A1		Day 4/Set 1 Total CK (CPK) Units
N	Value	Description
174		Missing
2707	.S	Legitimately skipped
0	1	1=ng/mL
40	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF28A2</i>		<i>Day 4/Set 1 Total CK (CPK) Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
171		Missing
2707	.S	Legitimately skipped
42	1	1=Range Set 1
1	2	2=Range Set 2

<i>MIF28A3</i>		<i>Day 4/Set 1 Total CK (CPK) Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
207		Missing
2707	.S	Legitimately skipped
3	1	1=absent/negative/normal
0	2	2=trace/weak positive
4	3	3=present/positive/abnormal

<i>MIF28B</i>		<i>Day 4/Set 1 CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
190		Missing
2707	.S	Legitimately skipped
24	Range	0.27 - 300 (median=2.65 mean=16.841 std=60.519)

<i>MIF28B1</i>		<i>Day 4/Set 1 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
193		Missing
2707	.S	Legitimately skipped
21	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF28B2</i>		<i>Day 4/Set 1 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
192		Missing
2707	.S	Legitimately skipped
21	1	1=Range Set 1
1	2	2=Range Set 2

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<i>MIF28B3</i>		<i>Day 4/Set 1 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
213		Missing
2707	.S	Legitimately skipped
1	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF28C</i>		<i>Day 4/Set 1 Total LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
211		Missing
2707	.S	Legitimately skipped
3	Range	278 - 897 (median=570 mean=581.7 std=309.7)

<i>MIF28C1</i>		<i>Day 4/Set 1 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
213		Missing
2707	.S	Legitimately skipped
0	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L

<i>MIF28C2</i>		<i>Day 4/Set 1 Total LDH Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
211		Missing
2707	.S	Legitimately skipped
3	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF28C3</i>		<i>Day 4/Set 1 Total LDH Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

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<i>MIF28D</i>		<i>Day 4/Set 1 LDH-1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
205		Missing
2707	.S	Legitimately skipped
9	Range	2 - 22 (median=13 mean=12.2 std=7.4)

<i>MIF28D1</i>		<i>Day 4/Set 1 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF28D2</i>		<i>Day 4/Set 1 LDH-1 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF28D3</i>		<i>Day 4/Set 1 LDH-1 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF28E</i>		<i>Day 4/Set 1 LDH-2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped

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<i>MIF28E1</i>		<i>Day 4/Set 1 LDH-2 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF28E2</i>		<i>Day 4/Set 1 LDH-2 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF28E3</i>		<i>Day 4/Set 1 LDH-2 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214		Missing
2707	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF28F</i>		<i>Day 4/Set 1 Troponin <</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
150		Missing
2707	.S	Legitimately skipped
51	0	0=n/a
13	1	1=<

<i>MIF28F0</i>		<i>Day 4/Set 1 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
147		Missing
2707	.S	Legitimately skipped
67	Range	0 - 5000 (median=0.112 mean=93.3868 std=621.6358)

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<i>MIF28F1</i>		<i>Day 4/Set 1 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
158		Missing
2707	.S	Legitimately skipped
56	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF28F2</i>		<i>Day 4/Set 1 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
151		Missing
2707	.S	Legitimately skipped
61	1	1=Range Set 1
2	2	2=Range Set 2

<i>MIF28F3</i>		<i>Day 4/Set 1 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
194		Missing
2707	.S	Legitimately skipped
9	1	1=absent/negative/normal
1	2	2=trace/weak positive
10	3	3=present/positive/abnormal

<i>MIF28F4</i>		<i>Day 4/Set 1 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
146		Missing
2707	.S	Legitimately skipped
6	1	1= Troponin, type not specified
33	2	2= Troponin I
25	3	3= Troponin T
4	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure

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MIF28G		second set of enzymes reported fourth day
N	Value	Description
1		Missing
2600	.S	Legitimately skipped
302	0	0=No/NR
18	1	1=Yes

MIF28Y		Year of MIF28
N	Value	Description
2833		Missing
88	Range	2008 - 2023 (median=2015 mean=2015.6 std=3.8)

MIF29_FUTIME		Follow-up time from baseline for MIF29 (Date of Day 4/Set 2)
N	Value	Description
2903		Missing
18	Range	777 - 4630 (median=2573.5 mean=2538.00 std=1297.02)

MIF29A		Day 4/Set 2 Total CK (CPK)
N	Value	Description
151		Missing
2768	.S	Legitimately skipped
2	Range	711 - 60661 (median=30686 mean=30686.0 std=42391.1)

MIF29A1		Day 4/Set 2 Total CK (CPK) Units
N	Value	Description
151		Missing
2768	.S	Legitimately skipped
0	1	1=ng/mL
2	2	2=Units/L
0	3	3=Åµg/L

MIF29A2		Day 4/Set 2 Total CK (CPK) Range Set
N	Value	Description
151		Missing
2768	.S	Legitimately skipped
2	1	1=Range Set 1
0	2	2=Range Set 2

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<i>MIF29A3</i>		<i>Day 4/Set 2 Total CK (CPK)Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2768	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
1	3	3=present/positive/abnormal

<i>MIF29B</i>		<i>Day 4/Set 2 CK-MB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2768	.S	Legitimately skipped
1	Range	3.19 - 3.19 (median=3.19 mean=3.190 std=.)

<i>MIF29B1</i>		<i>Day 4/Set 2 CK-MB Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2768	.S	Legitimately skipped
1	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF29B2</i>		<i>Day 4/Set 2 CK-MB Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2768	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF29B3</i>		<i>Day 4/Set 2 CK-MB Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

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<i>MIF29C</i>		<i>Day 4/Set 2 Total LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2768	.S	Legitimately skipped
1	Range	848 - 848 (median=848 mean=848.0 std=.)

<i>MIF29C1</i>		<i>Day 4/Set 2 Total LDH Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2768	.S	Legitimately skipped
0	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L

<i>MIF29C2</i>		<i>Day 4/Set 2 Total LDH Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
152		Missing
2768	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF29C3</i>		<i>Day 4/Set 2 Total LDH Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF29D</i>		<i>Day 4/Set 2 LDH-1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped

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<i>MIF29D1</i>		<i>Day 4/Set 2 LDH-1 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

<i>MIF29D2</i>		<i>Day 4/Set 2 LDH-1 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF29D3</i>		<i>Day 4/Set 2 LDH-1 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF29E</i>		<i>Day 4/Set 2 LDH-2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped

<i>MIF29E1</i>		<i>Day 4/Set 2 LDH-2 Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF29E2</i>		<i>Day 4/Set 2 LDH-2 Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2

<i>MIF29E3</i>		<i>Day 4/Set 2 LDH-2 Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2768	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal

<i>MIF29F</i>		<i>Day 4/Set 2 Troponin <</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
137		Missing
2768	.S	Legitimately skipped
12	0	0=n/a
4	1	1=<

<i>MIF29F0</i>		<i>Day 4/Set 2 Troponin Value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
137		Missing
2768	.S	Legitimately skipped
16	Range	0.01 - 597 (median=0.0685 mean=42.04044 std=149.07110)

<i>MIF29F1</i>		<i>Day 4/Set 2 Troponin Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
139		Missing
2768	.S	Legitimately skipped
14	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L

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<i>MIF29F2</i>		<i>Day 4/Set 2 Troponin Range Set</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
137		Missing
2768	.S	Legitimately skipped
15	1	1=Range Set 1
1	2	2=Range Set 2

<i>MIF29F3</i>		<i>Day 4/Set 2 Troponin Words Code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
146		Missing
2768	.S	Legitimately skipped
3	1	1=absent/negative/normal
1	2	2=trace/weak positive
3	3	3=present/positive/abnormal

<i>MIF29F4</i>		<i>Day 4/Set 2 Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
137		Missing
2768	.S	Legitimately skipped
0	1	1= Troponin, type not specified
7	2	2= Troponin I
7	3	3= Troponin T
2	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure

<i>MIF29Y</i>		<i>Year of MIF29</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2903		Missing
18	Range	2012 - 2023 (median=2016 mean=2016.3 std=3.5)

<i>MIF30</i>		<i>Mention of having trauma/surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1076	.S	Legitimately skipped
1746	0	0=No/NR
98	1	1=Yes

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MIF30A		Cardiac Procedure
N	Value	Description
1		Missing
2823	.S	Legitimately skipped
63	0	0=No
34	1	1=Yes

MIF30A1_FUTIME		Follow-up time from baseline for MIF30A1 (Date of Cardiac Procedure)
N	Value	Description
2888		Missing
33	Range	27 - 5064 (median=2057 mean=2157.8 std=1643.6)

MIF30A1Y		Year of MIF30A1
N	Value	Description
2888		Missing
33	Range	2010 - 2023 (median=2015 mean=2015.4 std=4.5)

MIF30B		CPR or cardioversion
N	Value	Description
3		Missing
2823	.S	Legitimately skipped
80	0	0=No
15	1	1=Yes

MIF30B1_FUTIME		Follow-up time from baseline for MIF30B1 (Date CPR or cardioversion)
N	Value	Description
2906		Missing
15	Range	193 - 4900 (median=2547 mean=2368.5 std=1671.7)

MIF30B1Y		Year of MIF30B1
N	Value	Description
2906		Missing
15	Range	2010 - 2023 (median=2016 mean=2016.3 std=4.5)

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<i>MIF30C</i>		<i>Other cardiac trauma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2823	.S	Legitimately skipped
93	0	0=No
3	1	1=Yes

<i>MIF30C1_FUTIME</i>		<i>Follow-up time from baseline for MIF30C1 (Date of Other cardiac trauma)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2918		Missing
3	Range	774 - 4900 (median=1065 mean=2246.3 std=2302.7)

<i>MIF30C1Y</i>		<i>Year of MIF30C1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2918		Missing
3	Range	2012 - 2023 (median=2012 mean=2015.7 std=6.4)

<i>MIF30C2</i>		<i>Specify type of cardiac trauma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2918	~	Legitimately skipped
2	Present	Text suppressed

<i>MIF30D</i>		<i>Rhabdomyolysis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2823	.S	Legitimately skipped
81	0	0=No
15	1	1=Yes

<i>MIF30D1_FUTIME</i>		<i>Follow-up time from baseline for MIF30D1 (Date of Rhabdomyolysis)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2916		Missing
5	Range	59 - 4715 (median=394 mean=1654.8 std=2032.4)

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<i>MIF30D1Y</i>		<i>Year of MIF30D1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2916		Missing
5	Range	2008 - 2022 (median=2012 mean=2013.8 std=6.0)

<i>MIF30E</i>		<i>Intramuscular Injection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2823	.S	Legitimately skipped
95	0	0=No
0	1	1=Yes

<i>MIF30E1_FUTIME</i>		<i>Follow-up time from baseline for MIF30E1 (Date of Intramuscular Injection)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2921		Missing

<i>MIF30E1Y</i>		<i>Year of MIF30E1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2921		Missing

<i>MIF30F</i>		<i>Non-cardiac procedure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2823	.S	Legitimately skipped
73	0	0=No
22	1	1=Yes

<i>MIF30F1_FUTIME</i>		<i>Follow-up time from baseline for MIF30F1 (Date of Non)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2902		Missing
19	Range	816 - 4787 (median=2850 mean=2904.4 std=1121.7)

<i>MIF30F1Y</i>		<i>Year of MIF30F1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2902		Missing
19	Range	2013 - 2022 (median=2018 mean=2017.8 std=2.8)

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<i>MIF30F2</i>		<i>Specify non-cardiac procedure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2899	~	Legitimately skipped
22	Present	Text suppressed

<i>MIF30G</i>		<i>Non-cardiac trauma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2823	.S	Legitimately skipped
83	0	0=No
12	1	1=Yes

<i>MIF30G1_FUTIME</i>		<i>Follow-up time from baseline for MIF30G1 (cardiac procedure)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2911		Missing
10	Range	262 - 4856 (median=1958 mean=2266.2 std=1576.6)

<i>MIF30G1Y</i>		<i>Year of MIF30G1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2911		Missing
10	Range	2010 - 2023 (median=2014.5 mean=2015.50 std=4.72)

<i>MIF31</i>		<i>Item number from biomarkers section</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
2823	.S	Legitimately skipped
89	Range	22 - 28 (median=22 mean=23.0 std=1.8)

<i>MIF32</i>		<i>Evidence of hemolytic disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2918	0	0=No/NR
2	1	1=Yes

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MIF33		Have active liver disease?
N	Value	Description
2801	0	0=No/NR
120	1	1=Yes

MIF34		Any 12 lead ECGs taken?
N	Value	Description
1		Missing
282	0	0=No
2151	1	1=Yes
487	9	9=NR

MIF34A_FUTIME		Follow-up time from baseline for MIF34A (Date of non)
N	Value	Description
835		Missing
2086	Range	5 - 5302 (median=2427 mean=2421.8 std=1380.1)

MIF34AY		Year of MIF34A
N	Value	Description
832		Missing
2089	Range	2008 - 2024 (median=2016 mean=2016.1 std=3.8)

MIF34B		Copy of first ECG enclosed?
N	Value	Description
10		Missing
824	.S	Legitimately skipped
1117	0	0=No
970	1	1=Yes

MIF34C_FUTIME		Follow-up time from baseline for MIF34C (cardiac trauma)
N	Value	Description
1933		Missing
988	Range	6 - 5202 (median=2101 mean=2235.6 std=1402.2)

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<i>MIF34CY</i>		<i>Year of MIF34C</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1932		Missing
989	Range	2008 - 2024 (median=2015 mean=2015.7 std=3.9)

<i>MIF34D</i>		<i>Copy of second ECG enclosed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72		Missing
1859	.S	Legitimately skipped
378	0	0=No
612	1	1=Yes

<i>MIF34E_FUTIME</i>		<i>Follow-up time from baseline for MIF34E (Date of first ECG)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2400		Missing
521	Range	6 - 5202 (median=2049 mean=2171.6 std=1408.7)

<i>MIF34EY</i>		<i>Year of MIF34E</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2400		Missing
521	Range	2008 - 2024 (median=2015 mean=2015.5 std=4.0)

<i>MIF34F</i>		<i>Copy of third ECG enclosed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
113		Missing
2285	.S	Legitimately skipped
176	0	0=No
347	1	1=Yes

<i>MIF34G_FUTIME</i>		<i>Follow-up time from baseline for MIF34G (Date of second ECG)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2581		Missing
340	Range	22 - 5203 (median=1673.5 mean=1949.40 std=1418.34)

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<i>MIF34GY</i>		<i>Year of MIF34G</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2581		Missing
340	Range	2008 - 2024 (median=2014 mean=2014.8 std=4.1)

<i>MIF34H</i>		<i>Copy of last ECG enclosed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117		Missing
2463	.S	Legitimately skipped
95	0	0=No
246	1	1=Yes

<i>MIF35</i>		<i>Transthoracic echocardiogram performed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1764	0	0=No/NR
1157	1	1=Yes

<i>MIF35A</i>		<i>LV ejection fraction TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
108		Missing
1764	.S	Legitimately skipped
1049	Range	5 - 85 (median=55 mean=51.7 std=15.3)

<i>MIF36</i>		<i>Nuclear Medicare Scan performed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2818	0	0=No/NR
103	1	1=Yes

<i>MIF36A</i>		<i>Ejection fraction of scan</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
2818	.S	Legitimately skipped
94	Range	10 - 85 (median=58 mean=56.1 std=17.0)

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<i>MIF36B</i>		<i>RV(%) of scan</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
103		Missing
2818	.S	Legitimately skipped

<i>MIF36C</i>		<i>Positive stress test for ischemia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2818	.S	Legitimately skipped
81	0	0=No/NR
22	1	1=Yes

<i>MIF37</i>		<i>Stress test performed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2740	0	0=No/NR
181	1	1=Yes

<i>MIF37A</i>		<i>Ejection fraction of Stress Test</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
89		Missing
2740	.S	Legitimately skipped
92	Range	17 - 96 (median=59.5 mean=56.75 std=17.67)

<i>MIF37B</i>		<i>Stress test positive for ischemia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2740	.S	Legitimately skipped
147	0	0=No/NR
31	1	1=Yes

<i>MIF37C</i>		<i>Greater than or equal to 1 mm ST?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2740	.S	Legitimately skipped
176	0	0=No/NR
2	1	1=Yes

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<i>MIF37D</i>		<i>Ischemic pain or equivalent occurred?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2740	.S	Legitimately skipped
175	0	0=No/NR
4	1	1=Yes

<i>MIF38</i>		<i>Was a coronary angiography performed?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2494	0	0=No/NR
427	1	1=Yes

<i>MIF38A_FUTIME</i>		<i>Follow-up time from baseline for MIF38A (Date of third ECG)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2509		Missing
412	Range	22 - 5213 (median=2062.5 mean=2172.05 std=1397.99)

<i>MIF38AY</i>		<i>Year of MIF38A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2509		Missing
412	Range	2008 - 2024 (median=2015 mean=2015.5 std=4.0)

<i>MIF38B</i>		<i>Ejection % of coronary angiography</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
258		Missing
2494	.S	Legitimately skipped
169	Range	10 - 70 (median=50 mean=48.0 std=14.4)

<i>MIF38C</i>		<i>70% or greater obstruction?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2494	.S	Legitimately skipped
123	0	0=No
278	1	1=Yes
23	9	9=NR

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<i>MIF38D</i>		<i>Coronary bypass present?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2494	.S	Legitimately skipped
290	0	0=No
40	1	1=Yes
94	9	9=NR

<i>MIF38D1</i>		<i>If yes, number of occluded grafts</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2881	.S	Legitimately skipped
38	Range	0 - 4 (median=0 mean=0.7 std=1.0)

<i>MIF39</i>		<i>Coronary reperfusion attempted?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2648	0	0=No/NR
273	1	1=Yes

<i>MIF39A</i>		<i>Approximate time onset to reperfusion</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2648	.S	Legitimately skipped
7	1	1= < 2 hours
9	2	2= 2 to 4 hours
4	3	3= 4 to 6 hours
4	4	4= 6 to 12 hours
8	5	5= 12 to 24 hours
130	6	6= > 24 hours
110	7	7= Not Sure

<i>MIF40A</i>		<i>Coronary artery bypass graft surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2844	0	0=No/NR
76	1	1=Yes

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<i>MIF40A1_FUTIME</i>		<i>Follow-up time from baseline for MIF40A1 (Date of last ECG)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2847		Missing
74	Range	154 - 5156 (median=1609 mean=2086.6 std=1433.8)

<i>MIF40A1Y</i>		<i>Year of MIF40A1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2847		Missing
74	Range	2009 - 2023 (median=2014.5 mean=2015.28 std=4.15)

<i>MIF40A2</i>		<i>Time of Coronary artery bypass</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
67		Missing
2844	.S	Legitimately skipped
10	Range	4:16:00 - 19:05:00

<i>MIF40A3</i>		<i>AM/PM of the coronary bypass</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7		Missing
2912	.S	Legitimately skipped
1	1	1=am
1	2	2=pm

<i>MIF40B</i>		<i>Coronary atherectomy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2918	0	0=No/NR
2	1	1=Yes

<i>MIF40B1_FUTIME</i>		<i>Follow-up time from baseline for MIF40B1 (Date of coronary angiography)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2919		Missing
2	Range	108 - 4752 (median=2430 mean=2430.0 std=3283.8)

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<i>MIF40B1Y</i>		<i>Year of MIF40B1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2919		Missing
2	Range	2009 - 2023 (median=2016 mean=2016.0 std=9.9)

<i>MIF40B2</i>		<i>Time of coronary atherectomy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2918	.S	Legitimately skipped
1	Range	9:30:00 - 9:30:00

<i>MIF40B3</i>		<i>AM/PM of coronary atherectomy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2920	.S	Legitimately skipped
0	1	1=am
0	2	2=pm

<i>MIF40C</i>		<i>Intra-arterial/intravenous thrombolytic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2915	0	0=No/NR
5	1	1=Yes

<i>MIF40C1_FUTIME</i>		<i>Follow-up time from baseline for MIF40C1 (Date of Coronary artery bypass)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2916		Missing
5	Range	134 - 1947 (median=235 mean=549.6 std=783.5)

<i>MIF40C1Y</i>		<i>Year of MIF40C1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2916		Missing
5	Range	2010 - 2015 (median=2011 mean=2011.6 std=1.9)

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<i>MIF40C2</i>		<i>Time of thrombolytic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
2915	.S	Legitimately skipped
2	Range	19:02:00 - 20:15:00

<i>MIF40C3</i>		<i>AM/PM of thrombolytic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2918	.S	Legitimately skipped
0	1	1=am
2	2	2=pm

<i>MIF40D</i>		<i>Coronary angioplasty w/o stent</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2867	0	0=No/NR
52	1	1=Yes

<i>MIF40D1_FUTIME</i>		<i>Follow-up time from baseline for MIF40D1 (Date of Coronary atherectomy)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2870		Missing
51	Range	141 - 5213 (median=3729 mean=3391.9 std=1299.6)

<i>MIF40D1Y</i>		<i>Year of MIF40D1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2870		Missing
51	Range	2010 - 2024 (median=2020 mean=2019.1 std=3.5)

<i>MIF40D2</i>		<i>Time of coronary angioplasty w/e stent</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
31		Missing
2868	.S	Legitimately skipped
22	Range	8:37:00 - 16:54:00

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<i>MIF40D3</i>		<i>AM/PM coronary angioplasty w/e stent</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2919	.S	Legitimately skipped
0	1	1=am
0	2	2=pm

<i>MIF40E</i>		<i>Coronary angioplasty with stent place</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2723	0	0=No/NR
197	1	1=Yes

<i>MIF40E1_FUTIME</i>		<i>Follow-up time from baseline for MIF40E1 (Date of thrombolytic)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2728		Missing
193	Range	22 - 4902 (median=2130 mean=2119.1 std=1365.2)

<i>MIF40E1Y</i>		<i>Year of MIF40E1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2728		Missing
193	Range	2009 - 2023 (median=2015 mean=2015.4 std=3.8)

<i>MIF40E2</i>		<i>Time of coronary angioplasty with stent</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
119		Missing
2723	.S	Legitimately skipped
79	Range	0:32:00 - 23:35:00

<i>MIF40E3</i>		<i>AM/PM of coronary angioplasty with stent</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
12		Missing
2901	.S	Legitimately skipped
4	1	1=am
4	2	2=pm

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<i>MIF40F</i>		<i>Valve surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2881	0	0=No/NR
40	1	1=Yes

<i>MIF40G</i>		<i>Non-cardiac surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2488	0	0=No/NR
432	1	1=Yes

<i>MIF40H</i>		<i>Aortic balloon pump</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2899	0	0=No/NR
22	1	1=Yes

<i>MIF40I</i>		<i>Pacemaker placement</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2868	0	0=No/NR
53	1	1=Yes

<i>MIF40J</i>		<i>Cardioversion or defibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2855	0	0=No/NR
66	1	1=Yes

<i>MIF40J1_FUTIME</i>		<i>Follow-up time from baseline for MIF40J1 (Date of coronary angioplasty w/o stent)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2871		Missing
50	Range	62 - 5036 (median=3091 mean=2759.9 std=1614.5)

<i>MIF40J1Y</i>		<i>Year of MIF40J1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2871		Missing
50	Range	2009 - 2024 (median=2018 mean=2017.1 std=4.5)

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<i>MIF40J2</i>		<i>Time of cardioversion or defibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
53		Missing
2855	.S	Legitimately skipped
13	Range	5:48:00 - 23:30:00

<i>MIF40J3</i>		<i>AM/PM of cardioversion or defibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6		Missing
2913	.S	Legitimately skipped
0	1	1=am
2	2	2=pm

<i>MIF40J4A</i>		<i>Ventricular Fibrillation/Flutter</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2855	.S	Legitimately skipped
55	0	0=No/NR
10	1	1=Yes

<i>MIF40J4B</i>		<i>Ventricular Tachycardia (VT)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2855	.S	Legitimately skipped
57	0	0=No/NR
8	1	1=Yes

<i>MIF40J4C</i>		<i>Asystole</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2855	.S	Legitimately skipped
57	0	0=No/NR
8	1	1=Yes

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<i>MIF40J4E</i>		<i>Atrial Fibrillation/Flutter</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2855	.S	Legitimately skipped
17	0	0=No/NR
48	1	1=Yes

<i>MIF40J4F</i>		<i>Pulseless Electrical Activity (PEA)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2855	.S	Legitimately skipped
57	0	0=No/NR
8	1	1=Yes

<i>MIF41A1</i>		<i>Admission Meds of Nitroglycerin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2211	0	0=No/NR
710	1	1=Yes

<i>MIF41A2</i>		<i>Discharge Meds of Nitroglycerin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2639	0	0=No/NR
281	1	1=Yes

<i>MIF41B1</i>		<i>Admission of Beta Blockers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1293	0	0=No/NR
1628	1	1=Yes

<i>MIF41B2</i>		<i>Discharge Meds of Beta Blockers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1702	0	0=No/NR
1218	1	1=Yes

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<i>MIF41C1</i>		<i>Admission Meds of Channel Blockers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2035	0	0=No/NR
885	1	1=Yes

<i>MIF41C2</i>		<i>Discharge Meds of Channel Blockers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2356	0	0=No/NR
563	1	1=Yes

<i>MIF41D1</i>		<i>Admission Meds of ACE or ARB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1444	0	0=No/NR
1476	1	1=Yes

<i>MIF41D2</i>		<i>Discharge Meds of ACE or ARB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
1923	0	0=No/NR
996	1	1=Yes

<i>MIF41E1</i>		<i>Admission Meds of scheduled aspirin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1356	0	0=No/NR
1564	1	1=Yes

<i>MIF41E2</i>		<i>Discharge Meds of scheduled aspirin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1761	0	0=No/NR
1159	1	1=Yes

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<i>MIF41F1</i>		<i>Admission Meds of Heparin/Enoxaparin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1789	0	0=No/NR
1132	1	1=Yes

<i>MIF41F2</i>		<i>Discharge Meds of of Heparin/Enoxaparin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2808	0	0=No/NR
111	1	1=Yes

<i>MIF41G1</i>		<i>Admission Meds of Coumadin,warfarin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2517	0	0=No/NR
404	1	1=Yes

<i>MIF41G2</i>		<i>Discharge Meds of Coumadin,warfarin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2596	0	0=No/NR
323	1	1=Yes

<i>MIF41H1</i>		<i>Admission Meds of Anti-platelet agents</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2239	0	0=No/NR
682	1	1=Yes

<i>MIF41H2</i>		<i>Discharge Meds of Anti-platelet agents</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2426	0	0=No/NR
493	1	1=Yes

<i>MIF41I1</i>		<i>Admission Meds of Statin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1269	0	0=No/NR
1652	1	1=Yes

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<i>MIF41I2</i>		<i>Discharge Meds of Statin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1596	0	0=No/NR
1324	1	1=Yes

<i>MIF42A</i>		<i>Treated with IV pressors</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2709	0	0=No/NR
212	1	1=Yes

<i>MIF42B</i>		<i>Treated with IV nitroglycerin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2820	0	0=No/NR
101	1	1=Yes

<i>MIF42C</i>		<i>Treated with inhibitors</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2893	0	0=No/NR
28	1	1=Yes

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

Myocardial Infarction Reviewer 1 Form: FORM CODE=MID1, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
2862	Present	Text suppressed

VISIT		Visit
N	Value	Description
187	1	Visit 1
218	2	Visit 2
258	3	Visit 3
254	4	Visit 4
254	5	Visit 5
201	6	Visit 6
211	7	Visit 7
264	8	Visit 8
217	9	Visit 9
191	10	Visit 10
251	11	Visit 11
149	12	Visit 12
142	13	Visit 13
60	14	Visit 14
5	15	Visit 15

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1934	1	Occurrence 1
521	2	Occurrence 2
192	3	Occurrence 3
84	4	Occurrence 4
50	5	Occurrence 5
29	6	Occurrence 6
14	7	Occurrence 7
9	8	Occurrence 8
8	9	Occurrence 9
9	10	Occurrence 10
4	11	Occurrence 11
4	12	Occurrence 12
1	13	Occurrence 13
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19

<i>MID10A_FUTIME</i>		<i>Follow-up time from baseline for MID10A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2859	Range	361 - 6141 (median=3633 mean=3775.4 std=918.0)

<i>MID10AY</i>		<i>Year of MID10A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2859	Range	2010 - 2025 (median=2019 mean=2019.8 std=2.5)

<i>MID10C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2862	Present	Text suppressed

<i>MID10D_FUTIME</i>		<i>Follow-up time from baseline for MID10D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2862	Range	-25 - 5416 (median=2345 mean=2376.3 std=1353.6)

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Section B: Myocardial Infarction Ascertainment

<i>MID10DY</i>		<i>Year of MID10D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2862	Range	2008 - 2024 (median=2016 mean=2016.0 std=3.7)

<i>MID11</i>		<i>Evidence of cardiac pain?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
1856	0	0=No
859	1	1=Yes
143	9	9=NR

<i>MID12</i>		<i>Level of cardiac biomarkers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
392	1	1= Abnormal (at least one value at 2 X the upper limit normal)
213	2	2= Equivocal (> normal, but < 2X the upper limit normal)
57	3	3= Incomplete
1073	4	4= Normal
1122	5	5= None recorded

<i>MID13</i>		<i>Interpretations of ECGs</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
5	1	1= Evolution of Major Q-Wave
56	2	2= Evolution of ST-T Elevation with or without Q-Wave
4	3	3= New LBBB
86	4	4= Evolution of ST-Depression/T wave inversion alone
1	5	5= Evolution of Minor Q-Wave alone
20	6	6= Single ECG with Major Q-wave
3	7	7= Single ECG with LBBB, described as new
2683	8	8= Absent, Uncodable or Other ECG

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Section B: Myocardial Infarction Ascertainment

<i>MID14</i>		<i>Myocardial infarction classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
157	1	1=Definite
193	2	2=Probable
2483	3	3=No MI
25	4	4=Unclassifiable

<i>MID14A</i>		<i>Type of MI?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
2509	.S	Legitimately skipped
69	0	0=Transmural
193	1	1=Subendocardial
86	9	9=Unsure/unknown

<i>MID14B</i>		<i>Location of MI?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
2509	.S	Legitimately skipped
35	1	1=Anterior
3	2	2=Posterior
35	3	3=Inferior
29	4	4=Lateral
2	5	5=Septal
26	6	6=More than one area
218	9	9=Unable to determine

<i>MID14C</i>		<i>Was the MI procedure-related?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
2509	.S	Legitimately skipped
5	1	1=Yes, cardiovascular procedure
5	2	2=Yes, non-cardiovascular procedure
338	3	3=No/Unsure

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<i>MID14D</i>		<i>Subclass of MI?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6		Missing
2509	.S	Legitimately skipped
176	1	1=Type 1
122	2	2=Type 2
0	3	3=Type 3
1	4	4=Type 4a
2	5	5=Type 4b
0	6	6=Type 5
46	7	7=Unsure

<i>MID15</i>		<i>aborted MI due to quick intervention</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1833		Missing
1029	Range	0 - 9 (median=0 mean=0.1 std=1.1)

<i>MID16</i>		<i>Comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2523		Missing
339	Present	Text suppressed

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Section B: Myocardial Infarction Ascertainment

Myocardial Infarction Reviewer 2 Form: FORM CODE=MID2, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
2862	Present	Text suppressed

VISIT		Visit
N	Value	Description
187	1	Visit 1
218	2	Visit 2
258	3	Visit 3
254	4	Visit 4
254	5	Visit 5
201	6	Visit 6
211	7	Visit 7
264	8	Visit 8
217	9	Visit 9
191	10	Visit 10
251	11	Visit 11
149	12	Visit 12
142	13	Visit 13
60	14	Visit 14
5	15	Visit 15

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1934	1	Occurrence 1
521	2	Occurrence 2
192	3	Occurrence 3
84	4	Occurrence 4
50	5	Occurrence 5
29	6	Occurrence 6
14	7	Occurrence 7
9	8	Occurrence 8
8	9	Occurrence 9
9	10	Occurrence 10
4	11	Occurrence 11
4	12	Occurrence 12
1	13	Occurrence 13
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19

<i>MID20A_FUTIME</i>		<i>Follow-up time from baseline for MID20A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2859	Range	1802 - 6133 (median=3652 mean=3811.3 std=939.0)

<i>MID20AY</i>		<i>Year of MID20A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2859	Range	2015 - 2025 (median=2019 mean=2019.9 std=2.6)

<i>MID20C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2862	Present	Text suppressed

<i>MID20D_FUTIME</i>		<i>Follow-up time from baseline for MID20D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6		Missing
2856	Range	-25 - 5416 (median=2350.5 mean=2380.21 std=1352.24)

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

<i>MID20DY</i>		<i>Year of MID20D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6		Missing
2856	Range	2008 - 2024 (median=2016 mean=2016.1 std=3.7)

<i>MID21</i>		<i>Evidence of cardiac pain?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
1808	0	0=No
869	1	1=Yes
181	9	9=NR

<i>MID22</i>		<i>Level of cardiac biomarkers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
377	1	1= Abnormal (at least one value at 2 X the upper limit normal)
219	2	2= Equivocal (> normal, but < 2X the upper limit normal)
78	3	3= Incomplete
1080	4	4= Normal
1104	5	5= None recorded

<i>MID23</i>		<i>Interpretations of ECGs</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
2	1	1= Evolution of Major Q-Wave
65	2	2= Evolution of ST-T Elevation with or without Q-Wave
2	3	3= New LBBB
92	4	4= Evolution of ST-Depression/T wave inversion alone
1	5	5= Evolution of Minor Q-Wave alone
21	6	6= Single ECG with Major Q-wave
3	7	7= Single ECG with LBBB, described as new
2672	8	8= Absent, Uncodable or Other ECG

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Section B: Myocardial Infarction Ascertainment

MID24		Myocardial infarction classification
N	Value	Description
4		Missing
151	1	1=Definite
222	2	2=Probable
2464	3	3=No MI
21	4	4=Unclassifiable

MID24A		Type of MI?
N	Value	Description
2		Missing
2487	.S	Legitimately skipped
70	0	0=Transmural
231	1	1=Subendocardial
72	9	9=Unsure/unknown

MID24B		Location of MI?
N	Value	Description
2		Missing
2487	.S	Legitimately skipped
33	1	1=Anterior
2	2	2=Posterior
39	3	3=Inferior
17	4	4=Lateral
4	5	5=Septal
34	6	6=More than one area
244	9	9=Unable to determine

MID24C		Was the MI procedure-related?
N	Value	Description
2		Missing
2487	.S	Legitimately skipped
4	1	1=Yes, cardiovascular procedure
6	2	2=Yes, non-cardiovascular procedure
363	3	3=No/Unsure

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Section B: Myocardial Infarction Ascertainment

<i>MID24D</i>		<i>Subclass of MI?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2487	.S	Legitimately skipped
162	1	1=Type 1
161	2	2=Type 2
2	3	3=Type 3
0	4	4=Type 4a
0	5	5=Type 4b
0	6	6=Type 5
47	7	7=Unsure

<i>MID25</i>		<i>aborted MI due to quick intervention</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1816		Missing
1046	Range	0 - 9 (median=0 mean=0.2 std=1.4)

<i>MID26</i>		<i>Comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2515		Missing
347	Present	Text suppressed

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

Myocardial Infarction Adjudication Reviewer Form: FORM CODE=MID3, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
696	Present	Text suppressed

VISIT		Visit
N	Value	Description
130	1	Visit 1
90	2	Visit 2
82	3	Visit 3
72	4	Visit 4
52	5	Visit 5
42	6	Visit 6
30	7	Visit 7
28	8	Visit 8
26	9	Visit 9
31	10	Visit 10
42	11	Visit 11
31	12	Visit 12
30	13	Visit 13
10	14	Visit 14

OCCURRENCE		Occurrence
N	Value	Description
473	1	Occurrence 1
127	2	Occurrence 2
40	3	Occurrence 3
17	4	Occurrence 4
12	5	Occurrence 5
10	6	Occurrence 6
5	7	Occurrence 7
4	8	Occurrence 8
3	9	Occurrence 9
3	10	Occurrence 10
1	13	Occurrence 13
1	19	Occurrence 19

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

<i>MID30A_FUTIME</i>		<i>Follow-up time from baseline for MID30A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
246		Missing
450	Range	2383 - 6169 (median=3795.5 mean=4094.61 std=983.99)

<i>MID30AY</i>		<i>Year of MID30A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
246		Missing
450	Range	2015 - 2025 (median=2020 mean=2020.7 std=2.7)

<i>MID30C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
696	Present	Text suppressed

<i>MID30D_FUTIME</i>		<i>Follow-up time from baseline for MID30D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
80		Missing
616	Range	5 - 5207 (median=1613.5 mean=1984.28 std=1422.12)

<i>MID30DY</i>		<i>Year of MID30D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
80		Missing
616	Range	2008 - 2024 (median=2014 mean=2014.9 std=4.0)

<i>MID31</i>		<i>Evidence of cardiac pain?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
246		Missing
174	0	0=No
245	1	1=Yes
31	9	9=NR

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

MID32		Level of cardiac biomarkers
N	Value	Description
246		Missing
197	1	1= Abnormal (at least one value at 2 X the upper limit normal)
111	2	2= Equivocal (> normal, but < 2X the upper limit normal)
23	3	3= Incomplete
55	4	4= Normal
64	5	5= None recorded

MID33		Interpretations of ECGs
N	Value	Description
246		Missing
0	1	1= Evolution of Major Q-Wave
31	2	2= Evolution of ST-T Elevation with or without Q-Wave
0	3	3= New LBBB
30	4	4= Evolution of ST-Depression/T wave inversion alone
1	5	5= Evolution of Minor Q-Wave alone
8	6	6= Single ECG with Major Q-wave
0	7	7= Single ECG with LBBB, described as new
380	8	8= Absent, Uncodable or Other ECG

MID34		Myocardial infarction classification
N	Value	Description
246		Missing
64	1	1=Definite
147	2	2=Probable
217	3	3=No MI
22	4	4=Unclassifiable

MID34A		Type of MI?
N	Value	Description
239		Missing
246	.S	Legitimately skipped
41	0	0=Transmural
132	1	1=Subendocardial
38	9	9=Unsure/unknown

Chapter 1: Endpoint Ascertainment

Section B: Myocardial Infarction Ascertainment

MID34B		Location of MI?
N	Value	Description
239		Missing
246	.S	Legitimately skipped
14	1	1=Anterior
1	2	2=Posterior
19	3	3=Inferior
12	4	4=Lateral
2	5	5=Septal
18	6	6=More than one area
145	9	9=Unable to determine

MID34C		Was the MI procedure-related?
N	Value	Description
239		Missing
246	.S	Legitimately skipped
2	1	1=Yes, cardiovascular procedure
8	2	2=Yes, non-cardiovascular procedure
201	3	3=No/Unsure

MID34D		Subclass of MI?
N	Value	Description
239		Missing
246	.S	Legitimately skipped
94	1	1=Type 1
82	2	2=Type 2
0	3	3=Type 3
1	4	4=Type 4a
0	5	5=Type 4b
0	6	6=Type 5
34	7	7=Unsure

MID35		aborted MI due to quick intervention
N	Value	Description
623		Missing
73	Range	0 - 9 (median=0 mean=1.6 std=3.5)

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<i>MID36</i>		<i>Comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
597		Missing
99	Present	Text suppressed

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Section C: Heart Failure Ascertainment

Heart Failure Abstraction Form: FORM CODE=HTF, VERSION=A

ID		HCHS Participant ID
N	Value	Description
1552	Present	Text suppressed

VISIT		Visit
N	Value	Description
65	1	Visit 1
98	2	Visit 2
114	3	Visit 3
136	4	Visit 4
141	5	Visit 5
105	6	Visit 6
104	7	Visit 7
135	8	Visit 8
116	9	Visit 9
128	10	Visit 10
189	11	Visit 11
95	12	Visit 12
88	13	Visit 13
35	14	Visit 14
3	15	Visit 15

Chapter 1: Endpoint Ascertainment

Section C: Heart Failure Ascertainment

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1009	1	Occurrence 1
273	2	Occurrence 2
123	3	Occurrence 3
54	4	Occurrence 4
30	5	Occurrence 5
17	6	Occurrence 6
11	7	Occurrence 7
8	8	Occurrence 8
5	9	Occurrence 9
8	10	Occurrence 10
3	11	Occurrence 11
4	12	Occurrence 12
2	13	Occurrence 13
1	14	Occurrence 14
1	15	Occurrence 15
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19

<i>HTF0A_FUTIME</i>		<i>Follow-up time from baseline for HTF0A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1552	Range	17 - 5962 (median=3518 mean=3568.0 std=1092.0)

<i>HTF0AY</i>		<i>Year of HTF0A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1552	Range	2010 - 2025 (median=2019 mean=2019.3 std=3.0)

<i>HTF0C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1552	Present	Text suppressed

<i>HTF0D_FUTIME</i>		<i>Follow-up time from baseline for HTF0D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1552	Range	10 - 5302 (median=2657 mean=2577.7 std=1330.4)

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<i>HTF0DY</i>		<i>Year of HTF0D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1552	Range	2008 - 2024 (median=2017 mean=2016.6 std=3.6)

<i>HTF1</i>		<i>Was the event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
90	1	1= In hospital only
3	2	2= Emergency Dept. visit only (ED)
1456	3	3= Both ED and in hospital
0	4	4= Observation care only
2	5	5= Both ED and observation care
1	6	6= Unsure/Unknown

<i>HTF2_FUTIME</i>		<i>Follow-up time from baseline for HTF2 (Date of arrival)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10		Missing
1542	Range	10 - 5302 (median=2669 mean=2584.3 std=1329.5)

<i>HTF2A</i>		<i>Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72		Missing
1480	Range	0:00:00 - 23:59:00

<i>HTF2A1</i>		<i>AM/PM of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
1483	.S	Legitimately skipped
25	1	1=AM
29	2	2=PM

<i>HTF2B_FUTIME</i>		<i>Follow-up time from baseline for HTF2B (Date of admission)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1551	Range	10 - 5302 (median=2652 mean=2576.3 std=1329.0)

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Section C: Heart Failure Ascertainment

<i>HTF2BY</i>		<i>Year of HTF2B</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1551	Range	2008 - 2024 (median=2017 mean=2016.6 std=3.6)

<i>HTF2Y</i>		<i>Year of HTF2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10		Missing
1542	Range	2008 - 2024 (median=2017 mean=2016.6 std=3.6)

<i>HTF3_FUTIME</i>		<i>Follow-up time from baseline for HTF3 (Date of discharge)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13		Missing
1539	Range	17 - 5306 (median=2653 mean=2582.6 std=1329.4)

<i>HTF3A</i>		<i>Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
662		Missing
890	Range	0:03:00 - 23:59:00

<i>HTF3A1</i>		<i>AM/PM of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
37		Missing
1483	.S	Legitimately skipped
8	1	1=AM
24	2	2=PM

<i>HTF3Y</i>		<i>Year of HTF3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13		Missing
1539	Range	2008 - 2024 (median=2017 mean=2016.6 std=3.6)

<i>HTF4</i>		<i>Admitting diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
228		Missing
1324	Present	Text suppressed

Chapter 1: Endpoint Ascertainment

Section C: Heart Failure Ascertainment

<i>HTF5</i>		<i>Discharge diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
201		Missing
1351	Present	Text suppressed

<i>HTF6</i>		<i>EMS transport</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1057	0	0=No/NR
495	1	1=Yes

<i>HTF7</i>		<i>Transferred from hosp</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1498	0	0=No/NR
54	1	1=Yes

<i>HTF8</i>		<i>Code status DNR</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1469	0	0=No/NR
83	1	1=Yes

<i>HTF9</i>		<i>Alive at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
96	0	0=No
1456	1	1=Yes

<i>HTF10A</i>		<i>PND at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
242	0	0=No
111	1	1=Yes
1199	9	9=NR

<i>HTF10B</i>		<i>Orthopnea at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
271	0	0=No
285	1	1=Yes
996	9	9=NR

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<i>HTF10C</i>		<i>Shortness of breath at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
323	0	0=No
1109	1	1=Yes
120	9	9=NR

<i>HTF10D</i>		<i>Edema at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
764	0	0=No
430	1	1=Yes
358	9	9=NR

<i>HTF10E</i>		<i>Hypoxia at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
282	0	0=No
449	1	1=Yes
821	9	9=NR

<i>HTF10F</i>		<i>Dyspnea (at rest) at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
472	0	0=No
531	1	1=Yes
549	9	9=NR

<i>HTF10G</i>		<i>Dyspnea (on exertion) at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
263	0	0=No
1019	1	1=Yes
270	9	9=NR

<i>HTF11</i>		<i>Exacerbation of HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1009	0	0=No
487	1	1=Yes
56	9	9=NR

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<i>HTF12</i>		<i>Onset/progression of HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
753	0	0=No
555	1	1=Yes
244	9	9=NR

<i>HTF13A</i>		<i>Diastolic heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1192	0	0=No/NR
360	1	1=Yes

<i>HTF13B</i>		<i>Systolic heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1194	0	0=No/NR
358	1	1=Yes

<i>HTF13C</i>		<i>Right-sided heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1523	0	0=No/NR
29	1	1=Yes

<i>HTF13D</i>		<i>Ischemic cardiomyopathy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1396	0	0=No/NR
156	1	1=Yes

<i>HTF13E</i>		<i>Idiopathic/dilated cardiomyopathy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1413	0	0=No/NR
139	1	1=Yes

<i>HTF13F</i>		<i>Myocarditis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1548	0	0=No/NR
4	1	1=Yes

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<i>HTF13G</i>		<i>Peripartum cardiomyopathy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1552	0	0=No/NR
0	1	1=Yes

<i>HTF13H</i>		<i>Other cardiomyopathy/HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1385	0	0=No/NR
167	1	1=Yes

<i>HTF13H1</i>		<i>Specify type cardiomyopathy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1385	~	Legitimately skipped
167	Present	Text suppressed

<i>HTF14</i>		<i>Cardiac imaging performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
993	0	0=No/NR
559	1	1=Yes

<i>HTF14A</i>		<i>Quantitative EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
993	.S	Legitimately skipped
58	0	0=No
501	1	1=Yes

<i>HTF14B</i>		<i>Lowest LV EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
1051	.S	Legitimately skipped
499	Range	10 - 86 (median=45 mean=44.0 std=17.9)

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<i>HTF14C</i>		<i>Qualitative EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1494	~	Legitimately skipped
1	D	D= Decreased moderately
0	M	M= Mildly reduced
28	N	N= Normal
2	O	O= None of the above
0	S	S= Severely reduced
26	U	U= Unsure-Not available

<i>HTF14D</i>		<i>Time since lowest EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
121		Missing
993	.S	Legitimately skipped
438	Range	0.1 - 99 (median=4.4 mean=10.46 std=15.26)

<i>HTF14E</i>		<i>Type of Imaging from EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
993	.S	Legitimately skipped
407	1	1= ECHO
0	2	2= MUGA
28	3	3= Catheterization with ventriculography
0	4	4= CT
0	5	5= MRI
22	6	6= Other
102	7	7= Unknown

<i>HTF15A</i>		<i>Diagnosis of HF prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
782	0	0=No/NR
770	1	1=Yes

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<i>HTF15B</i>		<i>Hospitalization for HF prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
782	.S	Legitimately skipped
566	0	0=No/NR
204	1	1=Yes

<i>HTF15C</i>		<i>Treatment for HF prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
782	.S	Legitimately skipped
406	0	0=No/NR
364	1	1=Yes

<i>HTF15D</i>		<i>Valvular HD prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1386	0	0=No/NR
166	1	1=Yes

<i>HTF15E</i>		<i>RHD prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1530	0	0=No/NR
22	1	1=Yes

<i>HTF15F</i>		<i>Congenital HD prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1544	0	0=No/NR
8	1	1=Yes

<i>HTF15G</i>		<i>Coronary HD (ever) prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
958	0	0=No/NR
594	1	1=Yes

<i>HTF15H</i>		<i>Coronary HD (within year) prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1446	0	0=No/NR
106	1	1=Yes

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<i>HTF15I</i>		<i>Angina prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1257	0	0=No/NR
295	1	1=Yes

<i>HTF15J</i>		<i>Myocardial infarction prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1288	0	0=No/NR
264	1	1=Yes

<i>HTF15K</i>		<i>Atrial fib/atrial flutter prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1290	0	0=No/NR
262	1	1=Yes

<i>HTF15L</i>		<i>Heart block/bradycardia prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1506	0	0=No/NR
46	1	1=Yes

<i>HTF15M</i>		<i>Ventricular fib or tachycard prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1514	0	0=No/NR
38	1	1=Yes

<i>HTF15N</i>		<i>Hypertension prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
275	0	0=No/NR
1277	1	1=Yes

<i>HTF15O</i>		<i>Diabetes prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
595	0	0=No/NR
957	1	1=Yes

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<i>HTF15P</i>		<i>COPD prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1211	0	0=No/NR
341	1	1=Yes

<i>HTF15Q</i>		<i>Cor pulmonale prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1486	0	0=No/NR
66	1	1=Yes

<i>HTF15R</i>		<i>Pulmonary hypertension prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1465	0	0=No/NR
87	1	1=Yes

<i>HTF15S</i>		<i>ESRD prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1366	0	0=No/NR
186	1	1=Yes

<i>HTF16A</i>		<i>CABG past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1346	0	0=No/NR
206	1	1=Yes

<i>HTF16B</i>		<i>PCI past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1320	0	0=No/NR
232	1	1=Yes

<i>HTF16C</i>		<i>Valve surgery past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1468	0	0=No/NR
84	1	1=Yes

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<i>HTF16D</i>		<i>Pacemaker past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1440	0	0=No/NR
112	1	1=Yes

<i>HTF16E</i>		<i>AICD past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1404	0	0=No/NR
148	1	1=Yes

<i>HTF16F</i>		<i>Ablation for arrhythmia past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1526	0	0=No/NR
26	1	1=Yes

<i>HTF16G</i>		<i>Cardiac transplant past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1549	0	0=No/NR
3	1	1=Yes

<i>HTF16H</i>		<i>VAD past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1550	0	0=No/NR
2	1	1=Yes

<i>HTF17A</i>		<i>Myocardial Infarction active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1403	0	0=No/NR
149	1	1=Yes

<i>HTF17B</i>		<i>Shock/Cardiogenic Shock active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1507	0	0=No/NR
45	1	1=Yes

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<i>HTF17C</i>		<i>Ventricular Fib active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1473	0	0=No/NR
79	1	1=Yes

<i>HTF17D</i>		<i>Ventricular Tachycardia active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1521	0	0=No/NR
31	1	1=Yes

<i>HTF17E</i>		<i>Atrial Fib/Flutter active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1390	0	0=No/NR
162	1	1=Yes

<i>HTF17F</i>		<i>COPD exacerbation active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1377	0	0=No/NR
175	1	1=Yes

<i>HTF17G</i>		<i>Cardiac Surgery active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1500	0	0=No/NR
52	1	1=Yes

<i>HTF17H</i>		<i>Non-cardiac surgery active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1429	0	0=No/NR
123	1	1=Yes

<i>HTF17I</i>		<i>Pulmonary Embolus active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1529	0	0=No/NR
23	1	1=Yes

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<i>HTF17J</i>		<i>Pneumonia active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1254	0	0=No/NR
298	1	1=Yes

<i>HTF17K</i>		<i>Renal failure/insufficiency active</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1400	0	0=No/NR
152	1	1=Yes

<i>HTF18</i>		<i>Weight or BMI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
461		Missing
1091	Range	17 - 575 (median=86 mean=103.5 std=55.8)

<i>HTF18A</i>		<i>Units Weight/BMI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
464		Missing
200	1	1= Weight in lbs
853	2	2= Weight in Kg
35	3	3= BMI

<i>HTF19A</i>		<i>JVD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
895	0	0=No
141	1	1=Yes
516	9	9=NR

<i>HTF19B</i>		<i>Crackles or rales</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
856	0	0=No
489	1	1=Yes
207	9	9=NR

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<i>HTF19C</i>		<i>Wheezing</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
989	0	0=No
327	1	1=Yes
236	9	9=NR

<i>HTF19D</i>		<i>Rhonchi</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
871	0	0=No
254	1	1=Yes
427	9	9=NR

<i>HTF19E</i>		<i>S3 gallop</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
823	0	0=No
33	1	1=Yes
696	9	9=NR

<i>HTF19F</i>		<i>Lower extremity edema-unilateral</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1368	0	0=No
18	1	1=Yes
166	9	9=NR

<i>HTF19G</i>		<i>Lower extremity edema-bilateral</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
962	0	0=No
432	1	1=Yes
158	9	9=NR

<i>HTF20</i>		<i>Chest X-ray performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
163	0	0=No/NR
1389	1	1=Yes

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<i>HTF20A</i>		<i>Pulmonary edema/CHF x-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
163	.S	Legitimately skipped
1060	0	0=No/NR
329	1	1=Yes

<i>HTF20B</i>		<i>Cardiomegaly/Cardiothoracic ratio x-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
163	.S	Legitimately skipped
794	0	0=No/NR
595	1	1=Yes

<i>HTF20C</i>		<i>Pulmonary vascular congestion x-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
163	.S	Legitimately skipped
1011	0	0=No/NR
378	1	1=Yes

<i>HTF20D</i>		<i>Pleural effusion x-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
163	.S	Legitimately skipped
1028	0	0=No/NR
361	1	1=Yes

<i>HTF21</i>		<i>CT scan or CT angiogram</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1126	0	0=No/NR
426	1	1=Yes

<i>HTF21A</i>		<i>Pulmonary edema CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1126	.S	Legitimately skipped
362	0	0=No/NR
64	1	1=Yes

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<i>HTF21B</i>		<i>Cardiomegaly CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1126	.S	Legitimately skipped
311	0	0=No/NR
115	1	1=Yes

<i>HTF21C</i>		<i>Pleural effusion CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1126	.S	Legitimately skipped
275	0	0=No/NR
151	1	1=Yes

<i>HTF21D</i>		<i>Enlarged vena cava CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1126	.S	Legitimately skipped
425	0	0=No/NR
1	1	1=Yes

<i>HTF21E</i>		<i>Enlarged Pulmonary arteries CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1126	.S	Legitimately skipped
402	0	0=No/NR
24	1	1=Yes

<i>HTF22</i>		<i>TTE performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
818	0	0=No/NR
734	1	1=Yes

<i>HTF22A_FUTIME</i>		<i>Follow-up time from baseline for HTF22A (Date of TTE)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
907		Missing
645	Range	11 - 5398 (median=2761 mean=2609.9 std=1338.9)

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<i>HTF22AY</i>		<i>Year of HTF22A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
903		Missing
649	Range	2008 - 2024 (median=2017 mean=2016.7 std=3.7)

<i>HTF22B</i>		<i>Left Ventricular Ejection fraction TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
48		Missing
818	.S	Legitimately skipped
686	Range	5 - 84 (median=55 mean=48.0 std=16.5)

<i>HTF22C</i>		<i>LVH TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10		Missing
818	.S	Legitimately skipped
172	0	0=None
51	1	1=Present
161	2	2=Mild
56	3	3=Mod
11	4	4=Severe
273	9	9=NR

<i>HTF22D</i>		<i>Impaired LV systolic function TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
818	.S	Legitimately skipped
291	0	0=None
39	1	1=Present
43	2	2=Mild
50	3	3=Mod
100	4	4=Severe
202	9	9=NR

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<i>HTF22E</i>		<i>Impaired RV systolic function TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10		Missing
818	.S	Legitimately skipped
380	0	0=None
20	1	1=Present
21	2	2=Mild
23	3	3=Mod
11	4	4=Severe
269	9	9=NR

<i>HTF22F</i>		<i>Pulmonary hypertension TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
818	.S	Legitimately skipped
82	0	0=None
19	1	1=Present
44	2	2=Mild
47	3	3=Mod
27	4	4=Severe
506	9	9=NR

<i>HTF22G</i>		<i>Diastolic dysfunction TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
818	.S	Legitimately skipped
60	0	0=None
195	1	1=Present
86	2	2=Mild
41	3	3=Mod
14	4	4=Severe
330	9	9=NR

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<i>HTF23</i>		<i>TEE performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1498	0	0=No/NR
53	1	1=Yes

<i>HTF23A_FUTIME</i>		<i>Follow-up time from baseline for HTF23A (Date of TEE)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1502		Missing
50	Range	407 - 5401 (median=2846 mean=2816.8 std=1309.0)

<i>HTF23AY</i>		<i>Year of HTF23A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1502		Missing
50	Range	2010 - 2023 (median=2017.5 mean=2017.26 std=3.39)

<i>HTF23B1</i>		<i>LV ejection fraction TEE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
28		Missing
1499	.S	Legitimately skipped
25	Range	20 - 75 (median=50 mean=47.9 std=13.4)

<i>HTF23B2</i>		<i>RV ejection fraction TEE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
53		Missing
1499	.S	Legitimately skipped

<i>HTF24</i>		<i>Coronary angiography performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1372	0	0=No/NR
180	1	1=Yes

<i>HTF24A_FUTIME</i>		<i>Follow-up time from baseline for HTF24A (Date of angiography)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1377		Missing
175	Range	22 - 5161 (median=2439 mean=2371.3 std=1377.1)

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<i>HTF24AY</i>		<i>Year of HTF24A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1376		Missing
176	Range	2008 - 2024 (median=2016 mean=2016.1 std=3.9)

<i>HTF24B</i>		<i>LV Ejection fraction angio</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
103		Missing
1372	.S	Legitimately skipped
77	Range	10 - 67 (median=42 mean=41.4 std=15.3)

<i>HTF24C</i>		<i>70% obstruction artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1372	.S	Legitimately skipped
71	0	0=No/NR
109	1	1=Yes

<i>HTF25</i>		<i>MUGA or RVG performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1548	0	0=No/NR
4	1	1=Yes

<i>HTF25A</i>		<i>LV ejection fraction MUGA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1548	.S	Legitimately skipped
4	Range	10 - 68 (median=62 mean=50.5 std=27.3)

<i>HTF25B</i>		<i>RV ejection fraction MUGA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
1548	.S	Legitimately skipped

<i>HTF26</i>		<i>MRI performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1550	0	0=No/NR
2	1	1=Yes

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<i>HTF26A</i>		<i>LV ejection fraction MRI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1550	.S	Legitimately skipped
1	Range	30 - 30 (median=30 mean=30.0 std=.)

<i>HTF26B</i>		<i>RV ejection fraction MRI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
1550	.S	Legitimately skipped

<i>HTF27A</i>		<i>LV ejection fraction during visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
818		Missing
734	Range	0 - 87 (median=51 mean=47.5 std=17.0)

<i>HTF27B</i>		<i>Stress test positive for ischemia visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1525	0	0=No/NR
27	1	1=Yes

<i>HTF27C</i>		<i>Regional wall abnormalities visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1306	0	0=No/NR
246	1	1=Yes

<i>HTF27D</i>		<i>Dilated left ventricle visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1392	0	0=No/NR
160	1	1=Yes

<i>HTF27E</i>		<i>Dilated right ventricle visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1452	0	0=No/NR
100	1	1=Yes

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<i>HTF27F</i>		<i>Impaired left systolic visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1308	0	0=No/NR
244	1	1=Yes

<i>HTF27G</i>		<i>Left diastolic dysfunction visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1231	0	0=No/NR
321	1	1=Yes

<i>HTF27H</i>		<i>VSD visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1548	0	0=No/NR
4	1	1=Yes

<i>HTF27I</i>		<i>ASD visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1551	0	0=No/NR
1	1	1=Yes

<i>HTF27J</i>		<i>PDA visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1552	0	0=No/NR
0	1	1=Yes

<i>HTF27K</i>		<i>Artificial heart valve visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1534	0	0=No/NR
18	1	1=Yes

<i>HTF27L</i>		<i>HOCM visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1549	0	0=No/NR
3	1	1=Yes

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<i>HTF27M</i>		<i>Valvular HD visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1503	0	0=No/NR
49	1	1=Yes

<i>HTF28A</i>		<i>Worst BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1218		Missing
334	Range	5 - 116000 (median=307.55 mean=1613.843 std=7086.795)

<i>HTF28B</i>		<i>Last BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1328		Missing
224	Range	5 - 116000 (median=318.05 mean=1809.852 std=8432.691)

<i>HTF28C</i>		<i>Upper Limit BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1336		Missing
216	Range	23 - 250 (median=100 mean=104.1 std=24.8)

<i>HTF29A</i>		<i>Worst ProBNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1119		Missing
433	Range	7 - 175001 (median=1561 mean=8793.4 std=20134.2)

<i>HTF29B</i>		<i>Last BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1269		Missing
283	Range	7 - 175001 (median=1576 mean=9403.6 std=22254.1)

<i>HTF29C</i>		<i>Upper Limit ProBNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1255		Missing
297	Range	99 - 1800 (median=177 mean=378.8 std=398.4)

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<i>HTF30</i>		<i>Troponin value available</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10		Missing
417	0	0=No
1125	1	1=Yes

<i>HTF30A1</i>		<i>Less than worst Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
719		Missing
408	<	
3	>	
422	~	Legitimately skipped

<i>HTF30A2</i>		<i>Worst Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
30		Missing
422	.S	Legitimately skipped
1100	Range	0 - 12000 (median=0.04 mean=70.549 std=719.409)

<i>HTF30B</i>		<i>Upper Limit Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
379		Missing
422	.S	Legitimately skipped
751	Range	0.01 - 100 (median=0.05 mean=2.771 std=8.934)

<i>HTF30C</i>		<i>Type of Troponin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7		Missing
422	.S	Legitimately skipped
116	1	1= Troponin, type not specified
579	2	2= Troponin I
327	3	3= Troponin T
96	4	4= High Sensitivity Troponin (HS)
5	5	5= Unsure

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<i>HTF31</i>		<i>Worst sodium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
97		Missing
1455	Range	107 - 161 (median=137 mean=136.1 std=4.5)

<i>HTF32</i>		<i>Worst serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
91		Missing
1461	Range	0.2 - 15.1 (median=1.1 mean=2.04 std=2.26)

<i>HTF33</i>		<i>Worst BUN</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
93		Missing
1459	Range	4 - 210 (median=24 mean=33.0 std=26.1)

<i>HTF34</i>		<i>Worst hemoglobin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
80		Missing
1472	Range	4.3 - 18.1 (median=11.5 mean=11.30 std=2.39)

<i>HTF35</i>		<i>Worst hematocrit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
107		Missing
1445	Range	12.2 - 54.9 (median=35.2 mean=34.49 std=6.89)

<i>HTF36A</i>		<i>Cardioversion or Defibrillation visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1515	0	0=No/NR
37	1	1=Yes

<i>HTF36B</i>		<i>Aortic balloon pump visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1533	0	0=No/NR
19	1	1=Yes

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<i>HTF36C</i>		<i>PCI visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1483	0	0=No/NR
69	1	1=Yes

<i>HTF36D</i>		<i>CPAP or BIPAP visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1359	0	0=No/NR
193	1	1=Yes

<i>HTF36E</i>		<i>Mechanical Ventilation visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1402	0	0=No/NR
150	1	1=Yes

<i>HTF36F</i>		<i>Thoracentesis visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1518	0	0=No/NR
34	1	1=Yes

<i>HTF36G</i>		<i>VAD visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1548	0	0=No/NR
4	1	1=Yes

<i>HTF36H</i>		<i>Heart transplant visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1551	0	0=No/NR
1	1	1=Yes

<i>HTF36I</i>		<i>Hemodialysis or hemofiltration visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1382	0	0=No/NR
170	1	1=Yes

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<i>HTF37A</i>		<i>Med list admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	0	0=No/NR
1390	1	1=Yes

<i>HTF37B</i>		<i>Med list discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	0	0=No/NR
1150	1	1=Yes

<i>HTF38A</i>		<i>ACE inhibitors admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
885	0	0=No/NR
505	1	1=Yes

<i>HTF38B</i>		<i>ACE inhibitors discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
772	0	0=No/NR
378	1	1=Yes

<i>HTF39A</i>		<i>Angiotensin admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
162	.S	Legitimately skipped
1072	0	0=No/NR
317	1	1=Yes

<i>HTF39B</i>		<i>Angiotensin discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
402	.S	Legitimately skipped
917	0	0=No/NR
232	1	1=Yes

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<i>HTF40A</i>		<i>Beta blockers admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
574	0	0=No/NR
816	1	1=Yes

<i>HTF40B</i>		<i>Beta blockers discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
474	0	0=No/NR
676	1	1=Yes

<i>HTF41A</i>		<i>Digitalis admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
1330	0	0=No/NR
60	1	1=Yes

<i>HTF41B</i>		<i>Digitalis discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
1106	0	0=No/NR
44	1	1=Yes

<i>HTF42A</i>		<i>Diuretics admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
717	0	0=No/NR
673	1	1=Yes

<i>HTF42B</i>		<i>Diuretics discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
613	0	0=No/NR
537	1	1=Yes

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<i>HTF43A</i>		<i>Aldosterone blocker admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
1279	0	0=No/NR
111	1	1=Yes

<i>HTF43B</i>		<i>Aldosterone blocker discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
402	.S	Legitimately skipped
1055	0	0=No/NR
94	1	1=Yes

<i>HTF44A</i>		<i>Lipid agents admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
520	0	0=No/NR
870	1	1=Yes

<i>HTF44B</i>		<i>Lipid agents discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
393	0	0=No/NR
757	1	1=Yes

<i>HTF45A</i>		<i>Nitrates admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
1120	0	0=No/NR
270	1	1=Yes

<i>HTF45B</i>		<i>Nitrates discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
1003	0	0=No/NR
147	1	1=Yes

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<i>HTF46A</i>		<i>Hydralazine admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
162	.S	Legitimately skipped
1221	0	0=No/NR
169	1	1=Yes

<i>HTF46B</i>		<i>Hydralazine discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
1040	0	0=No/NR
110	1	1=Yes

<i>HTF47A</i>		<i>IV inotropes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1513	0	0=No/NR
38	1	1=Yes

<i>HTF47B</i>		<i>IV diuretics</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1065	0	0=No/NR
486	1	1=Yes

<i>HTF47C</i>		<i>Oral diuretics</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1350	0	0=No/NR
201	1	1=Yes

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Heart Failure Reviewer 1 Form: FORM CODE=HFD1, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
1533	Present	Text suppressed

VISIT		Visit
N	Value	Description
65	1	Visit 1
98	2	Visit 2
114	3	Visit 3
134	4	Visit 4
141	5	Visit 5
104	6	Visit 6
104	7	Visit 7
135	8	Visit 8
116	9	Visit 9
127	10	Visit 10
184	11	Visit 11
94	12	Visit 12
83	13	Visit 13
32	14	Visit 14
2	15	Visit 15

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OCCURRENCE		Occurrence
N	Value	Description
996	1	Occurrence 1
270	2	Occurrence 2
122	3	Occurrence 3
54	4	Occurrence 4
29	5	Occurrence 5
16	6	Occurrence 6
11	7	Occurrence 7
8	8	Occurrence 8
5	9	Occurrence 9
8	10	Occurrence 10
3	11	Occurrence 11
4	12	Occurrence 12
2	13	Occurrence 13
1	14	Occurrence 14
1	15	Occurrence 15
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19

HFD10A_FUTIME		Follow-up time from baseline for HFD10A (Completion Date)
N	Value	Description
14		Missing
1519	Range	1895 - 6015 (median=3947 mean=3981.5 std=838.5)

HFD10AY		Year of HFD10A
N	Value	Description
14		Missing
1519	Range	2015 - 2025 (median=2021 mean=2020.4 std=2.2)

HFD10C		Event ID
N	Value	Description
1533	Present	Text suppressed

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<i>HFD10D_FUTIME</i>		<i>Follow-up time from baseline for HFD10D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1533	Range	10 - 5302 (median=2623 mean=2563.4 std=1327.2)

<i>HFD10DY</i>		<i>Year of HFD10D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1533	Range	2008 - 2024 (median=2017 mean=2016.6 std=3.6)

<i>HFD11</i>		<i>History of heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
758	0	0=No
687	1	1=Yes
74	9	9=NR

<i>HFD12</i>		<i>ADHF diagnosed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16		Missing
1051	0	0=No/NR
466	1	1=Yes

<i>HFD13</i>		<i>Pulmonary edema/congestion</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16		Missing
930	0	0=No
430	1	1=Yes
157	9	9=NR

<i>HFD14</i>		<i>Dilated ventricle</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25		Missing
520	0	0=No
31	1	1=Yes, history
177	2	2=Yes, current imaging
780	9	9=Unknown

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<i>HFD15</i>		<i>Poor LV function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25		Missing
506	0	0=No
141	1	1=Yes, history
307	2	2=Yes, current imaging
554	9	9=Unknown

<i>HFD16</i>		<i>Poor RV function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25		Missing
510	0	0=No
21	1	1=Yes, history
81	2	2=Yes, current imaging
896	9	9=Unknown

<i>HFD17</i>		<i>Diastolic dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25		Missing
91	0	0=No
81	1	1=Yes, history
316	2	2=Yes, current imaging
1020	9	9=Unknown

<i>HFD18</i>		<i>quantitative EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16		Missing
61	1	1=less than 20
94	2	2=20 to 29
95	3	3=30 to 39
116	4	4=40 to 49
489	5	5=greater than or equal to 50
662	6	6=Unknown

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<i>HFD19</i>		<i>Reviewer Classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
268	1	1=Definite
219	2	2=Probable
313	3	3=No, chronic stable HF
703	4	4=No, HF unlikely
16	5	5=Unclassifiable

<i>HFD19A</i>		<i>classify the severity</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
1032	.S	Legitimately skipped
70	1	1=Mild
301	2	2=Moderate
113	3	3=Severe
3	4	4=Unknown

<i>HFD19B</i>		<i>ADHF right-sided HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
1032	.S	Legitimately skipped
407	0	0=No
20	1	1=Yes
60	9	9=Unknown

<i>HFD110A1</i>		<i>a. Coronary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.5 std=0.5)

<i>HFD110A10</i>		<i>a. Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.0 std=0.2)

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<i>HFD110A11</i>		<i>a.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)

<i>HFD110A12</i>		<i>a.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)

<i>HFD110A13</i>		<i>a.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.0 std=0.1)

<i>HFD110A14</i>		<i>a.Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.0 std=0.2)

<i>HFD110A2</i>		<i>a.Hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=1 mean=0.7 std=0.4)

<i>HFD110A3</i>		<i>a.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)

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<i>HFD110A4</i>		<i>a.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.1 std=0.2)

<i>HFD110A5</i>		<i>a.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)

<i>HFD110A6</i>		<i>a.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.1 std=0.3)

<i>HFD110A7</i>		<i>a.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.1 std=0.4)

<i>HFD110A8</i>		<i>a.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.4 std=0.5)

<i>HFD110A9</i>		<i>a.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
387	.S	Legitimately skipped
172	Range	0 - 1 (median=0 mean=0.1 std=0.3)

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<i>HFD110B1</i>		<i>b. Coronary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
957		Missing
495	.S	Legitimately skipped
81	Range	0 - 1 (median=1 mean=0.6 std=0.5)

<i>HFD110B10</i>		<i>b. Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
573	.S	Legitimately skipped
4	Range	0 - 1 (median=0.5 mean=0.50 std=0.58)

<i>HFD110B11</i>		<i>b. Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
543	.S	Legitimately skipped
34	Range	0 - 1 (median=0.5 mean=0.50 std=0.51)

<i>HFD110B12</i>		<i>b. Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
957		Missing
550	.S	Legitimately skipped
26	Range	1 - 1 (median=1 mean=1.0 std=0.0)

<i>HFD110B13</i>		<i>b. Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
576	.S	Legitimately skipped
1	Range	1 - 1 (median=1 mean=1.0 std=.)

<i>HFD110B14</i>		<i>b. Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
957		Missing
571	.S	Legitimately skipped
5	Range	0 - 1 (median=1 mean=0.6 std=0.5)

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<i>HFD110B2</i>		<i>b.Hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
962		Missing
451	.S	Legitimately skipped
120	Range	0 - 1 (median=0 mean=0.3 std=0.5)

<i>HFD110B3</i>		<i>b.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
550	.S	Legitimately skipped
27	Range	0 - 1 (median=1 mean=0.8 std=0.4)

<i>HFD110B4</i>		<i>b.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
567	.S	Legitimately skipped
10	Range	0 - 1 (median=1 mean=0.6 std=0.5)

<i>HFD110B5</i>		<i>b.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
543	.S	Legitimately skipped
34	Range	0 - 1 (median=1 mean=0.7 std=0.5)

<i>HFD110B6</i>		<i>b.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
556	.S	Legitimately skipped
21	Range	0 - 1 (median=1 mean=1.0 std=0.2)

<i>HFD110B7</i>		<i>b.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
552	.S	Legitimately skipped
25	Range	0 - 1 (median=1 mean=0.9 std=0.3)

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<i>HFD110B8</i>		<i>b.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
957		Missing
514	.S	Legitimately skipped
62	Range	0 - 1 (median=1 mean=0.7 std=0.4)

<i>HFD110B9</i>		<i>b.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
556	.S	Legitimately skipped
21	Range	0 - 1 (median=0 mean=0.4 std=0.5)

<i>HFD111</i>		<i>Asymp LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
1092	0	0=No/NR
38	1	1=Yes
389	9	9=Unknown

<i>HFD112</i>		<i>Reviewer comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1208		Missing
325	Present	Text suppressed

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Heart Failure Reviewer 2 Form: FORM CODE=HFD2, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
1533	Present	Text suppressed

VISIT		Visit
N	Value	Description
65	1	Visit 1
98	2	Visit 2
114	3	Visit 3
134	4	Visit 4
141	5	Visit 5
104	6	Visit 6
104	7	Visit 7
135	8	Visit 8
116	9	Visit 9
127	10	Visit 10
184	11	Visit 11
94	12	Visit 12
83	13	Visit 13
32	14	Visit 14
2	15	Visit 15

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OCCURRENCE		Occurrence
N	Value	Description
996	1	Occurrence 1
270	2	Occurrence 2
122	3	Occurrence 3
54	4	Occurrence 4
29	5	Occurrence 5
16	6	Occurrence 6
11	7	Occurrence 7
8	8	Occurrence 8
5	9	Occurrence 9
8	10	Occurrence 10
3	11	Occurrence 11
4	12	Occurrence 12
2	13	Occurrence 13
1	14	Occurrence 14
1	15	Occurrence 15
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19

HFD20A_FUTIME		Follow-up time from baseline for HFD20A (Completion Date)
N	Value	Description
11		Missing
1522	Range	1886 - 6010 (median=3946 mean=3983.2 std=851.9)

HFD20AY		Year of HFD20A
N	Value	Description
11		Missing
1522	Range	2015 - 2025 (median=2021 mean=2020.4 std=2.3)

HFD20C		Event ID
N	Value	Description
1533	Present	Text suppressed

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<i>HFD20D_FUTIME</i>		<i>Follow-up time from baseline for HFD20D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1533	Range	10 - 5302 (median=2623 mean=2563.4 std=1327.2)

<i>HFD20DY</i>		<i>Year of HFD20D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1533	Range	2008 - 2024 (median=2017 mean=2016.6 std=3.6)

<i>HFD21</i>		<i>History of heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
786	0	0=No
651	1	1=Yes
85	9	9=NR

<i>HFD22</i>		<i>ADHF diagnosed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
12		Missing
1062	0	0=No/NR
459	1	1=Yes

<i>HFD23</i>		<i>Pulmonary edema/congestion</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13		Missing
834	0	0=No
486	1	1=Yes
200	9	9=NR

<i>HFD24</i>		<i>Dilated ventricle</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
510	0	0=No
39	1	1=Yes, history
186	2	2=Yes, current imaging
783	9	9=Unknown

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<i>HFD25</i>		<i>Poor LV function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
498	0	0=No
151	1	1=Yes, history
321	2	2=Yes, current imaging
548	9	9=Unknown

<i>HFD26</i>		<i>Poor RV function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
517	0	0=No
16	1	1=Yes, history
77	2	2=Yes, current imaging
908	9	9=Unknown

<i>HFD27</i>		<i>Diastolic dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
99	0	0=No
71	1	1=Yes, history
352	2	2=Yes, current imaging
996	9	9=Unknown

<i>HFD28</i>		<i>quantitative EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
55	1	1=less than 20
106	2	2=20 to 29
103	3	3=30 to 39
121	4	4=40 to 49
489	5	5=greater than or equal to 50
648	6	6=Unknown

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<i>HFD29</i>		<i>Reviewer Classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
300	1	1=Definite
180	2	2=Probable
267	3	3=No, chronic stable HF
756	4	4=No, HF unlikely
19	5	5=Unclassifiable

<i>HFD29A</i>		<i>classify the severity</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
1042	.S	Legitimately skipped
87	1	1=Mild
294	2	2=Moderate
94	3	3=Severe
5	4	4=Unknown

<i>HFD29B</i>		<i>ADHF right-sided HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
1042	.S	Legitimately skipped
422	0	0=No
13	1	1=Yes
45	9	9=Unknown

<i>HFD210A1</i>		<i>a. Coronary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
996		Missing
407	.S	Legitimately skipped
130	Range	0 - 1 (median=1 mean=0.6 std=0.5)

<i>HFD210A10</i>		<i>a. Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1029		Missing
407	.S	Legitimately skipped
97	Range	0 - 1 (median=0 mean=0.0 std=0.2)

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<i>HFD210A11</i>		<i>a.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1020		Missing
407	.S	Legitimately skipped
106	Range	0 - 1 (median=0 mean=0.3 std=0.5)

<i>HFD210A12</i>		<i>a.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1020		Missing
407	.S	Legitimately skipped
106	Range	0 - 1 (median=0 mean=0.2 std=0.4)

<i>HFD210A13</i>		<i>a.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1030		Missing
407	.S	Legitimately skipped
96	Range	0 - 1 (median=0 mean=0.0 std=0.1)

<i>HFD210A14</i>		<i>a.Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1030		Missing
407	.S	Legitimately skipped
96	Range	0 - 1 (median=0 mean=0.1 std=0.2)

<i>HFD210A2</i>		<i>a.Hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
976		Missing
407	.S	Legitimately skipped
150	Range	0 - 1 (median=1 mean=0.9 std=0.3)

<i>HFD210A3</i>		<i>a.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1016		Missing
407	.S	Legitimately skipped
110	Range	0 - 1 (median=0 mean=0.3 std=0.5)

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<i>HFD210A4</i>		<i>a.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1031		Missing
407	.S	Legitimately skipped
95	Range	0 - 1 (median=0 mean=0.0 std=0.2)

<i>HFD210A5</i>		<i>a.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1022		Missing
407	.S	Legitimately skipped
104	Range	0 - 1 (median=0 mean=0.3 std=0.5)

<i>HFD210A6</i>		<i>a.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1028		Missing
407	.S	Legitimately skipped
98	Range	0 - 1 (median=0 mean=0.1 std=0.3)

<i>HFD210A7</i>		<i>a.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1025		Missing
407	.S	Legitimately skipped
101	Range	0 - 1 (median=0 mean=0.2 std=0.4)

<i>HFD210A8</i>		<i>a.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
994		Missing
407	.S	Legitimately skipped
132	Range	0 - 1 (median=1 mean=0.5 std=0.5)

<i>HFD210A9</i>		<i>a.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1026		Missing
407	.S	Legitimately skipped
100	Range	0 - 1 (median=0 mean=0.2 std=0.4)

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<i>HFD210B1</i>		<i>b. Coronary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
974		Missing
504	.S	Legitimately skipped
55	Range	0 - 1 (median=1 mean=0.6 std=0.5)

<i>HFD210B10</i>		<i>b. Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
957		Missing
574	.S	Legitimately skipped
2	Range	0 - 1 (median=0.5 mean=0.50 std=0.71)

<i>HFD210B11</i>		<i>b. Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
963		Missing
540	.S	Legitimately skipped
30	Range	0 - 1 (median=0 mean=0.3 std=0.5)

<i>HFD210B12</i>		<i>b. Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
962		Missing
555	.S	Legitimately skipped
16	Range	0 - 1 (median=1 mean=0.7 std=0.5)

<i>HFD210B13</i>		<i>b. Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
957		Missing
575	.S	Legitimately skipped
1	Range	0 - 0 (median=0 mean=0.0 std=.)

<i>HFD210B14</i>		<i>b. Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
957		Missing
572	.S	Legitimately skipped
4	Range	0 - 1 (median=1 mean=0.8 std=0.5)

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<i>HFD210B2</i>		<i>b.Hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
998		Missing
443	.S	Legitimately skipped
92	Range	0 - 1 (median=0 mean=0.3 std=0.5)

<i>HFD210B3</i>		<i>b.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
965		Missing
544	.S	Legitimately skipped
24	Range	0 - 1 (median=1 mean=0.5 std=0.5)

<i>HFD210B4</i>		<i>b.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
573	.S	Legitimately skipped
4	Range	0 - 1 (median=0.5 mean=0.50 std=0.58)

<i>HFD210B5</i>		<i>b.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
960		Missing
543	.S	Legitimately skipped
30	Range	0 - 1 (median=1 mean=0.5 std=0.5)

<i>HFD210B6</i>		<i>b.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
956		Missing
567	.S	Legitimately skipped
10	Range	0 - 1 (median=1 mean=0.9 std=0.3)

<i>HFD210B7</i>		<i>b.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
958		Missing
560	.S	Legitimately skipped
15	Range	1 - 1 (median=1 mean=1.0 std=0.0)

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<i>HFD210B8</i>		<i>b.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
975		Missing
505	.S	Legitimately skipped
53	Range	0 - 1 (median=0 mean=0.5 std=0.5)

<i>HFD210B9</i>		<i>b.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
960		Missing
561	.S	Legitimately skipped
12	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>HFD211</i>		<i>Asymp LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
1149	0	0=No/NR
24	1	1=Yes
349	9	9=Unknown

<i>HFD212</i>		<i>Reviewer comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1269		Missing
264	Present	Text suppressed

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Section C: Heart Failure Ascertainment

Heart Failure Adjudication Reviewer Form: FORM CODE=HFD3, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
605	Present	Text suppressed

VISIT		Visit
N	Value	Description
43	1	Visit 1
58	2	Visit 2
54	3	Visit 3
61	4	Visit 4
56	5	Visit 5
35	6	Visit 6
41	7	Visit 7
52	8	Visit 8
44	9	Visit 9
37	10	Visit 10
56	11	Visit 11
31	12	Visit 12
32	13	Visit 13
4	14	Visit 14
1	15	Visit 15

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<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
366	1	Occurrence 1
110	2	Occurrence 2
61	3	Occurrence 3
28	4	Occurrence 4
12	5	Occurrence 5
9	6	Occurrence 6
6	7	Occurrence 7
2	8	Occurrence 8
2	9	Occurrence 9
5	10	Occurrence 10
1	11	Occurrence 11
1	12	Occurrence 12
1	18	Occurrence 18
1	19	Occurrence 19

<i>HFD30A_FUTIME</i>		<i>Follow-up time from baseline for HFD30A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
566	Range	2002 - 6049 (median=4130.5 mean=4289.41 std=791.95)

<i>HFD30AY</i>		<i>Year of HFD30A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
566	Range	2015 - 2025 (median=2020 mean=2021.2 std=2.0)

<i>HFD30C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
605	Present	Text suppressed

<i>HFD30D_FUTIME</i>		<i>Follow-up time from baseline for HFD30D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
605	Range	10 - 5116 (median=2228 mean=2288.6 std=1356.5)

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<i>HFD30DY</i>		<i>Year of HFD30D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
605	Range	2008 - 2023 (median=2016 mean=2015.8 std=3.7)

<i>HFD31</i>		<i>History of heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
184	0	0=No
359	1	1=Yes
23	9	9=NR

<i>HFD32</i>		<i>ADHF diagnosed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
292	0	0=No/NR
274	1	1=Yes

<i>HFD33</i>		<i>Pulmonary edema/congestion</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
246	0	0=No
252	1	1=Yes
68	9	9=NR

<i>HFD34</i>		<i>Dilated ventricle</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
182	0	0=No
14	1	1=Yes, history
93	2	2=Yes, current imaging
277	9	9=Unknown

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<i>HFD35</i>		<i>Poor LV function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
172	0	0=No
73	1	1=Yes, history
179	2	2=Yes, current imaging
142	9	9=Unknown

<i>HFD36</i>		<i>Poor RV function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
199	0	0=No
6	1	1=Yes, history
49	2	2=Yes, current imaging
312	9	9=Unknown

<i>HFD37</i>		<i>Diastolic dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
22	0	0=No
60	1	1=Yes, history
139	2	2=Yes, current imaging
345	9	9=Unknown

<i>HFD38</i>		<i>quantitative EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
27	1	1=less than 20
47	2	2=20 to 29
54	3	3=30 to 39
71	4	4=40 to 49
168	5	5=greater than or equal to 50
199	6	6=Unknown

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<i>HFD39</i>		<i>Reviewer Classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
138	1	1=Definite
167	2	2=Probable
150	3	3=No, chronic stable HF
96	4	4=No, HF unlikely
15	5	5=Unclassifiable

<i>HFD39A</i>		<i>classify the severity</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
36		Missing
264	.S	Legitimately skipped
31	1	1=Mild
204	2	2=Moderate
67	3	3=Severe
3	4	4=Unknown

<i>HFD39B</i>		<i>ADHF right-sided HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
36		Missing
264	.S	Legitimately skipped
274	0	0=No
7	1	1=Yes
24	9	9=Unknown

<i>HFD310A1</i>		<i>a. Coronary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0.5 mean=0.50 std=0.52)

<i>HFD310A10</i>		<i>a. Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.1 std=0.3)

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<i>HFD310A11</i>		<i>a.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.1 std=0.4)

<i>HFD310A12</i>		<i>a.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.1 std=0.4)

<i>HFD310A13</i>		<i>a.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>HFD310A14</i>		<i>a.Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>HFD310A2</i>		<i>a.Hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=1 mean=0.6 std=0.5)

<i>HFD310A3</i>		<i>a.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.1 std=0.4)

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<i>HFD310A4</i>		<i>a.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>HFD310A5</i>		<i>a.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.4 std=0.5)

<i>HFD310A6</i>		<i>a.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.1 std=0.4)

<i>HFD310A7</i>		<i>a.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>HFD310A8</i>		<i>a.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.1 std=0.3)

<i>HFD310A9</i>		<i>a.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
27	.S	Legitimately skipped
14	Range	0 - 1 (median=0 mean=0.2 std=0.4)

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<i>HFD310B1</i>		<i>b. Coronary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
46	.S	Legitimately skipped
7	Range	0 - 1 (median=1 mean=0.7 std=0.5)

<i>HFD310B10</i>		<i>b. Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
553		Missing
52	.S	Legitimately skipped

<i>HFD310B11</i>		<i>b. Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
51	.S	Legitimately skipped
2	Range	0 - 1 (median=0.5 mean=0.50 std=0.71)

<i>HFD310B12</i>		<i>b. Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
554		Missing
51	.S	Legitimately skipped

<i>HFD310B13</i>		<i>b. Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
53	.S	Legitimately skipped

<i>HFD310B14</i>		<i>b. Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
53	.S	Legitimately skipped

<i>HFD310B2</i>		<i>b. Hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
44	.S	Legitimately skipped
9	Range	0 - 1 (median=0 mean=0.1 std=0.3)

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<i>HFD310B3</i>		<i>b. Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
51	.S	Legitimately skipped
2	Range	1 - 1 (median=1 mean=1.0 std=0.0)

<i>HFD310B4</i>		<i>b. Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
53	.S	Legitimately skipped

<i>HFD310B5</i>		<i>b. Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
553		Missing
47	.S	Legitimately skipped
5	Range	0 - 1 (median=1 mean=0.6 std=0.5)

<i>HFD310B6</i>		<i>b. Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
553		Missing
51	.S	Legitimately skipped
1	Range	1 - 1 (median=1 mean=1.0 std=.)

<i>HFD310B7</i>		<i>b. Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
53	.S	Legitimately skipped

<i>HFD310B8</i>		<i>b. Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
52	.S	Legitimately skipped
1	Range	0 - 0 (median=0 mean=0.0 std=.)

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<i>HFD310B9</i>		<i>b. Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552		Missing
50	.S	Legitimately skipped
3	Range	0 - 1 (median=1 mean=0.7 std=0.6)

<i>HFD311</i>		<i>Asymp LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39		Missing
465	0	0=No/NR
17	1	1=Yes
84	9	9=Unknown

<i>HFD312</i>		<i>Reviewer comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
550		Missing
55	Present	Text suppressed

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Pulmonary Abstraction Form: FORM CODE=PUL, VERSION=B

ID		HCHS Participant ID
N	Value	Description
3015	Present	Text suppressed

VISIT		Visit
N	Value	Description
270	1	Visit 1
293	2	Visit 2
305	3	Visit 3
269	4	Visit 4
304	5	Visit 5
208	6	Visit 6
201	7	Visit 7
211	8	Visit 8
163	9	Visit 9
181	10	Visit 10
269	11	Visit 11
151	12	Visit 12
121	13	Visit 13
61	14	Visit 14
8	15	Visit 15

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<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2076	1	Occurrence 1
505	2	Occurrence 2
185	3	Occurrence 3
85	4	Occurrence 4
52	5	Occurrence 5
31	6	Occurrence 6
16	7	Occurrence 7
14	8	Occurrence 8
8	9	Occurrence 9
8	10	Occurrence 10
6	11	Occurrence 11
3	12	Occurrence 12
1	13	Occurrence 13
1	14	Occurrence 14
1	15	Occurrence 15
1	16	Occurrence 16
2	17	Occurrence 17
2	18	Occurrence 18
1	19	Occurrence 19
3	21	Occurrence 21
12	31	Occurrence 31
1	32	Occurrence 32
1	40	Occurrence 40

<i>PUL0A_FUTIME</i>		<i>Follow-up time from baseline for PUL0A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3015	Range	22 - 6094 (median=2958 mean=3231.1 std=1174.5)

<i>PUL0AY</i>		<i>Year of PUL0A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3015	Range	2009 - 2025 (median=2017 mean=2018.4 std=3.2)

<i>PUL0C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3015	Present	Text suppressed

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<i>PUL0D_FUTIME</i>		<i>Follow-up time from baseline for PUL0D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3015	Range	1 - 5556 (median=2028 mean=2200.6 std=1382.4)

<i>PUL0DY</i>		<i>Year of PUL0D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3015	Range	2008 - 2024 (median=2015 mean=2015.6 std=3.8)

<i>PUL1</i>		<i>Event type</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
249	1	1= In hospital only
635	2	2= Emergency Dept. visit only (ED)
2104	3	3= Both ED and in hospital
2	4	4= Observation care only
20	5	5= Both ED and observation care

<i>PUL2</i>		<i>Hospital stay less than 24 hrs</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
638	.S	Legitimately skipped
2293	0	0=No/NR
82	1	1=Yes

<i>PUL3A_FUTIME</i>		<i>Follow-up time from baseline for PUL3A (Date of arrival)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
3000	Range	1 - 5556 (median=2046.5 mean=2206.05 std=1384.55)

<i>PUL3AY</i>		<i>Year of PUL3A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
3000	Range	2008 - 2024 (median=2015 mean=2015.6 std=3.8)

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<i>PUL3B</i>		<i>Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
160		Missing
2855	Range	0:00:00 - 23:58:00

<i>PUL3B1</i>		<i>AM/PM of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
46		Missing
2530	.S	Legitimately skipped
195	1	1=AM
244	2	2=PM

<i>PUL3C_FUTIME</i>		<i>Follow-up time from baseline for PUL3C (Date of admission)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
647		Missing
2368	Range	2 - 5516 (median=2200 mean=2320.1 std=1390.4)

<i>PUL3CY</i>		<i>Year of PUL3C</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
647		Missing
2368	Range	2008 - 2024 (median=2016 mean=2015.9 std=3.9)

<i>PUL4A_FUTIME</i>		<i>Follow-up time from baseline for PUL4A (Date of discharge)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29		Missing
2986	Range	4 - 5556 (median=2040 mean=2207.1 std=1378.9)

<i>PUL4AY</i>		<i>Year of PUL4A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
28		Missing
2987	Range	2008 - 2024 (median=2015 mean=2015.6 std=3.8)

<i>PUL4B</i>		<i>Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
967		Missing
2048	Range	0:00:00 - 23:59:00

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<i>PUL4B1</i>		<i>AM/PM of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
148		Missing
2530	.S	Legitimately skipped
79	1	1=AM
258	2	2=PM

<i>PUL5</i>		<i>Primary admitting diagnosis code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
541		Missing
2474	Present	Text suppressed

<i>PUL6</i>		<i>Primary discharge diagnosis code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
362		Missing
2653	Present	Text suppressed

<i>PUL7</i>		<i>Transported in EMS unit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2193	0	0=No/NR
822	1	1=Yes

<i>PUL8</i>		<i>Transferred from another hospital</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2941	0	0=No/NR
74	1	1=Yes

<i>PUL9</i>		<i>Code status no-code or DNR</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2875	0	0=No/NR
140	1	1=Yes

<i>PUL10</i>		<i>Patient alive at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
199	0	0=No/NR
2814	1	1=Yes

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<i>PUL11A</i>		<i>Cough at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
697	0	0=No
1609	1	1=Yes
709	9	9=Not Recorded

<i>PUL11B</i>		<i>Sputum production at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
999	0	0=No
768	1	1=Yes
1248	9	9=Not Recorded

<i>PUL11C</i>		<i>Sputum purulence at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1088	0	0=No
397	1	1=Yes
1530	9	9=Not Recorded

<i>PUL11D</i>		<i>Wheezing at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
994	0	0=No
833	1	1=Yes
1188	9	9=Not Recorded

<i>PUL11E</i>		<i>Chest pain at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
1345	0	0=No
1111	1	1=Yes
558	9	9=Not Recorded

<i>PUL11F</i>		<i>Leg edema at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1505	0	0=No
357	1	1=Yes
1153	9	9=Not Recorded

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<i>PUL11G</i>		<i>Rescue bronchodilator at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
526	0	0=No
1197	1	1=Yes
1292	9	9=Not Recorded

<i>PUL11H</i>		<i>Dyspnea at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
594	0	0=No
2103	1	1=Yes
318	9	9=Not Recorded

<i>PUL11I</i>		<i>Dyspnea (at rest) at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
794	0	0=No
1078	1	1=Yes
1143	9	9=Not Recorded

<i>PUL11J</i>		<i>Dyspnea (walking/on exertion) at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
487	0	0=No
1693	1	1=Yes
835	9	9=Not Recorded

<i>PUL11K</i>		<i>Woken up at night at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
486	0	0=No
102	1	1=Yes
2427	9	9=Not Recorded

<i>PUL11L</i>		<i>Fever at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1967	0	0=No
650	1	1=Yes
398	9	9=Not Recorded

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<i>PUL11M</i>		<i>AMS at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2541	0	0=No
312	1	1=Yes
162	9	9=Not Recorded

<i>PUL12</i>		<i>Reason COPD/chronic bronchitis/emphysema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2584	0	0=No/NR
431	1	1=Yes

<i>PUL13</i>		<i>Reason exacerbation of asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2401	0	0=No/NR
614	1	1=Yes

<i>PUL14</i>		<i>New onset of exacerbation prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1985	0	0=No/NR
1030	1	1=Yes

<i>PUL15A</i>		<i>Asthma history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1493	0	0=No/NR
1522	1	1=Yes

<i>PUL15B</i>		<i>Chronic bronchitis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2922	0	0=No/NR
93	1	1=Yes

<i>PUL15C</i>		<i>Emphysema history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2878	0	0=No/NR
137	1	1=Yes

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<i>PUL15D</i>		<i>COPD history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2170	0	0=No/NR
845	1	1=Yes

<i>PUL15E</i>		<i>Pulmonary fibrosis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2958	0	0=No/NR
57	1	1=Yes

<i>PUL15F</i>		<i>Sarcoidosis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3013	0	0=No/NR
2	1	1=Yes

<i>PUL15G</i>		<i>Lung cancer history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2962	0	0=No/NR
53	1	1=Yes

<i>PUL15H</i>		<i>Lung resection or lobectomy history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3001	0	0=No/NR
14	1	1=Yes

<i>PUL15I</i>		<i>Home oxygen history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2860	0	0=No/NR
155	1	1=Yes

<i>PUL15J</i>		<i>Pulmonary embolus history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2967	0	0=No/NR
48	1	1=Yes

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<i>PUL15K</i>		<i>Pulmonary hypertension history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2918	0	0=No/NR
97	1	1=Yes

<i>PUL15L</i>		<i>Cor pulmonale history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2835	0	0=No/NR
180	1	1=Yes

<i>PUL15M</i>		<i>OSA history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2590	0	0=No/NR
425	1	1=Yes

<i>PUL15N</i>		<i>Coronary artery disease history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2386	0	0=No/NR
629	1	1=Yes

<i>PUL15O</i>		<i>Heart failure history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2510	0	0=No/NR
505	1	1=Yes

<i>PUL15P</i>		<i>Atrial fibrillation/atrial flutter history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2802	0	0=No/NR
213	1	1=Yes

<i>PUL15Q</i>		<i>Diabetes history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1611	0	0=No/NR
1404	1	1=Yes

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<i>PUL15R</i>		<i>Pulmonary Tuberculosis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2967	0	0=No/NR
48	1	1=Yes

<i>PUL15S</i>		<i>Bronchiectasis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2973	0	0=No/NR
42	1	1=Yes

<i>PUL16A</i>		<i>Pre-bronchodilator FEV1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2996		Missing
19	Range	19 - 99 (median=70 mean=61.6 std=25.6)

<i>PUL16B</i>		<i>Post-bronchodilator FEV1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2995		Missing
20	Range	18 - 150 (median=77 mean=75.3 std=34.2)

<i>PUL17</i>		<i>FEV1/FVC ratio</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2991		Missing
24	Range	52 - 93 (median=74.5 mean=73.85 std=12.46)

<i>PUL17A</i>		<i>FEV1/FVC ratio Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2990		Missing
0	1	1=proportion or decimal
25	2	2=percent (If percent, then assure not percent predicted for the ratio)

<i>PUL18A</i>		<i>Active Upper Respiratory Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2603	0	0=No/NR
412	1	1=Yes

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<i>PUL18B</i>		<i>Active Pneumonia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2354	0	0=No/NR
661	1	1=Yes

<i>PUL18C</i>		<i>Active Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2981	0	0=No/NR
34	1	1=Yes

<i>PUL18D</i>		<i>Active Myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2886	0	0=No/NR
129	1	1=Yes

<i>PUL18E</i>		<i>Active Heart failure exacerbation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2706	0	0=No/NR
309	1	1=Yes

<i>PUL18F</i>		<i>Active Atrial fibrillation/flutter</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2869	0	0=No/NR
146	1	1=Yes

<i>PUL18G</i>		<i>Active SVT/MAT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2992	0	0=No/NR
23	1	1=Yes

<i>PUL18H</i>		<i>Active CABG/Valvular Surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2971	0	0=No/NR
44	1	1=Yes

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<i>PUL18I</i>		<i>Active Non-cardiac surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2686	0	0=No/NR
329	1	1=Yes

<i>PUL19A</i>		<i>Heart rate arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
313		Missing
2702	Range	0 - 199 (median=89 mean=90.7 std=20.3)

<i>PUL19B</i>		<i>Respiration rate arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
369		Missing
2646	Range	0 - 55 (median=19 mean=19.9 std=4.3)

<i>PUL19C</i>		<i>Oxy sat arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564		Missing
2451	Range	0 - 160 (median=97 mean=95.5 std=6.6)

<i>PUL19C1</i>		<i>Oxy Sats on room air arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
439		Missing
199	0	0=No
1143	1	1=Yes
1234	9	9=NR

<i>PUL19C2</i>		<i>Level oxy arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
121		Missing
2741	.S	Legitimately skipped
153	Range	1 - 180 (median=3 mean=18.7 std=35.2)

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<i>PUL19C3</i>		<i>Level oxy Units arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
121		Missing
2741	.S	Legitimately skipped
126	1	1= Liters
27	2	2=Percent

<i>PUL19D</i>		<i>Weight arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1221		Missing
1794	Range	36.36 - 575 (median=85 mean=101.3 std=49.8)

<i>PUL19D1</i>		<i>Weight units arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1232		Missing
291	1	1= Lbs
1492	2	2=Kg

<i>PUL20A</i>		<i>Use of accessory muscles at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
679	0	0=No
117	1	1=Yes
2219	9	9=NR

<i>PUL20B</i>		<i>Cyanosis at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1495	0	0=No
21	1	1=Yes
1499	9	9=NR

<i>PUL20C</i>		<i>Clubbing at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1133	0	0=No
11	1	1=Yes
1871	9	9=NR

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<i>PUL20D</i>		<i>JVD at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1605	0	0=No
81	1	1=Yes
1329	9	9=NR

<i>PUL20E</i>		<i>Crackles/rales at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1693	0	0=No
645	1	1=Yes
677	9	9=NR

<i>PUL20F</i>		<i>Wheezing or rhonchi at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1444	0	0=No
1219	1	1=Yes
352	9	9=NR

<i>PUL20G</i>		<i>Dec unilateral breath at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1114	0	0=No
40	1	1=Yes
1861	9	9=NR

<i>PUL20H</i>		<i>Dec bilateral breath at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
487	0	0=No
703	1	1=Yes
1825	9	9=NR

<i>PUL20I</i>		<i>Prolonged expiratory time at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
219	0	0=No
84	1	1=Yes
2712	9	9=NR

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<i>PUL20J</i>		<i>Egophony at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
174	0	0=No
10	1	1=Yes
2831	9	9=NR

<i>PUL20K</i>		<i>Lower extremity edema at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2012	0	0=No
457	1	1=Yes
546	9	9=NR

<i>PUL21</i>		<i>Chest X-ray performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	0	0=No/NR
2402	1	1=Yes

<i>PUL22A</i>		<i>Hyperinflation on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2314	0	0=No/NR
88	1	1=Yes

<i>PUL22B</i>		<i>Flattened diaphragms on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2376	0	0=No/NR
26	1	1=Yes

<i>PUL22C</i>		<i>Consolidation or infiltrate on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
1663	0	0=No/NR
739	1	1=Yes

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<i>PUL22D</i>		<i>Scarring on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2335	0	0=No/NR
67	1	1=Yes

<i>PUL22E</i>		<i>Nodules on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2398	0	0=No/NR
4	1	1=Yes

<i>PUL22F</i>		<i>Masses on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2397	0	0=No/NR
5	1	1=Yes

<i>PUL22G</i>		<i>Pulmonary edema on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
1860	0	0=No/NR
542	1	1=Yes

<i>PUL22H</i>		<i>Bilateral pleural effusion on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2192	0	0=No/NR
210	1	1=Yes

<i>PUL22I</i>		<i>Unilateral pleural effusion on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2240	0	0=No/NR
162	1	1=Yes

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<i>PUL22J</i>		<i>Emphysema on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
2359	0	0=No/NR
43	1	1=Yes

<i>PUL22K</i>		<i>Cardiomegaly on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
613	.S	Legitimately skipped
1857	0	0=No/NR
545	1	1=Yes

<i>PUL23</i>		<i>Chest/lung CT scan or CTA performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2254	0	0=No/NR
760	1	1=Yes

<i>PUL24A</i>		<i>Emphysema on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
660	0	0=No/NR
100	1	1=Yes

<i>PUL24B</i>		<i>Nodules on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
737	0	0=No/NR
23	1	1=Yes

<i>PUL24C</i>		<i>Masses on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
747	0	0=No/NR
13	1	1=Yes

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<i>PUL24D</i>		<i>Lymphadenopathy on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
663	0	0=No/NR
97	1	1=Yes

<i>PUL24E</i>		<i>Ground glass changes on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
578	0	0=No/NR
182	1	1=Yes

<i>PUL24F</i>		<i>Pneumonia on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
624	0	0=No/NR
136	1	1=Yes

<i>PUL24G</i>		<i>Fibrosis or honeycombing on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
730	0	0=No/NR
30	1	1=Yes

<i>PUL24H</i>		<i>Filling defect-vascular (PE) on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
752	0	0=No/NR
8	1	1=Yes

<i>PUL24I</i>		<i>Filling defect-mucus plug on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
746	0	0=No/NR
14	1	1=Yes

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<i>PUL24J</i>		<i>Cysts or blebs on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
749	0	0=No/NR
11	1	1=Yes

<i>PUL24K</i>		<i>Atelectasis on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
533	0	0=No/NR
227	1	1=Yes

<i>PUL24L</i>		<i>Calcifications on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
647	0	0=No/NR
113	1	1=Yes

<i>PUL24M</i>		<i>Pulmonary embolus on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
734	0	0=No/NR
26	1	1=Yes

<i>PUL24N</i>		<i>Enlarged pulmonary artery on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
720	0	0=No/NR
40	1	1=Yes

<i>PUL24O</i>		<i>Bronchiectasis on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
712	0	0=No/NR
48	1	1=Yes

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<i>PUL24P</i>		<i>Pulmonary edema on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
675	0	0=No/NR
85	1	1=Yes

<i>PUL24Q</i>		<i>Cardiomegaly on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
642	0	0=No/NR
118	1	1=Yes

<i>PUL24R</i>		<i>Bilateral pleural effusion on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
638	0	0=No/NR
122	1	1=Yes

<i>PUL24S</i>		<i>Unilateral pleural effusion on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
703	0	0=No/NR
57	1	1=Yes

<i>PUL24T</i>		<i>Airway wall thickening on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2255	.S	Legitimately skipped
709	0	0=No/NR
51	1	1=Yes

<i>PUL25</i>		<i>Spirometry performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2989	0	0=No/NR
26	1	1=Yes

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<i>PUL25A1</i>		<i>FEV1 spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2989	.S	Legitimately skipped
24	Range	0.66 - 3.07 (median=1.6 mean=1.70 std=0.70)

<i>PUL25A2</i>		<i>FEV1 Percent Predicted spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2989	.S	Legitimately skipped
24	Range	19 - 118 (median=66.5 mean=66.50 std=23.25)

<i>PUL25B1</i>		<i>FVC spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2989	.S	Legitimately skipped
23	Range	0.66 - 4.91 (median=2.36 mean=2.437 std=1.087)

<i>PUL25B2</i>		<i>FVC Percent Predicted spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2989	.S	Legitimately skipped
23	Range	15 - 121 (median=75 mean=73.3 std=25.5)

<i>PUL25C1</i>		<i>FEV1/FVC ratio spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
2989	.S	Legitimately skipped
23	Range	0.93 - 88 (median=71 mean=69.1 std=17.6)

<i>PUL25C2</i>		<i>Units for FEV1/FVC ratio spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
2989	.S	Legitimately skipped
1	1	1=proportion or decimal
20	2	2=percent (If percent, then assure not percent predicted for the ratio)

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<i>PUL26</i>		<i>Post-bronchodilator spirometry performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3001	0	0=No/NR
14	1	1=Yes

<i>PUL26A1</i>		<i>FEV1 post</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3001	.S	Legitimately skipped
14	Range	0.63 - 3.28 (median=2.255 mean=2.0307 std=0.8454)

<i>PUL26A2</i>		<i>FEV1 Percent Predicted post</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
3001	.S	Legitimately skipped
10	Range	18 - 81 (median=63 mean=61.0 std=19.9)

<i>PUL26B1</i>		<i>FVC post</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
3001	.S	Legitimately skipped
10	Range	0.63 - 3.76 (median=2.18 mean=2.388 std=1.063)

<i>PUL26B2</i>		<i>FVC Percent Predicted post</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
3001	.S	Legitimately skipped
10	Range	15 - 94 (median=71 mean=67.0 std=23.0)

<i>PUL26C1</i>		<i>FEV1/FVC ratio post</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
3001	.S	Legitimately skipped
10	Range	55 - 89 (median=69.19 mean=71.538 std=10.549)

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<i>PUL26C2</i>		<i>Units for FEV1/FVC ratio post</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
3001	.S	Legitimately skipped
0	1	1=proportion or decimal
10	2	2=percent (If percent, then assure not percent predicted for the ratio)

<i>PUL27</i>		<i>PEFR or PEF obtained at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2810	0	0=No/NR
204	1	1=Yes

<i>PUL27A_FUTIME</i>		<i>Follow-up time from baseline for PUL27A (Date of first PEF(R) taken)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2819		Missing
196	Range	41 - 4763 (median=974.5 mean=1215.59 std=946.95)

<i>PUL27AY</i>		<i>Year of PUL27A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2819		Missing
196	Range	2008 - 2022 (median=2012 mean=2012.8 std=2.7)

<i>PUL27B</i>		<i>First PEF recording</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6		Missing
2811	.S	Legitimately skipped
198	Range	50 - 450 (median=200 mean=211.3 std=77.7)

<i>PUL27C</i>		<i>Worst recording</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
2811	.S	Legitimately skipped
196	Range	50 - 450 (median=200 mean=207.9 std=76.7)

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<i>PUL28</i>		<i>PERF or PEF obtained at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2856	0	0=No/NR
159	1	1=Yes

<i>PUL28A_FUTIME</i>		<i>Follow-up time from baseline for PUL28A (Date of last PEF(R) taken)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2865		Missing
150	Range	41 - 4768 (median=885 mean=1174.0 std=923.5)

<i>PUL28AY</i>		<i>Year of PUL28A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2865		Missing
150	Range	2008 - 2022 (median=2012 mean=2012.7 std=2.5)

<i>PUL28B</i>		<i>Last PEF recording</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2856	.S	Legitimately skipped
159	Range	120 - 510 (median=300 mean=300.3 std=84.4)

<i>PUL29</i>		<i>VQ Scan done</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2962	0	0=No/NR
53	1	1=Yes

<i>PUL29A</i>		<i>VQ Scan results</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
2962	.S	Legitimately skipped
2	1	1=High probability
1	2	2=Intermediate probability
21	3	3=Low probability
26	4	4=No evidence of Pulmonary Embolus
1	5	5=Indeterminate

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<i>PUL30</i>		<i>TTE or TEE performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2329	0	0=No/NR
686	1	1=Yes

<i>PUL30A</i>		<i>TTE/TEE Ejection fraction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
69		Missing
2329	.S	Legitimately skipped
617	Range	5 - 85 (median=56 mean=52.9 std=15.4)

<i>PUL30B</i>		<i>TTE/TEE RVSP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
338		Missing
2329	.S	Legitimately skipped
348	Range	5 - 104 (median=36.855 mean=38.5762 std=16.0103)

<i>PUL30C</i>		<i>Right Ventricular Hypertrophy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
2329	.S	Legitimately skipped
139	0	0=None
6	1	1=Present
3	2	2=Mild
1	3	3=Mod
0	4	4=Severe
528	9	9=NR

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<i>PUL30D</i>		<i>Impaired RV systolic function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
2329	.S	Legitimately skipped
377	0	0=None
15	1	1=Present
17	2	2=Mild
15	3	3=Mod
9	4	4=Severe
245	9	9=NR

<i>PUL30E</i>		<i>Pulmonary hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
2329	.S	Legitimately skipped
95	0	0=None
11	1	1=Present
34	2	2=Mild
35	3	3=Mod
27	4	4=Severe
476	9	9=NR

<i>PUL30F</i>		<i>Tricuspid Regurgitation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
2329	.S	Legitimately skipped
181	0	0=None
12	1	1=Present
219	2	2=Mild
69	3	3=Mod
20	4	4=Severe
177	9	9=NR

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<i>PUL30G</i>		<i>Diastolic dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
2329	.S	Legitimately skipped
422	0	0=No/Unknown/NR
256	1	1=Yes

<i>PUL31</i>		<i>White Blood Cell Count</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
562		Missing
2453	Range	0.1 - 174 (median=9.8 mean=11.09 std=6.52)

<i>PUL32A</i>		<i>First Hemoglobin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
556		Missing
2459	Range	3.8 - 18.7 (median=12.1 mean=11.78 std=2.29)

<i>PUL33A</i>		<i>First Hematocrit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
593		Missing
2422	Range	11.7 - 55 (median=36.7 mean=35.82 std=6.48)

<i>PUL34A</i>		<i>First Sodium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
600		Missing
2415	Range	101 - 162 (median=137 mean=136.7 std=4.5)

<i>PUL35A</i>		<i>First Serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
579		Missing
2436	Range	0.2 - 37 (median=0.9 mean=1.46 std=1.91)

<i>PUL36A</i>		<i>First BUN</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
597		Missing
2418	Range	1.1 - 179 (median=18 mean=24.6 std=20.6)

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<i>PUL37A</i>		<i>First Bicarbonate</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
657		Missing
2358	Range	4 - 91 (median=27 mean=27.1 std=4.8)

<i>PUL38A</i>		<i>First BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2613		Missing
402	Range	3.7 - 116000 (median=128.3 mean=976.27 std=6030.54)

<i>PUL38B</i>		<i>Upper limit normal BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2742		Missing
273	Range	23 - 250 (median=100 mean=103.8 std=22.0)

<i>PUL39A</i>		<i>First ProBNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2540		Missing
475	Range	4 - 175001 (median=442 mean=6002.3 std=17429.9)

<i>PUL39B</i>		<i>Upper limit normal ProBNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2675		Missing
340	Range	99 - 70000 (median=125 mean=550.1 std=3793.1)

<i>PUL40</i>		<i>ABGs obtained</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
2289	0	0=No/NR
711	1	1=Yes

<i>PUL40A1</i>		<i>First blood gas pH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42		Missing
2300	.S	Legitimately skipped
673	Range	6.59 - 7.69 (median=7.4 mean=7.37 std=0.13)

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<i>PUL40A2</i>		<i>First blood gas PaCO2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
44		Missing
2300	.S	Legitimately skipped
671	Range	10.6 - 184 (median=40 mean=43.9 std=17.2)

<i>PUL40A3</i>		<i>First blood gas PaO2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
46		Missing
2300	.S	Legitimately skipped
669	Range	16 - 623.3 (median=78.2 mean=108.22 std=90.04)

<i>PUL40A4</i>		<i>First blood gas O2 Sat</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
151		Missing
2300	.S	Legitimately skipped
564	Range	18 - 100 (median=95.65 mean=90.976 std=14.045)

<i>PUL40B1</i>		<i>Last blood gas pH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
406		Missing
2300	.S	Legitimately skipped
309	Range	6.838 - 7.58 (median=7.402 mean=7.3773 std=0.1104)

<i>PUL40B2</i>		<i>Last blood gas PaCO2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
407		Missing
2300	.S	Legitimately skipped
308	Range	21 - 111 (median=42 mean=44.6 std=14.7)

<i>PUL40B3</i>		<i>Last blood gas PaO2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
406		Missing
2300	.S	Legitimately skipped
309	Range	19.1 - 534 (median=86.4 mean=102.80 std=57.62)

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<i>PUL40B4</i>		<i>Last blood gas O2 Sat</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
450		Missing
2300	.S	Legitimately skipped
265	Range	31 - 100 (median=96.4 mean=93.79 std=10.09)

<i>PUL40C</i>		<i>Blood gas on room air</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
2300	.S	Legitimately skipped
288	0	0=No
190	1	1=Yes
226	9	9=Not Recorded

<i>PUL40C1</i>		<i>Level oxy blood gas</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14		Missing
2725	.S	Legitimately skipped
276	Range	0.6 - 100 (median=40 mean=52.4 std=34.7)

<i>PUL40C2</i>		<i>Level oxy Units blood gas</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
12		Missing
2725	.S	Legitimately skipped
44	1	1=Liters
234	2	2=Percent

<i>PUL41</i>		<i>Sputum culture done</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2665	0	0=No/NR
349	1	1=Yes

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<i>PUL41A</i>		<i>Sputum Culture Results</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2666	.S	Legitimately skipped
125	0	0=Neg
104	1	1=Pos
120	9	9=Not Recorded

<i>PUL41B1</i>		<i>Haemophilus Influenzae sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
102	0	0=No
1	1	1=Yes

<i>PUL41B10</i>		<i>Other sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
34		Missing
2911	~	Legitimately skipped
70	Present	Text suppressed

<i>PUL41B2</i>		<i>Moraxella Catarrhalis sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
103	0	0=No
0	1	1=Yes

<i>PUL41B3</i>		<i>Streptococcus pneumoniae sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
100	0	0=No
3	1	1=Yes

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<i>PUL41B4</i>		<i>Methicillin-resistant Staph sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
86	0	0=No
17	1	1=Yes

<i>PUL41B5</i>		<i>Staphylococcus aureus sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
98	0	0=No
5	1	1=Yes

<i>PUL41B6</i>		<i>Mycoplasma pneumoniae sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
101	0	0=No
2	1	1=Yes

<i>PUL41B7</i>		<i>Pseudomonas Aureginosa sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
93	0	0=No
10	1	1=Yes

<i>PUL41B8</i>		<i>Chlamydophila pneumoniae sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
103	0	0=No
0	1	1=Yes

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<i>PUL41B9</i>		<i>Oropharyngeal flora sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2911	.S	Legitimately skipped
83	0	0=No
20	1	1=Yes

<i>PUL42</i>		<i>Blood culture done</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2243	0	0=No/NR
772	1	1=Yes

<i>PUL42A</i>		<i>Culture Results</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2243	.S	Legitimately skipped
446	0	0=Neg
70	1	1=Pos
255	9	9=Not Recorded

<i>PUL42B1</i>		<i>Haemophilus Influenzae blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2945	.S	Legitimately skipped
69	0	0=No
0	1	1=Yes

<i>PUL42B2</i>		<i>Moraxella Catarrhalis blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2945	.S	Legitimately skipped
69	0	0=No
0	1	1=Yes

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<i>PUL42B3</i>		<i>Streptococcus pneumoniae blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2945	.S	Legitimately skipped
66	0	0=No
3	1	1=Yes

<i>PUL42B4</i>		<i>Methicillin-resistant Staph blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2945	.S	Legitimately skipped
57	0	0=No
12	1	1=Yes

<i>PUL42B5</i>		<i>Staphylococcus aureus blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2945	.S	Legitimately skipped
64	0	0=No
5	1	1=Yes

<i>PUL42B6</i>		<i>Other blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13		Missing
2945	~	Legitimately skipped
57	Present	Text suppressed

<i>PUL43</i>		<i>Influenza swab</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
21		Missing
443	0	0=Neg
69	1	1=Pos
2482	9	9=Not Recorded

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<i>PUL44</i>		<i>CPAP or BiPap</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2714	0	0=No/NR
301	1	1=Yes

<i>PUL45</i>		<i>Mechanical Ventilation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2716	0	0=No/NR
299	1	1=Yes

<i>PUL46</i>		<i>Inhaled short-acting beta-agonists</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1566	0	0=No/NR
1449	1	1=Yes

<i>PUL47</i>		<i>Inhaled short-acting anticholinergic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1851	0	0=No/NR
1164	1	1=Yes

<i>PUL48</i>		<i>Use of continuous nebulizers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1671	0	0=No/NR
1344	1	1=Yes

<i>PUL49</i>		<i>Magnesium injections in ED</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2840	0	0=No/NR
175	1	1=Yes

<i>PUL50</i>		<i>Oxygen</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1659	0	0=No/NR
1356	1	1=Yes

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<i>PUL51</i>		<i>IV Antibiotics</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1927	0	0=No/NR
1088	1	1=Yes

<i>PUL52</i>		<i>IV corticosteroid</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1855	0	0=No/NR
1160	1	1=Yes

<i>PUL53</i>		<i>IV lasix or furosemide</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2602	0	0=No/NR
413	1	1=Yes

<i>PUL54A</i>		<i>Antibiotics-oral onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2375	0	0=No/NR
640	1	1=Yes

<i>PUL54B</i>		<i>Antibiotics-oral discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2242	0	0=No/NR
773	1	1=Yes

<i>PUL55A</i>		<i>Systemic corticosteroid onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2051	0	0=No/NR
964	1	1=Yes

<i>PUL55B</i>		<i>Systemic corticosteroid discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2225	0	0=No/NR
790	1	1=Yes

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<i>PUL56A</i>		<i>short acting beta-agonists onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1341	0	0=No/NR
1674	1	1=Yes

<i>PUL56B</i>		<i>short acting beta-agonists discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1838	0	0=No/NR
1177	1	1=Yes

<i>PUL57A</i>		<i>long-acting beta-agonist onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2813	0	0=No/NR
202	1	1=Yes

<i>PUL57B</i>		<i>long-acting beta-agonist discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2847	0	0=No/NR
168	1	1=Yes

<i>PUL58A</i>		<i>short-acting anticholinergics onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1998	0	0=No/NR
1017	1	1=Yes

<i>PUL58B</i>		<i>short-acting anticholinergics discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2727	0	0=No/NR
288	1	1=Yes

<i>PUL59A</i>		<i>long-acting anticholinergics onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2850	0	0=No/NR
165	1	1=Yes

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<i>PUL59B</i>		<i>long-acting anticholinergics discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2862	0	0=No/NR
153	1	1=Yes

<i>PUL60A</i>		<i>Inhaled corticosteroids onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2374	0	0=No/NR
641	1	1=Yes

<i>PUL60B</i>		<i>Inhaled corticosteroids discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2503	0	0=No/NR
512	1	1=Yes

<i>PUL61A</i>		<i>Nebulized bronchodilators onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1791	0	0=No/NR
1224	1	1=Yes

<i>PUL61B</i>		<i>Nebulized bronchodilators discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2613	0	0=No/NR
402	1	1=Yes

<i>PUL62A</i>		<i>Leukotriene antagonist onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2562	0	0=No/NR
453	1	1=Yes

<i>PUL62B</i>		<i>Leukotriene antagonist discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2646	0	0=No/NR
369	1	1=Yes

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<i>PUL63A</i>		<i>Home oxygen onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2829	0	0=No/NR
186	1	1=Yes

<i>PUL63B</i>		<i>Home oxygen discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2810	0	0=No/NR
205	1	1=Yes

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Pulmonary Reviewer 1 Form: FORM CODE=PLD1, VERSION=A

ID		HCHS Participant ID
N	Value	Description
2949	Present	Text suppressed

VISIT		Visit
N	Value	Description
255	1	Visit 1
280	2	Visit 2
300	3	Visit 3
268	4	Visit 4
303	5	Visit 5
208	6	Visit 6
201	7	Visit 7
210	8	Visit 8
163	9	Visit 9
182	10	Visit 10
260	11	Visit 11
147	12	Visit 12
113	13	Visit 13
53	14	Visit 14
6	15	Visit 15

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<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2039	1	Occurrence 1
499	2	Occurrence 2
185	3	Occurrence 3
84	4	Occurrence 4
51	5	Occurrence 5
30	6	Occurrence 6
16	7	Occurrence 7
14	8	Occurrence 8
8	9	Occurrence 9
8	10	Occurrence 10
3	11	Occurrence 11
3	12	Occurrence 12
1	13	Occurrence 13
1	14	Occurrence 14
1	15	Occurrence 15
1	16	Occurrence 16
2	17	Occurrence 17
2	18	Occurrence 18
1	19	Occurrence 19

<i>PLD10A_FUTIME</i>		<i>Follow-up time from baseline for PLD10A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2747	Range	1289 - 6157 (median=3590 mean=3850.4 std=1148.8)

<i>PLD10AY</i>		<i>Year of PLD10A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2747	Range	2013 - 2025 (median=2019 mean=2020.1 std=3.2)

<i>PLD10C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2949	Present	Text suppressed

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<i>PLD10D_FUTIME</i>		<i>Follow-up time from baseline for PLD10D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2948	Range	1 - 5556 (median=2042 mean=2200.1 std=1367.4)

<i>PLD10DY</i>		<i>Year of PLD10D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
2948	Range	2008 - 2024 (median=2015 mean=2015.6 std=3.8)

<i>PLD11</i>		<i>Meet SOL criteria for CLRD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
203		Missing
231	1	1= Definite
56	2	2= Highly Probable
1532	3	3= Probable
927	4	4= NO/unknown

<i>PLD12A</i>		<i>Asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
928	.S	Legitimately skipped
68	1	1= Definite
174	2	2= Highly Probable
1088	3	3= Probable
7	4	4= Definitely Not
482	5	5= Probably not/Unknown

<i>PLD12B</i>		<i>COPD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
928	.S	Legitimately skipped
18	1	1= Definite
25	2	2= Highly Probable
705	3	3= Probable
10	4	4= Definitely Not
1061	5	5= Probably not/Unknown

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<i>PLD12C</i>		<i>Emphysema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
928	.S	Legitimately skipped
81	1	1= Definite
2	2	2= Highly Probable
69	3	3= Probable
56	4	4= Definitely Not
1611	5	5= Probably not/Unknown

<i>PLD12D</i>		<i>Chronic Bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
928	.S	Legitimately skipped
0	1	1= Definite
0	2	2= Highly Probable
15	3	3= Probable
9	4	4= Definitely Not
1795	5	5= Probably not/Unknown

<i>PLD13</i>		<i>Predominant CLRD subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
928	.S	Legitimately skipped
1136	1	1= Asthma
582	2	2= COPD
67	3	3= Emphysema
3	4	4= Chronic Bronchitis
31	5	5= Unsure

<i>PLD14</i>		<i>Evidence of other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
928	.S	Legitimately skipped
1685	0	0=No
134	1	1=Yes

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<i>PLD14A</i>		<i>Specify other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2613	~	Legitimately skipped
134	Present	Text suppressed

<i>PLD15</i>		<i>Exacerbation of CLRD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
928	.S	Legitimately skipped
234	1	1= Definite
564	2	2= Highly Probable
242	3	3= Probable
779	4	4= No/Unknown

<i>PLD15A</i>		<i>Evidence for another diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
1707	.S	Legitimately skipped
771	1	1= No, "LONE" CLRD exacerbation
269	2	2= Yes, "COMORBID" CLRD

<i>PLD15B1</i>		<i>Pneumonia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2478	.S	Legitimately skipped
158	1	1=Yes
111	2	2=No/unknown

<i>PLD15B2</i>		<i>Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2478	.S	Legitimately skipped
4	1	1=Yes
265	2	2=No/unknown

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<i>PLD15B3</i>		<i>Pneumothorax</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2478	.S	Legitimately skipped
4	1	1=Yes
265	2	2=No/unknown

<i>PLD15B4</i>		<i>Acute myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2478	.S	Legitimately skipped
11	1	1=Yes
258	2	2=No/unknown

<i>PLD15B5</i>		<i>Acute decompensated heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2478	.S	Legitimately skipped
87	1	1=Yes
182	2	2=No/unknown

<i>PLD15B5A</i>		<i>Type of ADHF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2660	.S	Legitimately skipped
52	1	1= Left-sided
7	2	2= Right-sided
28	3	3= Unknown

<i>PLD15B6</i>		<i>Other condition</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2478	.S	Legitimately skipped
39	1	1=Yes
230	2	2=No/unknown

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<i>PLD15B6A</i>		<i>Specify other condition</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
202		Missing
2708	~	Legitimately skipped
39	Present	Text suppressed

<i>PLD16</i>		<i>Comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1737		Missing
928	~	Legitimately skipped
284	Present	Text suppressed

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Pulmonary Reviewer 2 Form: FORM CODE=PLD2, VERSION=A

ID		HCHS Participant ID
N	Value	Description
2945	Present	Text suppressed

VISIT		Visit
N	Value	Description
253	1	Visit 1
278	2	Visit 2
300	3	Visit 3
268	4	Visit 4
303	5	Visit 5
208	6	Visit 6
201	7	Visit 7
210	8	Visit 8
163	9	Visit 9
182	10	Visit 10
260	11	Visit 11
147	12	Visit 12
113	13	Visit 13
53	14	Visit 14
6	15	Visit 15

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OCCURRENCE		Occurrence
N	Value	Description
2037	1	Occurrence 1
497	2	Occurrence 2
185	3	Occurrence 3
84	4	Occurrence 4
51	5	Occurrence 5
30	6	Occurrence 6
16	7	Occurrence 7
14	8	Occurrence 8
8	9	Occurrence 9
8	10	Occurrence 10
3	11	Occurrence 11
3	12	Occurrence 12
1	13	Occurrence 13
1	14	Occurrence 14
1	15	Occurrence 15
1	16	Occurrence 16
2	17	Occurrence 17
2	18	Occurrence 18
1	19	Occurrence 19

PLD20A_FUTIME		Follow-up time from baseline for PLD20A (Completion Date)
N	Value	Description
153		Missing
2792	Range	420 - 6207 (median=3620.5 mean=3871.45 std=1178.34)

PLD20AY		Year of PLD20A
N	Value	Description
153		Missing
2792	Range	2009 - 2025 (median=2019 mean=2020.2 std=3.3)

PLD20C		Event ID
N	Value	Description
2945	Present	Text suppressed

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<i>PLD20D_FUTIME</i>		<i>Follow-up time from baseline for PLD20D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2945	Range	1 - 5556 (median=2046 mean=2201.5 std=1367.2)

<i>PLD20DY</i>		<i>Year of PLD20D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2945	Range	2008 - 2024 (median=2015 mean=2015.6 std=3.8)

<i>PLD21</i>		<i>Meet SOL criteria for CLRD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
199	1	1= Definite
61	2	2= Highly Probable
1577	3	3= Probable
955	4	4= NO/unknown

<i>PLD22A</i>		<i>Asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
955	.S	Legitimately skipped
65	1	1= Definite
164	2	2= Highly Probable
1122	3	3= Probable
6	4	4= Definitely Not
480	5	5= Probably not/Unknown

<i>PLD22B</i>		<i>COPD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
955	.S	Legitimately skipped
11	1	1= Definite
15	2	2= Highly Probable
726	3	3= Probable
5	4	4= Definitely Not
1080	5	5= Probably not/Unknown

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<i>PLD22C</i>		<i>Emphysema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
955	.S	Legitimately skipped
66	1	1= Definite
4	2	2= Highly Probable
74	3	3= Probable
61	4	4= Definitely Not
1632	5	5= Probably not/Unknown

<i>PLD22D</i>		<i>Chronic Bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
955	.S	Legitimately skipped
0	1	1= Definite
2	2	2= Highly Probable
11	3	3= Probable
10	4	4= Definitely Not
1814	5	5= Probably not/Unknown

<i>PLD23</i>		<i>Predominant CLRD subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
955	.S	Legitimately skipped
1144	1	1= Asthma
581	2	2= COPD
57	3	3= Emphysema
7	4	4= Chronic Bronchitis
48	5	5= Unsure

<i>PLD24</i>		<i>Evidence of other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
955	.S	Legitimately skipped
1676	0	0=No
161	1	1=Yes

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<i>PLD24A</i>		<i>Specify other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2631	~	Legitimately skipped
161	Present	Text suppressed

<i>PLD25</i>		<i>Exacerbation of CLRD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
154		Missing
955	.S	Legitimately skipped
219	1	1= Definite
596	2	2= Highly Probable
237	3	3= Probable
784	4	4= No/Unknown

<i>PLD25A</i>		<i>Evidence for another diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
1740	.S	Legitimately skipped
776	1	1= No, "LONE" CLRD exacerbation
276	2	2= Yes, "COMORBID" CLRD

<i>PLD25B1</i>		<i>Pneumonia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2516	.S	Legitimately skipped
169	1	1=Yes
107	2	2=No/unknown

<i>PLD25B2</i>		<i>Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2516	.S	Legitimately skipped
3	1	1=Yes
273	2	2=No/unknown

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<i>PLD25B3</i>		<i>Pneumothorax</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2516	.S	Legitimately skipped
5	1	1=Yes
271	2	2=No/unknown

<i>PLD25B4</i>		<i>Acute myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2516	.S	Legitimately skipped
4	1	1=Yes
272	2	2=No/unknown

<i>PLD25B5</i>		<i>Acute decompensated heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2516	.S	Legitimately skipped
78	1	1=Yes
198	2	2=No/unknown

<i>PLD25B5A</i>		<i>Type of ADHF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2714	.S	Legitimately skipped
47	1	1= Left-sided
8	2	2= Right-sided
23	3	3= Unknown

<i>PLD25B6</i>		<i>Other condition</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2516	.S	Legitimately skipped
58	1	1=Yes
218	2	2=No/unknown

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<i>PLD25B6A</i>		<i>Specify other condition</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
153		Missing
2734	~	Legitimately skipped
58	Present	Text suppressed

<i>PLD26</i>		<i>Comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1734		Missing
955	~	Legitimately skipped
256	Present	Text suppressed

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Pulmonary Adjudication Reviewer Form: FORM CODE=PLD3, VERSION=A

ID		HCHS Participant ID
N	Value	Description
666	Present	Text suppressed

VISIT		Visit
N	Value	Description
81	1	Visit 1
80	2	Visit 2
77	3	Visit 3
60	4	Visit 4
61	5	Visit 5
55	6	Visit 6
55	7	Visit 7
60	8	Visit 8
48	9	Visit 9
40	10	Visit 10
35	11	Visit 11
11	12	Visit 12
3	13	Visit 13

OCCURRENCE		Occurrence
N	Value	Description
439	1	Occurrence 1
131	2	Occurrence 2
51	3	Occurrence 3
15	4	Occurrence 4
13	5	Occurrence 5
6	6	Occurrence 6
4	8	Occurrence 8
3	9	Occurrence 9
3	10	Occurrence 10
1	11	Occurrence 11

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<i>PLD30A_FUTIME</i>		<i>Follow-up time from baseline for PLD30A (Completion Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
556	Range	328 - 6147 (median=3762 mean=3927.4 std=968.7)

<i>PLD30AY</i>		<i>Year of PLD30A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
556	Range	2009 - 2025 (median=2019 mean=2020.1 std=2.8)

<i>PLD30C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
666	Present	Text suppressed

<i>PLD30D_FUTIME</i>		<i>Follow-up time from baseline for PLD30D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
666	Range	1 - 4792 (median=1777 mean=1827.0 std=1165.6)

<i>PLD30DY</i>		<i>Year of PLD30D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
666	Range	2008 - 2021 (median=2014 mean=2014.6 std=3.3)

<i>PLD31</i>		<i>Meet SOL criteria for CLRD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
117	1	1= Definite
36	2	2= Highly Probable
355	3	3= Probable
48	4	4= NO/unknown

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<i>PLD32A</i>		<i>Asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
48	.S	Legitimately skipped
30	1	1= Definite
71	2	2= Highly Probable
269	3	3= Probable
0	4	4= Definitely Not
138	5	5= Probably not/Unknown

<i>PLD32B</i>		<i>COPD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
48	.S	Legitimately skipped
10	1	1= Definite
13	2	2= Highly Probable
181	3	3= Probable
2	4	4= Definitely Not
302	5	5= Probably not/Unknown

<i>PLD32C</i>		<i>Emphysema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
48	.S	Legitimately skipped
35	1	1= Definite
2	2	2= Highly Probable
20	3	3= Probable
5	4	4= Definitely Not
446	5	5= Probably not/Unknown

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<i>PLD32D</i>		<i>Chronic Bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
48	.S	Legitimately skipped
0	1	1= Definite
0	2	2= Highly Probable
7	3	3= Probable
3	4	4= Definitely Not
498	5	5= Probably not/Unknown

<i>PLD33</i>		<i>Predominant CLRD subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
48	.S	Legitimately skipped
312	1	1= Asthma
156	2	2= COPD
23	3	3= Emphysema
2	4	4= Chronic Bronchitis
15	5	5= Unsure

<i>PLD34</i>		<i>Evidence of other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
48	.S	Legitimately skipped
463	0	0=No
45	1	1=Yes

<i>PLD34A</i>		<i>Specify other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
511	~	Legitimately skipped
45	Present	Text suppressed

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<i>PLD35</i>		<i>Exacerbation of CLRD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
48	.S	Legitimately skipped
112	1	1= Definite
152	2	2= Highly Probable
82	3	3= Probable
162	4	4= No/Unknown

<i>PLD35A</i>		<i>Evidence for another diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
210	.S	Legitimately skipped
240	1	1= No, "LONE" CLRD exacerbation
106	2	2= Yes, "COMORBID" CLRD

<i>PLD35B1</i>		<i>Pneumonia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
450	.S	Legitimately skipped
62	1	1=Yes
44	2	2=No/unknown

<i>PLD35B2</i>		<i>Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
450	.S	Legitimately skipped
2	1	1=Yes
104	2	2=No/unknown

<i>PLD35B3</i>		<i>Pneumothorax</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
450	.S	Legitimately skipped
4	1	1=Yes
102	2	2=No/unknown

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<i>PLD35B4</i>		<i>Acute myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
450	.S	Legitimately skipped
3	1	1=Yes
103	2	2=No/unknown

<i>PLD35B5</i>		<i>Acute decompensated heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
450	.S	Legitimately skipped
27	1	1=Yes
79	2	2=No/unknown

<i>PLD35B5A</i>		<i>Type of ADHF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
529	.S	Legitimately skipped
17	1	1= Left-sided
3	2	2= Right-sided
7	3	3= Unknown

<i>PLD35B6</i>		<i>Other condition</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
450	.S	Legitimately skipped
23	1	1=Yes
83	2	2=No/unknown

<i>PLD35B6A</i>		<i>Specify other condition</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
533	~	Legitimately skipped
23	Present	Text suppressed

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<i>PLD36</i>		<i>Comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
461		Missing
48	~	Legitimately skipped
157	Present	Text suppressed

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Section E: Stroke Ascertainment

Stroke Abstraction Form: FORM CODE=STR, VERSION=1

<i>ID</i>		<i>HCHS Participant ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Present	Text suppressed

<i>VISIT</i>		<i>Visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
50	1	Visit 1
57	2	Visit 2
55	3	Visit 3
54	4	Visit 4
43	5	Visit 5
36	6	Visit 6
43	7	Visit 7
44	8	Visit 8
44	9	Visit 9
25	10	Visit 10
55	11	Visit 11
25	12	Visit 12
18	13	Visit 13
10	14	Visit 14
2	15	Visit 15

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
412	1	Occurrence 1
100	2	Occurrence 2
26	3	Occurrence 3
7	4	Occurrence 4
10	5	Occurrence 5
3	6	Occurrence 6
2	7	Occurrence 7
1	10	Occurrence 10

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<i>STR0A_FUTIME</i>		<i>Follow-up time from baseline for STR0A (Completion date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Range	3330 - 5912 (median=4155 mean=4206.4 std=544.4)

<i>STR0AY</i>		<i>Year of STR0A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Range	2020 - 2025 (median=2020 mean=2021.0 std=1.3)

<i>STR0C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Present	Text suppressed

<i>STR0D_FUTIME</i>		<i>Follow-up time from baseline for STR0D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Range	14 - 5662 (median=2070 mean=2182.0 std=1379.9)

<i>STR0DY</i>		<i>Year of STR0D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Range	2008 - 2024 (median=2015 mean=2015.4 std=3.9)

<i>STR1_FUTIME</i>		<i>Follow-up time from baseline for STR1 (Date of arrival)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Range	14 - 5662 (median=2060 mean=2181.2 std=1379.9)

<i>STR1A</i>		<i>Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15		Missing
546	Range	0:00:00 - 23:59:00

<i>STR1Y</i>		<i>Year of STR1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
561	Range	2008 - 2024 (median=2015 mean=2015.4 std=3.9)

<i>STR2</i>		<i>Primary admitting code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
80		Missing
481	Present	Text suppressed

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<i>STR3_FUTIME</i>		<i>Follow-up time from baseline for STR3 (Date of discharge)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
560	Range	21 - 5666 (median=2113 mean=2198.4 std=1383.8)

<i>STR3Y</i>		<i>Year of STR3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
560	Range	2008 - 2024 (median=2015 mean=2015.5 std=3.9)

<i>STR4</i>		<i>Primary discharge code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
64		Missing
497	Present	Text suppressed

<i>STR5</i>		<i>Arrival mode</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
232	1	1=EMS
133	2	2=Private transport/walk-in
11	3	3=Transfer from another hospital
0	4	4=Mobile stroke unit
185	9	9=Unknown

<i>STR6</i>		<i>Transferred from this hospital to another</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
544	0	0=No
14	1	1=Yes
3	9	9=Unknown

<i>STR7</i>		<i>DNR/DNI present</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
201	0	0=No
48	1	1=Yes
312	9	9=Unknown

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STR8		<i>New neurological signs/symptoms</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
114	0	0=No
436	1	1=Yes
11	9	9=Unknown

STR9		<i>Stroke or TIA this hospitalization</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
531	0	0=No/NR
30	1	1=Yes

STR10		<i>event in setting of a procedure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
532	0	0=No/NR
29	1	1=Yes

STR10A1		<i>cerebral angiogram</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
532	.S	Legitimately skipped
26	0	0=No/NR
3	1	1=Yes

STR10A2		<i>coronary angiogram</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
532	.S	Legitimately skipped
24	0	0=No/NR
5	1	1=Yes

STR10A3		<i>cardiac surgery or procedures</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
532	.S	Legitimately skipped
24	0	0=No/NR
5	1	1=Yes

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<i>STR10A4</i>		<i>carotid endarterectomy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
532	.S	Legitimately skipped
29	0	0=No/NR
0	1	1=Yes

<i>STR10A5</i>		<i>carotid stenting</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
532	.S	Legitimately skipped
29	0	0=No/NR
0	1	1=Yes

<i>STR10A6</i>		<i>other procedure performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
532	.S	Legitimately skipped
8	0	0=No/NR
21	1	1=Yes

<i>STR10A6A</i>		<i>other procedure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
540	~	Legitimately skipped
21	Present	Text suppressed

<i>STR11A</i>		<i>symptom onset date known</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
67		Missing
89	0	0=No
405	1	1=Yes

<i>STR11B_FUTIME</i>		<i>Follow-up time from baseline for STR11B (symptom onset date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
157		Missing
404	Range	25 - 5662 (median=2058 mean=2154.9 std=1397.0)

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<i>STR11BY</i>		<i>Year of STR11B</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
156		Missing
405	Range	2008 - 2024 (median=2015 mean=2015.4 std=3.9)

<i>STR11C</i>		<i>symptom onset time known</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
156	.S	Legitimately skipped
229	0	0=No
175	1	1=Yes

<i>STR11D</i>		<i>symptom onset time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
386	.S	Legitimately skipped
175	Range	0:33:00 - 23:40:00

<i>STR12A</i>		<i>free of deficits date known</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
67		Missing
88	0	0=No
406	1	1=Yes

<i>STR12B_FUTIME</i>		<i>Follow-up time from baseline for STR12B (free of deficits date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
156		Missing
405	Range	25 - 5662 (median=2070 mean=2168.7 std=1396.0)

<i>STR12BY</i>		<i>Year of STR12B</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
155		Missing
406	Range	2008 - 2024 (median=2015 mean=2015.4 std=3.9)

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<i>STR12C</i>		<i>free of deficits time known</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
155	.S	Legitimately skipped
220	0	0=No
185	1	1=Yes

<i>STR12D</i>		<i>free of deficits time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
376	.S	Legitimately skipped
185	Range	0:33:00 - 23:40:00

<i>STR13</i>		<i>history of stroke</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
264	0	0=No
243	1	1=Yes
54	9	9=Unknown

<i>STR13A</i>		<i>previous stroke date known</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
318	.S	Legitimately skipped
111	0	0=No
131	1	1=Yes

<i>STR13B_FUTIME</i>		<i>Follow-up time from baseline for STR13B (date of most recent stroke)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
524		Missing
37	Range	-539 - 4774 (median=1015 mean=1432.0 std=1284.8)

<i>STR13BY</i>		<i>Year of STR13B</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
431		Missing
130	Range	1990 - 2022 (median=2012 mean=2011.1 std=6.4)

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<i>STR13C</i>		<i>Previous stroke type</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
318	.S	Legitimately skipped
193	1	1=Ischemic
9	2	2=Intracranial hemorrhage
8	3	3=Subarachnoid hemorrhage
33	9	9=Unknown

<i>STR13C1</i>		<i>history of hematoma evacuation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552	.S	Legitimately skipped
0	0	0=No
1	1	1=Yes
8	9	9=Unknown

<i>STR13C2</i>		<i>history of intra-hematoma tPA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
552	.S	Legitimately skipped
0	0	0=No
0	1	1=Yes
9	9	9=Unknown

<i>STR13C3</i>		<i>history of aneurysm</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
553	.S	Legitimately skipped
1	0	0=No
4	1	1=Yes
3	9	9=Unknown

<i>STR13C4</i>		<i>history of aneurysm coiling</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
553	.S	Legitimately skipped
2	0	0=No
0	1	1=Yes
6	9	9=Unknown

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<i>STR13C5</i>		<i>history of aneurysm clipping</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
553	.S	Legitimately skipped
0	0	0=No
3	1	1=Yes
5	9	9=Unknown

<i>STR14</i>		<i>history of TIA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
328	0	0=No
57	1	1=Yes
176	9	9=Unknown

<i>STR14A</i>		<i>previous TIA date known</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
504	.S	Legitimately skipped
43	0	0=No
14	1	1=Yes

<i>STR14B_FUTIME</i>		<i>Follow-up time from baseline for STR14B (previous TIA date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
556		Missing
5	Range	644 - 4543 (median=3480 mean=2705.0 std=1693.7)

<i>STR14BY</i>		<i>Year of STR14B</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
547		Missing
14	Range	2004 - 2023 (median=2013 mean=2013.5 std=5.3)

<i>STR15</i>		<i>history of CAD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
352	0	0=No
160	1	1=Yes
49	9	9=Unknown

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STR16		history of PVD
N	Value	Description
465	0	0=No
34	1	1=Yes
62	9	9=Unknown

STR17		history of carotid disease
N	Value	Description
452	0	0=No
43	1	1=Yes
66	9	9=Unknown

STR18		history of ESRD
N	Value	Description
473	0	0=No
29	1	1=Yes
59	9	9=Unknown

STR19		history of diabetes
N	Value	Description
211	0	0=No
311	1	1=Yes
39	9	9=Unknown

STR20		history of HTN
N	Value	Description
68	0	0=No
467	1	1=Yes
26	9	9=Unknown

STR21		history of dyslipidemia
N	Value	Description
165	0	0=No
317	1	1=Yes
79	9	9=Unknown

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STR22		<i>history of smoking</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
294	0	0=No
192	1	1=Yes
75	9	9=Unknown

STR23		<i>history of alcohol use disorder</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
393	0	0=No
65	1	1=Yes
103	9	9=Unknown

STR24		<i>history of illicit stimulant use four weeks prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
377	0	0=No
29	1	1=Yes
155	9	9=Unknown

STR25A		<i>myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
546	0	0=No/NR
15	1	1=Yes

STR25B		<i>intracardiac thrombus or intracardiac tumor</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
558	0	0=No/NR
3	1	1=Yes

STR25C		<i>atrial fibrillation/atrial flutter</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
503	0	0=No/NR
58	1	1=Yes

STR25C1		<i>24 hours of cardiac monitoring</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
277	0	0=No/NR
284	1	1=Yes

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<i>STR25D</i>		<i>left-sided atrial enlargement on echocardiogram</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
459	0	0=No/NR
102	1	1=Yes

<i>STR25E</i>		<i>rheumatic heart disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
556	0	0=No/NR
5	1	1=Yes

<i>STR25F</i>		<i>systemic embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
559	0	0=No/NR
2	1	1=Yes

<i>STR25G</i>		<i>hematologic abnormality (hypercoagulable state)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
555	0	0=No/NR
6	1	1=Yes

<i>STR25H</i>		<i>hematologic abnormality (hemorrhagic)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
555	0	0=No/NR
6	1	1=Yes

<i>STR25I</i>		<i>migraine headaches</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	0	0=No/NR
31	1	1=Yes

<i>STR25J</i>		<i>left-sided bioprosthetic valve</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
551	0	0=No/NR
10	1	1=Yes

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<i>STR25K</i>		<i>left-sided mechanical valve</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
553	0	0=No/NR
8	1	1=Yes

<i>STR26</i>		<i>time between onset and resolution of new neuro symptoms</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82	1	1=Less than 24 hours
78	2	2=24-48 hours
52	3	3=Greater than 48 hours
228	4	4=Unknown
121	5	5=Not applicable

<i>STR27</i>		<i>NIHSS recorded</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
427	0	0=No
133	1	1=Yes

<i>STR27A</i>		<i>NIHSS score</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
133	Range	0 - 27 (median=3 mean=5.2 std=6.2)

<i>STR27B1</i>		<i>LOCa</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
380	0	0
18	1	1
9	2	2
20	3	3

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<i>STR27B10</i>		<i>Sensory</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
354	0	0
67	1	1
6	2	2

<i>STR27B11</i>		<i>Best Language</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
370	0	0
34	1	1
10	2	2
13	3	3

<i>STR27B12</i>		<i>Dysarthria</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
346	0	0
73	1	1
8	2	2

<i>STR27B13</i>		<i>Extinction and Inattention</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
415	0	0
3	1	1
9	2	2

<i>STR27B14</i>		<i>Total Score</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
427	Range	0 - 38 (median=2 mean=3.2 std=5.2)

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<i>STR27B2</i>		<i>LOCb</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
340	0	0
50	1	1
37	2	2

<i>STR27B3</i>		<i>LOCc</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
369	0	0
22	1	1
36	2	2

<i>STR27B4</i>		<i>Best Gaze</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
396	0	0
26	1	1
5	2	2

<i>STR27B5</i>		<i>Visual Fields</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
400	0	0
12	1	1
10	2	2
5	3	3

<i>STR27B6</i>		<i>Facial Palsy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
348	0	0
74	1	1
2	2	2
3	3	3

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STR27B7A		Motor Left Arm
N	Value	Description
134	.S	Legitimately skipped
345	0	0
52	1	1
5	2	2
3	3	3
22	4	4

STR27B7B		Motor Right Arm
N	Value	Description
134	.S	Legitimately skipped
346	0	0
55	1	1
6	2	2
4	3	3
16	4	4

STR27B8A		Motor Left Leg
N	Value	Description
134	.S	Legitimately skipped
350	0	0
45	1	1
7	2	2
6	3	3
19	4	4

STR27B8B		Motor Right Leg
N	Value	Description
134	.S	Legitimately skipped
354	0	0
49	1	1
6	2	2
4	3	3
14	4	4

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<i>STR27B9</i>		<i>Limb Ataxia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
134	.S	Legitimately skipped
401	0	0
19	1	1
7	2	2

<i>STR28</i>		<i>asleep at time of event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
237	0	0=No
80	1	1=Yes
243	9	9=Unknown

<i>STR29</i>		<i>severe headache at onset or admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
264	0	0=No
56	1	1=Yes
241	9	9=Unknown

<i>STR30</i>		<i>Vomiting</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
245	0	0=No
86	1	1=Yes
230	9	9=Unknown

<i>STR31</i>		<i>Blurry Vision</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
159	0	0=No
49	1	1=Yes
353	9	9=Unknown

<i>STR32</i>		<i>Diplopia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
150	0	0=No
16	1	1=Yes
395	9	9=Unknown

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<i>STR33</i>		<i>Vertigo</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
104	0	0=No
34	1	1=Yes
423	9	9=Unknown

<i>STR34</i>		<i>Dizziness</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
34	.S	Legitimately skipped
112	0	0=No
112	1	1=Yes
303	9	9=Unknown

<i>STR35</i>		<i>Seizure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
136	0	0=No
26	1	1=Yes
399	9	9=Unknown

<i>STR35A</i>		<i>Seizure as first neurologic symptom</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
535	.S	Legitimately skipped
5	0	0=No
11	1	1=Yes
10	9	9=Unknown

<i>STR36</i>		<i>Level of Consciousness</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
90	0	0=Normal
39	1	1=Abnormal
3	2	2=Unconscious
1	9	9=Unknown

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STR37		<i>Aphasia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
76	0	0=No
34	1	1=Yes
23	9	9=Unknown

STR38		<i>Hemianopsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
64	0	0=No
14	1	1=Yes
55	9	9=Unknown

STR39		<i>abnormality of cranial nerves</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
72	0	0=No
15	1	1=Yes
46	9	9=Unknown

STR39A		<i>location of abnormality</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
546	.S	Legitimately skipped
4	1	1=Left
7	2	2=Right
2	3	3=Both
2	9	9=NR

STR40		<i>Horizontal gaze palsy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
72	0	0=No
12	1	1=Yes
49	9	9=Unknown

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STR40A		Location of Gaze Deviation
N	Value	Description
549	.S	Legitimately skipped
4	1	1=to Left
7	2	2=to Right
0	3	3=Both
1	9	9=NR

STR41		Dysphagia
N	Value	Description
428	.S	Legitimately skipped
54	0	0=No
12	1	1=Yes
67	9	9=Unknown

STR42		Dysarthria
N	Value	Description
428	.S	Legitimately skipped
65	0	0=No
56	1	1=Yes
12	9	9=Unknown

STR43		weakness
N	Value	Description
428	.S	Legitimately skipped
31	0	0=No
98	1	1=Yes
4	9	9=Unknown

STR43A		face weakness
N	Value	Description
463	.S	Legitimately skipped
31	1	1=Left
24	2	2=Right
5	3	3=Both
17	4	4=none
21	9	9=NR

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<i>STR43B</i>		<i>arm weakness</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
463	.S	Legitimately skipped
39	1	1=Left
31	2	2=Right
15	3	3=Both
6	4	4=none
7	9	9=NR

<i>STR43C</i>		<i>leg weakness</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
463	.S	Legitimately skipped
36	1	1=Left
25	2	2=Right
13	3	3=Both
14	4	4=none
10	9	9=NR

<i>STR44</i>		<i>sensory deficit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
72	0	0=No
46	1	1=Yes
15	9	9=Unknown

<i>STR44A</i>		<i>sensory deficit face</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
515	.S	Legitimately skipped
16	1	1=Left
7	2	2=Right
2	3	3=Both
11	4	4=none
10	9	9=NR

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<i>STR44B</i>		<i>sensory deficit arm</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
515	.S	Legitimately skipped
22	1	1=Left
12	2	2=Right
1	3	3=Both
6	4	4=none
5	9	9=NR

<i>STR44C</i>		<i>sensory deficit leg</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
515	.S	Legitimately skipped
20	1	1=Left
10	2	2=Right
3	3	3=Both
7	4	4=none
6	9	9=NR

<i>STR45</i>		<i>visual fields abnormality</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
81	0	0=No
18	1	1=Yes
34	9	9=Unknown

<i>STR46</i>		<i>ataxic gait</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
58	0	0=No
12	1	1=Yes
63	9	9=Unknown

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<i>STR47</i>		<i>hemiataxia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
428	.S	Legitimately skipped
55	0	0=No
9	1	1=Yes
69	9	9=Unknown

<i>STR48A1</i>		<i>first systolic blood pressure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76		Missing
485	Range	64 - 270 (median=149 mean=151.0 std=31.8)

<i>STR48A2</i>		<i>first diastolic blood pressure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
78		Missing
483	Range	31 - 160 (median=78 mean=80.7 std=18.5)

<i>STR49</i>		<i>first glucose level</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76		Missing
485	Range	49 - 1221 (median=129 mean=164.3 std=108.2)

<i>STR50</i>		<i>lumbar puncture performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
547	0	0=No/NR
14	1	1=Yes

<i>STR50A</i>		<i>lumbar puncture diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
547	.S	Legitimately skipped
5	0	0=Normal
2	1	1=Abnormal
7	9	9=Unknown

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<i>STR51</i>		<i>CT scan of head performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
129	0	0=No
298	1	1=Yes, 1 CT
130	2	2=Yes, 2 or more CTs
4	9	9=NR

<i>STR51A_FUTIME</i>		<i>Follow-up time from baseline for STR51A (date of first CT)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
147		Missing
414	Range	36 - 5662 (median=2211.5 mean=2236.15 std=1401.28)

<i>STR51AY</i>		<i>Year of STR51A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
142		Missing
419	Range	2008 - 2024 (median=2015 mean=2015.6 std=4.0)

<i>STR51B</i>		<i>First CT acute or subacute stroke diagnosed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
133	.S	Legitimately skipped
290	0	0=No
127	1	1=Yes
11	9	9=Unknown/NR

<i>STR51C_FUTIME</i>		<i>Follow-up time from baseline for STR51C (date of last CT)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
440		Missing
121	Range	52 - 5056 (median=1707 mean=2039.7 std=1398.1)

<i>STR51CY</i>		<i>Year of STR51C</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
434		Missing
127	Range	2009 - 2024 (median=2014 mean=2015.1 std=4.0)

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<i>STR51D</i>		<i>Last CT acute or subacute stroke or previously unseen stroke diagnosed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
431	.S	Legitimately skipped
58	0	0=No
62	1	1=Yes
10	9	9=Unknown/NR

<i>STR52</i>		<i>MRI including head performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
246	0	0=No
278	1	1=Yes
37	9	9=Unknown/NR

<i>STR52A</i>		<i>Acute or subacute stroke diagnosed on MRI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
283	.S	Legitimately skipped
133	0	0=No
132	1	1=Yes
13	9	9=Unknown/NR

<i>STR52B</i>		<i>DWI performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
283	.S	Legitimately skipped
19	0	0=No
189	1	1=Yes
70	9	9=Unknown/NR

<i>STR52C</i>		<i>DWI diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
372	.S	Legitimately skipped
84	0	0=Normal
94	1	1=Abnormal
11	9	9=Unknown/NR

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<i>STR53</i>		<i>cerebrovascular angiography performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
207	0	0=No
327	1	1=Yes
27	9	9=NR

<i>STR53A_FUTIME</i>		<i>Follow-up time from baseline for STR53A (cerebrovascular angiography date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
263		Missing
298	Range	52 - 5662 (median=2228.5 mean=2224.84 std=1371.98)

<i>STR53AY</i>		<i>Year of STR53A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
261		Missing
300	Range	2008 - 2024 (median=2015 mean=2015.5 std=4.0)

<i>STR53B</i>		<i>CTA head</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
168	0	0=No
137	1	1=Yes
22	9	9=NR

<i>STR53C</i>		<i>CTA neck</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
178	0	0=No
122	1	1=Yes
27	9	9=NR

<i>STR53D</i>		<i>MRA head</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
173	0	0=No
128	1	1=Yes
26	9	9=NR

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<i>STR53E</i>		<i>MRA neck</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
241	0	0=No
52	1	1=Yes
34	9	9=NR

<i>STR53F</i>		<i>Carotid doppler</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
147	0	0=No
160	1	1=Yes
20	9	9=NR

<i>STR53G</i>		<i>Transcranial Doppler</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
286	0	0=No
9	1	1=Yes
32	9	9=NR

<i>STR53H</i>		<i>Catheter Angiogram</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
282	0	0=No
10	1	1=Yes
35	9	9=NR

<i>STR53I</i>		<i>Stenosis - Right Internal Carotid Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
196	0	0=No
46	1	1=Yes
85	9	9=NR

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<i>STR53J</i>		<i>Stenosis - Left Internal Carotid Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
189	0	0=No
49	1	1=Yes
89	9	9=NR

<i>STR53K</i>		<i>Stenosis - Right Vertebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
150	0	0=No
5	1	1=Yes
172	9	9=NR

<i>STR53L</i>		<i>Stenosis - Left Vertebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
153	0	0=No
3	1	1=Yes
171	9	9=NR

<i>STR53M</i>		<i>Stenosis - Right Internal Carotid Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
170	0	0=No
36	1	1=Yes
121	9	9=NR

<i>STR53N</i>		<i>Stenosis - Left Internal Carotid Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
171	0	0=No
35	1	1=Yes
121	9	9=NR

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<i>STR53O</i>		<i>Stenosis - Right Vertebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
171	0	0=No
10	1	1=Yes
146	9	9=NR

<i>STR53P</i>		<i>Stenosis - Left Vertebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
174	0	0=No
5	1	1=Yes
148	9	9=NR

<i>STR53Q</i>		<i>Stenosis - Right Middle Cerebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
165	0	0=No
15	1	1=Yes
147	9	9=NR

<i>STR53R</i>		<i>Stenosis - Left Middle Cerebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
168	0	0=No
18	1	1=Yes
141	9	9=NR

<i>STR53S</i>		<i>Stenosis - Right Anterior Cerebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
175	0	0=No
4	1	1=Yes
148	9	9=NR

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<i>STR53T</i>		<i>Stenosis - Left Anterior Cerebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
178	0	0=No
6	1	1=Yes
143	9	9=NR

<i>STR53U</i>		<i>Stenosis - Right Posterior Cerebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
170	0	0=No
11	1	1=Yes
146	9	9=NR

<i>STR53V</i>		<i>Stenosis - Left Posterior Cerebral Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
172	0	0=No
8	1	1=Yes
147	9	9=NR

<i>STR53W</i>		<i>Stenosis - Basilar Artery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
234	.S	Legitimately skipped
167	0	0=No
9	1	1=Yes
151	9	9=NR

<i>STR54</i>		<i>transthoracic echocardiography study performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
279	0	0=No/NR
282	1	1=Yes

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STR54A		<i>transthoracic ejection fraction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
31		Missing
279	.S	Legitimately skipped
251	Range	20 - 84 (median=56 mean=56.4 std=12.5)

STR55		<i>transesophageal echocardiography study performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
531	0	0=No/NR
30	1	1=Yes

STR55A		<i>transesophageal ejection fraction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13		Missing
531	.S	Legitimately skipped
17	Range	20 - 70 (median=55 mean=50.6 std=15.7)

STR56A		<i>intracardiac thrombus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
84	0	0=No
3	1	1=Yes
205	9	9=NR

STR56B		<i>mitral stenosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
87	0	0=No
3	1	1=Yes
202	9	9=NR

STR56C		<i>mitral regurgitation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
96	0	0=No
126	1	1=Yes
70	9	9=NR

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<i>STR56D</i>		<i>aortic stenosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
121	0	0=No
13	1	1=Yes
158	9	9=NR

<i>STR56E</i>		<i>aortic regurgitation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
130	0	0=No
54	1	1=Yes
108	9	9=NR

<i>STR56F</i>		<i>poor ventricular function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
146	0	0=No
78	1	1=Yes
68	9	9=NR

<i>STR56G</i>		<i>dilated left ventricle</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
182	0	0=No
28	1	1=Yes
82	9	9=NR

<i>STR56H</i>		<i>ascending aortic arch atheroma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
58	0	0=No
7	1	1=Yes
227	9	9=NR

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<i>STR56H1</i>		<i>ascending aortic arch atheroma diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
554	.S	Legitimately skipped
4	1	1=Mild
0	2	2=Moderate
0	3	3=Severe
3	9	9=NR

<i>STR56I</i>		<i>atrioseptal aneurysm</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
21	0	0=No
3	1	1=Yes
268	9	9=NR

<i>STR56J</i>		<i>patent foramen ovale (PFO)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
35	0	0=No
10	1	1=Yes
247	9	9=NR

<i>STR56K</i>		<i>valve vegetations</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
70	0	0=No
6	1	1=Yes
216	9	9=NR

<i>STR56L</i>		<i>artificial valve</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
70	0	0=No
8	1	1=Yes
214	9	9=NR

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<i>STR56M</i>		<i>left atrial enlargement</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
269	.S	Legitimately skipped
113	0	0=No
99	1	1=Yes
80	9	9=NR

<i>STR57</i>		<i>first sodium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
63		Missing
498	Range	117 - 153 (median=139 mean=138.6 std=4.0)

<i>STR58</i>		<i>first creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
58		Missing
503	Range	0.3 - 38 (median=0.9 mean=1.32 std=2.24)

<i>STR59</i>		<i>first BUN</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
63		Missing
498	Range	5 - 160 (median=17 mean=21.9 std=17.7)

<i>STR60</i>		<i>first Hgb</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
53		Missing
508	Range	3.9 - 19 (median=13 mean=12.8 std=2.0)

<i>STR61</i>		<i>first Hct</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72		Missing
489	Range	19 - 53.3 (median=39.1 mean=38.44 std=5.55)

<i>STR62</i>		<i>first INR</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
211		Missing
350	Range	0.8 - 12 (median=1 mean=1.2 std=1.0)

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STR63		<i>first PTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
251		Missing
310	Range	10 - 81 (median=28 mean=29.6 std=7.2)

STR63A		<i>Activated PTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
251	.S	Legitimately skipped
95	0	0=No
170	1	1=Yes
45	9	9=NR

STR64		<i>first platelet count</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
71		Missing
490	Range	7 - 833 (median=235 mean=245.8 std=94.5)

STR65		<i>Total Cholesterol</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
390		Missing
171	Range	49 - 379 (median=170 mean=177.2 std=46.9)

STR66		<i>LDL</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
389		Missing
172	Range	8 - 214 (median=96.5 mean=99.85 std=38.41)

STR67		<i>HbA1c</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
401		Missing
160	Range	4.43 - 38 (median=6.75 mean=7.533 std=3.111)

STR68		<i>receive thrombolytic treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
492	0	0=No
21	1	1=Yes
48	9	9=NR

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STR68A		<i>symptomatic ICH post tPA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
540	.S	Legitimately skipped
16	0	0=No
0	1	1=Yes
5	9	9=NR

STR69		<i>endovascular therapy for ischemic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
510	0	0=No
14	1	1=Yes
37	9	9=NR

STR69A		<i>symptomatic ICH post EVT?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
547	.S	Legitimately skipped
8	0	0=No
1	1	1=Yes
5	9	9=NR

STR70		<i>osmotic/hypertonic treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
512	0	0=No
10	1	1=Yes
39	9	9=NR

STR71		<i>decompressive hemicraniectomy needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
520	0	0=No
11	1	1=Yes
30	9	9=NR

STR72		<i>alive at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39	0	0=No
522	1	1=Yes

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<i>STR72A</i>		<i>autopsy performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
522	.S	Legitimately skipped
30	0	0=No
2	1	1=Yes
7	9	9=NR

<i>STR72A1</i>		<i>autopsy diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
559	.S	Legitimately skipped
0	1	1=recent bleeding of saccular aneurysm/subarachnoid hemorrhage
0	2	2=hemorrhagic infarction of the brain
0	3	3=lacunar stroke
0	4	4=embolic stroke
0	7	7=other

<i>STR73</i>		<i>ambulate independently before hospitalization</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39	.S	Legitimately skipped
23	0	0=No
469	1	1=Yes
30	9	9=Unknown/NR

<i>STR74</i>		<i>ambulate independently at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
39	.S	Legitimately skipped
74	0	0=No
347	1	1=Yes
101	9	9=Unknown/NR

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STR75		where discharged to
N	Value	Description
39	.S	Legitimately skipped
371	1	1=Home
50	2	2=Acute Rehabilitation
61	3	3=Skilled Nursing Facility
3	4	4=Assisted Living
6	5	5=Hospice
31	6	6=Other/Unknown

STR76		ADLs required
N	Value	Description
39	.S	Legitimately skipped
233	0	0=No
137	1	1=Yes
152	9	9=Unknown/NR

STR77		modified rankin scale done at discharge
N	Value	Description
39	.S	Legitimately skipped
509	0	0=No
6	1	1=Yes
7	9	9=Unknown/NR

STR77A		modified rankin scale score
N	Value	Description
555	.S	Legitimately skipped
4	0	0
0	1	1
0	2	2
0	3	3
0	4	4
2	5	5
0	6	6

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STR77B		<i>discharged ADLs/walk capabilities</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
45	.S	Legitimately skipped
185	0	0=Back to Normal
67	1	1=Exhibiting S/S related to stroke but otherwise without disability
37	2	2=Able to ambulate independently with some disability
19	3	3=Dependent on others for some ADLs, but able to walk
25	4	4=Moderately severe disability
16	5	5=Bedridden (requires 24/7 care)
167	9	9=Unknown/NR

STR78A		<i>prior to admission anticoagulants</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
535	0	0=No/NR
26	1	1=Yes

STR78B		<i>prior to admission beta blockers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
378	0	0=No/NR
183	1	1=Yes

STR78C		<i>prior to admission CCB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
438	0	0=No/NR
123	1	1=Yes

STR78D		<i>prior to admission ACE or ARB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
284	0	0=No/NR
277	1	1=Yes

STR78E		<i>prior to admission scheduled aspirin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
305	0	0=No/NR
256	1	1=Yes

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STR78F		<i>prior to admission heparin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
548	0	0=No/NR
13	1	1=Yes

STR78G		<i>prior to admission Coumadin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
533	0	0=No/NR
28	1	1=Yes

STR78H		<i>prior to admission anti-platelet agent (non-aspirin)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
466	0	0=No/NR
95	1	1=Yes

STR78I		<i>prior to admission statin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
288	0	0=No/NR
273	1	1=Yes

STR79A		<i>during/after admission anticoagulants</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
527	0	0=No/NR
34	1	1=Yes

STR79B		<i>during/after admission beta blockers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
342	0	0=No/NR
219	1	1=Yes

STR79C		<i>during/after admission CCB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
365	0	0=No/NR
196	1	1=Yes

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<i>STR79D</i>		<i>during/after admission ACE or ARB</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
300	0	0=No/NR
261	1	1=Yes

<i>STR79E</i>		<i>during/after admission scheduled aspirin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
210	0	0=No/NR
351	1	1=Yes

<i>STR79F</i>		<i>during/after admission heparin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
396	0	0=No/NR
165	1	1=Yes

<i>STR79G</i>		<i>during/after admission Coumadin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
522	0	0=No/NR
39	1	1=Yes

<i>STR79H</i>		<i>during/after admission anti-platelet agent (non-aspirin)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
418	0	0=No/NR
143	1	1=Yes

<i>STR79I</i>		<i>during/after admission statin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
203	0	0=No/NR
358	1	1=Yes

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Stroke Reviewer 1 Form: FORM CODE=STD1, VERSION=1

ID		HCHS Participant ID
N	Value	Description
558	Present	Text suppressed

VISIT		Visit
N	Value	Description
50	1	Visit 1
57	2	Visit 2
55	3	Visit 3
54	4	Visit 4
43	5	Visit 5
36	6	Visit 6
43	7	Visit 7
44	8	Visit 8
44	9	Visit 9
25	10	Visit 10
54	11	Visit 11
23	12	Visit 12
18	13	Visit 13
10	14	Visit 14
2	15	Visit 15

OCCURRENCE		Occurrence
N	Value	Description
410	1	Occurrence 1
99	2	Occurrence 2
26	3	Occurrence 3
7	4	Occurrence 4
10	5	Occurrence 5
3	6	Occurrence 6
2	7	Occurrence 7
1	10	Occurrence 10

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<i>STD10A_FUTIME</i>		<i>Follow-up time from baseline for STD10A (Completion date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
556	Range	3434 - 6048 (median=4329 mean=4428.0 std=560.1)

<i>STD10AY</i>		<i>Year of STD10A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
556	Range	2020 - 2025 (median=2021 mean=2021.5 std=1.3)

<i>STD10C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
558	Present	Text suppressed

<i>STD10D_FUTIME</i>		<i>Follow-up time from baseline for STD10D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
558	Range	14 - 5662 (median=2065 mean=2178.5 std=1380.7)

<i>STD10DY</i>		<i>Year of STD10D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
558	Range	2008 - 2024 (median=2015 mean=2015.4 std=3.9)

<i>STD11</i>		<i>primary diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
270	1	1 = Not a TIA or Stroke
51	2	2 = TIA
235	3	3 = Stroke

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STD12		stroke type
N	Value	Description
2		Missing
321	.S	Legitimately skipped
173	1	1 = Brain Infarction
22	2	2 = Subarachnoid Hemorrhage
26	3	3 = Intracerebral Hemorrhage
8	4	4 = Other Stroke Type
6	5	5 = Unknown Stroke Type

STD12A		other stroke type specify
N	Value	Description
2		Missing
548	~	Legitimately skipped
8	Present	Text suppressed

STD12B		unknown stroke type specify
N	Value	Description
2		Missing
550	~	Legitimately skipped
6	Present	Text suppressed

STD13		brain infarct subtype
N	Value	Description
2		Missing
375	.S	Legitimately skipped
6	0	0 = N/A (not applicable)
12	1	1 = Large vessel extracranial atheroembolic
10	2	2 = Large vessel intracranial atheroembolic
19	3	3 = Cardioembolic
39	4	4 = Lacunar Infarction
10	5	5 = Acute ischemic stroke of other known etiology
25	6	6 = Ischemic stroke of unknown cause (no probable etiology despite complete workup)
16	7	7 = Ischemic stroke of unknown cause (more than one etiology)
44	8	8 = Ischemic stroke of unknown cause (workup is incomplete)

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STD13A		<i>acute ischemic other known specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
546	~	Legitimately skipped
10	Present	Text suppressed

STD13B		<i>ischemic unknown cause</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
540	~	Legitimately skipped
16	Present	Text suppressed

STD14		<i>Was the Brain Infarction Procedure-Related?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
381	.S	Legitimately skipped
164	0	0=No
8	1	1=Yes
3	9	9=Unknown

STD14A		<i>specify procedure related infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
548	~	Legitimately skipped
8	Present	Text suppressed

STD15		<i>subarachnoid hemorrhage subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
526	.S	Legitimately skipped
7	0	0=N/A (not applicable)
13	1	1=Aneurysmal
3	2	2=Sulcal
2	3	3=Perimesencephalic
3	4	4=Other
2	9	9=Unknown

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STD15A		<i>subarachnoid subtype other specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
553	~	Legitimately skipped
3	Present	Text suppressed

STD16		<i>subarachnoid intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
533	.S	Legitimately skipped
14	0	0=No
8	1	1=Yes
1	9	9=Unknown

STD17		<i>subarachnoid intraparenchymal extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
533	.S	Legitimately skipped
16	0	0=No
7	1	1=Yes
0	9	9=Unknown

STD18		<i>Intracerebral Hemorrhage Subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
522	.S	Legitimately skipped
6	0	0=N/A (not applicable)
15	1	1=Small Vessel Hypertensive Vasculopathy
0	2	2=Cerebral Amyloid
0	3	3=Vascular Malformation Arteriovenous
1	4	4=Vascular Malformation Cavernous
0	5	5=Endocarditis
3	6	6=Primary Intraventricular Hemorrhage
5	7	7=Other
4	9	9=Unknown

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STD18A		<i>intracerebral Hemorrhage Subtype other specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
551	~	Legitimately skipped
5	Present	Text suppressed

STD19		<i>intracerebral intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
528	.S	Legitimately skipped
19	0	0=No
8	1	1=Yes
1	9	9=Unknown

STD110		<i>core of hemorrhage description</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
528	.S	Legitimately skipped
8	1	1=Lobar
16	2	2=Deep
2	3	3=Cerebellar
2	9	9=Unknown

STD111		<i>hemorrhagic procedure-related</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
500	.S	Legitimately skipped
52	0	0=No or N/A
2	1	1=Yes
2	9	9=Unknown

STD111A		<i>hemorrhagic procedure-related specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
554	~	Legitimately skipped
2	Present	Text suppressed

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STD112		overall reviewer comments
N	Value	Description
243		Missing
315	Present	Text suppressed

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Stroke Reviewer 2 Form: FORM CODE=STD2, VERSION=1

ID		HCHS Participant ID
N	Value	Description
550	Present	Text suppressed

VISIT		Visit
N	Value	Description
50	1	Visit 1
57	2	Visit 2
55	3	Visit 3
54	4	Visit 4
43	5	Visit 5
36	6	Visit 6
43	7	Visit 7
44	8	Visit 8
44	9	Visit 9
25	10	Visit 10
54	11	Visit 11
18	12	Visit 12
15	13	Visit 13
10	14	Visit 14
2	15	Visit 15

OCCURRENCE		Occurrence
N	Value	Description
405	1	Occurrence 1
97	2	Occurrence 2
25	3	Occurrence 3
7	4	Occurrence 4
10	5	Occurrence 5
3	6	Occurrence 6
2	7	Occurrence 7
1	10	Occurrence 10

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<i>STD20A_FUTIME</i>		<i>Follow-up time from baseline for STD20A (Completion date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
548	Range	3425 - 6044 (median=4311 mean=4402.7 std=562.5)

<i>STD20AY</i>		<i>Year of STD20A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
548	Range	2020 - 2025 (median=2021 mean=2021.3 std=1.4)

<i>STD20C</i>		<i>Event ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
550	Present	Text suppressed

<i>STD20D_FUTIME</i>		<i>Follow-up time from baseline for STD20D (Event Date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
550	Range	14 - 5662 (median=2046 mean=2144.7 std=1361.4)

<i>STD20DY</i>		<i>Year of STD20D</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
550	Range	2008 - 2024 (median=2015 mean=2015.3 std=3.8)

<i>STD21</i>		<i>primary diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
269	1	1 = Not a TIA or Stroke
56	2	2 = TIA
223	3	3 = Stroke

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STD22		stroke type
N	Value	Description
2		Missing
325	.S	Legitimately skipped
165	1	1 = Brain Infarction
24	2	2 = Subarachnoid Hemorrhage
27	3	3 = Intracerebral Hemorrhage
3	4	4 = Other Stroke Type
4	5	5 = Unknown Stroke Type

STD22A		other stroke type specify
N	Value	Description
2		Missing
545	~	Legitimately skipped
3	Present	Text suppressed

STD22B		unknown stroke type specify
N	Value	Description
2		Missing
544	~	Legitimately skipped
4	Present	Text suppressed

STD23		brain infarct subtype
N	Value	Description
2		Missing
380	.S	Legitimately skipped
1	0	0 = N/A (not applicable)
6	1	1 = Large vessel extracranial atheroembolic
12	2	2 = Large vessel intracranial atheroembolic
21	3	3 = Cardioembolic
43	4	4 = Lacunar Infarction
6	5	5 = Acute ischemic stroke of other known etiology
29	6	6 = Ischemic stroke of unknown cause (no probable etiology despite complete workup)
9	7	7 = Ischemic stroke of unknown cause (more than one etiology)
41	8	8 = Ischemic stroke of unknown cause (workup is incomplete)

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STD23A		<i>acute ischemic other known specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
542	~	Legitimately skipped
6	Present	Text suppressed

STD23B		<i>ischemic unknown cause</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
539	~	Legitimately skipped
9	Present	Text suppressed

STD24		<i>Was the Brain Infarction Procedure-Related?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
381	.S	Legitimately skipped
157	0	0=No
7	1	1=Yes
3	9	9=Unknown

STD24A		<i>specify procedure related infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
541	~	Legitimately skipped
7	Present	Text suppressed

STD25		<i>subarachnoid hemorrhage subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
521	.S	Legitimately skipped
3	0	0=N/A (not applicable)
13	1	1=Aneurysmal
4	2	2=Sulcal
2	3	3=Perimesencephalic
3	4	4=Other
2	9	9=Unknown

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STD25A		<i>subarachnoid subtype other specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
545	~	Legitimately skipped
3	Present	Text suppressed

STD26		<i>subarachnoid intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
524	.S	Legitimately skipped
13	0	0=No
7	1	1=Yes
4	9	9=Unknown

STD27		<i>subarachnoid intraparenchymal extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
524	.S	Legitimately skipped
16	0	0=No
5	1	1=Yes
3	9	9=Unknown

STD28		<i>Intracerebral Hemorrhage Subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
518	.S	Legitimately skipped
3	0	0=N/A (not applicable)
16	1	1=Small Vessel Hypertensive Vasculopathy
0	2	2=Cerebral Amyloid
1	3	3=Vascular Malformation Arteriovenous
0	4	4=Vascular Malformation Cavernous
1	5	5=Endocarditis
1	6	6=Primary Intraventricular Hemorrhage
2	7	7=Other
6	9	9=Unknown

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STD28A		<i>intracerebral Hemorrhage Subtype other specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
546	~	Legitimately skipped
2	Present	Text suppressed

STD29		<i>intracerebral intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
521	.S	Legitimately skipped
17	0	0=No
10	1	1=Yes
0	9	9=Unknown

STD210		<i>core of hemorrhage description</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
521	.S	Legitimately skipped
9	1	1=Lobar
14	2	2=Deep
3	3	3=Cerebellar
1	9	9=Unknown

STD211		<i>hemorrhagic procedure-related</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
494	.S	Legitimately skipped
52	0	0=No or N/A
1	1	1=Yes
1	9	9=Unknown

STD211A		<i>hemorrhagic procedure-related specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
547	~	Legitimately skipped
1	Present	Text suppressed

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<i>STD212</i>		<i>overall reviewer comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
277		Missing
273	Present	Text suppressed

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Stroke Adjudication Reviewer Form: FORM CODE=STD3, VERSION=1

<i>ID</i>		<i>HCHS Participant ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79	Present	Text suppressed

<i>VISIT</i>		<i>Visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	1	Visit 1
9	2	Visit 2
11	3	Visit 3
13	4	Visit 4
5	5	Visit 5
3	6	Visit 6
7	7	Visit 7
6	8	Visit 8
4	9	Visit 9
1	10	Visit 10
7	11	Visit 11
3	12	Visit 12
2	14	Visit 14

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
50	1	Occurrence 1
24	2	Occurrence 2
5	3	Occurrence 3

<i>STD30A_FUTIME</i>		<i>Follow-up time from baseline for STD30A (Completion date)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79	Range	3759 - 6018 (median=4579 mean=4601.9 std=519.7)

<i>STD30AY</i>		<i>Year of STD30A</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79	Range	2021 - 2025 (median=2022 mean=2022.1 std=1.3)

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STD30C		Event ID
N	Value	Description
79	Present	Text suppressed

STD30D_FUTIME		Follow-up time from baseline for STD30D (Event Date)
N	Value	Description
79	Range	81 - 5066 (median=1543 mean=1889.8 std=1326.2)

STD30DY		Year of STD30D
N	Value	Description
79	Range	2009 - 2022 (median=2014 mean=2014.7 std=3.8)

STD31		primary diagnosis
N	Value	Description
46	1	1 = Not a TIA or Stroke
12	2	2 = TIA
21	3	3 = Stroke

STD32		stroke type
N	Value	Description
58	.S	Legitimately skipped
14	1	1 = Brain Infarction
2	2	2 = Subarachnoid Hemorrhage
2	3	3 = Intracerebral Hemorrhage
2	4	4 = Other Stroke Type
1	5	5 = Unknown Stroke Type

STD32A		other stroke type specify
N	Value	Description
77	~	Legitimately skipped
2	Present	Text suppressed

STD32B		unknown stroke type specify
N	Value	Description
78	~	Legitimately skipped
1	Present	Text suppressed

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STD33		<i>brain infarct subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
63	.S	Legitimately skipped
2	0	0 = N/A (not applicable)
1	1	1 = Large vessel extracranial atheroembolic
1	2	2 = Large vessel intracranial atheroembolic
1	3	3 = Cardioembolic
4	4	4 = Lacunar Infarction
0	5	5 = Acute ischemic stroke of other known etiology
1	6	6 = Ischemic stroke of unknown cause (no probable etiology despite complete workup)
1	7	7 = Ischemic stroke of unknown cause (more than one etiology)
5	8	8 = Ischemic stroke of unknown cause (workup is incomplete)

STD33A		<i>acute ischemic other known specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79	~	Legitimately skipped

STD33B		<i>ischemic unknown cause</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
78	~	Legitimately skipped
1	Present	Text suppressed

STD34		<i>Was the Brain Infarction Procedure-Related?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
65	.S	Legitimately skipped
13	0	0=No
0	1	1=Yes
1	9	9=Unknown

STD34A		<i>specify procedure related infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79	~	Legitimately skipped

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STD35		<i>subarachnoid hemorrhage subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	.S	Legitimately skipped
1	0	0=N/A (not applicable)
0	1	1=Aneurysmal
2	2	2=Sulcal
0	3	3=Perimesencephalic
0	4	4=Other
1	9	9=Unknown

STD35A		<i>subarachnoid subtype other specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79	~	Legitimately skipped

STD36		<i>subarachnoid intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	.S	Legitimately skipped
3	0	0=No
0	1	1=Yes
0	9	9=Unknown

STD37		<i>subarachnoid intraparenchymal extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	.S	Legitimately skipped
1	0	0=No
2	1	1=Yes
0	9	9=Unknown

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STD38		<i>Intracerebral Hemorrhage Subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	.S	Legitimately skipped
1	0	0=N/A (not applicable)
0	1	1=Small Vessel Hypertensive Vasculopathy
0	2	2=Cerebral Amyloid
1	3	3=Vascular Malformation Arteriovenous
0	4	4=Vascular Malformation Cavemous
0	5	5=Endocarditis
0	6	6=Primary Intraventricular Hemorrhage
1	7	7=Other
1	9	9=Unknown

STD38A		<i>intracerebral Hemorrhage Subtype other specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
78	~	Legitimately skipped
1	Present	Text suppressed

STD39		<i>intracerebral intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	.S	Legitimately skipped
2	0	0=No
1	1	1=Yes
0	9	9=Unknown

STD310		<i>core of hemorrhage description</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	.S	Legitimately skipped
2	1	1=Lobar
0	2	2=Deep
1	3	3=Cerebellar
0	9	9=Unknown

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<i>STD311</i>		<i>hemorrhagic procedure-related</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
73	.S	Legitimately skipped
5	0	0=No or N/A
0	1	1=Yes
1	9	9=Unknown

<i>STD311A</i>		<i>hemorrhagic procedure-related specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
79	~	Legitimately skipped

<i>STD312</i>		<i>overall reviewer comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
22		Missing
57	Present	Text suppressed

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Pregnancy Complication Abstraction Form: FORM CODE=PRC, VERSION=1

<i>ID</i>		<i>HCHS Participant ID</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Present	Text suppressed

<i>VISIT</i>		<i>Label information missing in the dataset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14	1	Visit 1
15	2	Visit 2
24	3	Visit 3
17	4	Visit 4
17	5	Visit 5
15	6	Visit 6
18	7	Visit 7
1	8	Visit 8
6	9	Visit 9

<i>OCCURRENCE</i>		<i>Label information missing in the dataset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
109	1	Occurrence 1
14	2	Occurrence 2
3	3	Occurrence 3
1	4	Occurrence 4

<i>PRC1</i>		<i>presenting complaint</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
93	1	1= Labor, scheduled induction, scheduled c-section
18	2	2= Preeclampsia symptoms
1	3	3= Decreased fetal movement
0	4	4= Hyperglycemia symptoms
0	5	5= Poor glucose control
14	6	6= Other

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<i>PRC2</i>		<i>24 hour stay</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	0	0=No/Not Recorded
0	1	1=Yes

<i>PRC3A</i>		<i>Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17		Missing
110	Range	0:37:00 - 23:31:00

<i>PRC3AHR</i>		<i>Hour of Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17		Missing
110	Range	0 - 23 (median=11 mean=11.5 std=5.7)

<i>PRC3AMN</i>		<i>Minute of Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17		Missing
110	Range	0 - 57 (median=28.5 mean=26.29 std=16.62)

<i>PRC3ASC</i>		<i>Second of Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52		Missing
75	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>PRC4A</i>		<i>Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
55		Missing
72	Range	1:00:00 - 23:59:00

<i>PRC4AHR</i>		<i>Hour of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
55		Missing
72	Range	1 - 23 (median=15 mean=15.0 std=4.1)

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<i>PRC4AMN</i>		<i>Minute of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
55		Missing
72	Range	0 - 59 (median=30 mean=27.2 std=18.4)

<i>PRC4ASC</i>		<i>Second of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
77		Missing
50	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>PRC5</i>		<i>Admitting diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25		Missing
102	Present	Text suppressed

<i>PRC6</i>		<i>Discharge diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18		Missing
109	Present	Text suppressed

<i>PRC7</i>		<i>patient referred</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
102	0	0=No/Not Recorded
25	1	1=Yes

<i>PRC8</i>		<i>patient transferred</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
125	0	0=No/Not Recorded
2	1	1=Yes

<i>PRC9</i>		<i>patient alive</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
0	0	0=No/Not Recorded
127	1	1=Yes

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<i>PRC11</i>		<i>EDC method</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14	1	1= Last Menstrual Period (LMP) only
14	2	2= LMP consistent with first trimester ultrasound (<14 weeks)
7	3	3= LMP consistent with second trimester ultrasound (14 0/7 - 27 6/7 weeks)
2	4	4= LMP consistent with third trimester ultrasound (>=28 weeks)
9	5	5= First trimester ultrasound only (<14 weeks)
7	6	6= Second trimester ultrasound only (14 0/7 - 27 6/7 weeks)
2	7	7= Third trimester ultrasound only (>=28 weeks)
4	8	8= Assisted reproductive technology (ART)
68	9	9= Not documented

<i>PRC12A</i>		<i>headache</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
111	0	0=No/Not Recorded
15	1	1=Yes

<i>PRC12B</i>		<i>blurry vision</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
122	0	0=No/Not Recorded
4	1	1=Yes

<i>PRC12C</i>		<i>right upper quadrant pain or epigastric pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
123	0	0=No/Not Recorded
3	1	1=Yes

<i>PRC12D</i>		<i>nausea, vomiting</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
124	0	0=No/Not Recorded
2	1	1=Yes

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<i>PRC12E</i>		<i>vaginal bleeding</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
123	0	0=No/Not Recorded
3	1	1=Yes

<i>PRC12F</i>		<i>seizure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
126	0	0=No/Not Recorded
0	1	1=Yes

<i>PRC13A</i>		<i>diabetes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
111	0	0=No/Not Recorded
15	1	1=Yes

<i>PRC13B</i>		<i>gestational diabetes in prior pregnancies</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
114	0	0=No/Not Recorded
12	1	1=Yes

<i>PRC13C</i>		<i>gestational diabetes in current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
60	0	0=No/Not Recorded
66	1	1=Yes

<i>PRC13D</i>		<i>hypertension prior to pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
101	0	0=No/Not Recorded
25	1	1=Yes

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<i>PRC13E</i>		<i>gestational hypertension in prior pregnancies</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
114	0	0=No/Not Recorded
12	1	1=Yes

<i>PRC13F</i>		<i>gestational hypertension in current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
85	0	0=No/Not Recorded
41	1	1=Yes

<i>PRC13G</i>		<i>preeclampsia in prior pregnancies</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
122	0	0=No/Not Recorded
4	1	1=Yes

<i>PRC13H</i>		<i>preeclampsia in current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
98	0	0=No/Not Recorded
28	1	1=Yes

<i>PRC13I</i>		<i>thyroid disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
116	0	0=No/Not Recorded
10	1	1=Yes

<i>PRC13J</i>		<i>smoking prior to current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
122	0	0=No/Not Recorded
4	1	1=Yes

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<i>PRC13K</i>		<i>smoking during pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
121	0	0=No/Not Recorded
5	1	1=Yes

<i>PRC13L</i>		<i>known seizure disorder</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
124	0	0=No/Not Recorded
2	1	1=Yes

<i>PRC14</i>		<i>qualifying BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
84	0	0=No/Not Recorded
42	1	1=Yes

<i>PRC14A2</i>		<i>Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18		Missing
85	.S	Legitimately skipped
24	Range	1:02:00 - 20:53:00

<i>PRC14A2HR</i>		<i>Hour of Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
103		Missing
24	Range	1 - 20 (median=15 mean=13.3 std=5.2)

<i>PRC14A2MN</i>		<i>Minute of Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
103		Missing
24	Range	0 - 53 (median=30.5 mean=27.79 std=18.97)

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<i>PRC14A2SC</i>		<i>Second of Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
112		Missing
15	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>PRC14A3</i>		<i>initial systolic BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
42	Range	120 - 200 (median=150 mean=155.0 std=16.6)

<i>PRC14A4</i>		<i>initial diastolic BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
85	.S	Legitimately skipped
34	Range	72 - 116 (median=93 mean=92.8 std=11.0)

<i>PRC14B2</i>		<i>confirming BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
32		Missing
85	.S	Legitimately skipped
10	Range	1:51:00 - 21:55:00

<i>PRC14B2HR</i>		<i>Hour of confirming BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117		Missing
10	Range	1 - 21 (median=15.5 mean=13.70 std=7.73)

<i>PRC14B2MN</i>		<i>Minute of confirming BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117		Missing
10	Range	0 - 55 (median=22.5 mean=23.00 std=19.84)

<i>PRC14B2SC</i>		<i>Second of confirming BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
120		Missing
7	Range	0 - 0 (median=0 mean=0.0 std=0.0)

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<i>PRC14B3</i>		<i>confirming systolic BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24		Missing
85	.S	Legitimately skipped
18	Range	140 - 186 (median=149.5 mean=153.72 std=12.02)

<i>PRC14B4</i>		<i>confirming diastolic BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
26		Missing
85	.S	Legitimately skipped
16	Range	58 - 120 (median=92 mean=92.6 std=14.9)

<i>PRC15</i>		<i>anti-hypertensive meds</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
17	0	0= No
11	1	1= Yes
98	9	9=NR

<i>PRC16</i>		<i>vaginal exam</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
6	0	0= No
74	1	1= Yes
46	9	9=NR

<i>PRC16A</i>		<i>dilation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
53	.S	Legitimately skipped
72	Range	0 - 10 (median=3 mean=3.2 std=2.9)

<i>PRC16B</i>		<i>effacement</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
12		Missing
53	.S	Legitimately skipped
62	Range	0 - 100 (median=65 mean=56.7 std=36.8)

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<i>PRC17</i>		<i>prenatal records</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
51	0	0= No/ NR
75	1	1= Yes

<i>PRC17A</i>		<i>baseline urine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
69	0	0= No/ NR
6	1	1= Yes

<i>PRC17A2</i>		<i>total urine volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
121	.S	Legitimately skipped
1	Range	2375 - 2375 (median=2375 mean=2375.0 std=.)

<i>PRC17A3</i>		<i>total protein</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
121	.S	Legitimately skipped
5	Range	168 - 432 (median=349 mean=325.4 std=97.7)

<i>PRC17A3A</i>		<i>protein units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
121	.S	Legitimately skipped
3	1	1=mg/24 hours
1	2	2=mg/dL

<i>PRC17B</i>		<i>glucose loading test</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
51	0	0= No/ NR
24	1	1= Yes

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<i>PRC17B1</i>		<i>glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
103	.S	Legitimately skipped
19	Range	50 - 220 (median=148 mean=143.8 std=46.4)

<i>PRC17C</i>		<i>type of OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
53	0	0=Not performed/Not Recorded
2	1	1=100 g load
0	2	2=75g load
20	3	3=Test performed, but type of glucose load not documented

<i>PRC17C1A</i>		<i>OGTT discrete values</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
105	.S	Legitimately skipped
6	0	0= No
16	1	1= Yes

<i>PRC17C1B</i>		<i>fasting glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
111	.S	Legitimately skipped
13	Range	84 - 150 (median=95 mean=96.8 std=16.7)

<i>PRC17C1C</i>		<i>1 hr glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
111	.S	Legitimately skipped
16	Range	103 - 270 (median=183.5 mean=177.94 std=43.04)

<i>PRC17C1E</i>		<i>3 hr glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
111	.S	Legitimately skipped
14	Range	83 - 176 (median=128 mean=124.3 std=29.8)

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<i>PRC17C2</i>		<i>OGTT number of values reported</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
121	.S	Legitimately skipped
6	0	0= No
0	1	1= Yes

<i>PRC17C2A</i>		<i>number of abnormal values</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	0	0=0/4
0	1	1=1/4
0	2	2=2/4
0	3	3=3/4
0	4	4=4/4 (4 or more tests)

<i>PRC17C3</i>		<i>OGTT documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
121	.S	Legitimately skipped
0	0	0=no result documented
4	1	1=abnormal OGTT
2	2	2=normal OGTT

<i>PRC17D</i>		<i>prenatal Hgb A1c</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
62	0	0= No/ NR
13	1	1= Yes

<i>PRC17D1A</i>		<i>Hgb A1c value less than or equal to 20</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6		Missing
114	.S	Legitimately skipped
7	Range	5.2 - 9.3 (median=5.8 mean=6.41 std=1.41)

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<i>PRC17D2A</i>		<i>Hgb A1c greater than 20 value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
114	.S	Legitimately skipped
9	Range	5 - 8.4 (median=5.5 mean=5.71 std=1.03)

<i>PRC17E</i>		<i>outpatient pressure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
43	0	0= No/ NR
32	1	1= Yes

<i>PRC17E1A</i>		<i>highest systolic less than or equal to 20 value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24		Missing
95	.S	Legitimately skipped
8	Range	100 - 143 (median=135 mean=126.1 std=17.9)

<i>PRC17E2A</i>		<i>highest systolic greater than 20 value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
95	.S	Legitimately skipped
28	Range	108 - 160 (median=128 mean=130.7 std=13.9)

<i>PRC17E3A</i>		<i>highest diastolic less than or equal to 20 date</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24		Missing
95	.S	Legitimately skipped
8	Range	70 - 90 (median=76.5 mean=77.50 std=6.95)

<i>PRC17E4A</i>		<i>highest diastolic greater than 20 date</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4		Missing
95	.S	Legitimately skipped
28	Range	54 - 100 (median=77.5 mean=78.46 std=10.19)

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<i>PRC17F</i>		<i>weight gain available</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
42	0	0= No/ NR
33	1	1= Yes

<i>PRC17F1</i>		<i>pre weight</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
22		Missing
94	.S	Legitimately skipped
11	Range	72.7 - 230 (median=96.3 mean=120.01 std=51.27)

<i>PRC17F1A</i>		<i>pre weight units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16		Missing
94	.S	Legitimately skipped
7	1	1= kg
4	2	2= lb
6	9	9= NR

<i>PRC17F2</i>		<i>first weight</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
28		Missing
94	.S	Legitimately skipped
5	Range	63.1 - 173.5 (median=151 mean=135.5 std=43.3)

<i>PRC17F2A</i>		<i>first weight units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
23		Missing
94	.S	Legitimately skipped
1	1	1= kg
4	2	2= lb
5	9	9= NR

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<i>PRC17F3</i>		<i>last weight</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
94	.S	Legitimately skipped
25	Range	62.6 - 237 (median=120.6 mean=135.90 std=54.40)

<i>PRC17F3A</i>		<i>last weight units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
94	.S	Legitimately skipped
14	1	1= kg
11	2	2= lb
0	9	9= NR

<i>PRC17G</i>		<i>outpatient diabetes tx</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
33	0	0= No/ NR
42	1	1= Yes

<i>PRC17G1</i>		<i>outpatient insulin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
25	0	0= No/ NR
17	1	1= Yes

<i>PRC17G2</i>		<i>outpatient glyburide</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
24	0	0= No/ NR
18	1	1= Yes

<i>PRC17G3</i>		<i>outpatient metformin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
36	0	0= No/ NR
6	1	1= Yes

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<i>PRC17G4</i>		<i>outpatient other med</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
35	0	0= No/ NR
7	1	1= Yes

<i>PRC17G4A</i>		<i>outpatient med name</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
120	~	Legitimately skipped
7	Present	Text suppressed

<i>PRC18A</i>		<i>12 or 24 hour urine test</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
114	0	0= No/ NR
12	1	1= Yes

<i>PRC18A2</i>		<i>urine total volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
115	.S	Legitimately skipped
1	Range	2375 - 2375 (median=2375 mean=2375.0 std=.)

<i>PRC18A3</i>		<i>urine total protein</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
115	.S	Legitimately skipped
10	Range	0 - 2142 (median=356 mean=599.0 std=622.0)

<i>PRC18A4</i>		<i>24 hour creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
115	.S	Legitimately skipped
1	Range	2.1 - 2.1 (median=2.1 mean=2.10 std=.)

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<i>PRC18A5</i>		<i>urine duration</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
115	.S	Legitimately skipped
0	1	1=12h
10	2	2=24h

<i>PRC18B</i>		<i>last urine protein</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
118	0	0= No/ NR
8	1	1= Yes

<i>PRC18B2</i>		<i>last urine protein value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
119	.S	Legitimately skipped
8	Range	0.43 - 1211 (median=0.715 mean=200.7238 std=430.3331)

<i>PRC19A</i>		<i>platelet count</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
41	0	0= No/ NR
85	1	1= Yes

<i>PRC19A1</i>		<i>lowest platelet value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	.S	Legitimately skipped
85	Range	104 - 493 (median=198 mean=214.1 std=67.9)

<i>PRC19B</i>		<i>AST</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
84	0	0= No/ NR
42	1	1= Yes

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<i>PRC19B1</i>		<i>highest AST value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
42	Range	14 - 408 (median=23.5 mean=47.60 std=80.82)

<i>PRC19B2</i>		<i>lab limit SGOT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
10	0	0= No/ NR
32	1	1= Yes

<i>PRC19B2A</i>		<i>Lab limit SGOT value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
95	.S	Legitimately skipped
32	Range	32 - 60 (median=41 mean=41.3 std=6.7)

<i>PRC19C</i>		<i>ALT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
84	0	0= No/ NR
42	1	1= Yes

<i>PRC19C1</i>		<i>highest ALT value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
42	Range	7 - 402 (median=19.5 mean=40.71 std=71.39)

<i>PRC19C2</i>		<i>lab limit ALT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	.S	Legitimately skipped
10	0	0= No/ NR
32	1	1= Yes

<i>PRC19C2A</i>		<i>lab limit ALT value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
95	.S	Legitimately skipped
32	Range	24 - 78 (median=52 mean=51.0 std=10.2)

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<i>PRC19D</i>		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
83	0	0= No/ NR
43	1	1= Yes

<i>PRC19D1</i>		<i>highest serum creatinine value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
84	.S	Legitimately skipped
43	Range	0.39 - 2.13 (median=0.56 mean=0.618 std=0.271)

<i>PRC19E</i>		<i>uric acid</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
97	0	0= No/ NR
29	1	1= Yes

<i>PRC19E1</i>		<i>highest acid value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
98	.S	Legitimately skipped
29	Range	2.9 - 10.6 (median=4.9 mean=5.13 std=1.51)

<i>PRC19F</i>		<i>LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
104	0	0= No/ NR
22	1	1= Yes

<i>PRC19F1</i>		<i>highest LDH value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
105	.S	Legitimately skipped
22	Range	147 - 792 (median=212.5 mean=279.50 std=167.45)

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<i>PRC19G</i>		<i>admission hematocrit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
49	0	0= No/ NR
77	1	1= Yes

<i>PRC19G1</i>		<i>admission hematocrit value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
50	.S	Legitimately skipped
77	Range	25.2 - 44.8 (median=35.8 mean=35.11 std=3.83)

<i>PRC19H</i>		<i>post delivery hematocrit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
46	0	0= No/ NR
80	1	1= Yes

<i>PRC19H1</i>		<i>lowest hematocrit value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
47	.S	Legitimately skipped
80	Range	19.8 - 39.6 (median=30.9 mean=30.71 std=4.19)

<i>PRC19I</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
126	0	0= No/ NR
0	1	1= Yes

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<i>PRC20A</i>		<i>magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
4	1	1=None
13	2	2=Yes, for preeclampsia
1	3	3=Yes, for preeclampsia and other indication
0	4	4=Yes, for tocolysis
0	5	5=Yes, for fetal neuroprotection (choose neuroprotection if given for tocolysis and neuroprotection)
1	6	6=Yes, other indication
3	7	7=Yes, indication not documented
104	9	9=Unsure/NR

<i>PRC20B</i>		<i>insulin (subcutaneous)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
100	0	0=No/Not Recorded
26	1	1=Yes

<i>PRC20C</i>		<i>insulin (IV or drip)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
125	0	0=No/Not Recorded
1	1	1=Yes

<i>PRC20D</i>		<i>antihypertensives IV</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
122	0	0=No/Not Recorded
4	1	1=Yes

<i>PRC20E</i>		<i>antihypertensives PO</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
112	0	0=No/Not Recorded
14	1	1=Yes

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<i>PRC20F</i>		<i>betamethason/dexamethasone IM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
110	0	0=No/Not Recorded
16	1	1=Yes

<i>PRC20F1</i>		<i>prematurity</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
111	.S	Legitimately skipped
3	0	0=No/Not Recorded
13	1	1=Yes

<i>PRC20G</i>		<i>tocolysis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
122	0	0=No/Not Recorded
4	1	1=Yes

<i>PRC21</i>		<i>patient deliver</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
9	0	0=No
117	1	1=Yes
0	9	9=NR

<i>PRC21B</i>		<i>Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42		Missing
10	.S	Legitimately skipped
75	Range	0:02:00 - 23:48:00

<i>PRC21BHR</i>		<i>Hour of Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52		Missing
75	Range	0 - 23 (median=11 mean=11.4 std=6.7)

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<i>PRC21BMN</i>		<i>Minute of Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52		Missing
75	Range	0 - 59 (median=28 mean=28.4 std=19.0)

<i>PRC21BSC</i>		<i>Second of Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
77		Missing
50	Range	0 - 0 (median=0 mean=0.0 std=0.0)

<i>PRC22A</i>		<i>gestional weeks</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
10	.S	Legitimately skipped
114	Range	27 - 41 (median=39 mean=37.8 std=2.5)

<i>PRC22B</i>		<i>gestional day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5		Missing
10	.S	Legitimately skipped
112	Range	0 - 7 (median=3 mean=2.7 std=2.2)

<i>PRC23</i>		<i>type of labor</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
10	.S	Legitimately skipped
38	1	1= No labor
32	2	2= Spontaneous
6	3	3= Spontaneous, augmented
38	4	4= Induced
2	9	9= Unsure/NR

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<i>PRC24</i>		<i>scheduled delivery reason</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
50	.S	Legitimately skipped
1	1	1= IUFD (intrauterine fetal demise)
0	2	2= Prior stillbirth
0	3	3=Congenital malformation / fetal anomaly
2	4	4= IUGR (intrauterine growth restriction)
6	5	5= Oligohydramnious
1	6	6= Non-reassuring fetal status
0	7	7= Abrupton
26	8	8= Hypertension / preeclampsia
10	9	9= Diabetes
0	10	10= Chorioamnionitis
0	11	11= PROM
2	12	12= Macrosomia
1	13	13= Postterm
0	14	14= Cholestasis
17	15	15= Elective
3	16	16= Not documented
8	17	17= Other
0	18	18= N/A

<i>PRC24A</i>		<i>scheduled delivery other</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
119	~	Legitimately skipped
8	Present	Text suppressed

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<i>PRC25</i>		<i>c-section indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
0	1	1= CPD
8	2	2= Failed induction / FTP
3	3	3= Fetal distress
0	4	4= Cord prolapsed
7	5	5= Malpresentation Twin A / Singleton
0	6	6= Malpresentation Twin B
1	7	7= Abruption
1	8	8= Previa
3	9	9= Preeclampsia / hypertension
0	10	10= Multiple gestation
0	11	11= Communicable disease
1	12	12= Previous C Section
2	13	13= Macrosomia
23	14	14= Elective
3	15	15= Not documented
4	16	16= Other
61	17	17= N/A

<i>PRC25A</i>		<i>c-section specify reason</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
123	~	Legitimately skipped
4	Present	Text suppressed

<i>PRC26</i>		<i>skin incision</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
71	.S	Legitimately skipped
3	1	1=Vertical
29	2	2= Pfannensteil
24	9	9= Not Recorded

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<i>PRC27</i>		<i>uterine incision</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
71	.S	Legitimately skipped
0	1	1= Low vertical
2	2	2= Classical, other vertical
44	3	3= Low Transverse
0	4	4= T or J incision
10	9	9= Not Recorded

<i>PRC28A</i>		<i>placental abruption</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
115	0	0=No/Not Recorded
2	1	1=Yes

<i>PRC28B</i>		<i>postpartum hemorrhage</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
113	0	0=No/Not Recorded
4	1	1=Yes

<i>PRC28C</i>		<i>third degree laceration</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
116	0	0=No/Not Recorded
1	1	1=Yes

<i>PRC28D</i>		<i>fourth degree laceration</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
117	0	0=No/Not Recorded
0	1	1=Yes

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<i>PRC28E</i>		<i>injury to viscera</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
117	0	0=No/Not Recorded
0	1	1=Yes

<i>PRC28F</i>		<i>intubation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
117	0	0=No/Not Recorded
0	1	1=Yes

<i>PRC28G</i>		<i>ICU admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
115	0	0=No/Not Recorded
2	1	1=Yes

<i>PRC28H</i>		<i>transfusion</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
113	0	0=No/Not Recorded
4	1	1=Yes

<i>PRC28H1</i>		<i>platelets</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
116	0	0=No/Not Recorded
1	1	1=Yes

<i>PRC28H2</i>		<i>cryoprecipitate</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
117	0	0=No/Not Recorded
0	1	1=Yes

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<i>PRC28H3</i>		<i>frozen plasma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
116	0	0=No/Not Recorded
1	1	1=Yes

<i>PRC28I</i>		<i>tubal ligation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
97	0	0=No/Not Recorded
20	1	1=Yes

<i>PRC28J</i>		<i>dilation and curettage</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
117	0	0=No/Not Recorded
0	1	1=Yes

<i>PRC28K</i>		<i>hysterectomy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
116	0	0=No/Not Recorded
1	1	1=Yes

<i>PRC28L</i>		<i>intrapartum IV/IM narcotic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
116	0	0=No/Not Recorded
1	1	1=Yes

<i>PRC28M</i>		<i>intrapartum regional epidural</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
67	0	0=No/Not Recorded
50	1	1=Yes

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<i>PRC28N</i>		<i>intrapartum regional spinal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
76	0	0=No/Not Recorded
41	1	1=Yes

<i>PRC28O</i>		<i>intrapartum general anesthesia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
115	0	0=No/Not Recorded
2	1	1=Yes

<i>PRC29</i>		<i>infants born</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	.S	Legitimately skipped
112	1	1=1
3	2	2=2
0	3	3=3
0	4	4=4
2	9	9=Not Recorded

<i>PRC30A1</i>		<i>grams 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42		Missing
12	.S	Legitimately skipped
73	Range	810 - 4730 (median=3255 mean=3161.7 std=664.1)

<i>PRC30A2</i>		<i>pounds 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
93		Missing
12	.S	Legitimately skipped
22	Range	3 - 10 (median=7 mean=6.7 std=1.8)

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<i>PRC30A3</i>		<i>ounces 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
95		Missing
12	.S	Legitimately skipped
20	Range	1 - 15 (median=11 mean=10.5 std=3.8)

<i>PRC30B</i>		<i>one min apgar 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
23		Missing
12	.S	Legitimately skipped
92	Range	0 - 9 (median=9 mean=8.1 std=1.6)

<i>PRC30C</i>		<i>five min apgar 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
23		Missing
12	.S	Legitimately skipped
92	Range	0 - 9 (median=9 mean=8.8 std=1.0)

<i>PRC30D</i>		<i>shoulder dystocia 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
12	.S	Legitimately skipped
21	0	0= No
2	1	1= Yes
89	9	9= NR

<i>PRC30E</i>		<i>birth trauma 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
12	.S	Legitimately skipped
13	0	0= No
0	1	1= Yes
101	9	9= NR

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<i>PRC30F</i>		<i>stillbirth 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
12	.S	Legitimately skipped
110	0	0= No
2	1	1= Yes
2	9	9= NR

<i>PRC30G</i>		<i>type of delivery 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
12	.S	Legitimately skipped
49	1	1= SVD
1	2	2= FAVD
6	3	3= VAVD
0	4	4= Breech vaginal
55	5	5= C-section
3	9	9= NR/Other

<i>PRC30H</i>		<i>infant sex 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
12	.S	Legitimately skipped
40	1	1= Male
51	2	2= Female
23	9	9= NR

<i>PRC31A1</i>		<i>grams 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
124	.S	Legitimately skipped
1	Range	1320 - 1320 (median=1320 mean=1320.0 std=.)

<i>PRC31A2</i>		<i>pounds 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
124	.S	Legitimately skipped

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<i>PRC31A3</i>		<i>ounces 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3		Missing
124	.S	Legitimately skipped

<i>PRC31B</i>		<i>one min apgar 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
124	.S	Legitimately skipped
2	Range	5 - 6 (median=5.5 mean=5.50 std=0.71)

<i>PRC31C</i>		<i>five min apgar 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
124	.S	Legitimately skipped
2	Range	8 - 8 (median=8 mean=8.0 std=0.0)

<i>PRC31D</i>		<i>shoulder dystocia 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
124	.S	Legitimately skipped
1	0	0= No
0	1	1= Yes
2	9	9= NR

<i>PRC31E</i>		<i>birth trauma 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
124	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
3	9	9= NR

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<i>PRC31F</i>		<i>stillbirth 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
124	.S	Legitimately skipped
3	0	0= No
0	1	1= Yes
0	9	9= NR

<i>PRC31G</i>		<i>type of delivery 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
124	.S	Legitimately skipped
1	1	1= SVD
0	2	2= FAVD
0	3	3= VAVD
0	4	4= Breech vaginal
2	5	5= C-section
0	9	9= NR/Other

<i>PRC31H</i>		<i>infant sex 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
124	.S	Legitimately skipped
2	1	1= Male
1	2	2= Female
0	9	9= NR

<i>PRC32A1</i>		<i>grams 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC32A2</i>		<i>pounds 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC32A3</i>		<i>ounces 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

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<i>PRC32B</i>		<i>one min apgar 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC32C</i>		<i>five min apgar 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC32D</i>		<i>shoulder dystocia 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
0	9	9= NR

<i>PRC32E</i>		<i>birth trauma 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
0	9	9= NR

<i>PRC32F</i>		<i>stillbirth</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
0	9	9= NR

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<i>PRC32G</i>		<i>type of delivery 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	1	1= SVD
0	2	2= FAVD
0	3	3= VAVD
0	4	4= Breech vaginal
0	5	5= C-section
0	9	9= NR/Other

<i>PRC32H</i>		<i>infant sex 3</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	1	1= Male
0	2	2= Female
0	9	9= NR

<i>PRC33A1</i>		<i>grams 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC33A2</i>		<i>pounds 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC33A3</i>		<i>ounces 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC33B</i>		<i>one min apgar 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC33C</i>		<i>five min apgar 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

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<i>PRC33D</i>		<i>shoulder dystocia 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
0	9	9= NR

<i>PRC33E</i>		<i>birth trauma 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
0	9	9= NR

<i>PRC33F</i>		<i>stillbirth 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
0	9	9= NR

<i>PRC33G</i>		<i>type of delivery 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	1	1= SVD
0	2	2= FAVD
0	3	3= VAVD
0	4	4= Breech vaginal
0	5	5= C-section
0	9	9= NR/Other

<i>PRC33H</i>		<i>infant sex 4</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped
0	1	1= Male
0	2	2= Female
0	9	9= NR

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<i>PRC0A_FUTIME</i>		<i>Futime from Randomization date for (PRC0A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	2093 - 3703 (median=2756 mean=2708.0 std=372.0)

<i>PRC0D_FUTIME</i>		<i>Futime from Randomization date for (PRC0D)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	212 - 3405 (median=1333 mean=1433.4 std=803.4)

<i>PRC3_FUTIME</i>		<i>Futime from Randomization date for (PRC3)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
126	Range	212 - 3405 (median=1330 mean=1421.1 std=806.4)

<i>PRC3B_FUTIME</i>		<i>Futime from Randomization date for (PRC3B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	212 - 3405 (median=1333 mean=1433.4 std=803.4)

<i>PRC4_FUTIME</i>		<i>Futime from Randomization date for (PRC4)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
126	Range	242 - 3427 (median=1350 mean=1445.5 std=806.8)

<i>PRC10_FUTIME</i>		<i>Futime from Randomization date for (PRC10)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
26		Missing
101	Range	252 - 3493 (median=1489 mean=1501.5 std=844.5)

<i>PRC14A1_FUTIME</i>		<i>Futime from Randomization date for (PRC14A1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
86		Missing
41	Range	240 - 3405 (median=1674 mean=1555.3 std=911.6)

<i>PRC14B1_FUTIME</i>		<i>Futime from Randomization date for (PRC14B1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110		Missing
17	Range	240 - 3405 (median=1771 mean=1652.8 std=860.1)

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<i>PRC17A1_FUTIME</i>		<i>Futime from Randomization date for (PRC17A1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
123		Missing
4	Range	273 - 1174 (median=958.5 mean=841.00 std=404.61)

<i>PRC17B2_FUTIME</i>		<i>Futime from Randomization date for (PRC17B2)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
120		Missing
7	Range	99 - 2631 (median=903 mean=1116.6 std=1038.9)

<i>PRC17C1_FUTIME</i>		<i>Futime from Randomization date for (PRC17C1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117		Missing
10	Range	129 - 2640 (median=1188.5 mean=1193.90 std=954.45)

<i>PRC17D1B_FUTIME</i>		<i>Futime from Randomization date for (PRC17D1B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
123		Missing
4	Range	1546 - 2631 (median=2045 mean=2066.8 std=450.4)

<i>PRC17D2B_FUTIME</i>		<i>Futime from Randomization date for (PRC17D2B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
122		Missing
5	Range	185 - 2210 (median=1532 mean=1377.2 std=775.7)

<i>PRC17E1B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E1B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
123		Missing
4	Range	338 - 1893 (median=1222 mean=1168.8 std=725.7)

<i>PRC17E2B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E2B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
102		Missing
25	Range	279 - 3350 (median=1602 mean=1617.3 std=787.4)

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<i>PRC17E3B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E3B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
123		Missing
4	Range	338 - 1886 (median=1255 mean=1183.5 std=738.5)

<i>PRC17E4B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E4B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
101		Missing
26	Range	279 - 3350 (median=1608.5 mean=1627.15 std=771.02)

<i>PRC17F2B_FUTIME</i>		<i>Futime from Randomization date for (PRC17F2B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
123		Missing
4	Range	134 - 1880 (median=552.5 mean=779.75 std=780.99)

<i>PRC17F3B_FUTIME</i>		<i>Futime from Randomization date for (PRC17F3B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
104		Missing
23	Range	276 - 2404 (median=1583 mean=1374.1 std=687.1)

<i>PRC18A1_FUTIME</i>		<i>Futime from Randomization date for (PRC18A1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
120		Missing
7	Range	273 - 1675 (median=1081 mean=939.4 std=495.8)

<i>PRC18B1_FUTIME</i>		<i>Futime from Randomization date for (PRC18B1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
119		Missing
8	Range	1944 - 3405 (median=2250 mean=2470.5 std=562.7)

<i>PRC21A_FUTIME</i>		<i>Futime from Randomization date for (PRC21A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18		Missing
109	Range	240 - 3417 (median=1338 mean=1451.5 std=807.2)

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Pregnancy Complication Reviewer 1 Form: FORM CODE=PCD1, VERSION=1

ID		HCHS Participant ID
N	Value	Description
127	Present	Text suppressed

VISIT		Label information missing in the dataset
N	Value	Description
14	1	Visit 1
15	2	Visit 2
24	3	Visit 3
17	4	Visit 4
17	5	Visit 5
15	6	Visit 6
18	7	Visit 7
1	8	Visit 8
6	9	Visit 9

OCCURRENCE		Label information missing in the dataset
N	Value	Description
109	1	Occurrence 1
14	2	Occurrence 2
3	3	Occurrence 3
1	4	Occurrence 4

PCD11		PCD reviews required
N	Value	Description
53	1	1= Preeclampsia/Eclampsia
59	2	2= Gestational Diabetes
15	3	3= Both Preeclampsia/Eclampsia and Gestational Diabetes

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<i>PCD12</i>		<i>prior hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
17	0	0= No
12	1	1= Yes
0	9	9=NR

<i>PCD12A1</i>		<i>SBP Prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
34	.S	Legitimately skipped
6	0	0= No
5	1	1= Yes
1	9	9=NR/unsure

<i>PCD12A2</i>		<i>self-report hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
34	.S	Legitimately skipped
6	0	0= No
5	1	1= Yes
1	9	9=NR/unsure

<i>PCD12A3</i>		<i>documentation of hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
34	.S	Legitimately skipped
3	0	0= No
9	1	1= Yes
0	9	9=NR/unsure

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<i>PCD13</i>		<i>SBP less than 20 weeks</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
83		Missing
17	.S	Legitimately skipped
3	0	0= No
2	1	1= Yes
22	9	9=NR/unsure

<i>PCD13A</i>		<i>documentation of treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
84		Missing
17	.S	Legitimately skipped
4	0	0= No
1	1	1= Yes
21	9	9=NR/unsure

<i>PCD14</i>		<i>hypertensive blood pressures 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
83		Missing
17	.S	Legitimately skipped
9	0	0= No
4	1	1= Yes
14	9	9=Unknown

<i>PCD15</i>		<i>proteinuria documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
17	.S	Legitimately skipped
2	1	1= >= 0.3g / 24h urine
7	2	2= >=1+ urine dipstick
2	3	3=urine protein:creatinine ratio documented as "positive" by local lab criteria
1	4	4=more than one of the lab criteria of 1-3
2	5	5= Presence of proteinuria documented in notes, but specific lab test not noted
8	6	6= Negative test for proteinuria
6	9	9= NR/unsure

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<i>PCD16A</i>		<i>two occasions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
12	0	0= No
6	1	1= Yes
11	9	9=NR/unsure

<i>PCD16A1</i>		<i>more than 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
40	.S	Legitimately skipped
1	0	0= No
3	1	1= Yes
2	9	9=NR/unsure

<i>PCD16B</i>		<i>anti-hypertensive medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
13	0	0= No
6	1	1= Yes
10	9	9=NR/unsure

<i>PCD16C</i>		<i>elevated liver enzymes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
20	0	0= No
1	1	1= Yes
8	9	9=NR/unsure

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<i>PCD16D</i>		<i>persistent pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
21	0	0= No
0	1	1= Yes
8	9	9=NR/unsure

<i>PCD16E</i>		<i>thrombocytopenia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
26	0	0= No
0	1	1= Yes
3	9	9=NR/unsure

<i>PCD16F</i>		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
20	0	0= No
0	1	1= Yes
9	9	9=NR/unsure

<i>PCD16G</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
4	0	0= No
1	1	1= Yes
24	9	9=NR/unsure

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<i>PCD16H</i>		<i>new onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
18	0	0= No
3	1	1= Yes
8	9	9=NR/unsure

<i>PCD17</i>		<i>intrapartum magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
6	0	0= No
5	1	1= Yes
18	9	9=NR/Unsure

<i>PCD17A</i>		<i>magnesium indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
41	.S	Legitimately skipped
1	0	0= No
4	1	1= Yes
0	9	9=NR/Unsure

<i>PCD18</i>		<i>grand mal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
0	1	1= A grand mal seizure in the absence of any known causes for seizure
0	2	2= A grand mal seizure in the setting of a brain lesion or diagnosis of epilepsy
0	3	3= A grand mal seizure without specifics to code as 1 or 2
28	4	4= No grand mal seizure noted
0	5	5= Seizure, but unsure if grand mal
1	9	9= Unsure

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<i>PCD19</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
2	1	1= Definite
4	2	2= Probable
6	3	3= Possible
14	4	4= Unlikely
3	5	5= Unclassifiable

<i>PCD110</i>		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
3	1	1= Definite
8	2	2= Probable
3	3	3= Possible
13	4	4= Unlikely
2	5	5= Unclassifiable

<i>PCD111</i>		<i>severe features</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
1	1	1= Definite
3	2	2= Probable
1	3	3= Possible
24	4	4= Unlikely
0	5	5= Unclassifiable

Chapter 1: Endpoint Ascertainment

Section F: Pregnancy Complication Ascertainment

<i>PCD112</i>		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
81		Missing
17	.S	Legitimately skipped
0	1	1= Definite
0	2	2= Probable
0	3	3= Possible
29	4	4= Unlikely
0	5	5= Unclassifiable

<i>PCD113</i>		<i>pregestatal DM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
21	.S	Legitimately skipped
16	0	0= No
8	1	1= Yes
0	9	9=Unknown

<i>PCD114A</i>		<i>fasting glucose ?126</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
37	.S	Legitimately skipped
5	0	0= No
3	1	1= Yes
0	9	9=NR/unsure

<i>PCD114B</i>		<i>2h glucose ? 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
37	.S	Legitimately skipped
2	0	0= No
2	1	1= Yes
4	9	9=NR/unsure

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<i>PCD114C</i>		<i>prior HbA1c ? 6.5</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
37	.S	Legitimately skipped
3	0	0= No
4	1	1= Yes
1	9	9=NR/unsure

<i>PCD114D</i>		<i>pregestational DM documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
83		Missing
37	.S	Legitimately skipped
4	0	0= No
3	1	1= Yes
0	9	9=NR/unsure

<i>PCD114E</i>		<i>DM self-report</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
37	.S	Legitimately skipped
4	0	0= No
3	1	1= Yes
1	9	9=NR/unsure

<i>PCD115</i>		<i>glucose intolerance present</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
21	.S	Legitimately skipped
2	0	0= No
7	1	1= Yes
15	9	9=NR

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<i>PCD115A1</i>		<i>3 hour OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
38	.S	Legitimately skipped
0	0	0= No
3	1	1= Yes
4	9	9=NR/unsure

<i>PCD115A2</i>		<i>2h 75g OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
38	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
7	9	9=NR/unsure

<i>PCD115A3</i>		<i>50g GLT > 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
38	.S	Legitimately skipped
2	0	0= No
0	1	1= Yes
5	9	9=NR/unsure

<i>PCD115A4</i>		<i>two or more glucose > 125</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
38	.S	Legitimately skipped
0	0	0= No
1	1	1= Yes
6	9	9=NR/unsure

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<i>PCD115A5</i>		<i>abnormal or elevated OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
38	.S	Legitimately skipped
1	0	0= No
1	1	1= Yes
5	9	9=NR/unsure

<i>PCD115B</i>		<i>Non-diagnostic glucose intolerance</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
28	.S	Legitimately skipped
2	0	0= No
1	1	1= Yes
14	9	9=NR/unsure

<i>PCD115B1</i>		<i>elevated 1 hour GLT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
44	.S	Legitimately skipped
0	0	0= No
0	1	1= Yes
1	9	9=NR/unsure

<i>PCD115B2</i>		<i>evidence of abnormal glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
44	.S	Legitimately skipped
0	0	0= No
1	1	1= Yes
0	9	9=NR/unsure

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<i>PCD116</i>		<i>treatment with insulin or oral</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
21	.S	Legitimately skipped
2	0	0= No
20	1	1= Yes
2	9	9=NR

<i>PCD117</i>		<i>GDM diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
21	.S	Legitimately skipped
6	0	0= No
18	1	1= Yes
0	9	9=NR

<i>PCD117A</i>		<i>physician mention GDM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
21	.S	Legitimately skipped
6	0	0= No
18	1	1= Yes
0	9	9=NR

<i>PCD118</i>		<i>Gestational Diabetes classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
21	.S	Legitimately skipped
13	1	1=Definite
2	2	2=Probable
1	3	3=Possible
2	4	4=Unlikely
1	5	5=Unclassifiable
5	6	6=Pregestational

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<i>PCD119</i>		<i>Comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
105		Missing
22	Present	Text suppressed

<i>PCD10A_FUTIME</i>		<i>Futime from Randomization date for (PCD10A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
82		Missing
45	Range	2563 - 3896 (median=3173 mean=3217.3 std=331.3)

<i>PCD10D_FUTIME</i>		<i>Futime from Randomization date for (PCD10D)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	289 - 3197 (median=2476 mean=2484.0 std=424.3)

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Pregnancy Complication Reviewer 2 Form: FORM CODE=PCD2, VERSION=1

ID		HCHS Participant ID
N	Value	Description
127	Present	Text suppressed

VISIT		Label information missing in the dataset
N	Value	Description
14	1	Visit 1
15	2	Visit 2
24	3	Visit 3
17	4	Visit 4
17	5	Visit 5
15	6	Visit 6
18	7	Visit 7
1	8	Visit 8
6	9	Visit 9

OCCURRENCE		Label information missing in the dataset
N	Value	Description
109	1	Occurrence 1
14	2	Occurrence 2
3	3	Occurrence 3
1	4	Occurrence 4

PCD219		Comments
N	Value	Description
93		Missing
34	Present	Text suppressed

PCD21		PCD reviews required
N	Value	Description
53	1	1= Preeclampsia/Eclampsia
59	2	2= Gestational Diabetes
15	3	3= Both Preeclampsia/Eclampsia and Gestational Diabetes

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<i>PCD22</i>		<i>prior hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
39	0	0= No
17	1	1= Yes
3	9	9=NR

<i>PCD22A1</i>		<i>SBP Prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
101	.S	Legitimately skipped
11	0	0= No
4	1	1= Yes
3	9	9=NR/unsure

<i>PCD22A2</i>		<i>self-report hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
101	.S	Legitimately skipped
8	0	0= No
4	1	1= Yes
6	9	9=NR/unsure

<i>PCD22A3</i>		<i>documentation of hypertensions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
101	.S	Legitimately skipped
3	0	0= No
13	1	1= Yes
2	9	9=NR/unsure

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<i>PCD23</i>		<i>SBP less than 20 weeks</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
22	0	0= No
0	1	1= Yes
38	9	9=NR/unsure

<i>PCD23A</i>		<i>documentation of treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10		Missing
59	.S	Legitimately skipped
24	0	0= No
2	1	1= Yes
32	9	9=NR/unsure

<i>PCD24</i>		<i>hypertensive blood pressures 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
11		Missing
59	.S	Legitimately skipped
14	0	0= No
12	1	1= Yes
31	9	9=Unknown

<i>PCD25</i>		<i>proteinuria documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
3	1	1= >= 0.3g / 24h urine
7	2	2= >=1+ urine dipstick
4	3	3=urine protein:creatinine ratio documented as "positive" by local lab criteria
3	4	4=more than one of the lab criteria of 1-3
2	5	5= Presence of proteinuria documented in notes, but specific lab test not noted
16	6	6= Negative test for proteinuria
24	9	9= NR/unsure

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<i>PCD26A</i>		<i>two occasions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
26	0	0= No
6	1	1= Yes
28	9	9=NR/unsure

<i>PCD26A1</i>		<i>more than 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
113	.S	Legitimately skipped
2	0	0= No
4	1	1= Yes
0	9	9=NR/unsure

<i>PCD26B</i>		<i>anti-hypertensive medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
30	0	0= No
9	1	1= Yes
21	9	9=NR/unsure

<i>PCD26C</i>		<i>elevated liver enzymes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
40	0	0= No
3	1	1= Yes
17	9	9=NR/unsure

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<i>PCD26D</i>		<i>persistent pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
44	0	0= No
0	1	1= Yes
16	9	9=NR/unsure

<i>PCD26E</i>		<i>thrombocytopenia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
43	0	0= No
0	1	1= Yes
17	9	9=NR/unsure

<i>PCD26F</i>		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
41	0	0= No
1	1	1= Yes
18	9	9=NR/unsure

<i>PCD26G</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
38	0	0= No
1	1	1= Yes
21	9	9=NR/unsure

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<i>PCD26H</i>		<i>new onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
59	.S	Legitimately skipped
42	0	0= No
7	1	1= Yes
11	9	9=NR/unsure

<i>PCD27</i>		<i>intrapartum magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
17	0	0= No
13	1	1= Yes
29	9	9=NR/Unsure

<i>PCD27A</i>		<i>magnesium indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
113	.S	Legitimately skipped
2	0	0= No
3	1	1= Yes
8	9	9=NR/Unsure

<i>PCD28</i>		<i>grand mal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
0	1	1= A grand mal seizure in the absence of any known causes for seizure
0	2	2= A grand mal seizure in the setting of a brain lesion or diagnosis of epilepsy
0	3	3= A grand mal seizure without specifics to code as 1 or 2
56	4	4= No grand mal seizure noted
0	5	5= Seizure, but unsure if grand mal
3	9	9= Unsure

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<i>PCD29</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
6	1	1= Definite
11	2	2= Probable
12	3	3= Possible
21	4	4= Unlikely
9	5	5= Unclassifiable

<i>PCD210</i>		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
7	1	1= Definite
13	2	2= Probable
12	3	3= Possible
21	4	4= Unlikely
6	5	5= Unclassifiable

<i>PCD211</i>		<i>severe features</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
3	1	1= Definite
2	2	2= Probable
4	3	3= Possible
47	4	4= Unlikely
3	5	5= Unclassifiable

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<i>PCD212</i>		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
59	.S	Legitimately skipped
0	1	1= Definite
0	2	2= Probable
1	3	3= Possible
57	4	4= Unlikely
1	5	5= Unclassifiable

<i>PCD213</i>		<i>pregestatal DM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
53	.S	Legitimately skipped
50	0	0= No
12	1	1= Yes
4	9	9=Unknown

<i>PCD214A</i>		<i>fasting glucose ?126</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
107	.S	Legitimately skipped
3	0	0= No
3	1	1= Yes
6	9	9=NR/unsure

<i>PCD214B</i>		<i>2h glucose ? 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
107	.S	Legitimately skipped
2	0	0= No
2	1	1= Yes
8	9	9=NR/unsure

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<i>PCD214C</i>		<i>prior HbA1c ? 6.5</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
107	.S	Legitimately skipped
3	0	0= No
4	1	1= Yes
5	9	9=NR/unsure

<i>PCD214D</i>		<i>pregestational DM documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
107	.S	Legitimately skipped
0	0	0= No
9	1	1= Yes
2	9	9=NR/unsure

<i>PCD214E</i>		<i>DM self-report</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
107	.S	Legitimately skipped
3	0	0= No
4	1	1= Yes
5	9	9=NR/unsure

<i>PCD215</i>		<i>glucose intolerance present</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
53	.S	Legitimately skipped
5	0	0= No
24	1	1= Yes
36	9	9=NR

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<i>PCD215A1</i>		<i>3 hour OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
94	.S	Legitimately skipped
4	0	0= No
7	1	1= Yes
14	9	9=NR/unsure

<i>PCD215A2</i>		<i>2h 75g OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
94	.S	Legitimately skipped
3	0	0= No
5	1	1= Yes
17	9	9=NR/unsure

<i>PCD215A3</i>		<i>50g GLT > 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
94	.S	Legitimately skipped
8	0	0= No
1	1	1= Yes
16	9	9=NR/unsure

<i>PCD215A4</i>		<i>two or more glucose > 125</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
94	.S	Legitimately skipped
4	0	0= No
0	1	1= Yes
21	9	9=NR/unsure

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<i>PCD215A5</i>		<i>abnormal or elevated OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
94	.S	Legitimately skipped
8	0	0= No
0	1	1= Yes
17	9	9=NR/unsure

<i>PCD215B</i>		<i>Non-diagnostic glucose intolerance</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
77	.S	Legitimately skipped
6	0	0= No
7	1	1= Yes
29	9	9=NR/unsure

<i>PCD215B1</i>		<i>elevated 1 hour GLT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
112	.S	Legitimately skipped
0	0	0= No
1	1	1= Yes
6	9	9=NR/unsure

<i>PCD215B2</i>		<i>evidence of abnormal glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
112	.S	Legitimately skipped
0	0	0= No
3	1	1= Yes
4	9	9=NR/unsure

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<i>PCD216</i>		<i>treatment with insulin or oral</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
53	.S	Legitimately skipped
11	0	0= No
39	1	1= Yes
15	9	9=NR

<i>PCD217</i>		<i>GDM diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
53	.S	Legitimately skipped
7	0	0= No
55	1	1= Yes
3	9	9=NR

<i>PCD217A</i>		<i>physician mention GDM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10		Missing
53	.S	Legitimately skipped
7	0	0= No
54	1	1= Yes
3	9	9=NR

<i>PCD218</i>		<i>Gestational Diabetes classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8		Missing
53	.S	Legitimately skipped
25	1	1=Definite
20	2	2=Probable
3	3	3=Possible
6	4	4=Unlikely
6	5	5=Unclassifiable
6	6	6=Pregestational

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<i>PCD20A_FUTIME</i>		<i>Futime from Randomization date for (PCD20A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
26		Missing
101	Range	2585 - 3868 (median=3147 mean=3192.2 std=293.5)

<i>PCD20D_FUTIME</i>		<i>Futime from Randomization date for (PCD20D)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	289 - 3197 (median=2476 mean=2499.9 std=405.2)

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Pregnancy Complication Adjudication Reviewer Form: FORM CODE=PCD3, VERSION=1

ID		HCHS Participant ID
N	Value	Description
124	Present	Text suppressed

VISIT		Label information missing in the dataset
N	Value	Description
13	1	Visit 1
15	2	Visit 2
24	3	Visit 3
17	4	Visit 4
17	5	Visit 5
15	6	Visit 6
17	7	Visit 7
1	8	Visit 8
5	9	Visit 9

OCCURRENCE		Label information missing in the dataset
N	Value	Description
107	1	Occurrence 1
13	2	Occurrence 2
3	3	Occurrence 3
1	4	Occurrence 4

PCD319		Comments
N	Value	Description
53		Missing
71	Present	Text suppressed

PCD31		PCD reviews required
N	Value	Description
48	1	1= Preeclampsia/Eclampsia
52	2	2= Gestational Diabetes
24	3	3= Both Preeclampsia/Eclampsia and Gestational Diabetes

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<i>PCD32</i>		<i>prior hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
44	0	0= No
27	1	1= Yes
1	9	9=NR

<i>PCD32A1</i>		<i>SBP Prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
97	.S	Legitimately skipped
19	0	0= No
7	1	1= Yes
0	9	9=NR/unsure

<i>PCD32A2</i>		<i>self-report hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
97	.S	Legitimately skipped
19	0	0= No
7	1	1= Yes
0	9	9=NR/unsure

<i>PCD32A3</i>		<i>documentation of hypertensions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
97	.S	Legitimately skipped
6	0	0= No
21	1	1= Yes
0	9	9=NR/unsure

<i>PCD33</i>		<i>SBP less than 20 weeks</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
52	.S	Legitimately skipped
5	0	0= No
1	1	1= Yes
64	9	9=NR/unsure

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<i>PCD33A</i>		<i>documentation of treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9		Missing
52	.S	Legitimately skipped
3	0	0= No
0	1	1= Yes
60	9	9=NR/unsure

<i>PCD34</i>		<i>hypertensive blood pressures 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
10	0	0= No
25	1	1= Yes
37	9	9=Unknown

<i>PCD35</i>		<i>proteinuria documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
52	.S	Legitimately skipped
6	1	1= >= 0.3g / 24h urine
10	2	2= >=1+ urine dipstick
8	3	3=urine protein:creatinine ratio documented as "positive" by local lab criteria
2	4	4=more than one of the lab criteria of 1-3
2	5	5= Presence of proteinuria documented in notes, but specific lab test not noted
20	6	6= Negative test for proteinuria
23	9	9= NR/unsure

<i>PCD36A</i>		<i>two occasions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
26	0	0= No
8	1	1= Yes
38	9	9=NR/unsure

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<i>PCD36A1</i>		<i>more than 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
116	.S	Legitimately skipped
2	0	0= No
3	1	1= Yes
2	9	9=NR/unsure

<i>PCD36B</i>		<i>anti-hypertensive medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
27	0	0= No
8	1	1= Yes
37	9	9=NR/unsure

<i>PCD36C</i>		<i>elevated liver enzymes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
42	0	0= No
3	1	1= Yes
27	9	9=NR/unsure

<i>PCD36D</i>		<i>persistent pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
31	0	0= No
2	1	1= Yes
39	9	9=NR/unsure

<i>PCD36E</i>		<i>thrombocytopenia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
48	0	0= No
0	1	1= Yes
24	9	9=NR/unsure

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<i>PCD36F</i>		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
41	0	0= No
2	1	1= Yes
29	9	9=NR/unsure

<i>PCD36G</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
5	0	0= No
1	1	1= Yes
66	9	9=NR/unsure

<i>PCD36H</i>		<i>new onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
27	0	0= No
11	1	1= Yes
34	9	9=NR/unsure

<i>PCD37</i>		<i>intrapartum magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
18	0	0= No
17	1	1= Yes
37	9	9=NR/Unsure

<i>PCD37A</i>		<i>magnesium indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2		Missing
107	.S	Legitimately skipped
0	0	0= No
15	1	1= Yes
0	9	9=NR/Unsure

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<i>PCD38</i>		<i>grand mal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
0	1	1= A grand mal seizure in the absence of any known causes for seizure
0	2	2= A grand mal seizure in the setting of a brain lesion or diagnosis of epilepsy
0	3	3= A grand mal seizure without specifics to code as 1 or 2
72	4	4= No grand mal seizure noted
0	5	5= Seizure, but unsure if grand mal
0	9	9= Unsure

<i>PCD39</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
2	1	1= Definite
11	2	2= Probable
10	3	3= Possible
44	4	4= Unlikely
5	5	5= Unclassifiable

<i>PCD310</i>		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
14	1	1= Definite
11	2	2= Probable
6	3	3= Possible
33	4	4= Unlikely
8	5	5= Unclassifiable

<i>PCD311</i>		<i>severe features</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
7	1	1= Definite
5	2	2= Probable
2	3	3= Possible
50	4	4= Unlikely
8	5	5= Unclassifiable

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<i>PCD312</i>		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
0	1	1= Definite
0	2	2= Probable
0	3	3= Possible
72	4	4= Unlikely
0	5	5= Unclassifiable

<i>PCD313</i>		<i>pregestatal DM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
48	.S	Legitimately skipped
58	0	0= No
18	1	1= Yes
0	9	9=Unknown

<i>PCD314A</i>		<i>fasting glucose ?126</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
106	.S	Legitimately skipped
11	0	0= No
7	1	1= Yes
0	9	9=NR/unsure

<i>PCD314B</i>		<i>2h glucose ? 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
106	.S	Legitimately skipped
6	0	0= No
6	1	1= Yes
6	9	9=NR/unsure

<i>PCD314C</i>		<i>prior HbA1c ? 6.5</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
106	.S	Legitimately skipped
9	0	0= No
9	1	1= Yes
0	9	9=NR/unsure

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<i>PCD314D</i>		<i>pregestational DM documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
106	.S	Legitimately skipped
6	0	0= No
11	1	1= Yes
1	9	9=NR/unsure

<i>PCD314E</i>		<i>DM self-report</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
106	.S	Legitimately skipped
11	0	0= No
6	1	1= Yes
1	9	9=NR/unsure

<i>PCD315</i>		<i>glucose intolerance present</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
48	.S	Legitimately skipped
7	0	0= No
14	1	1= Yes
55	9	9=NR

<i>PCD315A1</i>		<i>3 hour OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
110	.S	Legitimately skipped
6	0	0= No
6	1	1= Yes
1	9	9=NR/unsure

<i>PCD315A2</i>		<i>2h 75g OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
110	.S	Legitimately skipped
11	0	0= No
0	1	1= Yes
2	9	9=NR/unsure

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<i>PCD315A3</i>		<i>50g GLT > 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
110	.S	Legitimately skipped
9	0	0= No
3	1	1= Yes
1	9	9=NR/unsure

<i>PCD315A4</i>		<i>two or more glucose > 125</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
110	.S	Legitimately skipped
11	0	0= No
0	1	1= Yes
2	9	9=NR/unsure

<i>PCD315A5</i>		<i>abnormal or elevated OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
110	.S	Legitimately skipped
8	0	0= No
4	1	1= Yes
2	9	9=NR/unsure

<i>PCD315B</i>		<i>Non-diagnostic glucose intolerance</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
62	.S	Legitimately skipped
3	0	0= No
7	1	1= Yes
51	9	9=NR/unsure

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<i>PCD315B1</i>		<i>elevated 1 hour GLT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
116	.S	Legitimately skipped
0	0	0= No
5	1	1= Yes
2	9	9=NR/unsure

<i>PCD315B2</i>		<i>evidence of abnormal glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1		Missing
116	.S	Legitimately skipped
4	0	0= No
2	1	1= Yes
1	9	9=NR/unsure

<i>PCD316</i>		<i>treatment with insulin or oral</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
48	.S	Legitimately skipped
24	0	0= No
43	1	1= Yes
9	9	9=NR

<i>PCD317</i>		<i>GDM diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
48	.S	Legitimately skipped
19	0	0= No
57	1	1= Yes
0	9	9=NR

<i>PCD317A</i>		<i>physician mention GDM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
48	.S	Legitimately skipped
20	0	0= No
56	1	1= Yes
0	9	9=NR

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<i>PCD318</i>		<i>Gestational Diabetes classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
48	.S	Legitimately skipped
27	1	1=Definite
21	2	2=Probable
1	3	3=Possible
7	4	4=Unlikely
3	5	5=Unclassifiable
17	6	6=Pregestational

<i>PCD30A_FUTIME</i>		<i>Futime from Randomization date for (PCD30A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
124	Range	2855 - 3981 (median=3261 mean=3317.1 std=293.0)

<i>PCD30D_FUTIME</i>		<i>Futime from Randomization date for (PCD30D)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
124	Range	212 - 3405 (median=1346.5 mean=1449.21 std=782.42)

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HCHS incident MI Event Derived Variable file 2008-2021: FORM CODE=INCIDENT_MI_EVNT, VERSION=2008_2021_INV1

<i>INCIDENT_MI_2021</i>		<i>Incident Myocardial Infarction (MI) Event by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16171	0	No
244	1	Yes

<i>RECUR_MI_2021_1</i>		<i>MI as 1st Recur MI 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16383	0	No
32	1	Yes

<i>RECUR_MI_2021_2</i>		<i>MI as 2nd Recur MI 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16407	0	No
8	1	Yes

<i>RECUR_MI_2021_3</i>		<i>MI as 3rd Recur MI 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

<i>FUTIME_MI</i>		<i>Follow up time of MI event, censor or the end of follow-up by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4270 mean=4146.8 std=739.7)

<i>FUTIME_MI_1</i>		<i>Follow up time 1st Recur MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4275 mean=4168.9 std=700.0)

<i>FUTIME_MI_2</i>		<i>Follow up time 2nd Recur MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.3 std=698.0)

<i>FUTIME_MI_3</i>		<i>Follow up time 3rd Recur MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

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<i>PRECHD_NO_ANGINA</i>		<i>Prevalent Coronary Heart Disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
77		Missing
15480	0	No
858	1	Yes

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HCHS incident HF Event Derived Variable file 2008-2021: FORM CODE=INCIDENT_HF_EVNT, VERSION=2008_2021_INV1

<i>INCIDENT_HF_2021</i>		<i>Incident Heart Failure (HF) Event by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16155	0	No
260	1	Yes

<i>RECUR_HF_2021_1</i>		<i>HF as 1st Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16330	0	No
85	1	Yes

<i>RECUR_HF_2021_2</i>		<i>HF as 2nd Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16372	0	No
43	1	Yes

<i>RECUR_HF_2021_3</i>		<i>HF as 3rd Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16392	0	No
23	1	Yes

<i>RECUR_HF_2021_4</i>		<i>HF as 4th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16403	0	No
12	1	Yes

<i>RECUR_HF_2021_5</i>		<i>HF as 5th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16409	0	No
6	1	Yes

<i>RECUR_HF_2021_6</i>		<i>HF as 6th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16411	0	No
4	1	Yes

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<i>RECUR_HF_2021_7</i>		<i>HF as 7th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

<i>RECUR_HF_2021_8</i>		<i>HF as 8th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

<i>RECUR_HF_2021_9</i>		<i>HF as 9th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>RECUR_HF_2021_10</i>		<i>HF as 10th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>RECUR_HF_2021_11</i>		<i>HF as 11th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>RECUR_HF_2021_12</i>		<i>HF as 12th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>RECUR_HF_2021_13</i>		<i>HF as 13th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

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<i>FUTIME_HF</i>		<i>Follow up time of HF event, censor or the end of follow-up by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4270 mean=4147.8 std=739.3)

<i>FUTIME_HF_1</i>		<i>Follow up time 1st Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4273 mean=4163.4 std=710.3)

<i>FUTIME_HF_2</i>		<i>Follow up time 2nd Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4275 mean=4167.8 std=702.3)

<i>FUTIME_HF_3</i>		<i>Follow up time 3rd Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4169.3 std=699.8)

<i>FUTIME_HF_4</i>		<i>Follow up time 4th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.3 std=698.2)

<i>FUTIME_HF_5</i>		<i>Follow up time 5th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.4 std=697.9)

<i>FUTIME_HF_6</i>		<i>Follow up time 6th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.5 std=697.7)

<i>FUTIME_HF_7</i>		<i>Follow up time 7th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

<i>FUTIME_HF_8</i>		<i>Follow up time 8th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

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<i>FUTIME_HF_9</i>		<i>Follow up time 9th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_10</i>		<i>Follow up time 10th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_11</i>		<i>Follow up time 11th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_12</i>		<i>Follow up time 12th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_13</i>		<i>Follow up time 13th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>HFPEF</i>		<i>HFpEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16250	0	No
165	1	Yes

<i>HFrEF</i>		<i>HFrEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16250	0	No
165	1	Yes

<i>MHEA5</i>		<i>Has a doctor ever said that you had heart failure?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
88		Missing
10	.Q	DK/Refuse
16008	0	No
309	1	Yes

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<i>PRECHD_NO_ANGINA</i>		<i>Prevalent Coronary Heart Disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
77		Missing
15480	0	No
858	1	Yes

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Section A: Incident Event Derived Variable File

HCHS incident PUL Event Derived Variable file 2008-2018: FORM CODE=INCIDENT_PUL_EVNT, VERSION=2008_2018_INV1

ASTHMA_PLD		PLD define or highly probable Asthma by 2018
N	Value	Description
16285		Missing
0	0	No
130	1	Yes

COPD_PLD		PLD define or highly probable COPD by 2018
N	Value	Description
16394		Missing
0	0	No
21	1	Yes

CLRD_2018		Chronic Lower Respiratory Disease by 2018
N	Value	Description
16269	0	No
146	1	Yes

CLRD_ASTHMA_2018		CLRD due to Asthma by 2018
N	Value	Description
16285	0	No
130	1	Yes

CLRD_COPD_2018		CLRD due to COPD by 2018
N	Value	Description
16394	0	No
21	1	Yes

EXACERBATION_CLRD		Exacerbation of CLRD by 2018
N	Value	Description
16307	0	No
108	1	Yes

FUTIME_CLRD		Follow up time of CLRD event, censor or the end of follow-up by 2018
N	Value	Description
16415	Range	0 - 3954 (median=3190 mean=3144.4 std=528.2)

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<i>FUTIME_CLRD_ASTH MA</i>		<i>Follow up time of ASTHMA event, censor or the end of follow-up by 2018</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 3954 (median=3190 mean=3145.7 std=526.1)

<i>FUTIME_CLRD_COPD</i>		<i>Follow up time of COPD event, censor or the end of follow-up by 2018</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 3954 (median=3195 mean=3160.7 std=491.3)

<i>ASTHMA_EVER</i>		<i>Ever had Asthma (self-report)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
272		Missing
13439	0	No
2637	1	Yes
67	9	Unknown

<i>COPD_EVER</i>		<i>Self-report of ever had COPD/Emph or Chronic Bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
339		Missing
14987	0	No
1089	1	Yes

<i>PREVALENCE_CLRD</i>		<i>Prevalence of CLRD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13166	0	No
3249	1	Yes

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Section A: Incident Event Derived Variable File

HCHS incident STR Event Derived Variable file 2008-2021: FORM CODE=INCIDENT_STR_EVNT, VERSION=2008_2021_INV1

<i>INCIDENT_STR_2021</i>		<i>Incident Stroke (STR) Event by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16247	0	No
168	1	Yes

<i>RECUR_STR_2021_1</i>		<i>STR as Recur STR 2021 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16390	0	No
25	1	Yes

<i>RECUR_STR_2021_2</i>		<i>STR as Recur STR 2021 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

<i>FUTIME_STR</i>		<i>Follow up time of STR event, censor or the end of follow-up by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4271 mean=4151.1 std=733.9)

<i>FUTIME_STR_1</i>		<i>Recur FUTIME STR 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4275 mean=4168.4 std=702.1)

<i>FUTIME_STR_2</i>		<i>Recur FUTIME STR 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

<i>MHEA10</i>		<i>Has a doctor ever said that you had a stroke?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
87		Missing
2	.Q	DK/Refuse
16074	0	No
252	1	Yes

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Section A: Incident Event Derived Variable File

HCHS incident PRC Event Derived Variable file all years: FORM CODE=INCIDENT_PRC_EVNT, VERSION=ALL_INV1

<i>PRE_GHTN</i>		<i>Prevalence of Pregnancy only HBP or GHTN (Preeclampsia or Eclampsia) reported by Visit 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8408		Missing
7385	0	No
622	1	Yes

<i>PRE_GDM</i>		<i>Prevalence GDM reported by Visit 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
9377		Missing
6349	0	No
689	1	Yes

<i>PRE_PCD</i>		<i>Prevalence of Pregnancy Complications Event(Pregnancy only HBP GHTN GDM) by Visit 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8148		Missing
7059	0	No
1208	1	Yes

<i>INCIDENT_GHTN</i>		<i>Pregnancy Complications Diagnosis of GHTN (Preeclampsia to Eclampsia) Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16390	0	No
25	1	Yes

<i>INCIDENT_GDM</i>		<i>Pregnancy Complications Diagnosis of GDM Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16368	0	No
47	1	Yes

<i>INCIDENT_PCD</i>		<i>Pregnancy Complications Diagnosis (PCD) Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16301	0	No
114	1	Yes

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Section B: Combined Event Derived Variable File for All Incident Events
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HCHS Combined Incident Event Derived Variable file 2008-2021: FORM CODE=EVENT_PART_DERV, VERSION=2008_2021_INV1

ASTHMA_PLD		PLD define or highly probable Asthma by 2018
N	Value	Description
16285		Missing
0	0	No
130	1	Yes

COPD_PLD		PLD define or highly probable COPD by 2018
N	Value	Description
16394		Missing
0	0	No
21	1	Yes

CLRD_2018		Chronic Lower Respiratory Disease by 2018
N	Value	Description
16269	0	No
146	1	Yes

CLRD_ASTHMA_2018		CLRD due to Asthma by 2018
N	Value	Description
16285	0	No
130	1	Yes

CLRD_COPD_2018		CLRD due to COPD by 2018
N	Value	Description
16394	0	No
21	1	Yes

EXACERBATION_CLRD		Exacerbation of CLRD by 2018
N	Value	Description
16307	0	No
108	1	Yes

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<i>FUTIME_CLRD</i>		<i>Follow up time of CLRD event, censor or the end of follow-up by 2018</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 3954 (median=3190 mean=3144.4 std=528.2)

<i>FUTIME_CLRD_ASTH MA</i>		<i>Follow up time of ASTHMA event, censor or the end of follow-up by 2018</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 3954 (median=3190 mean=3145.7 std=526.1)

<i>FUTIME_CLRD_COPD</i>		<i>Follow up time of COPD event, censor or the end of follow-up by 2018</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 3954 (median=3195 mean=3160.7 std=491.3)

<i>INCIDENT_MI_2021</i>		<i>Incident Myocardial Infarction (MI) Event by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16171	0	No
244	1	Yes

<i>RECUR_MI_2021_1</i>		<i>MI as 1st Recur MI 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16383	0	No
32	1	Yes

<i>RECUR_MI_2021_2</i>		<i>MI as 2nd Recur MI 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16407	0	No
8	1	Yes

<i>RECUR_MI_2021_3</i>		<i>MI as 3rd Recur MI 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

<i>FUTIME_MI</i>		<i>Follow up time of MI event, censor or the end of follow-up by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4270 mean=4146.8 std=739.7)

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<i>FUTIME_MI_1</i>		<i>Follow up time 1st Recur MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4275 mean=4168.9 std=700.0)

<i>FUTIME_MI_2</i>		<i>Follow up time 2nd Recur MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.3 std=698.0)

<i>FUTIME_MI_3</i>		<i>Follow up time 3rd Recur MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

<i>INCIDENT_HF_2021</i>		<i>Incident Heart Failure (HF) Event by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16155	0	No
260	1	Yes

<i>RECUR_HF_2021_1</i>		<i>HF as 1st Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16330	0	No
85	1	Yes

<i>RECUR_HF_2021_2</i>		<i>HF as 2nd Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16372	0	No
43	1	Yes

<i>RECUR_HF_2021_3</i>		<i>HF as 3rd Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16392	0	No
23	1	Yes

<i>RECUR_HF_2021_4</i>		<i>HF as 4th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16403	0	No
12	1	Yes

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<i>RECUR_HF_2021_5</i> HF as 5th Recur HF 2021		
<i>N</i>	<i>Value</i>	<i>Description</i>
16409	0	No
6	1	Yes

<i>RECUR_HF_2021_6</i> HF as 6th Recur HF 2021		
<i>N</i>	<i>Value</i>	<i>Description</i>
16411	0	No
4	1	Yes

<i>RECUR_HF_2021_7</i> HF as 7th Recur HF 2021		
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

<i>RECUR_HF_2021_8</i> HF as 8th Recur HF 2021		
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

<i>RECUR_HF_2021_9</i> HF as 9th Recur HF 2021		
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>RECUR_HF_2021_10</i> HF as 10th Recur HF 2021		
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>RECUR_HF_2021_11</i> HF as 11th Recur HF 2021		
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

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Section B: Combined Event Derived Variable File for All Incident Events (N=16,415)

<i>RECUR_HF_2021_12</i>		<i>HF as 12th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>RECUR_HF_2021_13</i>		<i>HF as 13th Recur HF 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16414	0	No
1	1	Yes

<i>FUTIME_HF</i>		<i>Follow up time of HF event, censor or the end of follow-up by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4270 mean=4147.8 std=739.3)

<i>FUTIME_HF_1</i>		<i>Follow up time 1st Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4273 mean=4163.4 std=710.3)

<i>FUTIME_HF_2</i>		<i>Follow up time 2nd Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4275 mean=4167.8 std=702.3)

<i>FUTIME_HF_3</i>		<i>Follow up time 3rd Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4169.3 std=699.8)

<i>FUTIME_HF_4</i>		<i>Follow up time 4th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.3 std=698.2)

<i>FUTIME_HF_5</i>		<i>Follow up time 5th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.4 std=697.9)

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<i>FUTIME_HF_6</i>		<i>Follow up time 6th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.5 std=697.7)

<i>FUTIME_HF_7</i>		<i>Follow up time 7th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

<i>FUTIME_HF_8</i>		<i>Follow up time 8th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

<i>FUTIME_HF_9</i>		<i>Follow up time 9th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_10</i>		<i>Follow up time 10th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_11</i>		<i>Follow up time 11th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_12</i>		<i>Follow up time 12th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>FUTIME_HF_13</i>		<i>Follow up time 13th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.8 std=697.2)

<i>HFPEF</i>		<i>HFpEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16250	0	No
165	1	Yes

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<i>HFREF</i>		<i>HFrEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16250	0	No
165	1	Yes

<i>INCIDENT_GHTN</i>		<i>Pregnancy Complications Diagnosis of GHTN (Preeclampsia to Eclampsia) Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16393	0	No
22	1	Yes

<i>INCIDENT_GDM</i>		<i>Pregnancy Complications Diagnosis of GDM Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16376	0	No
39	1	Yes

<i>INCIDENT_PCD</i>		<i>Pregnancy Complications Diagnosis (PCD) Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16322	0	No
93	1	Yes

<i>INCIDENT_STR_2021</i>		<i>Incident Stroke (STR) Event by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16247	0	No
168	1	Yes

<i>RECUR_STR_2021_1</i>		<i>STR as Recur STR 2021 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16390	0	No
25	1	Yes

<i>RECUR_STR_2021_2</i>		<i>STR as Recur STR 2021 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16412	0	No
3	1	Yes

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<i>FUTIME_STR</i>		<i>Follow up time of STR event, censor or the end of follow-up by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4271 mean=4151.1 std=733.9)

<i>FUTIME_STR_1</i>		<i>Recur FUTIME STR 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4275 mean=4168.4 std=702.1)

<i>FUTIME_STR_2</i>		<i>Recur FUTIME STR 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4276 mean=4170.7 std=697.3)

<i>INCIDENT_CVD_2021</i>		<i>Incident CVD (MI, HF or Stroke) Event by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15865	0	No
550	1	Yes

<i>FUTIME_CVD</i>		<i>Follow up time of CVD (MI, HF or Stroke) event, censor or the end of follow-up by 2021</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	0 - 5050 (median=4262 mean=4112.9 std=797.4)