



Codebook for Incident Events 2008-2019 January 2024

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

HCHS/SOL Incident Events 2008-2019 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>
7420	Present	Text suppressed

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5511	1	Occurrence 1
1113	2	Occurrence 2
408	3	Occurrence 3
164	4	Occurrence 4
90	5	Occurrence 5
50	6	Occurrence 6
29	7	Occurrence 7
19	8	Occurrence 8
9	9	Occurrence 9
8	10	Occurrence 10
5	11	Occurrence 11
4	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>EEF1</i>		<i>Event location</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1610	1	1= In hospital
797	2	2= Emergency Dept. (ED)
4920	3	3= Both ED and in hospital
87	4	4= other
6		Missing

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Section A: Events Eligibility Ascertainment

EEF1A		Other event location
N	Value	Description
87	Present	Text suppressed
7332	~	Legitimately skipped
1		Missing

EEF4		Vital status at discharge
N	Value	Description
156	0	0= Dead
7245	1	1= Alive
19		Missing

EEF5A		ICD discharge code
N	Value	Description
6610	Present	Text suppressed
810		Missing

EEF5H		ICD discharge code
N	Value	Description
3199	Present	Text suppressed
4221		Missing

EEF5O		ICD discharge code
N	Value	Description
1126	Present	Text suppressed
6294		Missing

EEF5B		ICD discharge code
N	Value	Description
6190	Present	Text suppressed
1230		Missing

EEF5I		ICD discharge code
N	Value	Description
2794	Present	Text suppressed
4626		Missing

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EEF5P		ICD discharge code
N	Value	Description
956	Present	Text suppressed
6464		Missing

EEF5C		ICD discharge code
N	Value	Description
5747	Present	Text suppressed
1673		Missing

EEF5J		ICD discharge code
N	Value	Description
2418	Present	Text suppressed
5002		Missing

EEF5Q		ICD discharge code
N	Value	Description
798	Present	Text suppressed
6622		Missing

EEF5D		ICD discharge code
N	Value	Description
5237	Present	Text suppressed
2183		Missing

EEF5K		ICD discharge code
N	Value	Description
2051	Present	Text suppressed
5369		Missing

EEF5R		ICD discharge code
N	Value	Description
711	Present	Text suppressed
6709		Missing

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EEF5E		ICD discharge code
N	Value	Description
4692	Present	Text suppressed
2728		Missing

EEF5L		ICD discharge code
N	Value	Description
1776	Present	Text suppressed
5644		Missing

EEF5S		ICD discharge code
N	Value	Description
598	Present	Text suppressed
6822		Missing

EEF5F		ICD discharge code
N	Value	Description
4172	Present	Text suppressed
3248		Missing

EEF5M		ICD discharge code
N	Value	Description
1514	Present	Text suppressed
5906		Missing

EEF5T		ICD discharge code
N	Value	Description
506	Present	Text suppressed
6914		Missing

EEF5G		ICD discharge code
N	Value	Description
3688	Present	Text suppressed
3732		Missing

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Section A: Events Eligibility Ascertainment

EEF5N		ICD discharge code
N	Value	Description
1323	Present	Text suppressed
6097		Missing

EEF6A		ICD procedure code
N	Value	Description
3447	Present	Text suppressed
3973		Missing

EEF6K		ICD procedure code
N	Value	Description
65	Present	Text suppressed
7355		Missing

EEF6B		ICD procedure code
N	Value	Description
2197	Present	Text suppressed
5223		Missing

EEF6L		ICD procedure code
N	Value	Description
45	Present	Text suppressed
7375		Missing

EEF6C		ICD procedure code
N	Value	Description
1401	Present	Text suppressed
6019		Missing

EEF6M		ICD procedure code
N	Value	Description
34	Present	Text suppressed
7386		Missing

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Section A: Events Eligibility Ascertainment

EEF6D		ICD procedure code
N	Value	Description
902	Present	Text suppressed
6518		Missing

EEF6N		ICD procedure code
N	Value	Description
21	Present	Text suppressed
7399		Missing

EEF6E		ICD procedure code
N	Value	Description
556	Present	Text suppressed
6864		Missing

EEF6O		ICD procedure code
N	Value	Description
16	Present	Text suppressed
7404		Missing

EEF6F		ICD procedure code
N	Value	Description
384	Present	Text suppressed
7036		Missing

EEF6P		ICD procedure code
N	Value	Description
13	Present	Text suppressed
7407		Missing

EEF6G		ICD procedure code
N	Value	Description
267	Present	Text suppressed
7153		Missing

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Section A: Events Eligibility Ascertainment

EEF6Q		ICD procedure code
N	Value	Description
11	Present	Text suppressed
7409		Missing

EEF6H		ICD procedure code
N	Value	Description
186	Present	Text suppressed
7234		Missing

EEF6R		ICD procedure code
N	Value	Description
8	Present	Text suppressed
7412		Missing

EEF6I		ICD procedure code
N	Value	Description
130	Present	Text suppressed
7290		Missing

EEF6S		ICD procedure code
N	Value	Description
7	Present	Text suppressed
7413		Missing

EEF6J		ICD procedure code
N	Value	Description
98	Present	Text suppressed
7322		Missing

EEF6T		ICD procedure code
N	Value	Description
5	Present	Text suppressed
7415		Missing

Chapter 1: Endpoint Ascertainment

Section A: Events Eligibility Ascertainment

<i>EEF7</i>		<i>discharge summary mentions cardiac</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6176	0	0=No
1241	1	1=Yes
3		Missing

<i>EEF7A</i>		<i>MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
987	0	0=No
251	1	1=Yes
3		Missing

<i>EEF7O</i>		<i>acute PE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1158	0	0=No
81	1	1=Yes
2		Missing

<i>EEF7B</i>		<i>angina</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1144	0	0=No
96	1	1=Yes
1		Missing

<i>EEF7P</i>		<i>aortic aneurysm</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1236	0	0=No
2	1	1=Yes
3		Missing

Chapter 1: Endpoint Ascertainment

Section A: Events Eligibility Ascertainment

EEF7C		ischemic heart disease
N	Value	Description
6179	.S	Legitimately skipped
1197	0	0=No
42	1	1=Yes
2		Missing

EEF7Q		CCU
N	Value	Description
6179	.S	Legitimately skipped
1147	0	0=No
92	1	1=Yes
2		Missing

EEF7D		CHD
N	Value	Description
6179	.S	Legitimately skipped
740	0	0=No
499	1	1=Yes
2		Missing

EEF7R		cardiac cath
N	Value	Description
6179	.S	Legitimately skipped
903	0	0=No
335	1	1=Yes
3		Missing

EEF7E		unstable angina
N	Value	Description
6179	.S	Legitimately skipped
1181	0	0=No
58	1	1=Yes
2		Missing

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Section A: Events Eligibility Ascertainment

EEF7S		CABG
N	Value	Description
6179	.S	Legitimately skipped
1089	0	0=No
150	1	1=Yes
2		Missing

EEF7F		coronary insufficiency
N	Value	Description
6179	.S	Legitimately skipped
1235	0	0=No
4	1	1=Yes
2		Missing

EEF7T		coronary stent
N	Value	Description
6179	.S	Legitimately skipped
1017	0	0=No
221	1	1=Yes
3		Missing

EEF7G		cardiac arrest
N	Value	Description
6179	.S	Legitimately skipped
1159	0	0=No
80	1	1=Yes
2		Missing

EEF7U		elevated CK-MB
N	Value	Description
6179	.S	Legitimately skipped
1014	0	0=No
225	1	1=Yes
2		Missing

Chapter 1: Endpoint Ascertainment

Section A: Events Eligibility Ascertainment

<i>EEF7H</i>		<i>AHD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1113	0	0=No
126	1	1=Yes
2		Missing

<i>EEF7V</i>		<i>coronary angiography</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1073	0	0=No
166	1	1=Yes
2		Missing

<i>EEF7I</i>		<i>heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1067	0	0=No
171	1	1=Yes
3		Missing

<i>EEF7W</i>		<i>cardiac angioplasty</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1184	0	0=No
55	1	1=Yes
2		Missing

<i>EEF7J</i>		<i>CHF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
814	0	0=No
425	1	1=Yes
2		Missing

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Section A: Events Eligibility Ascertainment

EEF7X		<i>leg amputation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1229	0	0=No
10	1	1=Yes
2		Missing

EEF7K		<i>cardiomyopathy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1045	0	0=No
194	1	1=Yes
2		Missing

EEF7Y		<i>leg angioplasty</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1227	0	0=No
12	1	1=Yes
2		Missing

EEF7L		<i>atherosclerosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1224	0	0=No
15	1	1=Yes
2		Missing

EEF7Z		<i>leg revascularization</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1215	0	0=No
24	1	1=Yes
2		Missing

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Section A: Events Eligibility Ascertainment

<i>EEF7M</i>		<i>PVD</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1113	0	0=No
126	1	1=Yes
2		Missing

<i>EEF7AA</i>		<i>pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1200	0	0=No
40	1	1=Yes
1		Missing

<i>EEF7N</i>		<i>claudication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6179	.S	Legitimately skipped
1215	0	0=No
24	1	1=Yes
2		Missing

<i>EEF8</i>		<i>discharge summary mentions stroke</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7098	0	0=No
320	1	1=Yes
2		Missing

<i>EEF8A</i>		<i>TIA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
246	0	0=No
74	1	1=Yes

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Section A: Events Eligibility Ascertainment

<i>EEF8I</i>		<i>carotid endarterectomy or carotid ste</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
310	0	0=No
10	1	1=Yes

<i>EEF8B</i>		<i>stroke</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
159	0	0=No
161	1	1=Yes

<i>EEF8J</i>		<i>arteriovenous malformation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
315	0	0=No
5	1	1=Yes

<i>EEF8C</i>		<i>cerebral infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
227	0	0=No
93	1	1=Yes

<i>EEF8K</i>		<i>cerebral aneurysm</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
305	0	0=No
15	1	1=Yes

<i>EEF8D</i>		<i>lacunar (syndrome or infarction)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
307	0	0=No
13	1	1=Yes

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EEF8L		cerebral, intracerebral, brain hemorr
N	Value	Description
7100	.S	Legitimately skipped
256	0	0=No
64	1	1=Yes

EEF8E		cerebral embolus
N	Value	Description
7100	.S	Legitimately skipped
314	0	0=No
6	1	1=Yes

EEF8M		subarachnoid hemorrhage
N	Value	Description
7100	.S	Legitimately skipped
287	0	0=No
33	1	1=Yes

EEF8F		cerebral thrombosis
N	Value	Description
7100	.S	Legitimately skipped
316	0	0=No
4	1	1=Yes

EEF8N		cerebrovascular disease
N	Value	Description
7100	.S	Legitimately skipped
307	0	0=No
13	1	1=Yes

EEF8G		central or cerebral venous thrombosis
N	Value	Description
7100	.S	Legitimately skipped
320	0	0=No
0	1	1=Yes

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Section A: Events Eligibility Ascertainment

<i>EEF8O</i>		<i>CT/MRI scan showing new cerebrovascul</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
184	0	0=No
136	1	1=Yes

<i>EEF8H</i>		<i>intracranial stenting</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7100	.S	Legitimately skipped
320	0	0=No
0	1	1=Yes

<i>EEF9</i>		<i>discharge summary mentions HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6668	0	0=No
750	1	1=Yes
2		Missing

<i>EEF9A</i>		<i>heart failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6670	.S	Legitimately skipped
564	0	0=No
185	1	1=Yes
1		Missing

<i>EEF9J</i>		<i>low ejection fraction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4870	.S	Legitimately skipped
513	0	0=No
123	1	1=Yes
1914		Missing

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Section A: Events Eligibility Ascertainment

EEF9B		CHF
N	Value	Description
6670	.S	Legitimately skipped
303	0	0=No
446	1	1=Yes
1		Missing

EEF9K		ejection fraction less than 40
N	Value	Description
4870	.S	Legitimately skipped
509	0	0=No
127	1	1=Yes
1914		Missing

EEF9C		pump failure
N	Value	Description
6670	.S	Legitimately skipped
749	0	0=No
0	1	1=Yes
1		Missing

EEF9L		cardiomegaly
N	Value	Description
6670	.S	Legitimately skipped
568	0	0=No
181	1	1=Yes
1		Missing

EEF9D		cardiomyopathy
N	Value	Description
6670	.S	Legitimately skipped
558	0	0=No
191	1	1=Yes
1		Missing

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Section A: Events Eligibility Ascertainment

<i>EEF9M</i>		<i>impaired systolic function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6670	.S	Legitimately skipped
642	0	0=No
107	1	1=Yes
1		Missing

<i>EEF9E</i>		<i>ventricular failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6670	.S	Legitimately skipped
749	0	0=No
0	1	1=Yes
1		Missing

<i>EEF9N</i>		<i>impaired diastolic function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6670	.S	Legitimately skipped
574	0	0=No
175	1	1=Yes
1		Missing

<i>EEF9F</i>		<i>LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6670	.S	Legitimately skipped
683	0	0=No
66	1	1=Yes
1		Missing

<i>EEF9O</i>		<i>pulmonary edema device</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6670	.S	Legitimately skipped
730	0	0=No
19	1	1=Yes
1		Missing

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Section A: Events Eligibility Ascertainment

EEF9G		jugular venous distension
N	Value	Description
6670	.S	Legitimately skipped
713	0	0=No
36	1	1=Yes
1		Missing

EEF9P		heart biopsy
N	Value	Description
6670	.S	Legitimately skipped
749	0	0=No
0	1	1=Yes
1		Missing

EEF9H		orthopnea
N	Value	Description
6670	.S	Legitimately skipped
646	0	0=No
104	1	1=Yes

EEF9Q		AICD check
N	Value	Description
6670	.S	Legitimately skipped
732	0	0=No
17	1	1=Yes
1		Missing

EEF9I		paroxysmal nocturnal dyspnea
N	Value	Description
6670	.S	Legitimately skipped
705	0	0=No
44	1	1=Yes
1		Missing

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Section A: Events Eligibility Ascertainment

EEF9R		implantation of cardiac resync therapy device
N	Value	Description
6670	.S	Legitimately skipped
741	0	0=No
8	1	1=Yes
1		Missing

EEF10		discharge summary mentions pulmonary
N	Value	Description
6164	0	0=No
1254	1	1=Yes
2		Missing

EEF10A		acute bronchitis
N	Value	Description
6166	.S	Legitimately skipped
994	0	0=No
257	1	1=Yes
3		Missing

EEF10J		pneumonia with underlying COPD
N	Value	Description
6166	.S	Legitimately skipped
1197	0	0=No
55	1	1=Yes
2		Missing

EEF10B		acute bronchitis with exacerbation of asthma
N	Value	Description
6166	.S	Legitimately skipped
1224	0	0=No
27	1	1=Yes
3		Missing

Chapter 1: Endpoint Ascertainment

Section A: Events Eligibility Ascertainment

EEF10K		<i>emphysema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1208	0	0=No
43	1	1=Yes
3		Missing

EEF10C		<i>obstructive chronic bronchitis with acute bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1230	0	0=No
21	1	1=Yes
3		Missing

EEF10L		<i>bronchiectasis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1225	0	0=No
26	1	1=Yes
3		Missing

EEF10D		<i>acute exacerbation of asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
756	0	0=No
497	1	1=Yes
1		Missing

EEF10M		<i>extrinsic allergic alveolitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1251	0	0=No
0	1	1=Yes
3		Missing

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Section A: Events Eligibility Ascertainment

EEF10E		chronic obstruction asthma with acute exacerbation
N	Value	Description
6166	.S	Legitimately skipped
998	0	0=No
253	1	1=Yes
3		Missing

EEF10N		acute respiratory failure
N	Value	Description
6166	.S	Legitimately skipped
967	0	0=No
284	1	1=Yes
3		Missing

EEF10F		chronic obstructive asthma
N	Value	Description
6166	.S	Legitimately skipped
947	0	0=No
304	1	1=Yes
3		Missing

EEF10O		acute and chronic respiratory failure
N	Value	Description
6166	.S	Legitimately skipped
1207	0	0=No
44	1	1=Yes
3		Missing

EEF10G		status asthmaticus
N	Value	Description
6166	.S	Legitimately skipped
1239	0	0=No
12	1	1=Yes
3		Missing

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<i>EEF10P</i>		<i>positive spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1250	0	0=No
0	1	1=Yes
4		Missing

<i>EEF10H</i>		<i>chronic airway obstruction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1234	0	0=No
17	1	1=Yes
3		Missing

<i>EEF10Q</i>		<i>pulmonary insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1243	0	0=No
6	1	1=Yes
5		Missing

<i>EEF10I</i>		<i>chronic bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6166	.S	Legitimately skipped
1227	0	0=No
24	1	1=Yes
3		Missing

<i>EEF11</i>		<i>discharge summary mentions pregnancy-related complications</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4963	0	0=No
126	1	1=Yes
2331		Missing

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EEF11A		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
90	0	0=No
36	1	1=Yes
1851		Missing

EEF11J		<i>chronic hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
107	0	0=No
18	1	1=Yes
1852		Missing

EEF11B		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
122	0	0=No
3	1	1=Yes
1852		Missing

EEF11K		<i>hyperglycemia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
124	0	0=No
1	1	1=Yes
1852		Missing

EEF11C		<i>PEC</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
123	0	0=No
2	1	1=Yes
1852		Missing

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<i>EEF11L</i>		<i>gestational diabetes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
58	0	0=No
67	1	1=Yes
1852		Missing

<i>EEF11D</i>		<i>PreX</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
125	0	0=No
0	1	1=Yes
1852		Missing

<i>EEF11M</i>		<i>insulin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
106	0	0=No
19	1	1=Yes
1852		Missing

<i>EEF11E</i>		<i>PET</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
125	0	0=No
0	1	1=Yes
1852		Missing

<i>EEF11N</i>		<i>HELLP syndrome</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
124	0	0=No
1	1	1=Yes
1852		Missing

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<i>EEF11F</i>		<i>Toxemia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
125	0	0=No
0	1	1=Yes
1852		Missing

<i>EEF11O</i>		<i>seizure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
125	0	0=No
0	1	1=Yes
1852		Missing

<i>EEF11G</i>		<i>pregnancy-induced hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
116	0	0=No
9	1	1=Yes
1852		Missing

<i>EEF11P</i>		<i>renal failure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
125	0	0=No
0	1	1=Yes
1852		Missing

<i>EEF11H</i>		<i>pregestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
123	0	0=No
2	1	1=Yes
1852		Missing

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<i>EEF11Q</i>		<i>proteinuria</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
116	0	0=No
9	1	1=Yes
1852		Missing

<i>EEF11I</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
96	0	0=No
29	1	1=Yes
1852		Missing

<i>EEF11R</i>		<i>oliguria</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5443	.S	Legitimately skipped
125	0	0=No
0	1	1=Yes
1852		Missing

<i>EEF12A</i>		<i>no additional form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3692	N	
3332	Y	
396	~	Legitimately skipped

<i>EEF12B</i>		<i>MI form needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4687	N	
2258	Y	
474	~	Legitimately skipped
1		Missing

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EEF12C		Stroke form needed
N	Value	Description
6393	N	
452	Y	
574	~	Legitimately skipped
1		Missing

EEF12D		HF form needed
N	Value	Description
5702	N	
1135	Y	
567	~	Legitimately skipped
16		Missing

EEF12E		Pulmonary form needed
N	Value	Description
4575	N	
2356	Y	
489	~	Legitimately skipped

EEF12F		Pregnancy-Related Complications form needed
N	Value	Description
1139	Hn	
23	Hy	
3664	N	
126	Y	
618	~	Legitimately skipped
1850		Missing

EEF12G		Death form needed
N	Value	Description
4785	N	
156	Y	
629	~	Legitimately skipped
1850		Missing

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EEF0A_FUTIME		Futime from Randomization date for (EEF0A)
N	Value	Description
7419	Range	-116 - 5557 (median=2420 mean=2560.0 std=889.9)
1		Missing

EEF0D_FUTIME		Futime from Randomization date for (EEF0D)
N	Value	Description
7417	Range	-297 - 4256 (median=1448 mean=1612.7 std=1054.7)
3		Missing

EEF2_FUTIME		Futime from Randomization date for (EEF2)
N	Value	Description
6840	Range	-196 - 4256 (median=1536.5 mean=1661.10 std=1044.46)
580		Missing

EEF3_FUTIME		Futime from Randomization date for (EEF3)
N	Value	Description
7206	Range	-297 - 4261 (median=1479.5 mean=1637.43 std=1047.72)
214		Missing

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

ID		HCHS Participant ID
N	Value	Description
2278	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
1516	1	Occurrence 1
412	2	Occurrence 2
162	3	Occurrence 3
74	4	Occurrence 4
40	5	Occurrence 5
24	6	Occurrence 6
12	7	Occurrence 7
7	8	Occurrence 8
5	9	Occurrence 9
6	10	Occurrence 10
3	11	Occurrence 11
3	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

MIF1		Type of Event
N	Value	Description
335	1	1= In hospital only
2	2	2= Emergency Dept. visit only (ED)
1939	3	3= Both ED and in hospital
2		Missing

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MIF2A		Time of arrival
N	Value	Description
2057	Range	0:00:00 - 23:59:00
221		Missing

MIF2AHR		Hour of Time of arrival
N	Value	Description
2057	Range	0 - 23 (median=13 mean=12.7 std=5.8)
221		Missing

MIF2AMN		Minute of Time of arrival
N	Value	Description
2057	Range	0 - 59 (median=29 mean=28.8 std=17.2)
221		Missing

MIF2ASC		Second of Time of arrival
N	Value	Description
944	Range	0 - 0 (median=0 mean=0.0 std=0.0)
1334		Missing

MIF2A1		AM/PM of arrival
N	Value	Description
2032	.S	Legitimately skipped
93	1	1=AM
110	2	2=PM
43		Missing

MIF3A		Time of discharge
N	Value	Description
1225	Range	0:08:00 - 23:59:00
1053		Missing

MIF3AHR		Hour of Time of discharge
N	Value	Description
1225	Range	0 - 23 (median=16 mean=15.7 std=4.0)
1053		Missing

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MIF3AMN		Minute of Time of discharge
N	Value	Description
1225	Range	0 - 59 (median=30 mean=29.4 std=19.3)
1053		Missing

MIF3ASC		Second of Time of discharge
N	Value	Description
551	Range	0 - 0 (median=0 mean=0.0 std=0.0)
1727		Missing

MIF3A1		AM/PM of discharge
N	Value	Description
2032	.S	Legitimately skipped
15	1	1=AM
99	2	2=PM
132		Missing

MIF4		Primary admitting diagnosis code
N	Value	Description
2003	Present	Text suppressed
275		Missing

MIF5		Primary discharge diagnosis code
N	Value	Description
2081	Present	Text suppressed
197		Missing

MIF6		EMS unit transport patient to hospital
N	Value	Description
1713	0	0=No/NR
565	1	1=Yes

MIF7		Patient transferred to this hospital?
N	Value	Description
2194	0	0=No/NR
84	1	1=Yes

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MIF8		Patient's code ever 'no-code' or 'DNR'
N	Value	Description
2205	0	0=No/NR
73	1	1=Yes

MIF9		Patient alive at discharge?
N	Value	Description
96	0	0=No/NR
2182	1	1=Yes

MIF9A		Patient dead on arrival?
N	Value	Description
2182	.S	Legitimately skipped
96	0	0=No
0	1	1=Yes

MIF9B		Patient die in emergency department?
N	Value	Description
2182	.S	Legitimately skipped
95	0	0=No
1	1	1=Yes

MIF9C		Was an autopsy performed?
N	Value	Description
2182	.S	Legitimately skipped
79	0	0=No
4	1	1=Yes
13		Missing

MIF10		Acute episode occur prior to admission
N	Value	Description
46	0	0=No
2194	1	1=Yes
37	9	9=NR
1		Missing

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MIF10A		Time estimation of acute episode
N	Value	Description
84	.S	Legitimately skipped
44	1	1= (< 1 hr)
95	2	2= (>= 1 - < 3 hrs)
69	3	3= (>= 3 - < 6 hrs)
56	4	4= (>= 6 - < 12 hrs)
65	5	5= (>= 12 - < 24 hrs)
1237	6	6= (>= 24 hrs)
627	7	7= Unsure
1		Missing

MIF11		Mention of acute CHD at hospital
N	Value	Description
2040	0	0=No
63	1	1=Yes
174	9	9=NR
1		Missing

MIF12		Acute episode within 72 hours to arrival
N	Value	Description
987	0	0=No
850	1	1=Yes
441	9	9=NR

MIF12A		Pain/discomfort involve the chest?
N	Value	Description
1428	.S	Legitimately skipped
13	0	0=No
836	1	1=Yes
1	9	9=NR

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MIF12B		Pain crescendo over time?
N	Value	Description
1428	.S	Legitimately skipped
328	0	0=No
83	1	1=Yes
439	9	9=NR

MIF12C		Pain/discomfort have non-cardiac origin?
N	Value	Description
1428	.S	Legitimately skipped
410	0	0=No
107	1	1=Yes
333	9	9=NR

MIF13		Nausea/vomiting related?
N	Value	Description
1055	0	0=No
563	1	1=Yes
660	9	9=NR

MIF14		Diaphoresis associated?
N	Value	Description
869	0	0=No
175	1	1=Yes
1234	9	9=NR

MIF15		Fatigue/malaise associated ?
N	Value	Description
341	0	0=No
409	1	1=Yes
1528	9	9=NR

MIF16A1		Sys. blood pressure at arrival
N	Value	Description
2014	Range	0 - 270 (median=138 mean=140.1 std=30.2)
264		Missing

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MIF16A2		<i>Dia. blood pressure at arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2008	Range	0 - 191 (median=75 mean=76.4 std=17.1)
270		Missing

MIF16B		<i>Heart rate (bpm)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1981	Range	0 - 234 (median=83 mean=86.1 std=22.3)
297		Missing

MIF17A		<i>Myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1933	0	0=No/NR
344	1	1=Yes
1		Missing

MIF17A1		<i>If history of MI, MI within 4 weeks?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1934	.S	Legitimately skipped
323	0	0=No/NR
20	1	1=Yes
1		Missing

MIF17B		<i>Angina</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2039	0	0=No/NR
239	1	1=Yes

MIF17C		<i>Percutaneous coronary intervention</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1948	0	0=No/NR
330	1	1=Yes

MIF17D		<i>CABG</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1988	0	0=No/NR
290	1	1=Yes

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MIF17E		Coronary artery disease
N	Value	Description
1379	0	0=No/NR
899	1	1=Yes

MIF17F		Heart failure
N	Value	Description
1769	0	0=No/NR
509	1	1=Yes

MIF17G		Arrhythmia
N	Value	Description
1928	0	0=No/NR
349	1	1=Yes
1		Missing

MIF17G1		Atrial Fibrillation/Flutter
N	Value	Description
1928	.S	Legitimately skipped
78	0	0=No/NR
270	1	1=Yes
2		Missing

MIF17G2		Ventricular Fibrillation/Tachycardia
N	Value	Description
1928	.S	Legitimately skipped
312	0	0=No/NR
36	1	1=Yes
2		Missing

MIF17G3		Other arrhythmia
N	Value	Description
1928	.S	Legitimately skipped
295	0	0=No/NR
53	1	1=Yes
2		Missing

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MIF18A		Indicated Angina
N	Value	Description
1902	0	0=No/NR
376	1	1=Yes

MIF18B		Indicated Acute myocardial Infarction
N	Value	Description
2072	0	0=No/NR
206	1	1=Yes

MIF18C		Indicated ST elevation not on ECG
N	Value	Description
2262	0	0=No/NR
16	1	1=Yes

MIF18D		Indicated congestive heart failure
N	Value	Description
1880	0	0=No/NR
398	1	1=Yes

MIF18D1		If yes, heart failure/pulmonary edema?
N	Value	Description
1880	.S	Legitimately skipped
340	0	0=No/NR
58	1	1=Yes

MIF18E		Indicated shock or cardiogenic shock
N	Value	Description
2240	0	0=No/NR
38	1	1=Yes

MIF18E1		If yes, shock occur within 24 hours
N	Value	Description
2240	.S	Legitimately skipped
28	0	0=No/NR
10	1	1=Yes

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MIF18F		Indicated ventricular fibrillation/asystole
N	Value	Description
2195	0	0=No/NR
83	1	1=Yes

MIF18F1		If yes, arrest occur within 24 hours
N	Value	Description
2195	.S	Legitimately skipped
64	0	0=No/NR
19	1	1=Yes

MIF18G		Indicated Ventricular Tachycardia
N	Value	Description
2233	0	0=No/NR
45	1	1=Yes

MIF18H		Indicated Atrial fibrillation or atrial flutter
N	Value	Description
2060	0	0=No/NR
218	1	1=Yes

MIF19		Cardiac enzymes reported within days
N	Value	Description
880	0	0=No/NR
1396	1	1=Yes
2		Missing

MIF19A		Cardiac enzymes reported arrival day
N	Value	Description
774	.S	Legitimately skipped
192	0	0=No/NR
1311	1	1=Yes
1		Missing

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MIF19B		Cardiac enzymes reported day after arrival
N	Value	Description
774	.S	Legitimately skipped
914	0	0=No/NR
589	1	1=Yes
1		Missing

MIF19C		Cardiac enzymes reported third day
N	Value	Description
774	.S	Legitimately skipped
1344	0	0=No/NR
159	1	1=Yes
1		Missing

MIF19D		Cardiac enzymes reported fourth day
N	Value	Description
774	.S	Legitimately skipped
1432	0	0=No/NR
71	1	1=Yes
1		Missing

MIF20A1		Set 1 CK Upper limit of normal
N	Value	Description
551	Range	100 - 398 (median=200 mean=198.4 std=53.0)
881	.S	Legitimately skipped
846		Missing

MIF20A2		Set 1 CK units
N	Value	Description
881	.S	Legitimately skipped
1	1	1=ng/mL
593	2	2=Units/L
0	3	3=Åµg/L
803		Missing

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MIF20B1		Set 1 CK-MB upper limit of normal
N	Value	Description
422	Range	0.5 - 10 (median=5 mean=5.4 std=1.8)
881	.S	Legitimately skipped
975		Missing

MIF20B2		Set 1 CK-MB units
N	Value	Description
881	.S	Legitimately skipped
426	1	1=ng/mL
0	2	2=Units/L
1	3	3=Åµg/L
970		Missing

MIF20C1		Set 1 Total LDH upper limit of normal
N	Value	Description
50	Range	179 - 618 (median=210 mean=250.0 std=124.5)
881	.S	Legitimately skipped
1347		Missing

MIF20C2		Set 1 Total LDH units
N	Value	Description
881	.S	Legitimately skipped
0	1	1=ng/mL
51	2	2=Units/L
0	3	3=Åµg/L
1346		Missing

MIF20D1		Set 1 LDH-1 upper limit of normal
N	Value	Description
1	Range	100 - 100 (median=100 mean=100.0 std=)
881	.S	Legitimately skipped
1396		Missing

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MIF20D2		Set 1 LDH-1 Units
N	Value	Description
881	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
1397		Missing

MIF20E1		Set 1 LDH-2 upper limit of normal
N	Value	Description
1	Range	100 - 100 (median=100 mean=100.0 std=)
881	.S	Legitimately skipped
1396		Missing

MIF20E2		Set 1 LDH-2 units
N	Value	Description
881	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
1397		Missing

MIF20F1		Set 1 Troponin upper limit of normal
N	Value	Description
926	Range	0.01 - 22 (median=0.05 mean=0.462 std=2.453)
881	.S	Legitimately skipped
471		Missing

MIF20F2		Set 1 Troponin units
N	Value	Description
881	.S	Legitimately skipped
970	1	1=ng/mL
0	2	2=Units/L
14	3	3=Åµg/L
413		Missing

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MIF20F3		Set 1 Type of Troponin
N	Value	Description
881	.S	Legitimately skipped
120	1	1= Troponin, type not specified
621	2	2= Troponin I
459	3	3= Troponin T
10	4	4= High Sensitivity Troponin (HS)
2	5	5= Unsure
185		Missing

MIF20G		second set of normal values
N	Value	Description
774	.S	Legitimately skipped
1444	0	0=No/NR
39	1	1=Yes
21		Missing

MIF21A1		Set 2 Total CK upper limit of normal
N	Value	Description
8	Range	41 - 374 (median=250 mean=238.1 std=102.1)
2102	.S	Legitimately skipped
168		Missing

MIF21A2		Set 2 Total CK units
N	Value	Description
2102	.S	Legitimately skipped
0	1	1=ng/mL
8	2	2=Units/L
0	3	3=Åµg/L
168		Missing

MIF21B1		Set 2 CK-MB
N	Value	Description
6	Range	4.2 - 10 (median=5.85 mean=6.800 std=2.612)
2102	.S	Legitimately skipped
170		Missing

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MIF21B2		Set 2 CK-MB Units
N	Value	Description
2102	.S	Legitimately skipped
5	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
171		Missing

MIF21C1		Set 2 Total LDH Upper limit of normal
N	Value	Description
2102	.S	Legitimately skipped
176		Missing

MIF21C2		Set 2 Total LDH Units
N	Value	Description
2102	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
176		Missing

MIF21D1		Set 2 LDH-1 upper limit of normal
N	Value	Description
2102	.S	Legitimately skipped
176		Missing

MIF21D2		Set 2 LDH-1 Units
N	Value	Description
2102	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
176		Missing

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MIF21E1		Set 2 LDH-2 upper limit of normal
N	Value	Description
2102	.S	Legitimately skipped
176		Missing

MIF21E2		Set 2 LDH-2 units
N	Value	Description
2102	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
176		Missing

MIF21F1		Set 2 Troponin upper limit of normal
N	Value	Description
34	Range	0.01 - 0.48 (median=0.053 mean=0.0828 std=0.1038)
2102	.S	Legitimately skipped
142		Missing

MIF21F2		Set 2 Troponin units
N	Value	Description
2102	.S	Legitimately skipped
34	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
142		Missing

MIF21F3		Set 2 Type of Troponin
N	Value	Description
2102	.S	Legitimately skipped
4	1	1= Troponin, type not specified
28	2	2= Troponin I
5	3	3= Troponin T
0	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure
139		Missing

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MIF22A		Day 1/Set 1 Total CK (CPK)
N	Value	Description
603	Range	11 - 18460 (median=103 mean=326.5 std=1323.4)
960	.S	Legitimately skipped
715		Missing

MIF22A1		Day 1/Set 1 Total CK (CPK) Units
N	Value	Description
960	.S	Legitimately skipped
3	1	1=ng/mL
510	2	2=Units/L
0	3	3=Åµg/L
805		Missing

MIF22A2		Day 1/Set 1 Total CK Range Set
N	Value	Description
960	.S	Legitimately skipped
539	1	1=Range Set 1
0	2	2=Range Set 2
779		Missing

MIF22A3		Day 1/Set 1 Total CK Words Code
N	Value	Description
960	.S	Legitimately skipped
100	1	1=absent/negative/normal
2	2	2=trace/weak positive
24	3	3=present/positive/abnormal
1192		Missing

MIF22B		Day 1/Set 1 CK-MB
N	Value	Description
426	Range	0.22 - 300 (median=2.6 mean=6.94 std=24.49)
960	.S	Legitimately skipped
892		Missing

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MIF22B1		Day 1/Set 1 CK-MB Units
N	Value	Description
960	.S	Legitimately skipped
350	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
968		Missing

MIF22B2		Day 1/Set 1 CK-MB Range Set
N	Value	Description
960	.S	Legitimately skipped
356	1	1=Range Set 1
5	2	2=Range Set 2
957		Missing

MIF22B3		Day 1/Set 1 CK-MB Words Code
N	Value	Description
960	.S	Legitimately skipped
68	1	1=absent/negative/normal
1	2	2=trace/weak positive
6	3	3=present/positive/abnormal
1243		Missing

MIF22C		Day 1/Set 1 Total LDH
N	Value	Description
51	Range	108 - 3840 (median=208 mean=376.9 std=586.7)
960	.S	Legitimately skipped
1267		Missing

MIF22C1		Day 1/Set 1 Total LDH Units
N	Value	Description
960	.S	Legitimately skipped
0	1	1=ng/mL
38	2	2=Units/L
0	3	3=Åµg/L
1280		Missing

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MIF22C2		Day 1/Set 1 Total LDH Range Set
N	Value	Description
960	.S	Legitimately skipped
43	1	1=Range Set 1
1	2	2=Range Set 2
1274		Missing

MIF22C3		Day 1/Set 1 Total LDH Words Code
N	Value	Description
960	.S	Legitimately skipped
5	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
1313		Missing

MIF22D1		Day 1/Set 1 LDH-1 Units
N	Value	Description
960	.S	Legitimately skipped
0	1	1=ng/mL
2	2	2=Units/L
0	3	3=Åµg/L
1316		Missing

MIF22D2		Day 1/Set 1 LDH-1 Range Set
N	Value	Description
960	.S	Legitimately skipped
2	1	1=Range Set 1
0	2	2=Range Set 2
1316		Missing

MIF22D3		Day 1/Set 1 LDH-1 Words Code
N	Value	Description
960	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
1318		Missing

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MIF22E		Day 1/Set 1 LDH-2
N	Value	Description
2	Range	26 - 32.7 (median=29.35 mean=29.350 std=4.738)
960	.S	Legitimately skipped
1316		Missing

MIF22E1		Day 1/Set 1 LDH-2 Units
N	Value	Description
960	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
1318		Missing

MIF22E2		Day 1/Set 1 LDH-2 Range Set
N	Value	Description
960	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
1318		Missing

MIF22E3		Day 1/Set 1 LDH-2 Words Code
N	Value	Description
960	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
1318		Missing

MIF22F		Day 1/Set 1 Troponin <
N	Value	Description
960	.S	Legitimately skipped
621	0	0=n/a
580	1	1=<
117		Missing

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MIF22F0		Day 1/Set 1 Troponin Value
N	Value	Description
1214	Range	0 - 4097 (median=0.03 mean=7.654 std=146.044)
960	.S	Legitimately skipped
104		Missing

MIF22F1		Day 1/Set 1 Troponin Units
N	Value	Description
960	.S	Legitimately skipped
950	1	1=ng/mL
1	2	2=Units/L
10	3	3=Åµg/L
357		Missing

MIF22F2		Day 1/Set 1 Troponin Range Set
N	Value	Description
960	.S	Legitimately skipped
1093	1	1=Range Set 1
4	2	2=Range Set 2
221		Missing

MIF22F3		Day 1/Set 1 Troponin Words Code
N	Value	Description
960	.S	Legitimately skipped
391	1	1=absent/negative/normal
26	2	2=trace/weak positive
152	3	3=present/positive/abnormal
749		Missing

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MIF22F4		Day 1/Set 1 Type of Troponin
N	Value	Description
960	.S	Legitimately skipped
165	1	1= Troponin, type not specified
620	2	2= Troponin I
450	3	3= Troponin T
10	4	4= High Sensitivity Troponin (HS)
9	5	5= Unsure
64		Missing

MIF22G		second set of enzymes reported the first day
N	Value	Description
853	.S	Legitimately skipped
899	0	0=No/NR
526	1	1=Yes

MIF23A		Day 1/Set 2 Total CK (CPK)
N	Value	Description
224	Range	20 - 12110 (median=108 mean=395.9 std=1132.1)
1678	.S	Legitimately skipped
376		Missing

MIF23A1		Day 1/Set 2 Total CK Units
N	Value	Description
1678	.S	Legitimately skipped
2	1	1=ng/mL
206	2	2=Units/L
0	3	3=Åµg/L
392		Missing

MIF23A2		Day 1/Set 2 Total CK Range Set
N	Value	Description
1678	.S	Legitimately skipped
205	1	1=Range Set 1
2	2	2=Range Set 2
393		Missing

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MIF23A3		Day 1/Set 2 Total CK Words Code
N	Value	Description
1678	.S	Legitimately skipped
34	1	1=absent/negative/normal
0	2	2=trace/weak positive
3	3	3=present/positive/abnormal
563		Missing

MIF23B		Day 1/ Set 2 CK-MB
N	Value	Description
170	Range	0.28 - 962.4 (median=2.6 mean=25.87 std=94.75)
1678	.S	Legitimately skipped
430		Missing

MIF23B1		Day 1/ Set 2 CK-MB Units
N	Value	Description
1678	.S	Legitimately skipped
151	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
449		Missing

MIF23B2		Day 1/ Set 2 CK-MB Range Set
N	Value	Description
1678	.S	Legitimately skipped
149	1	1=Range Set 1
4	2	2=Range Set 2
447		Missing

MIF23B3		Day 1/ Set 2 CK-MB Words Code
N	Value	Description
1678	.S	Legitimately skipped
25	1	1=absent/negative/normal
1	2	2=trace/weak positive
2	3	3=present/positive/abnormal
572		Missing

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MIF23C		Day 1/Set 2 Total LDH
N	Value	Description
7	Range	140 - 256 (median=165 mean=177.3 std=40.7)
1678	.S	Legitimately skipped
593		Missing

MIF23C1		Day 1/Set 2 Total LDH Units
N	Value	Description
1678	.S	Legitimately skipped
0	1	1=ng/mL
7	2	2=Units/L
0	3	3=Åµg/L
593		Missing

MIF23C2		Day 1/Set 2 Total LDH Range Set
N	Value	Description
1678	.S	Legitimately skipped
8	1	1=Range Set 1
0	2	2=Range Set 2
592		Missing

MIF23C3		Day 1/Set 2 Total LDH Words Code
N	Value	Description
1678	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
600		Missing

MIF23D1		Day 1/Set 2 LDH-1 Units
N	Value	Description
1678	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
600		Missing

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MIF23D2		Day 1/Set 2 LDH-1 Range Set
N	Value	Description
1678	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2
599		Missing

MIF23D3		Day 1/Set 2 LDH-1 Words Code
N	Value	Description
1678	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
600		Missing

MIF23E		Day 1/Set 2 LDH-2
N	Value	Description
1	Range	30.4 - 30.4 (median=30.4 mean=30.40 std=)
1678	.S	Legitimately skipped
599		Missing

MIF23E1		Day 1/Set 2 LDH-2 Units
N	Value	Description
1678	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Âµg/L
600		Missing

MIF23E2		Day 1/Set 2 LDH-2 Range Set
N	Value	Description
1678	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2
599		Missing

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MIF23E3		Day 1/Set 2 LDH-2 Words Code
N	Value	Description
1678	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
600		Missing

MIF23F		Day 1/Set 2 Troponin <
N	Value	Description
1678	.S	Legitimately skipped
290	0	0=n/a
201	1	1=<
109		Missing

MIF23F0		Day 1/Set 2 Troponin Value
N	Value	Description
497	Range	0 - 4815 (median=0.042 mean=21.4096 std=278.4749)
1678	.S	Legitimately skipped
103		Missing

MIF23F1		Day 1/Set 2 Troponin Units
N	Value	Description
1678	.S	Legitimately skipped
414	1	1=ng/mL
0	2	2=Units/L
5	3	3=Åµg/L
181		Missing

MIF23F2		Day 1/Set 2 Troponin Range Set
N	Value	Description
1678	.S	Legitimately skipped
443	1	1=Range Set 1
18	2	2=Range Set 2
139		Missing

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MIF23F3		Day 1/Set 2 Troponin Words Code
N	Value	Description
1678	.S	Legitimately skipped
146	1	1=absent/negative/normal
7	2	2=trace/weak positive
84	3	3=present/positive/abnormal
363		Missing

MIF23F4		Day 1/Set 2 Type of Troponin
N	Value	Description
1678	.S	Legitimately skipped
56	1	1= Troponin, type not specified
282	2	2= Troponin I
169	3	3= Troponin T
6	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure
87		Missing

MIF24A		Day 2/Set 1 Total CK (CPK)
N	Value	Description
301	Range	18 - 20001 (median=95 mean=508.0 std=1895.0)
1616	.S	Legitimately skipped
361		Missing

MIF24A1		Day 2/Set 1 Total CK (CPK) Units
N	Value	Description
1616	.S	Legitimately skipped
1	1	1=ng/mL
268	2	2=Units/L
0	3	3=Åµg/L
393		Missing

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MIF24A2		Day 2/Set 1 Total CK (CPK) Range Set
N	Value	Description
1616	.S	Legitimately skipped
273	1	1=Range Set 1
2	2	2=Range Set 2
387		Missing

MIF24A3		Day 2/Set 1 Total CK (CPK) Words Code
N	Value	Description
1616	.S	Legitimately skipped
33	1	1=absent/negative/normal
0	2	2=trace/weak positive
7	3	3=present/positive/abnormal
622		Missing

MIF24B		Day 2/Set 1 CK-MB
N	Value	Description
232	Range	0 - 613 (median=2.41 mean=15.893 std=56.336)
1616	.S	Legitimately skipped
430		Missing

MIF24B1		Day 2/Set 1 CK-MB Units
N	Value	Description
1616	.S	Legitimately skipped
207	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
455		Missing

MIF24B2		Day 2/Set 1 CK-MB Range Set
N	Value	Description
1616	.S	Legitimately skipped
206	1	1=Range Set 1
5	2	2=Range Set 2
451		Missing

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MIF24B3		Day 2/Set 1 CK-MB Words Code
N	Value	Description
1616	.S	Legitimately skipped
32	1	1=absent/negative/normal
0	2	2=trace/weak positive
3	3	3=present/positive/abnormal
627		Missing

MIF24C		Day 2/Set 1 Total LDH
N	Value	Description
14	Range	131 - 1487 (median=201 mean=489.8 std=498.5)
1616	.S	Legitimately skipped
648		Missing

MIF24C1		Day 2/Set 1 Total LDH Units
N	Value	Description
1616	.S	Legitimately skipped
0	1	1=ng/mL
13	2	2=Units/L
0	3	3=Åµg/L
649		Missing

MIF24C2		Day 2/Set 1 Total LDH Range Set
N	Value	Description
1616	.S	Legitimately skipped
14	1	1=Range Set 1
0	2	2=Range Set 2
648		Missing

MIF24C3		Day 2/Set 1 Total LDH Words Code
N	Value	Description
1616	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
662		Missing

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MIF24D1		Day 2/Set 1 LDH-1 Units
N	Value	Description
1616	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
662		Missing

MIF24D2		Day 2/Set 1 LDH-1 Range Set
N	Value	Description
1616	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2
661		Missing

MIF24D3		Day 2/Set 1 LDH-1 Words Code
N	Value	Description
1616	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
662		Missing

MIF24E		Day 2/Set 1 LDH-2
N	Value	Description
2	Range	28.9 - 195 (median=111.95 mean=111.950 std=117.450)
1616	.S	Legitimately skipped
660		Missing

MIF24E1		Day 2/Set 1 LDH-2 Units
N	Value	Description
1616	.S	Legitimately skipped
0	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L
661		Missing

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MIF24E2		Day 2/Set 1 LDH-2 Range Set
N	Value	Description
1616	.S	Legitimately skipped
2	1	1=Range Set 1
0	2	2=Range Set 2
660		Missing

MIF24E3		Day 2/Set 1 LDH-2 Words Code
N	Value	Description
1616	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
662		Missing

MIF24F		Day 2/Set 1 Troponin <
N	Value	Description
1616	.S	Legitimately skipped
325	0	0=n/a
213	1	1=<
124		Missing

MIF24F0		Day 2/Set 1 Troponin Value
N	Value	Description
547	Range	0 - 8229 (median=0.04 mean=18.556 std=352.402)
1616	.S	Legitimately skipped
115		Missing

MIF24F1		Day 2/Set 1 Troponin Units
N	Value	Description
1616	.S	Legitimately skipped
466	1	1=ng/mL
0	2	2=Units/L
5	3	3=Åµg/L
191		Missing

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MIF24F2		Day 2/Set 1 Troponin Range Set
N	Value	Description
1616	.S	Legitimately skipped
489	1	1=Range Set 1
13	2	2=Range Set 2
160		Missing

MIF24F3		Day 2/Set 1 Troponin Words Code
N	Value	Description
1616	.S	Legitimately skipped
134	1	1=absent/negative/normal
6	2	2=trace/weak positive
85	3	3=present/positive/abnormal
437		Missing

MIF24F4		Day 2/Set 1 Type of Troponin
N	Value	Description
1616	.S	Legitimately skipped
30	1	1= Troponin, type not specified
283	2	2= Troponin I
231	3	3= Troponin T
4	4	4= High Sensitivity Troponin (HS)
4	5	5= Unsure
110		Missing

MIF24G		second set of enzymes reported second day
N	Value	Description
1509	.S	Legitimately skipped
522	0	0=No/NR
246	1	1=Yes
1		Missing

MIF25A		Day 2/Set 2 Total CK (CPK)
N	Value	Description
133	Range	16 - 12326 (median=95 mean=468.0 std=1385.3)
1915	.S	Legitimately skipped
230		Missing

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MIF25A1		Day 2/Set 2 Total CK (CPK) Units
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=ng/mL
127	2	2=Units/L
0	3	3=Åµg/L
236		Missing

MIF25A2		Day 2/Set 2 Total CK (CPK) Range Set
N	Value	Description
1915	.S	Legitimately skipped
127	1	1=Range Set 1
3	2	2=Range Set 2
233		Missing

MIF25A3		Day 2/Set 2 Total CK (CPK) Words Code
N	Value	Description
1915	.S	Legitimately skipped
21	1	1=absent/negative/normal
0	2	2=trace/weak positive
4	3	3=present/positive/abnormal
338		Missing

MIF25B		Day 2/Set 2 CK-MB
N	Value	Description
104	Range	0.06 - 207.3 (median=2.15 mean=12.935 std=35.164)
1915	.S	Legitimately skipped
259		Missing

MIF25B1		Day 2/Set 2 CK-MB Units
N	Value	Description
1915	.S	Legitimately skipped
100	1	1=ng/mL
2	2	2=Units/L
0	3	3=Åµg/L
261		Missing

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MIF25B2		Day 2/Set 2 CK-MB Range Set
N	Value	Description
1915	.S	Legitimately skipped
98	1	1=Range Set 1
5	2	2=Range Set 2
260		Missing

MIF25B3		Day 2/Set 2 CK-MB Words Code
N	Value	Description
1915	.S	Legitimately skipped
18	1	1=absent/negative/normal
0	2	2=trace/weak positive
2	3	3=present/positive/abnormal
343		Missing

MIF25C		Day 2/Set 2 Total LDH
N	Value	Description
6	Range	110 - 268 (median=165.5 mean=178.17 std=55.50)
1915	.S	Legitimately skipped
357		Missing

MIF25C1		Day 2/Set 2 Total LDH Units
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=ng/mL
6	2	2=Units/L
0	3	3=Âµg/L
357		Missing

MIF25C2		Day 2/Set 2 Total LDH Range Set
N	Value	Description
1915	.S	Legitimately skipped
6	1	1=Range Set 1
0	2	2=Range Set 2
357		Missing

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MIF25C3		Day 2/Set 2 Total LDH Words Code
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
363		Missing

MIF25D1		Day 2/Set 2 LDH-1 Units
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
363		Missing

MIF25D2		Day 2/Set 2 LDH-1 Range Set
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
363		Missing

MIF25D3		Day 2/Set 2 LDH-1 Words Code
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
363		Missing

MIF25E		Day 2/Set 2 LDH-2
N	Value	Description
1915	.S	Legitimately skipped
363		Missing

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MIF25E1		Day 2/Set 2 LDH-2 Units
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
363		Missing

MIF25E2		Day 2/Set 2 LDH-2 Range Set
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
363		Missing

MIF25E3		Day 2/Set 2 LDH-2 Words Code
N	Value	Description
1915	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
363		Missing

MIF25F		Day 2/Set 2 Troponin <
N	Value	Description
1915	.S	Legitimately skipped
147	0	0=n/a
71	1	1=<
145		Missing

MIF25F0		Day 2/Set 2 Troponin Value
N	Value	Description
224	Range	0 - 11000 (median=0.065 mean=51.5741 std=734.9034)
1915	.S	Legitimately skipped
139		Missing

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MIF25F1		Day 2/Set 2 Troponin Units
N	Value	Description
1915	.S	Legitimately skipped
199	1	1=ng/mL
0	2	2=Units/L
3	3	3=Åµg/L
161		Missing

MIF25F2		Day 2/Set 2 Troponin Range Set
N	Value	Description
1915	.S	Legitimately skipped
209	1	1=Range Set 1
6	2	2=Range Set 2
148		Missing

MIF25F3		Day 2/Set 2 Troponin Words Code
N	Value	Description
1915	.S	Legitimately skipped
49	1	1=absent/negative/normal
1	2	2=trace/weak positive
46	3	3=present/positive/abnormal
267		Missing

MIF25F4		Day 2/Set 2 Type of Troponin
N	Value	Description
1915	.S	Legitimately skipped
11	1	1= Troponin, type not specified
128	2	2= Troponin I
85	3	3= Troponin T
2	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure
137		Missing

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MIF26A		Day 3/Set 1 Total CK (CPK)
N	Value	Description
75	Range	24 - 14206 (median=173 mean=914.0 std=2320.9)
1991	.S	Legitimately skipped
212		Missing

MIF26A1		Day 3/Set 1 Total CK (CPK) Units
N	Value	Description
1991	.S	Legitimately skipped
0	1	1=ng/mL
68	2	2=Units/L
0	3	3=Åµg/L
219		Missing

MIF26A2		Day 3/Set 1 Total CK (CPK) Range Set
N	Value	Description
1991	.S	Legitimately skipped
68	1	1=Range Set 1
1	2	2=Range Set 2
218		Missing

MIF26A3		Day 3/Set 1 Total CK (CPK) Words Code
N	Value	Description
1991	.S	Legitimately skipped
4	1	1=absent/negative/normal
0	2	2=trace/weak positive
5	3	3=present/positive/abnormal
278		Missing

MIF26B		Day 3/Set 1 CK-MB
N	Value	Description
57	Range	0.22 - 300 (median=2.7 mean=13.44 std=41.51)
1991	.S	Legitimately skipped
230		Missing

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MIF26B1		Day 3/Set 1 CK-MB Units
N	Value	Description
1991	.S	Legitimately skipped
53	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L
233		Missing

MIF26B2		Day 3/Set 1 CK-MB Range Set
N	Value	Description
1991	.S	Legitimately skipped
54	1	1=Range Set 1
1	2	2=Range Set 2
232		Missing

MIF26B3		Day 3/Set 1 CK-MB Words Code
N	Value	Description
1991	.S	Legitimately skipped
7	1	1=absent/negative/normal
0	2	2=trace/weak positive
1	3	3=present/positive/abnormal
279		Missing

MIF26C		Day 3/Set 1 Total LDH
N	Value	Description
9	Range	138 - 1138 (median=305 mean=478.8 std=382.9)
1991	.S	Legitimately skipped
278		Missing

MIF26C1		Day 3/Set 1 Total LDH Units
N	Value	Description
1991	.S	Legitimately skipped
0	1	1=ng/mL
6	2	2=Units/L
0	3	3=Åµg/L
281		Missing

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MIF26C2		Day 3/Set 1 Total LDH Range Set
N	Value	Description
1991	.S	Legitimately skipped
7	1	1=Range Set 1
0	2	2=Range Set 2
280		Missing

MIF26C3		Day 3/Set 1 Total LDH Words Code
N	Value	Description
1991	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
287		Missing

MIF26D1		Day 3/Set 1 LDH-1 Units
N	Value	Description
1991	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
287		Missing

MIF26D2		Day 3/Set 1 LDH-1 Range Set
N	Value	Description
1991	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2
286		Missing

MIF26D3		Day 3/Set 1 LDH-1 Words Code
N	Value	Description
1991	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
287		Missing

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MIF26E		Day 3/Set 1 LDH-2
N	Value	Description
1	Range	29.8 - 29.8 (median=29.8 mean=29.80 std=)
1991	.S	Legitimately skipped
286		Missing

MIF26E1		Day 3/Set 1 LDH-2 Units
N	Value	Description
1991	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
287		Missing

MIF26E2		Day 3/Set 1 LDH-2 Range Set
N	Value	Description
1991	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2
286		Missing

MIF26E3		Day 3/Set 1 LDH-2 Words Code
N	Value	Description
1991	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
287		Missing

MIF26F		Day 3/Set 1 Troponin <
N	Value	Description
1991	.S	Legitimately skipped
91	0	0=n/a
35	1	1=<
161		Missing

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MIF26F0		Day 3/Set 1 Troponin Value
N	Value	Description
129	Range	0 - 11631 (median=0.11 mean=94.130 std=1023.843)
1991	.S	Legitimately skipped
158		Missing

MIF26F1		Day 3/Set 1 Troponin Units
N	Value	Description
1991	.S	Legitimately skipped
115	1	1=ng/mL
0	2	2=Units/L
3	3	3=Åµg/L
169		Missing

MIF26F2		Day 3/Set 1 Troponin Range Set
N	Value	Description
1991	.S	Legitimately skipped
119	1	1=Range Set 1
5	2	2=Range Set 2
163		Missing

MIF26F3		Day 3/Set 1 Troponin Words Code
N	Value	Description
1991	.S	Legitimately skipped
19	1	1=absent/negative/normal
0	2	2=trace/weak positive
24	3	3=present/positive/abnormal
244		Missing

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MIF26F4		Day 3/Set 1 Type of Troponin
N	Value	Description
1991	.S	Legitimately skipped
9	1	1= Troponin, type not specified
72	2	2= Troponin I
46	3	3= Troponin T
2	4	4= High Sensitivity Troponin (HS)
2	5	5= Unsure
156		Missing

MIF26G		second set of enzymes reported third day
N	Value	Description
1884	.S	Legitimately skipped
364	0	0=No/NR
29	1	1=Yes
1		Missing

MIF27A		Day 3/Set 2 Total CK (CPK)
N	Value	Description
14	Range	57 - 10891 (median=194 mean=1252.7 std=2933.0)
2113	.S	Legitimately skipped
151		Missing

MIF27A1		Day 3/Set 2 Total CK (CPK) Units
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=ng/mL
14	2	2=Units/L
0	3	3=Åµg/L
151		Missing

MIF27A2		Day 3/Set 2 Total CK (CPK) Range Set
N	Value	Description
2113	.S	Legitimately skipped
14	1	1=Range Set 1
0	2	2=Range Set 2
151		Missing

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MIF27A3		Day 3/Set 2 Total CK (CPK) Words Code
N	Value	Description
2113	.S	Legitimately skipped
2	1	1=absent/negative/normal
0	2	2=trace/weak positive
3	3	3=present/positive/abnormal
160		Missing

MIF27B		Day 3/Set 2 CK-MB
N	Value	Description
11	Range	0.83 - 7.1 (median=2.7 mean=3.62 std=2.37)
2113	.S	Legitimately skipped
154		Missing

MIF27B1		Day 3/Set 2 CK-MB Units
N	Value	Description
2113	.S	Legitimately skipped
11	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
154		Missing

MIF27B2		Day 3/Set 2 CK-MB Range Set
N	Value	Description
2113	.S	Legitimately skipped
11	1	1=Range Set 1
0	2	2=Range Set 2
154		Missing

MIF27B3		Day 3/Set 2 CK-MB Words Code
N	Value	Description
2113	.S	Legitimately skipped
2	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
163		Missing

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MIF27C		Day 3/Set 2 Total LDH
N	Value	Description
2113	.S	Legitimately skipped
165		Missing

MIF27C1		Day 3/Set 2 Total LDH Units
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
165		Missing

MIF27C2		Day 3/Set 2 Total LDH Range Set
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
165		Missing

MIF27C3		Day 3/Set 2 Total LDH Words Code
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
165		Missing

MIF27D1		Day 3/Set 2 LDH-1 Units
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
165		Missing

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MIF27D2		Day 3/Set 2 LDH-1 Range Set
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
165		Missing

MIF27D3		Day 3/Set 2 LDH-1 Words Code
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
165		Missing

MIF27E		Day 3/Set 2 LDH-2
N	Value	Description
2113	.S	Legitimately skipped
165		Missing

MIF27E1		Day 3/Set 2 LDH-2 Units
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
165		Missing

MIF27E2		Day 3/Set 2 LDH-2 Range Set
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
165		Missing

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MIF27E3		Day 3/Set 2 LDH-2 Words Code
N	Value	Description
2113	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
165		Missing

MIF27F		Day 3/Set 2 Troponin <
N	Value	Description
2113	.S	Legitimately skipped
19	0	0=n/a
7	1	1=<
139		Missing

MIF27F0		Day 3/Set 2 Troponin Value
N	Value	Description
24	Range	0.01 - 24.7 (median=0.056 mean=1.4940 std=5.0669)
2113	.S	Legitimately skipped
141		Missing

MIF27F1		Day 3/Set 2 Troponin Units
N	Value	Description
2113	.S	Legitimately skipped
22	1	1=ng/mL
0	2	2=Units/L
1	3	3=Åµg/L
142		Missing

MIF27F2		Day 3/Set 2 Troponin Range Set
N	Value	Description
2113	.S	Legitimately skipped
23	1	1=Range Set 1
0	2	2=Range Set 2
142		Missing

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MIF27F3		Day 3/Set 2 Troponin Words Code
N	Value	Description
2113	.S	Legitimately skipped
5	1	1=absent/negative/normal
0	2	2=trace/weak positive
4	3	3=present/positive/abnormal
156		Missing

MIF27F4		Day 3/Set 2 Type of Troponin
N	Value	Description
2113	.S	Legitimately skipped
1	1	1= Troponin, type not specified
13	2	2= Troponin I
10	3	3= Troponin T
0	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure
141		Missing

MIF28A		Day 4/Set 1 Total CK (CPK)
N	Value	Description
33	Range	20 - 44034 (median=134 mean=1965.2 std=7775.1)
2077	.S	Legitimately skipped
168		Missing

MIF28A1		Day 4/Set 1 Total CK (CPK) Units
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=ng/mL
31	2	2=Units/L
0	3	3=Åµg/L
170		Missing

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MIF28A2		Day 4/Set 1 Total CK (CPK) Range Set
N	Value	Description
2077	.S	Legitimately skipped
32	1	1=Range Set 1
1	2	2=Range Set 2
168		Missing

MIF28A3		Day 4/Set 1 Total CK (CPK) Words Code
N	Value	Description
2077	.S	Legitimately skipped
2	1	1=absent/negative/normal
0	2	2=trace/weak positive
3	3	3=present/positive/abnormal
196		Missing

MIF28B		Day 4/Set 1 CK-MB
N	Value	Description
24	Range	0.27 - 300 (median=2.65 mean=16.841 std=60.519)
2077	.S	Legitimately skipped
177		Missing

MIF28B1		Day 4/Set 1 CK-MB Units
N	Value	Description
2077	.S	Legitimately skipped
21	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
180		Missing

MIF28B2		Day 4/Set 1 CK-MB Range Set
N	Value	Description
2077	.S	Legitimately skipped
21	1	1=Range Set 1
1	2	2=Range Set 2
179		Missing

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MIF28B3		Day 4/Set 1 CK-MB Words Code
N	Value	Description
2077	.S	Legitimately skipped
1	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
200		Missing

MIF28C		Day 4/Set 1 Total LDH
N	Value	Description
2	Range	570 - 897 (median=733.5 mean=733.50 std=231.22)
2077	.S	Legitimately skipped
199		Missing

MIF28C1		Day 4/Set 1 Total LDH Units
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L
200		Missing

MIF28C2		Day 4/Set 1 Total LDH Range Set
N	Value	Description
2077	.S	Legitimately skipped
2	1	1=Range Set 1
0	2	2=Range Set 2
199		Missing

MIF28C3		Day 4/Set 1 Total LDH Words Code
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
201		Missing

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MIF28D1		Day 4/Set 1 LDH-1 Units
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
201		Missing

MIF28D2		Day 4/Set 1 LDH-1 Range Set
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
201		Missing

MIF28D3		Day 4/Set 1 LDH-1 Words Code
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
201		Missing

MIF28E		Day 4/Set 1 LDH-2
N	Value	Description
2077	.S	Legitimately skipped
201		Missing

MIF28E1		Day 4/Set 1 LDH-2 Units
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
201		Missing

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MIF28E2		Day 4/Set 1 LDH-2 Range Set
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
201		Missing

MIF28E3		Day 4/Set 1 LDH-2 Words Code
N	Value	Description
2077	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
201		Missing

MIF28F		Day 4/Set 1 Troponin <
N	Value	Description
2077	.S	Legitimately skipped
43	0	0=n/a
12	1	1=<
146		Missing

MIF28F0		Day 4/Set 1 Troponin Value
N	Value	Description
57	Range	0 - 5000 (median=0.11 mean=88.895 std=662.117)
2077	.S	Legitimately skipped
144		Missing

MIF28F1		Day 4/Set 1 Troponin Units
N	Value	Description
2077	.S	Legitimately skipped
50	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
151		Missing

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MIF28F2		Day 4/Set 1 Troponin Range Set
N	Value	Description
2077	.S	Legitimately skipped
52	1	1=Range Set 1
1	2	2=Range Set 2
148		Missing

MIF28F3		Day 4/Set 1 Troponin Words Code
N	Value	Description
2077	.S	Legitimately skipped
8	1	1=absent/negative/normal
1	2	2=trace/weak positive
6	3	3=present/positive/abnormal
186		Missing

MIF28F4		Day 4/Set 1 Type of Troponin
N	Value	Description
2077	.S	Legitimately skipped
6	1	1= Troponin, type not specified
27	2	2= Troponin I
25	3	3= Troponin T
0	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure
143		Missing

MIF28G		second set of enzymes reported fourth day
N	Value	Description
1970	.S	Legitimately skipped
292	0	0=No/NR
15	1	1=Yes
1		Missing

MIF29A		Day 4/Set 2 Total CK (CPK)
N	Value	Description
2	Range	711 - 60661 (median=30686 mean=30686.0 std=42391.1)
2124	.S	Legitimately skipped
152		Missing

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MIF29A1		Day 4/Set 2 Total CK (CPK) Units
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=ng/mL
2	2	2=Units/L
0	3	3=Åµg/L
152		Missing

MIF29A2		Day 4/Set 2 Total CK (CPK) Range Set
N	Value	Description
2124	.S	Legitimately skipped
2	1	1=Range Set 1
0	2	2=Range Set 2
152		Missing

MIF29A3		Day 4/Set 2 Total CK (CPK) Words Code
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
1	3	3=present/positive/abnormal
153		Missing

MIF29B		Day 4/Set 2 CK-MB
N	Value	Description
1	Range	3.19 - 3.19 (median=3.19 mean=3.190 std=)
2124	.S	Legitimately skipped
153		Missing

MIF29B1		Day 4/Set 2 CK-MB Units
N	Value	Description
2124	.S	Legitimately skipped
1	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
153		Missing

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MIF29B2		Day 4/Set 2 CK-MB Range Set
N	Value	Description
2124	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2
153		Missing

MIF29B3		Day 4/Set 2 CK-MB Words Code
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
154		Missing

MIF29C		Day 4/Set 2 Total LDH
N	Value	Description
1	Range	848 - 848 (median=848 mean=848.0 std=)
2124	.S	Legitimately skipped
153		Missing

MIF29C1		Day 4/Set 2 Total LDH Units
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=ng/mL
1	2	2=Units/L
0	3	3=Åµg/L
153		Missing

MIF29C2		Day 4/Set 2 Total LDH Range Set
N	Value	Description
2124	.S	Legitimately skipped
1	1	1=Range Set 1
0	2	2=Range Set 2
153		Missing

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MIF29C3		Day 4/Set 2 Total LDH Words Code
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
154		Missing

MIF29D1		Day 4/Set 2 LDH-1 Units
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
154		Missing

MIF29D2		Day 4/Set 2 LDH-1 Range Set
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
154		Missing

MIF29D3		Day 4/Set 2 LDH-1 Words Code
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
154		Missing

MIF29E		Day 4/Set 2 LDH-2
N	Value	Description
2124	.S	Legitimately skipped
154		Missing

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MIF29E1		Day 4/Set 2 LDH-2 Units
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
154		Missing

MIF29E2		Day 4/Set 2 LDH-2 Range Set
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=Range Set 1
0	2	2=Range Set 2
154		Missing

MIF29E3		Day 4/Set 2 LDH-2 Words Code
N	Value	Description
2124	.S	Legitimately skipped
0	1	1=absent/negative/normal
0	2	2=trace/weak positive
0	3	3=present/positive/abnormal
154		Missing

MIF29F		Day 4/Set 2 Troponin <
N	Value	Description
2124	.S	Legitimately skipped
9	0	0=n/a
4	1	1=<
141		Missing

MIF29F0		Day 4/Set 2 Troponin Value
N	Value	Description
13	Range	0.01 - 1.61 (median=0.068 mean=0.1913 std=0.4326)
2124	.S	Legitimately skipped
141		Missing

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MIF29F1		Day 4/Set 2 Troponin Units
N	Value	Description
2124	.S	Legitimately skipped
13	1	1=ng/mL
0	2	2=Units/L
0	3	3=Åµg/L
141		Missing

MIF29F2		Day 4/Set 2 Troponin Range Set
N	Value	Description
2124	.S	Legitimately skipped
13	1	1=Range Set 1
0	2	2=Range Set 2
141		Missing

MIF29F3		Day 4/Set 2 Troponin Words Code
N	Value	Description
2124	.S	Legitimately skipped
3	1	1=absent/negative/normal
1	2	2=trace/weak positive
0	3	3=present/positive/abnormal
150		Missing

MIF29F4		Day 4/Set 2 Type of Troponin
N	Value	Description
2124	.S	Legitimately skipped
0	1	1= Troponin, type not specified
6	2	2= Troponin I
7	3	3= Troponin T
0	4	4= High Sensitivity Troponin (HS)
0	5	5= Unsure
141		Missing

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MIF30		Mention of having trauma/surgery
N	Value	Description
881	.S	Legitimately skipped
1319	0	0=No/NR
77	1	1=Yes
1		Missing

MIF30A		Cardiac Procedure
N	Value	Description
2201	.S	Legitimately skipped
49	0	0=No
27	1	1=Yes
1		Missing

MIF30B		CPR or cardioversion
N	Value	Description
2201	.S	Legitimately skipped
64	0	0=No
10	1	1=Yes
3		Missing

MIF30C		Other cardiac trauma
N	Value	Description
2201	.S	Legitimately skipped
73	0	0=No
2	1	1=Yes
2		Missing

MIF30C2		Specify type of cardiac trauma
N	Value	Description
1	Present	Text suppressed
2276	~	Legitimately skipped
1		Missing

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<i>MIF30D</i>		<i>Rhabdomyolysis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2201	.S	Legitimately skipped
62	0	0=No
13	1	1=Yes
2		Missing

<i>MIF30E</i>		<i>Intramuscular Injection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2201	.S	Legitimately skipped
74	0	0=No
0	1	1=Yes
3		Missing

<i>MIF30F</i>		<i>Non-cardiac procedure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2201	.S	Legitimately skipped
57	0	0=No
17	1	1=Yes
3		Missing

<i>MIF30F2</i>		<i>Specify non-cardiac procedure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17	Present	Text suppressed
2261	~	Legitimately skipped

<i>MIF30G</i>		<i>Non-cardiac trauma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2201	.S	Legitimately skipped
64	0	0=No
10	1	1=Yes
3		Missing

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MIF31		Item number from biomarkers section
N	Value	Description
69	Range	22 - 28 (median=22 mean=23.2 std=1.9)
2201	.S	Legitimately skipped
8		Missing

MIF32		Evidence of hemolytic disease
N	Value	Description
2275	0	0=No/NR
2	1	1=Yes
1		Missing

MIF33		Have active liver disease?
N	Value	Description
2185	0	0=No/NR
93	1	1=Yes

MIF34		Any 12 lead ECGs taken?
N	Value	Description
192	0	0=No
1643	1	1=Yes
442	9	9=NR
1		Missing

MIF34B		Copy of first ECG enclosed?
N	Value	Description
676	.S	Legitimately skipped
756	0	0=No
836	1	1=Yes
10		Missing

MIF34D		Copy of second ECG enclosed?
N	Value	Description
1419	.S	Legitimately skipped
242	0	0=No
542	1	1=Yes
75		Missing

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MIF34F		Copy of third ECG enclosed?
N	Value	Description
1748	.S	Legitimately skipped
111	0	0=No
303	1	1=Yes
116		Missing

MIF34H		Copy of last ECG enclosed?
N	Value	Description
1875	.S	Legitimately skipped
64	0	0=No
218	1	1=Yes
121		Missing

MIF35		Transthoracic echocardiogram performed?
N	Value	Description
1409	0	0=No/NR
869	1	1=Yes

MIF35A		LV ejection fraction TTE
N	Value	Description
794	Range	5 - 81 (median=55 mean=52.0 std=15.3)
1409	.S	Legitimately skipped
75		Missing

MIF36		Nuclear Medicare Scan performed?
N	Value	Description
2186	0	0=No/NR
92	1	1=Yes

MIF36A		Ejection fraction of scan
N	Value	Description
83	Range	10 - 85 (median=58 mean=55.4 std=17.2)
2186	.S	Legitimately skipped
9		Missing

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MIF36B		RV(%) of scan
N	Value	Description
2186	.S	Legitimately skipped
92		Missing

MIF36C		Positive stress test for ischemia
N	Value	Description
2186	.S	Legitimately skipped
71	0	0=No/NR
21	1	1=Yes

MIF37		Stress test performed?
N	Value	Description
2118	0	0=No/NR
160	1	1=Yes

MIF37A		Ejection fraction of Stress Test
N	Value	Description
75	Range	17 - 96 (median=60 mean=57.0 std=17.5)
2118	.S	Legitimately skipped
85		Missing

MIF37B		Stress test positive for ischemia
N	Value	Description
2118	.S	Legitimately skipped
131	0	0=No/NR
26	1	1=Yes
3		Missing

MIF37C		Greater than or equal to 1 mm ST?
N	Value	Description
2118	.S	Legitimately skipped
155	0	0=No/NR
2	1	1=Yes
3		Missing

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MIF37D		Ischemic pain or equivalent occurred?
N	Value	Description
2118	.S	Legitimately skipped
154	0	0=No/NR
4	1	1=Yes
2		Missing

MIF38		Was a coronary angiography performed?
N	Value	Description
1931	0	0=No/NR
347	1	1=Yes

MIF38B		Ejection % of coronary angiography
N	Value	Description
155	Range	10 - 70 (median=50 mean=48.1 std=14.6)
1931	.S	Legitimately skipped
192		Missing

MIF38C		70% or greater obstruction?
N	Value	Description
1931	.S	Legitimately skipped
97	0	0=No
230	1	1=Yes
18	9	9=NR
2		Missing

MIF38D		Coronary bypass present?
N	Value	Description
1931	.S	Legitimately skipped
226	0	0=No
35	1	1=Yes
84	9	9=NR
2		Missing

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MIF38D1		If yes, number of occluded grafts
N	Value	Description
33	Range	0 - 3 (median=0 mean=0.5 std=0.8)
2243	.S	Legitimately skipped
2		Missing

MIF39		Coronary reperfusion attempted?
N	Value	Description
2049	0	0=No/NR
229	1	1=Yes

MIF39A		Approximate time onset to reperfusion
N	Value	Description
2049	.S	Legitimately skipped
4	1	1= < 2 hours
7	2	2= 2 to 4 hours
3	3	3= 4 to 6 hours
4	4	4= 6 to 12 hours
7	5	5= 12 to 24 hours
113	6	6= > 24 hours
90	7	7= Not Sure
1		Missing

MIF40A		Coronary artery bypass graft surgery
N	Value	Description
2216	0	0=No/NR
61	1	1=Yes
1		Missing

MIF40A2		Time of Coronary artery bypass
N	Value	Description
9	Range	4:16:00 - 19:05:00
2216	.S	Legitimately skipped
53		Missing

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MIF40A2HR		Hour of Time of Coronary artery bypass
N	Value	Description
9	Range	4 - 19 (median=12 mean=11.6 std=4.2)
2269		Missing

MIF40A2MN		Minute of Time of Coronary artery bypass
N	Value	Description
9	Range	5 - 55 (median=36 mean=32.2 std=20.4)
2269		Missing

MIF40A2SC		Second of Time of Coronary artery bypass
N	Value	Description
4	Range	0 - 0 (median=0 mean=0.0 std=0.0)
2274		Missing

MIF40A3		AM/PM of the coronary bypass
N	Value	Description
2269	.S	Legitimately skipped
1	1	1=am
1	2	2=pm
7		Missing

MIF40B		Coronary atherectomy
N	Value	Description
2276	0	0=No/NR
1	1	1=Yes
1		Missing

MIF40B2		Time of coronary atherectomy
N	Value	Description
2276	.S	Legitimately skipped
2		Missing

MIF40B2HR		Hour of Time of coronary atherectomy
N	Value	Description
2278		Missing

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MIF40B2MN		Minute of Time of coronary atherectomy
N	Value	Description
2278		Missing

MIF40B2SC		Second of Time of coronary atherectomy
N	Value	Description
2278		Missing

MIF40B3		AM/PM of coronary atherectomy
N	Value	Description
2277	.S	Legitimately skipped
0	1	1=am
0	2	2=pm
1		Missing

MIF40C		Intra-arterial/intravenous thrombolytic
N	Value	Description
2272	0	0=No/NR
5	1	1=Yes
1		Missing

MIF40C2		Time of thrombolytic
N	Value	Description
2	Range	19:02:00 - 20:15:00
2272	.S	Legitimately skipped
4		Missing

MIF40C2HR		Hour of Time of thrombolytic
N	Value	Description
2	Range	19 - 20 (median=19.5 mean=19.50 std=0.71)
2276		Missing

MIF40C2MN		Minute of Time of thrombolytic
N	Value	Description
2	Range	2 - 15 (median=8.5 mean=8.50 std=9.19)
2276		Missing

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MIF40C2SC		Second of Time of thrombolytic
N	Value	Description
2	Range	0 - 0 (median=0 mean=0.0 std=0.0)
2276		Missing

MIF40C3		AM/PM of thrombolytic
N	Value	Description
2275	.S	Legitimately skipped
0	1	1=am
2	2	2=pm
1		Missing

MIF40D		Coronary angioplasty w/o stent
N	Value	Description
2253	0	0=No/NR
23	1	1=Yes
2		Missing

MIF40D2		Time of coronary angioplasty w/e stent
N	Value	Description
10	Range	8:41:00 - 16:17:00
2254	.S	Legitimately skipped
14		Missing

MIF40D2HR		Hour of Time of coronary angioplasty w/e stent
N	Value	Description
10	Range	8 - 16 (median=11.5 mean=12.00 std=3.06)
2268		Missing

MIF40D2MN		Minute of Time of coronary angioplasty w/e stent
N	Value	Description
10	Range	0 - 45 (median=17 mean=23.8 std=18.0)
2268		Missing

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MIF40D2SC		Second of Time of coronary angioplasty w/e stent
N	Value	Description
2	Range	0 - 0 (median=0 mean=0.0 std=0.0)
2276		Missing

MIF40D3		AM/PM coronary angioplasty w/e stent
N	Value	Description
2276	.S	Legitimately skipped
0	1	1=am
0	2	2=pm
2		Missing

MIF40E		Coronary angioplasty with stent place
N	Value	Description
2110	0	0=No/NR
167	1	1=Yes
1		Missing

MIF40E2		Time of coronary angioplasty with stent
N	Value	Description
64	Range	1:02:00 - 19:34:00
2110	.S	Legitimately skipped
104		Missing

MIF40E2HR		Hour of Time of coronary angioplasty with stent
N	Value	Description
64	Range	1 - 19 (median=12 mean=12.1 std=3.7)
2214		Missing

MIF40E2MN		Minute of Time of coronary angioplasty with stent
N	Value	Description
64	Range	0 - 59 (median=26.5 mean=27.52 std=17.64)
2214		Missing

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MIF40E2SC		Second of Time of coronary angioplasty with stent
N	Value	Description
33	Range	0 - 0 (median=0 mean=0.0 std=0.0)
2245		Missing

MIF40E3		AM/PM of coronary angioplasty with stent
N	Value	Description
2258	.S	Legitimately skipped
4	1	1=am
4	2	2=pm
12		Missing

MIF40F		Valve surgery
N	Value	Description
2244	0	0=No/NR
34	1	1=Yes

MIF40G		Non-cardiac surgery
N	Value	Description
1940	0	0=No/NR
337	1	1=Yes
1		Missing

MIF40H		Aortic balloon pump
N	Value	Description
2259	0	0=No/NR
19	1	1=Yes

MIF40I		Pacemaker placement
N	Value	Description
2235	0	0=No/NR
43	1	1=Yes

MIF40J		Cardioversion or defibrillation
N	Value	Description
2238	0	0=No/NR
40	1	1=Yes

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MIF40J2		Time of cardioversion or defibrillation
N	Value	Description
6	Range	9:00:00 - 23:30:00
2238	.S	Legitimately skipped
34		Missing

MIF40J2HR		Hour of Time of cardioversion or defibrillation
N	Value	Description
6	Range	9 - 23 (median=10 mean=13.7 std=6.5)
2272		Missing

MIF40J2MN		Minute of Time of cardioversion or defibrillation
N	Value	Description
6	Range	0 - 30 (median=13.5 mean=13.83 std=12.62)
2272		Missing

MIF40J2SC		Second of Time of cardioversion or defibrillation
N	Value	Description
3	Range	0 - 0 (median=0 mean=0.0 std=0.0)
2275		Missing

MIF40J3		AM/PM of cardioversion or defibrillation
N	Value	Description
2270	.S	Legitimately skipped
0	1	1=am
2	2	2=pm
6		Missing

MIF40J4A		Ventricular Fibrillation/Flutter
N	Value	Description
2238	.S	Legitimately skipped
32	0	0=No/NR
7	1	1=Yes
1		Missing

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MIF40J4B		Ventricular Tachycardia (VT)
N	Value	Description
2238	.S	Legitimately skipped
32	0	0=No/NR
7	1	1=Yes
1		Missing

MIF40J4C		Asystole
N	Value	Description
2238	.S	Legitimately skipped
34	0	0=No/NR
5	1	1=Yes
1		Missing

MIF40J4D		Complete AV Block (3 HB)
N	Value	Description
2238	.S	Legitimately skipped
39	0	0=No/NR
0	1	1=Yes
1		Missing

MIF40J4E		Atrial Fibrillation/Flutter
N	Value	Description
2238	.S	Legitimately skipped
11	0	0=No/NR
28	1	1=Yes
1		Missing

MIF40J4F		Pulseless Electrical Activity (PEA)
N	Value	Description
2238	.S	Legitimately skipped
35	0	0=No/NR
4	1	1=Yes
1		Missing

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MIF41A1		Admission Meds of Nitroglycerin
N	Value	Description
1698	0	0=No/NR
580	1	1=Yes

MIF41A2		Discharge Meds of Nitroglycerin
N	Value	Description
2051	0	0=No/NR
226	1	1=Yes
1		Missing

MIF41B1		Admission of Beta Blockers
N	Value	Description
994	0	0=No/NR
1284	1	1=Yes

MIF41B2		Discharge Meds of Beta Blockers
N	Value	Description
1333	0	0=No/NR
944	1	1=Yes
1		Missing

MIF41C1		Admission Meds of Channel Blockers
N	Value	Description
1620	0	0=No/NR
657	1	1=Yes
1		Missing

MIF41C2		Discharge Meds of Channel Blockers
N	Value	Description
1862	0	0=No/NR
414	1	1=Yes
2		Missing

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MIF41D1		Admission Meds of ACE or ARB
N	Value	Description
1144	0	0=No/NR
1133	1	1=Yes
1		Missing

MIF41D2		Discharge Meds of ACE or ARB
N	Value	Description
1533	0	0=No/NR
743	1	1=Yes
2		Missing

MIF41E1		Admission Meds of scheduled aspirin
N	Value	Description
1057	0	0=No/NR
1220	1	1=Yes
1		Missing

MIF41E2		Discharge Meds of scheduled aspirin
N	Value	Description
1386	0	0=No/NR
891	1	1=Yes
1		Missing

MIF41F1		Admission Meds of Heparin/Enoxaparin
N	Value	Description
1475	0	0=No/NR
803	1	1=Yes

MIF41F2		Discharge Meds of of Heparin/Enoxaparin
N	Value	Description
2185	0	0=No/NR
91	1	1=Yes
2		Missing

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MIF41G1		Admission Meds of Coumadin,warfarin
N	Value	Description
1976	0	0=No/NR
302	1	1=Yes

MIF41G2		Discharge Meds of Coumadin,warfarin
N	Value	Description
2043	0	0=No/NR
233	1	1=Yes
2		Missing

MIF41H1		Admission Meds of Anti-platelet agents
N	Value	Description
1736	0	0=No/NR
542	1	1=Yes

MIF41H2		Discharge Meds of Anti-platelet agents
N	Value	Description
1897	0	0=No/NR
379	1	1=Yes
2		Missing

MIF41I1		Admission Meds of Statin
N	Value	Description
1047	0	0=No/NR
1231	1	1=Yes

MIF41I2		Discharge Meds of Statin
N	Value	Description
1311	0	0=No/NR
966	1	1=Yes
1		Missing

MIF42A		Treated with IV pressors
N	Value	Description
2151	0	0=No/NR
127	1	1=Yes

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MIF42B		Treated with IV nitroglycerin
N	Value	Description
2190	0	0=No/NR
88	1	1=Yes

MIF42C		Treated with inhibitors
N	Value	Description
2252	0	0=No/NR
26	1	1=Yes

MIF0A_FUTIME		Futime from Randomization date for (MIF0A)
N	Value	Description
2274	Range	594 - 5319 (median=2836.5 mean=2945.17 std=830.07)
4		Missing

MIF0D_FUTIME		Futime from Randomization date for (MIF0D)
N	Value	Description
2278	Range	5 - 4206 (median=1801 mean=1859.0 std=1052.5)

MIF2_FUTIME		Futime from Randomization date for (MIF2)
N	Value	Description
2212	Range	5 - 4206 (median=1819 mean=1870.8 std=1057.6)
66		Missing

MIF2B_FUTIME		Futime from Randomization date for (MIF2B)
N	Value	Description
2277	Range	5 - 4460 (median=1803 mean=1861.7 std=1053.5)
1		Missing

MIF3_FUTIME		Futime from Randomization date for (MIF3)
N	Value	Description
2266	Range	9 - 4227 (median=1801 mean=1865.2 std=1050.4)
12		Missing

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MIF22_FUTIME		Futime from Randomization date for (MIF22)
N	Value	Description
1308	Range	-78 - 4130 (median=1861.5 mean=1873.55 std=1064.41)
970		Missing

MIF23_FUTIME		Futime from Randomization date for (MIF23)
N	Value	Description
523	Range	-78 - 4033 (median=1852 mean=1871.8 std=1100.5)
1755		Missing

MIF24_FUTIME		Futime from Randomization date for (MIF24)
N	Value	Description
587	Range	6 - 3991 (median=1761 mean=1787.1 std=1025.3)
1691		Missing

MIF25_FUTIME		Futime from Randomization date for (MIF25)
N	Value	Description
245	Range	6 - 3657 (median=1822 mean=1806.0 std=971.4)
2033		Missing

MIF26_FUTIME		Futime from Randomization date for (MIF26)
N	Value	Description
159	Range	-108 - 3862 (median=1805 mean=1781.9 std=992.9)
2119		Missing

MIF27_FUTIME		Futime from Randomization date for (MIF27)
N	Value	Description
28	Range	504 - 3172 (median=1794 mean=1733.1 std=765.0)
2250		Missing

MIF28_FUTIME		Futime from Randomization date for (MIF28)
N	Value	Description
71	Range	50 - 4039 (median=1678 mean=1744.1 std=1057.7)
2207		Missing

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MIF29_FUTIME		Futime from Randomization date for (MIF29)
N	Value	Description
15	Range	777 - 3993 (median=1933 mean=2155.0 std=1043.6)
2263		Missing

MIF30A1_FUTIME		Futime from Randomization date for (MIF30A1)
N	Value	Description
26	Range	27 - 3988 (median=1273 mean=1529.5 std=1219.6)
2252		Missing

MIF30B1_FUTIME		Futime from Randomization date for (MIF30B1)
N	Value	Description
10	Range	193 - 3138 (median=1025 mean=1427.1 std=1116.3)
2268		Missing

MIF30C1_FUTIME		Futime from Randomization date for (MIF30C1)
N	Value	Description
2	Range	774 - 1065 (median=919.5 mean=919.50 std=205.77)
2276		Missing

MIF30D1_FUTIME		Futime from Randomization date for (MIF30D1)
N	Value	Description
4	Range	59 - 2778 (median=361 mean=889.8 std=1267.1)
2274		Missing

MIF30E1_FUTIME		Futime from Randomization date for (MIF30E1)
N	Value	Description
2278		Missing

MIF30F1_FUTIME		Futime from Randomization date for (MIF30F1)
N	Value	Description
14	Range	816 - 3886 (median=2349 mean=2479.4 std=963.3)
2264		Missing

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MIF30G1_FUTIME		Futime from Randomization date for (MIF30G1)
N	Value	Description
8	Range	262 - 3860 (median=1556 mean=1754.0 std=1270.1)
2270		Missing

MIF34A_FUTIME		Futime from Randomization date for (MIF34A)
N	Value	Description
1591	Range	5 - 4130 (median=1795 mean=1865.9 std=1050.9)
687		Missing

MIF34C_FUTIME		Futime from Randomization date for (MIF34C)
N	Value	Description
782	Range	6 - 4016 (median=1605.5 mean=1711.77 std=1048.01)
1496		Missing

MIF34E_FUTIME		Futime from Randomization date for (MIF34E)
N	Value	Description
412	Range	6 - 3972 (median=1454 mean=1633.8 std=1028.4)
1866		Missing

MIF34G_FUTIME		Futime from Randomization date for (MIF34G)
N	Value	Description
281	Range	22 - 3973 (median=1315 mean=1476.5 std=1040.1)
1997		Missing

MIF38A_FUTIME		Futime from Randomization date for (MIF38A)
N	Value	Description
335	Range	22 - 3949 (median=1546 mean=1695.4 std=1060.7)
1943		Missing

MIF40A1_FUTIME		Futime from Randomization date for (MIF40A1)
N	Value	Description
60	Range	154 - 3970 (median=1194 mean=1580.0 std=1050.4)
2218		Missing

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MIF40B1_FUTIME		Futime from Randomization date for (MIF40B1)
N	Value	Description
1	Range	108 - 108 (median=108 mean=108.0 std=)
2277		Missing

MIF40C1_FUTIME		Futime from Randomization date for (MIF40C1)
N	Value	Description
5	Range	134 - 1947 (median=235 mean=549.6 std=783.5)
2273		Missing

MIF40D1_FUTIME		Futime from Randomization date for (MIF40D1)
N	Value	Description
23	Range	141 - 3769 (median=2531 mean=2302.8 std=1092.9)
2255		Missing

MIF40E1_FUTIME		Futime from Randomization date for (MIF40E1)
N	Value	Description
163	Range	22 - 3901 (median=1638 mean=1737.4 std=1103.9)
2115		Missing

MIF40J1_FUTIME		Futime from Randomization date for (MIF40J1)
N	Value	Description
30	Range	62 - 3607 (median=1490 mean=1709.5 std=1167.0)
2248		Missing

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

ID		HCHS Participant ID
N	Value	Description
2238	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
1494	1	Occurrence 1
407	2	Occurrence 2
161	3	Occurrence 3
75	4	Occurrence 4
39	5	Occurrence 5
24	6	Occurrence 6
11	7	Occurrence 7
6	8	Occurrence 8
5	9	Occurrence 9
6	10	Occurrence 10
3	11	Occurrence 11
3	12	Occurrence 12
1	13	Occurrence 13
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19

MID16		Comments
N	Value	Description
234	Present	Text suppressed
2004		Missing

MID11		Evidence of cardiac pain?
N	Value	Description
1424	0	0=No
681	1	1=Yes
117	9	9=NR
16		Missing

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MID12		Level of cardiac biomarkers
N	Value	Description
268	1	1= Abnormal (at least one value at 2 X the upper limit normal)
149	2	2= Equivocal (> normal, but < 2X the upper limit normal)
45	3	3= Incomplete
857	4	4= Normal
902	5	5= None recorded
17		Missing

MID13		Interpretations of ECGs
N	Value	Description
5	1	1= Evolution of Major Q-Wave
46	2	2= Evolution of ST-T Elevation with or without Q-Wave
4	3	3= New LBBB
68	4	4= Evolution of ST-Depression/T wave inversion alone
1	5	5= Evolution of Minor Q-Wave alone
17	6	6= Single ECG with Major Q-wave
2	7	7= Single ECG with LBBB, described as new
2079	8	8= Absent, Uncodable or Other ECG
16		Missing

MID14		Myocardial infarction classification
N	Value	Description
117	1	1=Definite
119	2	2=Probable
1972	3	3=No MI
14	4	4=Unclassifiable
16		Missing

MID14A		Type of MI?
N	Value	Description
1987	.S	Legitimately skipped
58	0	0=Transmural
128	1	1=Subendocardial
48	9	9=Unsure/unknown
17		Missing

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Section B: Myocardial Infarction Ascertainment

MID14B		Location of MI?
N	Value	Description
1987	.S	Legitimately skipped
29	1	1=Anterior
3	2	2=Posterior
30	3	3=Inferior
24	4	4=Lateral
2	5	5=Septal
22	6	6=More than one area
124	9	9=Unable to determine
17		Missing

MID14C		Was the MI procedure-related?
N	Value	Description
1987	.S	Legitimately skipped
3	1	1=Yes, cardiovascular procedure
4	2	2=Yes, non-cardiovascular procedure
227	3	3=No/Unsure
17		Missing

MID14D		Subclass of MI?
N	Value	Description
1987	.S	Legitimately skipped
137	1	1=Type 1
57	2	2=Type 2
0	3	3=Type 3
0	4	4=Type 4a
2	5	5=Type 4b
0	6	6=Type 5
37	7	7=Unsure
18		Missing

MID15		aborted MI due to quick intervention
N	Value	Description
1029	Range	0 - 9 (median=0 mean=0.1 std=1.1)
1209		Missing

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MID10A_FUTIME		Futime from Randomization date for (MID10A)
N	Value	Description
2223	Range	361 - 5436 (median=3376 mean=3421.9 std=686.0)
15		Missing

MID10D_FUTIME		Futime from Randomization date for (MID10D)
N	Value	Description
2238	Range	-25 - 4206 (median=1810 mean=1871.5 std=1043.2)

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

ID		HCHS Participant ID
N	Value	Description
2238	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
1494	1	Occurrence 1
407	2	Occurrence 2
161	3	Occurrence 3
75	4	Occurrence 4
39	5	Occurrence 5
24	6	Occurrence 6
11	7	Occurrence 7
6	8	Occurrence 8
5	9	Occurrence 9
6	10	Occurrence 10
3	11	Occurrence 11
3	12	Occurrence 12
1	13	Occurrence 13
1	17	Occurrence 17
1	18	Occurrence 18
1	19	Occurrence 19

MID26		Comments
N	Value	Description
267	Present	Text suppressed
1971		Missing

MID21		Evidence of cardiac pain?
N	Value	Description
1344	0	0=No
696	1	1=Yes
162	9	9=NR
36		Missing

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MID22		Level of cardiac biomarkers
N	Value	Description
255	1	1= Abnormal (at least one value at 2 X the upper limit normal)
145	2	2= Equivocal (> normal, but < 2X the upper limit normal)
70	3	3= Incomplete
855	4	4= Normal
877	5	5= None recorded
36		Missing

MID23		Interpretations of ECGs
N	Value	Description
2	1	1= Evolution of Major Q-Wave
56	2	2= Evolution of ST-T Elevation with or without Q-Wave
2	3	3= New LBBB
70	4	4= Evolution of ST-Depression/T wave inversion alone
1	5	5= Evolution of Minor Q-Wave alone
16	6	6= Single ECG with Major Q-wave
3	7	7= Single ECG with LBBB, described as new
2052	8	8= Absent, Uncodable or Other ECG
36		Missing

MID24		Myocardial infarction classification
N	Value	Description
117	1	1=Definite
139	2	2=Probable
1932	3	3=No MI
14	4	4=Unclassifiable
36		Missing

MID24A		Type of MI?
N	Value	Description
1949	.S	Legitimately skipped
63	0	0=Transmural
151	1	1=Subendocardial
42	9	9=Unsure/unknown
33		Missing

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Section B: Myocardial Infarction Ascertainment

MID24B		Location of MI?
N	Value	Description
1949	.S	Legitimately skipped
29	1	1=Anterior
2	2	2=Posterior
32	3	3=Inferior
16	4	4=Lateral
4	5	5=Septal
31	6	6=More than one area
142	9	9=Unable to determine
33		Missing

MID24C		Was the MI procedure-related?
N	Value	Description
1949	.S	Legitimately skipped
3	1	1=Yes, cardiovascular procedure
5	2	2=Yes, non-cardiovascular procedure
248	3	3=No/Unsure
33		Missing

MID24D		Subclass of MI?
N	Value	Description
1949	.S	Legitimately skipped
124	1	1=Type 1
95	2	2=Type 2
1	3	3=Type 3
0	4	4=Type 4a
0	5	5=Type 4b
0	6	6=Type 5
35	7	7=Unsure
34		Missing

MID25		aborted MI due to quick intervention
N	Value	Description
1046	Range	0 - 9 (median=0 mean=0.2 std=1.4)
1192		Missing

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MID20A_FUTIME		Futime from Randomization date for (MID20A)
N	Value	Description
2204	Range	1802 - 5440 (median=3384 mean=3452.8 std=728.4)
34		Missing

MID20D_FUTIME		Futime from Randomization date for (MID20D)
N	Value	Description
2232	Range	-25 - 4206 (median=1815 mean=1875.2 std=1042.2)
6		Missing

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

ID		HCHS Participant ID
N	Value	Description
564	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
380	1	Occurrence 1
102	2	Occurrence 2
35	3	Occurrence 3
16	4	Occurrence 4
10	5	Occurrence 5
7	6	Occurrence 6
4	7	Occurrence 7
4	8	Occurrence 8
2	9	Occurrence 9
2	10	Occurrence 10
1	13	Occurrence 13
1	19	Occurrence 19

MID36		Comments
N	Value	Description
50	Present	Text suppressed
514		Missing

MID31		Evidence of cardiac pain?
N	Value	Description
108	0	0=No
176	1	1=Yes
19	9	9=NR
261		Missing

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MID32		Level of cardiac biomarkers
N	Value	Description
108	1	1= Abnormal (at least one value at 2 X the upper limit normal)
78	2	2= Equivocal (> normal, but < 2X the upper limit normal)
19	3	3= Incomplete
48	4	4= Normal
50	5	5= None recorded
261		Missing

MID33		Interpretations of ECGs
N	Value	Description
0	1	1= Evolution of Major Q-Wave
24	2	2= Evolution of ST-T Elevation with or without Q-Wave
0	3	3= New LBBB
18	4	4= Evolution of ST-Depression/T wave inversion alone
1	5	5= Evolution of Minor Q-Wave alone
3	6	6= Single ECG with Major Q-wave
0	7	7= Single ECG with LBBB, described as new
257	8	8= Absent, Uncodable or Other ECG
261		Missing

MID34		Myocardial infarction classification
N	Value	Description
42	1	1=Definite
81	2	2=Probable
169	3	3=No MI
11	4	4=Unclassifiable
261		Missing

MID34A		Type of MI?
N	Value	Description
185	.S	Legitimately skipped
27	0	0=Transmural
70	1	1=Subendocardial
26	9	9=Unsure/unknown
256		Missing

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Section B: Myocardial Infarction Ascertainment

MID34B		Location of MI?
N	Value	Description
185	.S	Legitimately skipped
7	1	1=Anterior
1	2	2=Posterior
16	3	3=Inferior
9	4	4=Lateral
2	5	5=Septal
12	6	6=More than one area
76	9	9=Unable to determine
256		Missing

MID34C		Was the MI procedure-related?
N	Value	Description
185	.S	Legitimately skipped
1	1	1=Yes, cardiovascular procedure
4	2	2=Yes, non-cardiovascular procedure
118	3	3=No/Unsure
256		Missing

MID34D		Subclass of MI?
N	Value	Description
185	.S	Legitimately skipped
58	1	1=Type 1
40	2	2=Type 2
0	3	3=Type 3
0	4	4=Type 4a
0	5	5=Type 4b
0	6	6=Type 5
25	7	7=Unsure
256		Missing

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

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MID30A_FUTIME		Futime from Randomization date for (MID30A)
N	Value	Description
303	Range	2383 - 5307 (median=3435 mean=3506.3 std=557.6)
261		Missing

MID30D_FUTIME		Futime from Randomization date for (MID30D)
N	Value	Description
483	Range	5 - 3949 (median=1253 mean=1423.3 std=991.1)
81		Missing

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

ID		HCHS Participant ID
N	Value	Description
1134	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
725	1	Occurrence 1
204	2	Occurrence 2
97	3	Occurrence 3
43	4	Occurrence 4
21	5	Occurrence 5
11	6	Occurrence 6
8	7	Occurrence 7
5	8	Occurrence 8
2	9	Occurrence 9
6	10	Occurrence 10
2	11	Occurrence 11
3	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

HTF1		Was the event
N	Value	Description
72	1	1= In hospital only
2	2	2= Emergency Dept. visit only (ED)
1057	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

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HTF2A		Time of arrival
N	Value	Description
1073	Range	0:02:00 - 23:59:00
61		Missing

HTF2AHR		Hour of Time of arrival
N	Value	Description
1073	Range	0 - 23 (median=13 mean=12.9 std=6.0)
61		Missing

HTF2AMN		Minute of Time of arrival
N	Value	Description
1073	Range	0 - 59 (median=29 mean=28.8 std=17.2)
61		Missing

HTF2ASC		Second of Time of arrival
N	Value	Description
459	Range	0 - 0 (median=0 mean=0.0 std=0.0)
675		Missing

HTF2A1		AM/PM of arrival
N	Value	Description
1062	.S	Legitimately skipped
26	1	1=AM
31	2	2=PM
15		Missing

HTF3A		Time of discharge
N	Value	Description
620	Range	0:08:00 - 23:59:00
514		Missing

HTF3AHR		Hour of Time of discharge
N	Value	Description
620	Range	0 - 23 (median=16 mean=15.5 std=4.2)
514		Missing

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<i>HTF3AMN</i>		<i>Minute of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
620	Range	0 - 59 (median=29.5 mean=28.39 std=19.11)
514		Missing

<i>HTF3ASC</i>		<i>Second of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
266	Range	0 - 0 (median=0 mean=0.0 std=0.0)
868		Missing

<i>HTF3A1</i>		<i>AM/PM of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1062	.S	Legitimately skipped
8	1	1=AM
25	2	2=PM
39		Missing

<i>HTF4</i>		<i>Admitting diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
983	Present	Text suppressed
151		Missing

<i>HTF5</i>		<i>Discharge diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1004	Present	Text suppressed
130		Missing

<i>HTF6</i>		<i>EMS transport</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
794	0	0=No/NR
340	1	1=Yes

<i>HTF7</i>		<i>Transferred from hosp</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1094	0	0=No/NR
40	1	1=Yes

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HTF8		Code status DNR
N	Value	Description
1086	0	0=No/NR
48	1	1=Yes

HTF9		Alive at discharge
N	Value	Description
56	0	0=No
1078	1	1=Yes

HTF10A		PND at event
N	Value	Description
191	0	0=No
98	1	1=Yes
845	9	9=NR

HTF10B		Orthopnea at event
N	Value	Description
203	0	0=No
232	1	1=Yes
699	9	9=NR

HTF10C		Shortness of breath at event
N	Value	Description
229	0	0=No
809	1	1=Yes
96	9	9=NR

HTF10D		Edema at event
N	Value	Description
555	0	0=No
342	1	1=Yes
237	9	9=NR

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HTF10E		Hypoxia at event
N	Value	Description
190	0	0=No
287	1	1=Yes
657	9	9=NR

HTF10F		Dyspnea (at rest) at event
N	Value	Description
304	0	0=No
348	1	1=Yes
482	9	9=NR

HTF10G		Dyspnea (on exertion) at event
N	Value	Description
173	0	0=No
716	1	1=Yes
245	9	9=NR

HTF11		Exacerbation of HF
N	Value	Description
737	0	0=No
352	1	1=Yes
45	9	9=NR

HTF12		Onset/progression of HF
N	Value	Description
596	0	0=No
402	1	1=Yes
136	9	9=NR

HTF13A		Diastolic heart failure
N	Value	Description
866	0	0=No/NR
268	1	1=Yes

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HTF13B		Systolic heart failure
N	Value	Description
874	0	0=No/NR
260	1	1=Yes

HTF13C		Right-sided heart failure
N	Value	Description
1114	0	0=No/NR
20	1	1=Yes

HTF13D		Ischemic cardiomyopathy
N	Value	Description
1015	0	0=No/NR
119	1	1=Yes

HTF13E		Idiopathic/dilated cardiomyopathy
N	Value	Description
1040	0	0=No/NR
94	1	1=Yes

HTF13F		Myocarditis
N	Value	Description
1130	0	0=No/NR
4	1	1=Yes

HTF13G		Peripartum cardiomyopathy
N	Value	Description
1134	0	0=No/NR
0	1	1=Yes

HTF13H		Other cardiomyopathy/HF
N	Value	Description
982	0	0=No/NR
152	1	1=Yes

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HTF13H1		Specify type cardiomyopathy
N	Value	Description
152	Present	Text suppressed
982	~	Legitimately skipped

HTF14		Cardiac imaging performed
N	Value	Description
739	0	0=No/NR
395	1	1=Yes

HTF14A		Quantitative EF
N	Value	Description
739	.S	Legitimately skipped
42	0	0=No
353	1	1=Yes

HTF14B		Lowest LV EF
N	Value	Description
352	Range	10 - 86 (median=45 mean=44.3 std=17.9)
781	.S	Legitimately skipped
1		Missing

HTF14C		Qualitative EF
N	Value	Description
1	D	D= Decreased moderately
0	M	M= Mildly reduced
23	N	N= Normal
1	O	O= None of the above
0	S	S= Severely reduced
16	U	U= Unsure-Not available
1092	~	Legitimately skipped
1		Missing

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HTF14D		Time since lowest EF
N	Value	Description
314	Range	0.1 - 84 (median=5 mean=10.0 std=13.6)
739	.S	Legitimately skipped
81		Missing

HTF14E		Type of Imaging from EF
N	Value	Description
739	.S	Legitimately skipped
290	1	1= ECHO
0	2	2= MUGA
23	3	3= Catheterization with ventriculography
0	4	4= CT
0	5	5= MRI
15	6	6= Other
67	7	7= Unknown

HTF15A		Diagnosis of HF prior
N	Value	Description
572	0	0=No/NR
562	1	1=Yes

HTF15B		Hospitalization for HF prior
N	Value	Description
572	.S	Legitimately skipped
419	0	0=No/NR
143	1	1=Yes

HTF15C		Treatment for HF prior
N	Value	Description
572	.S	Legitimately skipped
291	0	0=No/NR
271	1	1=Yes

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HTF15D		Valvular HD prior
N	Value	Description
1021	0	0=No/NR
113	1	1=Yes

HTF15E		RHD prior
N	Value	Description
1118	0	0=No/NR
16	1	1=Yes

HTF15F		Congenital HD prior
N	Value	Description
1128	0	0=No/NR
6	1	1=Yes

HTF15G		Coronary HD (ever) prior
N	Value	Description
694	0	0=No/NR
440	1	1=Yes

HTF15H		Coronary HD (within year) prior
N	Value	Description
1055	0	0=No/NR
79	1	1=Yes

HTF15I		Angina prior
N	Value	Description
903	0	0=No/NR
231	1	1=Yes

HTF15J		Myocardial infarction prior
N	Value	Description
936	0	0=No/NR
198	1	1=Yes

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<i>HTF15K</i>		<i>Atrial fib/atrial flutter prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
966	0	0=No/NR
168	1	1=Yes

<i>HTF15L</i>		<i>Heart block/bradycardia prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1103	0	0=No/NR
31	1	1=Yes

<i>HTF15M</i>		<i>Ventricular fib or tachycard prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1099	0	0=No/NR
35	1	1=Yes

<i>HTF15N</i>		<i>Hypertension prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
204	0	0=No/NR
930	1	1=Yes

<i>HTF15O</i>		<i>Diabetes prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
437	0	0=No/NR
697	1	1=Yes

<i>HTF15P</i>		<i>COPD prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
893	0	0=No/NR
241	1	1=Yes

<i>HTF15Q</i>		<i>Cor pulmonale prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1073	0	0=No/NR
61	1	1=Yes

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<i>HTF15R</i>		<i>Pulmonary hypertension prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1078	0	0=No/NR
56	1	1=Yes

<i>HTF15S</i>		<i>ESRD prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
993	0	0=No/NR
141	1	1=Yes

<i>HTF16A</i>		<i>CABG past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
980	0	0=No/NR
154	1	1=Yes

<i>HTF16B</i>		<i>PCI past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
970	0	0=No/NR
164	1	1=Yes

<i>HTF16C</i>		<i>Valve surgery past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1068	0	0=No/NR
66	1	1=Yes

<i>HTF16D</i>		<i>Pacemaker past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1058	0	0=No/NR
76	1	1=Yes

<i>HTF16E</i>		<i>AICD past</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1015	0	0=No/NR
119	1	1=Yes

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HTF16F		Ablation for arrhythmia past
N	Value	Description
1114	0	0=No/NR
20	1	1=Yes

HTF16G		Cardiac transplant past
N	Value	Description
1131	0	0=No/NR
3	1	1=Yes

HTF16H		VAD past
N	Value	Description
1132	0	0=No/NR
2	1	1=Yes

HTF17A		Myocardial Infarction active
N	Value	Description
1036	0	0=No/NR
98	1	1=Yes

HTF17B		Shock/Cardiogenic Shock active
N	Value	Description
1102	0	0=No/NR
32	1	1=Yes

HTF17C		Ventricular Fib active
N	Value	Description
1085	0	0=No/NR
49	1	1=Yes

HTF17D		Ventricular Tachycardia active
N	Value	Description
1103	0	0=No/NR
31	1	1=Yes

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HTF17E		Atrial Fib/Flutter active
N	Value	Description
1020	0	0=No/NR
114	1	1=Yes

HTF17F		COPD exacerbation active
N	Value	Description
993	0	0=No/NR
141	1	1=Yes

HTF17G		Cardiac Surgery active
N	Value	Description
1095	0	0=No/NR
39	1	1=Yes

HTF17H		Non-cardiac surgery active
N	Value	Description
1043	0	0=No/NR
91	1	1=Yes

HTF17I		Pulmonary Embolus active
N	Value	Description
1120	0	0=No/NR
14	1	1=Yes

HTF17J		Pneumonia active
N	Value	Description
968	0	0=No/NR
166	1	1=Yes

HTF17K		Renal failure/insufficiency active
N	Value	Description
1008	0	0=No/NR
126	1	1=Yes

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HTF18		Weight or BMI
N	Value	Description
753	Range	20 - 575 (median=88 mean=106.1 std=56.0)
381		Missing

HTF18A		Units Weight/BMI
N	Value	Description
147	1	1= Weight in lbs
588	2	2= Weight in Kg
14	3	3= BMI
385		Missing

HTF19A		JVD
N	Value	Description
656	0	0=No
119	1	1=Yes
359	9	9=NR

HTF19B		Crackles or rales
N	Value	Description
627	0	0=No
358	1	1=Yes
149	9	9=NR

HTF19C		Wheezing
N	Value	Description
699	0	0=No
266	1	1=Yes
169	9	9=NR

HTF19D		Rhonchi
N	Value	Description
639	0	0=No
200	1	1=Yes
295	9	9=NR

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HTF19E		S3 gallop
N	Value	Description
566	0	0=No
28	1	1=Yes
540	9	9=NR

HTF19F		Lower extremity edema-unilateral
N	Value	Description
998	0	0=No
12	1	1=Yes
124	9	9=NR

HTF19G		Lower extremity edema-bilateral
N	Value	Description
691	0	0=No
327	1	1=Yes
116	9	9=NR

HTF20		Chest X-ray performed
N	Value	Description
127	0	0=No/NR
1007	1	1=Yes

HTF20A		Pulmonary edema/CHF x-ray
N	Value	Description
127	.S	Legitimately skipped
745	0	0=No/NR
262	1	1=Yes

HTF20B		Cardiomegaly/Cardiothoracic ratio x-ray
N	Value	Description
127	.S	Legitimately skipped
575	0	0=No/NR
432	1	1=Yes

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HTF20C		Pulmonary vascular congestion x-ray
N	Value	Description
127	.S	Legitimately skipped
737	0	0=No/NR
270	1	1=Yes

HTF20D		Pleural effusion x-ray
N	Value	Description
127	.S	Legitimately skipped
711	0	0=No/NR
296	1	1=Yes

HTF21		CT scan or CT angiogram
N	Value	Description
826	0	0=No/NR
308	1	1=Yes

HTF21A		Pulmonary edema CT
N	Value	Description
826	.S	Legitimately skipped
262	0	0=No/NR
46	1	1=Yes

HTF21B		Cardiomegaly CT
N	Value	Description
826	.S	Legitimately skipped
223	0	0=No/NR
85	1	1=Yes

HTF21C		Pleural effusion CT
N	Value	Description
826	.S	Legitimately skipped
187	0	0=No/NR
121	1	1=Yes

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HTF21D		Enlarged vena cava CT
N	Value	Description
826	.S	Legitimately skipped
307	0	0=No/NR
1	1	1=Yes

HTF21E		Enlarged Pulmonary arteries CT
N	Value	Description
826	.S	Legitimately skipped
294	0	0=No/NR
14	1	1=Yes

HTF22		TTE performed
N	Value	Description
607	0	0=No/NR
527	1	1=Yes

HTF22B		Left Ventricular Ejection fraction TTE
N	Value	Description
497	Range	5 - 84 (median=53 mean=47.9 std=16.6)
607	.S	Legitimately skipped
30		Missing

HTF22C		LVH TTE
N	Value	Description
607	.S	Legitimately skipped
124	0	0=None
35	1	1=Present
128	2	2=Mild
48	3	3=Mod
10	4	4=Severe
175	9	9=NR
7		Missing

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<i>HTF22D</i>		<i>Impaired LV systolic function TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
607	.S	Legitimately skipped
205	0	0=None
36	1	1=Present
34	2	2=Mild
35	3	3=Mod
76	4	4=Severe
135	9	9=NR
6		Missing

<i>HTF22E</i>		<i>Impaired RV systolic function TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
607	.S	Legitimately skipped
264	0	0=None
15	1	1=Present
16	2	2=Mild
16	3	3=Mod
6	4	4=Severe
203	9	9=NR
7		Missing

<i>HTF22F</i>		<i>Pulmonary hypertension TTE</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
607	.S	Legitimately skipped
56	0	0=None
14	1	1=Present
32	2	2=Mild
38	3	3=Mod
20	4	4=Severe
360	9	9=NR
7		Missing

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HTF22G		Diastolic dysfunction TTE
N	Value	Description
607	.S	Legitimately skipped
43	0	0=None
138	1	1=Present
56	2	2=Mild
33	3	3=Mod
6	4	4=Severe
245	9	9=NR
6		Missing

HTF23		TEE performed
N	Value	Description
1096	0	0=No/NR
37	1	1=Yes
1		Missing

HTF23B1		LV ejection fraction TEE
N	Value	Description
18	Range	20 - 75 (median=50 mean=49.2 std=13.5)
1097	.S	Legitimately skipped
19		Missing

HTF23B2		RV ejection fraction TEE
N	Value	Description
1097	.S	Legitimately skipped
37		Missing

HTF24		Coronary angiography performed
N	Value	Description
992	0	0=No/NR
142	1	1=Yes

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HTF24B		LV Ejection fraction angio
N	Value	Description
69	Range	10 - 67 (median=40 mean=41.2 std=15.4)
992	.S	Legitimately skipped
73		Missing

HTF24C		70% obstruction artery
N	Value	Description
992	.S	Legitimately skipped
54	0	0=No/NR
88	1	1=Yes

HTF25		MUGA or RVG performed
N	Value	Description
1130	0	0=No/NR
4	1	1=Yes

HTF25A		LV ejection fraction MUGA
N	Value	Description
4	Range	10 - 68 (median=62 mean=50.5 std=27.3)
1130	.S	Legitimately skipped

HTF25B		RV ejection fraction MUGA
N	Value	Description
1130	.S	Legitimately skipped
4		Missing

HTF26		MRI performed
N	Value	Description
1132	0	0=No/NR
2	1	1=Yes

HTF26A		LV ejection fraction MRI
N	Value	Description
1	Range	30 - 30 (median=30 mean=30.0 std=)
1132	.S	Legitimately skipped
1		Missing

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HTF26B		RV ejection fraction MRI
N	Value	Description
1132	.S	Legitimately skipped
2		Missing

HTF27A		LV ejection fraction during visit
N	Value	Description
538	Range	0 - 87 (median=50 mean=47.4 std=17.0)
596		Missing

HTF27B		Stress test positive for ischemia visit
N	Value	Description
1112	0	0=No/NR
22	1	1=Yes

HTF27C		Regional wall abnormalities visit
N	Value	Description
944	0	0=No/NR
190	1	1=Yes

HTF27D		Dilated left ventricle visit
N	Value	Description
1012	0	0=No/NR
122	1	1=Yes

HTF27E		Dilated right ventricle visit
N	Value	Description
1065	0	0=No/NR
69	1	1=Yes

HTF27F		Impaired left systolic visit
N	Value	Description
942	0	0=No/NR
192	1	1=Yes

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<i>HTF27G</i>		<i>Left diastolic dysfunction visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
908	0	0=No/NR
226	1	1=Yes

<i>HTF27H</i>		<i>VSD visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1131	0	0=No/NR
3	1	1=Yes

<i>HTF27I</i>		<i>ASD visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1134	0	0=No/NR
0	1	1=Yes

<i>HTF27J</i>		<i>PDA visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1134	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>
1119	0	0=No/NR
15	1	1=Yes

<i>HTF27L</i>		<i>HOCM visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1133	0	0=No/NR
1	1	1=Yes

<i>HTF27M</i>		<i>Valvular HD visit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1097	0	0=No/NR
37	1	1=Yes

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HTF28A		Worst BNP
N	Value	Description
235	Range	5 - 33000 (median=309 mean=1407.0 std=3779.2)
899		Missing

HTF28B		Last BNP
N	Value	Description
169	Range	5 - 33000 (median=333 mean=1408.3 std=3927.1)
965		Missing

HTF28C		Upper Limit BNP
N	Value	Description
147	Range	23 - 126 (median=100 mean=99.4 std=9.5)
987		Missing

HTF29A		Worst ProBNP
N	Value	Description
290	Range	18 - 130000 (median=1922 mean=9283.5 std=17354.6)
844		Missing

HTF29B		Last BNP
N	Value	Description
223	Range	18 - 130000 (median=1958 mean=8617.4 std=16324.2)
911		Missing

HTF29C		Upper Limit ProBNP
N	Value	Description
223	Range	99 - 1800 (median=299 mean=426.5 std=418.6)
911		Missing

HTF30		Troponin value available
N	Value	Description
315	0	0=No
809	1	1=Yes
10		Missing

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HTF30A1		Less than worst Troponin
N	Value	Description
330	<	
3	>	
320	~	Legitimately skipped
481		Missing

HTF30A2		Worst Troponin
N	Value	Description
796	Range	0 - 11631 (median=0.03 mean=29.711 std=471.520)
320	.S	Legitimately skipped
18		Missing

HTF30B		Upper Limit Troponin
N	Value	Description
560	Range	0.01 - 100 (median=0.035 mean=0.8048 std=5.1675)
320	.S	Legitimately skipped
254		Missing

HTF30C		Type of Troponin
N	Value	Description
320	.S	Legitimately skipped
92	1	1= Troponin, type not specified
412	2	2= Troponin I
290	3	3= Troponin T
10	4	4= High Sensitivity Troponin (HS)
3	5	5= Unsure
7		Missing

HTF31		Worst sodium
N	Value	Description
1055	Range	117 - 161 (median=137 mean=136.4 std=4.4)
79		Missing

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HTF32		Worst serum creatinine
N	Value	Description
1060	Range	0.2 - 15.1 (median=1.16 mean=2.087 std=2.293)
74		Missing

HTF33		Worst BUN
N	Value	Description
1060	Range	6 - 194 (median=24 mean=33.0 std=25.6)
74		Missing

HTF34		Worst hemoglobin
N	Value	Description
1068	Range	4.4 - 18.1 (median=11.5 mean=11.34 std=2.38)
66		Missing

HTF35		Worst hematocrit
N	Value	Description
1046	Range	13.6 - 54.9 (median=35 mean=34.4 std=6.9)
88		Missing

HTF36A		Cardioversion or Defibrillation visit
N	Value	Description
1111	0	0=No/NR
23	1	1=Yes

HTF36B		Aortic balloon pump visit
N	Value	Description
1118	0	0=No/NR
16	1	1=Yes

HTF36C		PCI visit
N	Value	Description
1080	0	0=No/NR
54	1	1=Yes

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HTF36D		CPAP or BIPAP visit
N	Value	Description
993	0	0=No/NR
141	1	1=Yes

HTF36E		Mechanical Ventilation visit
N	Value	Description
1031	0	0=No/NR
103	1	1=Yes

HTF36F		Thoracentesis visit
N	Value	Description
1105	0	0=No/NR
29	1	1=Yes

HTF36G		VAD visit
N	Value	Description
1130	0	0=No/NR
4	1	1=Yes

HTF36H		Heart transplant visit
N	Value	Description
1134	0	0=No/NR
0	1	1=Yes

HTF36I		Hemodialysis or hemofiltration visit
N	Value	Description
1005	0	0=No/NR
129	1	1=Yes

HTF37A		Med list admission
N	Value	Description
133	0	0=No/NR
1001	1	1=Yes

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HTF37B		Med list discharge
N	Value	Description
332	0	0=No/NR
802	1	1=Yes

HTF38A		ACE inhibitors admission
N	Value	Description
133	.S	Legitimately skipped
611	0	0=No/NR
390	1	1=Yes

HTF38B		ACE inhibitors discharge
N	Value	Description
332	.S	Legitimately skipped
500	0	0=No/NR
302	1	1=Yes

HTF39A		Angiotensin admission
N	Value	Description
133	.S	Legitimately skipped
812	0	0=No/NR
188	1	1=Yes
1		Missing

HTF39B		Angiotensin discharge
N	Value	Description
332	.S	Legitimately skipped
675	0	0=No/NR
126	1	1=Yes
1		Missing

HTF40A		Beta blockers admission
N	Value	Description
133	.S	Legitimately skipped
416	0	0=No/NR
585	1	1=Yes

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HTF40B		Beta blockers discharge
N	Value	Description
332	.S	Legitimately skipped
327	0	0=No/NR
475	1	1=Yes

HTF41A		Digitalis admission
N	Value	Description
133	.S	Legitimately skipped
953	0	0=No/NR
48	1	1=Yes

HTF41B		Digitalis discharge
N	Value	Description
332	.S	Legitimately skipped
763	0	0=No/NR
39	1	1=Yes

HTF42A		Diuretics admission
N	Value	Description
133	.S	Legitimately skipped
533	0	0=No/NR
468	1	1=Yes

HTF42B		Diuretics discharge
N	Value	Description
332	.S	Legitimately skipped
436	0	0=No/NR
366	1	1=Yes

HTF43A		Aldosterone blocker admission
N	Value	Description
133	.S	Legitimately skipped
928	0	0=No/NR
73	1	1=Yes

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HTF43B		Aldosterone blocker discharge
N	Value	Description
332	.S	Legitimately skipped
742	0	0=No/NR
59	1	1=Yes
1		Missing

HTF44A		Lipid agents admission
N	Value	Description
133	.S	Legitimately skipped
401	0	0=No/NR
600	1	1=Yes

HTF44B		Lipid agents discharge
N	Value	Description
332	.S	Legitimately skipped
294	0	0=No/NR
508	1	1=Yes

HTF45A		Nitrates admission
N	Value	Description
133	.S	Legitimately skipped
795	0	0=No/NR
206	1	1=Yes

HTF45B		Nitrates discharge
N	Value	Description
332	.S	Legitimately skipped
689	0	0=No/NR
113	1	1=Yes

HTF46A		Hydralazine admission
N	Value	Description
133	.S	Legitimately skipped
874	0	0=No/NR
127	1	1=Yes

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HTF46B		Hydralazine discharge
N	Value	Description
332	.S	Legitimately skipped
714	0	0=No/NR
88	1	1=Yes

HTF47A		IV inotropes
N	Value	Description
1110	0	0=No/NR
23	1	1=Yes
1		Missing

HTF47B		IV diuretics
N	Value	Description
789	0	0=No/NR
344	1	1=Yes
1		Missing

HTF47C		Oral diuretics
N	Value	Description
1002	0	0=No/NR
131	1	1=Yes
1		Missing

HTF0A_FUTIME		Futime from Randomization date for (HTF0A)
N	Value	Description
1134	Range	17 - 5054 (median=2946.5 mean=3077.77 std=819.14)

HTF0D_FUTIME		Futime from Randomization date for (HTF0D)
N	Value	Description
1134	Range	10 - 4206 (median=1966.5 mean=1995.03 std=1035.49)

HTF2_FUTIME		Futime from Randomization date for (HTF2)
N	Value	Description
1125	Range	10 - 4206 (median=1968 mean=2000.5 std=1035.7)
9		Missing

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HTF2B_FUTIME		Futime from Randomization date for (HTF2B)
N	Value	Description
1132	Range	10 - 4207 (median=1968.5 mean=1997.13 std=1035.67)
2		Missing

HTF3_FUTIME		Futime from Randomization date for (HTF3)
N	Value	Description
1124	Range	17 - 4227 (median=1965.5 mean=1998.52 std=1032.37)
10		Missing

HTF22A_FUTIME		Futime from Randomization date for (HTF22A)
N	Value	Description
471	Range	11 - 4111 (median=2082 mean=2036.1 std=1058.2)
663		Missing

HTF23A_FUTIME		Futime from Randomization date for (HTF23A)
N	Value	Description
35	Range	407 - 3847 (median=2315 mean=2182.8 std=999.3)
1099		Missing

HTF24A_FUTIME		Futime from Randomization date for (HTF24A)
N	Value	Description
138	Range	22 - 3949 (median=1776 mean=1879.4 std=1083.4)
996		Missing

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Heart Failure Reviewer 1 Form: FORM CODE=HFD1, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
1123	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
716	1	Occurrence 1
204	2	Occurrence 2
96	3	Occurrence 3
43	4	Occurrence 4
20	5	Occurrence 5
11	6	Occurrence 6
8	7	Occurrence 7
5	8	Occurrence 8
2	9	Occurrence 9
6	10	Occurrence 10
2	11	Occurrence 11
3	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

HFD112		Reviewer comments
N	Value	Description
260	Present	Text suppressed
863		Missing

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HFD11		History of heart failure
N	Value	Description
550	0	0=No
506	1	1=Yes
60	9	9=NR
7		Missing

HFD12		ADHF diagnosed
N	Value	Description
762	0	0=No/NR
352	1	1=Yes
9		Missing

HFD13		Pulmonary edema/congestion
N	Value	Description
677	0	0=No
314	1	1=Yes
123	9	9=NR
9		Missing

HFD14		Dilated ventricle
N	Value	Description
385	0	0=No
25	1	1=Yes, history
146	2	2=Yes, current imaging
549	9	9=Unknown
18		Missing

HFD15		Poor LV function
N	Value	Description
371	0	0=No
88	1	1=Yes, history
231	2	2=Yes, current imaging
415	9	9=Unknown
18		Missing

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HFD16		Poor RV function
N	Value	Description
371	0	0=No
17	1	1=Yes, history
57	2	2=Yes, current imaging
660	9	9=Unknown
18		Missing

HFD17		Diastolic dysfunction
N	Value	Description
71	0	0=No
69	1	1=Yes, history
218	2	2=Yes, current imaging
747	9	9=Unknown
18		Missing

HFD18		quantitative EF
N	Value	Description
45	1	1=less than 20
67	2	2=20 to 29
62	3	3=30 to 39
93	4	4=40 to 49
354	5	5=greater than or equal to 50
493	6	6=Unknown
9		Missing

HFD19		Reviewer Classification
N	Value	Description
199	1	1=Definite
166	2	2=Probable
228	3	3=No, chronic stable HF
510	4	4=No, HF unlikely
13	5	5=Unclassifiable
7		Missing

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HFD19A		classify the severity
N	Value	Description
751	.S	Legitimately skipped
50	1	1=Mild
227	2	2=Moderate
86	3	3=Severe
2	4	4=Unknown
7		Missing

HFD19B		ADHF right-sided HF
N	Value	Description
751	.S	Legitimately skipped
296	0	0=No
14	1	1=Yes
55	9	9=Unknown
7		Missing

HFD110A1		a. Coronary disease
N	Value	Description
172	Range	0 - 1 (median=0 mean=0.5 std=0.5)
387	.S	Legitimately skipped
564		Missing

HFD110B1		b. Coronary disease
N	Value	Description
81	Range	0 - 1 (median=1 mean=0.6 std=0.5)
495	.S	Legitimately skipped
547		Missing

HFD110A2		a. Hypertension
N	Value	Description
172	Range	0 - 1 (median=1 mean=0.7 std=0.4)
387	.S	Legitimately skipped
564		Missing

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HFD110B2		<i>b.Hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
120	Range	0 - 1 (median=0 mean=0.3 std=0.5)
451	.S	Legitimately skipped
552		Missing

HFD110A3		<i>a.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)
387	.S	Legitimately skipped
564		Missing

HFD110B3		<i>b.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
27	Range	0 - 1 (median=1 mean=0.8 std=0.4)
550	.S	Legitimately skipped
546		Missing

HFD110A4		<i>a.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.1 std=0.2)
387	.S	Legitimately skipped
564		Missing

HFD110B4		<i>b.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	Range	0 - 1 (median=1 mean=0.6 std=0.5)
567	.S	Legitimately skipped
546		Missing

HFD110A5		<i>a.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)
387	.S	Legitimately skipped
564		Missing

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HFD110B5		<i>b.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
34	Range	0 - 1 (median=1 mean=0.7 std=0.5)
543	.S	Legitimately skipped
546		Missing

HFD110A6		<i>a.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.1 std=0.3)
387	.S	Legitimately skipped
564		Missing

HFD110B6		<i>b.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
21	Range	0 - 1 (median=1 mean=1.0 std=0.2)
556	.S	Legitimately skipped
546		Missing

HFD110A7		<i>a.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.1 std=0.4)
387	.S	Legitimately skipped
564		Missing

HFD110B7		<i>b.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	0 - 1 (median=1 mean=0.9 std=0.3)
552	.S	Legitimately skipped
546		Missing

HFD110A8		<i>a.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.4 std=0.5)
387	.S	Legitimately skipped
564		Missing

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HFD110B8		<i>b.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
62	Range	0 - 1 (median=1 mean=0.7 std=0.4)
514	.S	Legitimately skipped
547		Missing

HFD110A9		<i>a.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.1 std=0.3)
387	.S	Legitimately skipped
564		Missing

HFD110B9		<i>b.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
21	Range	0 - 1 (median=0 mean=0.4 std=0.5)
556	.S	Legitimately skipped
546		Missing

HFD110A10		<i>a.Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.0 std=0.2)
387	.S	Legitimately skipped
564		Missing

HFD110B10		<i>b.Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	0 - 1 (median=0.5 mean=0.50 std=0.58)
573	.S	Legitimately skipped
546		Missing

HFD110A11		<i>a.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)
387	.S	Legitimately skipped
564		Missing

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HFD110B11		<i>b.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
34	Range	0 - 1 (median=0.5 mean=0.50 std=0.51)
543	.S	Legitimately skipped
546		Missing

HFD110A12		<i>a.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.2 std=0.4)
387	.S	Legitimately skipped
564		Missing

HFD110B12		<i>b.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
26	Range	1 - 1 (median=1 mean=1.0 std=0.0)
550	.S	Legitimately skipped
547		Missing

HFD110A13		<i>a.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.0 std=0.1)
387	.S	Legitimately skipped
564		Missing

HFD110B13		<i>b.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	1 - 1 (median=1 mean=1.0 std=)
576	.S	Legitimately skipped
546		Missing

HFD110A14		<i>a.Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
172	Range	0 - 1 (median=0 mean=0.0 std=0.2)
387	.S	Legitimately skipped
564		Missing

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HFD110B14		<i>b. Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5	Range	0 - 1 (median=1 mean=0.6 std=0.5)
571	.S	Legitimately skipped
547		Missing

HFD111		<i>Asymp LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
773	0	0=No/NR
30	1	1=Yes
313	9	9=Unknown
7		Missing

HFD10A_FUTIME		<i>Futime from Randomization date for (HFD10A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1116	Range	1895 - 5136 (median=3584 mean=3626.2 std=643.5)
7		Missing

HFD10D_FUTIME		<i>Futime from Randomization date for (HFD10D)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1123	Range	10 - 4206 (median=1962 mean=1987.9 std=1035.4)

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Heart Failure Reviewer 2 Form: FORM CODE=HFD2, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
1123	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
716	1	Occurrence 1
204	2	Occurrence 2
96	3	Occurrence 3
43	4	Occurrence 4
20	5	Occurrence 5
11	6	Occurrence 6
8	7	Occurrence 7
5	8	Occurrence 8
2	9	Occurrence 9
6	10	Occurrence 10
2	11	Occurrence 11
3	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

HFD212		Reviewer comments
N	Value	Description
184	Present	Text suppressed
939		Missing

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HFD21		History of heart failure
N	Value	Description
575	0	0=No
473	1	1=Yes
69	9	9=NR
6		Missing

HFD22		ADHF diagnosed
N	Value	Description
772	0	0=No/NR
344	1	1=Yes
7		Missing

HFD23		Pulmonary edema/congestion
N	Value	Description
608	0	0=No
356	1	1=Yes
151	9	9=NR
8		Missing

HFD24		Dilated ventricle
N	Value	Description
374	0	0=No
32	1	1=Yes, history
157	2	2=Yes, current imaging
550	9	9=Unknown
10		Missing

HFD25		Poor LV function
N	Value	Description
362	0	0=No
97	1	1=Yes, history
246	2	2=Yes, current imaging
408	9	9=Unknown
10		Missing

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HFD26		Poor RV function
N	Value	Description
381	0	0=No
9	1	1=Yes, history
52	2	2=Yes, current imaging
671	9	9=Unknown
10		Missing

HFD27		Diastolic dysfunction
N	Value	Description
81	0	0=No
54	1	1=Yes, history
248	2	2=Yes, current imaging
730	9	9=Unknown
10		Missing

HFD28		quantitative EF
N	Value	Description
41	1	1=less than 20
76	2	2=20 to 29
68	3	3=30 to 39
94	4	4=40 to 49
355	5	5=greater than or equal to 50
483	6	6=Unknown
6		Missing

HFD29		Reviewer Classification
N	Value	Description
230	1	1=Definite
128	2	2=Probable
182	3	3=No, chronic stable HF
560	4	4=No, HF unlikely
17	5	5=Unclassifiable
6		Missing

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HFD29A		classify the severity
N	Value	Description
759	.S	Legitimately skipped
65	1	1=Mild
222	2	2=Moderate
67	3	3=Severe
4	4	4=Unknown
6		Missing

HFD29B		ADHF right-sided HF
N	Value	Description
759	.S	Legitimately skipped
310	0	0=No
10	1	1=Yes
38	9	9=Unknown
6		Missing

HFD210A1		a. Coronary disease
N	Value	Description
130	Range	0 - 1 (median=1 mean=0.6 std=0.5)
407	.S	Legitimately skipped
586		Missing

HFD210B1		b. Coronary disease
N	Value	Description
55	Range	0 - 1 (median=1 mean=0.6 std=0.5)
504	.S	Legitimately skipped
564		Missing

HFD210A2		a. Hypertension
N	Value	Description
150	Range	0 - 1 (median=1 mean=0.9 std=0.3)
407	.S	Legitimately skipped
566		Missing

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HFD210B2		b.Hypertension
N	Value	Description
92	Range	0 - 1 (median=0 mean=0.3 std=0.5)
443	.S	Legitimately skipped
588		Missing

HFD210A3		a.Atrial fibrillation
N	Value	Description
110	Range	0 - 1 (median=0 mean=0.3 std=0.5)
407	.S	Legitimately skipped
606		Missing

HFD210B3		b.Atrial fibrillation
N	Value	Description
24	Range	0 - 1 (median=1 mean=0.5 std=0.5)
544	.S	Legitimately skipped
555		Missing

HFD210A4		a.Other arrhythmia
N	Value	Description
95	Range	0 - 1 (median=0 mean=0.0 std=0.2)
407	.S	Legitimately skipped
621		Missing

HFD210B4		b.Other arrhythmia
N	Value	Description
4	Range	0 - 1 (median=0.5 mean=0.50 std=0.58)
573	.S	Legitimately skipped
546		Missing

HFD210A5		a.Valvular disease
N	Value	Description
104	Range	0 - 1 (median=0 mean=0.3 std=0.5)
407	.S	Legitimately skipped
612		Missing

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HFD210B5		b.Valvular disease
N	Value	Description
30	Range	0 - 1 (median=1 mean=0.5 std=0.5)
543	.S	Legitimately skipped
550		Missing

HFD210A6		a.Fluid or volume
N	Value	Description
98	Range	0 - 1 (median=0 mean=0.1 std=0.3)
407	.S	Legitimately skipped
618		Missing

HFD210B6		b.Fluid or volume
N	Value	Description
10	Range	0 - 1 (median=1 mean=0.9 std=0.3)
567	.S	Legitimately skipped
546		Missing

HFD210A7		a.Medication
N	Value	Description
101	Range	0 - 1 (median=0 mean=0.2 std=0.4)
407	.S	Legitimately skipped
615		Missing

HFD210B7		b.Medication
N	Value	Description
15	Range	1 - 1 (median=1 mean=1.0 std=0.0)
560	.S	Legitimately skipped
548		Missing

HFD210A8		a.Renal Insufficiency
N	Value	Description
132	Range	0 - 1 (median=1 mean=0.5 std=0.5)
407	.S	Legitimately skipped
584		Missing

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HFD210B8		<i>b.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
53	Range	0 - 1 (median=0 mean=0.5 std=0.5)
505	.S	Legitimately skipped
565		Missing

HFD210A9		<i>a.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
100	Range	0 - 1 (median=0 mean=0.2 std=0.4)
407	.S	Legitimately skipped
616		Missing

HFD210B9		<i>b.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
12	Range	0 - 0 (median=0 mean=0.0 std=0.0)
561	.S	Legitimately skipped
550		Missing

HFD210A10		<i>a.Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
97	Range	0 - 1 (median=0 mean=0.0 std=0.2)
407	.S	Legitimately skipped
619		Missing

HFD210B10		<i>b.Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2	Range	0 - 1 (median=0.5 mean=0.50 std=0.71)
574	.S	Legitimately skipped
547		Missing

HFD210A11		<i>a.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
106	Range	0 - 1 (median=0 mean=0.3 std=0.5)
407	.S	Legitimately skipped
610		Missing

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HFD210B11		<i>b.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
30	Range	0 - 1 (median=0 mean=0.3 std=0.5)
540	.S	Legitimately skipped
553		Missing

HFD210A12		<i>a.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
106	Range	0 - 1 (median=0 mean=0.2 std=0.4)
407	.S	Legitimately skipped
610		Missing

HFD210B12		<i>b.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16	Range	0 - 1 (median=1 mean=0.7 std=0.5)
555	.S	Legitimately skipped
552		Missing

HFD210A13		<i>a.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
96	Range	0 - 1 (median=0 mean=0.0 std=0.1)
407	.S	Legitimately skipped
620		Missing

HFD210B13		<i>b.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	0 - 0 (median=0 mean=0.0 std=)
575	.S	Legitimately skipped
547		Missing

HFD210A14		<i>a.Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
96	Range	0 - 1 (median=0 mean=0.1 std=0.2)
407	.S	Legitimately skipped
620		Missing

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HFD210B14		<i>b. Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	0 - 1 (median=1 mean=0.8 std=0.5)
572	.S	Legitimately skipped
547		Missing

HFD211		<i>Asymp LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
826	0	0=No/NR
21	1	1=Yes
270	9	9=Unknown
6		Missing

HFD20A_FUTIME		<i>Futime from Randomization date for (HFD20A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1117	Range	1886 - 5333 (median=3539 mean=3624.7 std=661.8)
6		Missing

HFD20D_FUTIME		<i>Futime from Randomization date for (HFD20D)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1123	Range	10 - 4206 (median=1962 mean=1987.9 std=1035.4)

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Heart Failure Adjudication Reviewer Form: FORM CODE=HFD3, VERSION=A,B

ID		HCHS Participant ID
N	Value	Description
473	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
278	1	Occurrence 1
91	2	Occurrence 2
49	3	Occurrence 3
26	4	Occurrence 4
8	5	Occurrence 5
6	6	Occurrence 6
5	7	Occurrence 7
2	8	Occurrence 8
4	10	Occurrence 10
1	11	Occurrence 11
1	12	Occurrence 12
1	18	Occurrence 18
1	19	Occurrence 19

HFD312		Reviewer comments
N	Value	Description
32	Present	Text suppressed
441		Missing

HFD31		History of heart failure
N	Value	Description
146	0	0=No
259	1	1=Yes
12	9	9=NR
56		Missing

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HFD32		ADHF diagnosed
N	Value	Description
226	0	0=No/NR
191	1	1=Yes
56		Missing

HFD33		Pulmonary edema/congestion
N	Value	Description
176	0	0=No
179	1	1=Yes
62	9	9=NR
56		Missing

HFD34		Dilated ventricle
N	Value	Description
137	0	0=No
11	1	1=Yes, history
71	2	2=Yes, current imaging
198	9	9=Unknown
56		Missing

HFD35		Poor LV function
N	Value	Description
123	0	0=No
48	1	1=Yes, history
130	2	2=Yes, current imaging
116	9	9=Unknown
56		Missing

HFD36		Poor RV function
N	Value	Description
147	0	0=No
3	1	1=Yes, history
33	2	2=Yes, current imaging
234	9	9=Unknown
56		Missing

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<i>HFD37</i>		<i>Diastolic dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17	0	0=No
40	1	1=Yes, history
99	2	2=Yes, current imaging
261	9	9=Unknown
56		Missing

<i>HFD38</i>		<i>quantitative EF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17	1	1=less than 20
33	2	2=20 to 29
39	3	3=30 to 39
55	4	4=40 to 49
121	5	5=greater than or equal to 50
152	6	6=Unknown
56		Missing

<i>HFD39</i>		<i>Reviewer Classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
95	1	1=Definite
119	2	2=Probable
108	3	3=No, chronic stable HF
80	4	4=No, HF unlikely
15	5	5=Unclassifiable
56		Missing

<i>HFD39A</i>		<i>classify the severity</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
206	.S	Legitimately skipped
18	1	1=Mild
147	2	2=Moderate
47	3	3=Severe
2	4	4=Unknown
53		Missing

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HFD39B		ADHF right-sided HF
N	Value	Description
206	.S	Legitimately skipped
189	0	0=No
5	1	1=Yes
20	9	9=Unknown
53		Missing

HFD310A1		a. Coronary disease
N	Value	Description
13	Range	0 - 1 (median=0 mean=0.5 std=0.5)
27	.S	Legitimately skipped
433		Missing

HFD310B1		b. Coronary disease
N	Value	Description
6	Range	0 - 1 (median=1 mean=0.7 std=0.5)
46	.S	Legitimately skipped
421		Missing

HFD310A2		a. Hypertension
N	Value	Description
13	Range	0 - 1 (median=1 mean=0.7 std=0.5)
27	.S	Legitimately skipped
433		Missing

HFD310B2		b. Hypertension
N	Value	Description
9	Range	0 - 1 (median=0 mean=0.1 std=0.3)
43	.S	Legitimately skipped
421		Missing

HFD310A3		a. Atrial fibrillation
N	Value	Description
13	Range	0 - 1 (median=0 mean=0.2 std=0.4)
27	.S	Legitimately skipped
433		Missing

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HFD310B3		<i>b.Atrial fibrillation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2	Range	1 - 1 (median=1 mean=1.0 std=0.0)
50	.S	Legitimately skipped
421		Missing

HFD310A4		<i>a.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 0 (median=0 mean=0.0 std=0.0)
27	.S	Legitimately skipped
433		Missing

HFD310B4		<i>b.Other arrhythmia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
421		Missing

HFD310A5		<i>a.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 1 (median=0 mean=0.5 std=0.5)
27	.S	Legitimately skipped
433		Missing

HFD310B5		<i>b.Valvular disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5	Range	0 - 1 (median=1 mean=0.6 std=0.5)
46	.S	Legitimately skipped
422		Missing

HFD310A6		<i>a.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 1 (median=0 mean=0.1 std=0.3)
27	.S	Legitimately skipped
433		Missing

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HFD310B6		<i>b.Fluid or volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
51	.S	Legitimately skipped
422		Missing

HFD310A7		<i>a.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 0 (median=0 mean=0.0 std=0.0)
27	.S	Legitimately skipped
433		Missing

HFD310B7		<i>b.Medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
421		Missing

HFD310A8		<i>a.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 1 (median=0 mean=0.1 std=0.3)
27	.S	Legitimately skipped
433		Missing

HFD310B8		<i>b.Renal Insufficiency</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	0 - 0 (median=0 mean=0.0 std=)
51	.S	Legitimately skipped
421		Missing

HFD310A9		<i>a.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 1 (median=0 mean=0.2 std=0.4)
27	.S	Legitimately skipped
433		Missing

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HFD310B9		<i>b.Cardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2	Range	0 - 1 (median=0.5 mean=0.50 std=0.71)
50	.S	Legitimately skipped
421		Missing

HFD310A10		<i>a.Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 1 (median=0 mean=0.1 std=0.3)
27	.S	Legitimately skipped
433		Missing

HFD310B10		<i>b.Noncardiovascular</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
51	.S	Legitimately skipped
422		Missing

HFD310A11		<i>a.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 1 (median=0 mean=0.2 std=0.4)
27	.S	Legitimately skipped
433		Missing

HFD310B11		<i>b.Pulmonary disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2	Range	0 - 1 (median=0.5 mean=0.50 std=0.71)
50	.S	Legitimately skipped
421		Missing

HFD310A12		<i>a.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 1 (median=0 mean=0.2 std=0.4)
27	.S	Legitimately skipped
433		Missing

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HFD310B12		<i>b.Pulmonary Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
50	.S	Legitimately skipped
423		Missing

HFD310A13		<i>a.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 0 (median=0 mean=0.0 std=0.0)
27	.S	Legitimately skipped
433		Missing

HFD310B13		<i>b.Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
421		Missing

HFD310A14		<i>a.Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 0 (median=0 mean=0.0 std=0.0)
27	.S	Legitimately skipped
433		Missing

HFD310B14		<i>b.Toxins</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
52	.S	Legitimately skipped
421		Missing

HFD311		<i>Asymp LV dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
334	0	0=No/NR
11	1	1=Yes
72	9	9=Unknown
56		Missing

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HFD30A_FUTIME		Futime from Randomization date for (HFD30A)
N	Value	Description
417	Range	2002 - 5354 (median=3830 mean=3931.2 std=549.7)
56		Missing

HFD30D_FUTIME		Futime from Randomization date for (HFD30D)
N	Value	Description
473	Range	10 - 4130 (median=1654 mean=1786.4 std=1052.9)

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Pulmonary Abstraction Form: FORM CODE=PUL, VERSION=B

ID		HCHS Participant ID
N	Value	Description
2372	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
1605	1	Occurrence 1
407	2	Occurrence 2
153	3	Occurrence 3
69	4	Occurrence 4
44	5	Occurrence 5
24	6	Occurrence 6
13	7	Occurrence 7
11	8	Occurrence 8
5	9	Occurrence 9
7	10	Occurrence 10
6	11	Occurrence 11
2	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

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<i>PUL1</i>		<i>Event type</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
211	1	1= In hospital only
572	2	2= Emergency Dept. visit only (ED)
1566	3	3= Both ED and in hospital
2	4	4= Observation care only
16	5	5= Both ED and observation care
5		Missing

<i>PUL2</i>		<i>Hospital stay less than 24 hrs</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
575	.S	Legitimately skipped
1727	0	0=No/NR
68	1	1=Yes
2		Missing

<i>PUL3B</i>		<i>Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2217	Range	0:00:00 - 23:58:00
155		Missing

<i>PUL3BHR</i>		<i>Hour of Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2217	Range	0 - 23 (median=13 mean=12.7 std=6.0)
155		Missing

<i>PUL3BMN</i>		<i>Minute of Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2217	Range	0 - 59 (median=29 mean=28.8 std=17.3)
155		Missing

<i>PUL3BSC</i>		<i>Second of Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1267	Range	0 - 0 (median=0 mean=0.0 std=0.0)
1105		Missing

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<i>PUL3B1</i>		<i>AM/PM of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1883	.S	Legitimately skipped
196	1	1=AM
247	2	2=PM
46		Missing

<i>PUL4B</i>		<i>Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1592	Range	0:00:00 - 23:59:00
780		Missing

<i>PUL4BHR</i>		<i>Hour of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1592	Range	0 - 23 (median=15 mean=14.7 std=5.3)
780		Missing

<i>PUL4BMN</i>		<i>Minute of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1592	Range	0 - 59 (median=29 mean=27.4 std=18.5)
780		Missing

<i>PUL4BSC</i>		<i>Second of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
933	Range	0 - 0 (median=0 mean=0.0 std=0.0)
1439		Missing

<i>PUL4B1</i>		<i>AM/PM of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1883	.S	Legitimately skipped
80	1	1=AM
260	2	2=PM
149		Missing

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<i>PUL5</i>		<i>Primary admitting diagnosis code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1949	Present	Text suppressed
423		Missing

<i>PUL6</i>		<i>Primary discharge diagnosis code</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2127	Present	Text suppressed
245		Missing

<i>PUL7</i>		<i>Transported in EMS unit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1766	0	0=No/NR
606	1	1=Yes

<i>PUL8</i>		<i>Transferred from another hospital</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2322	0	0=No/NR
50	1	1=Yes

<i>PUL9</i>		<i>Code status no-code or DNR</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2281	0	0=No/NR
91	1	1=Yes

<i>PUL10</i>		<i>Patient alive at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
122	0	0=No/NR
2248	1	1=Yes
2		Missing

<i>PUL11A</i>		<i>Cough at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
560	0	0=No
1232	1	1=Yes
580	9	9=Not Recorded

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<i>PUL11B</i>		<i>Sputum production at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
739	0	0=No
647	1	1=Yes
986	9	9=Not Recorded

<i>PUL11C</i>		<i>Sputum purulence at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
821	0	0=No
351	1	1=Yes
1200	9	9=Not Recorded

<i>PUL11D</i>		<i>Wheezing at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
828	0	0=No
746	1	1=Yes
798	9	9=Not Recorded

<i>PUL11E</i>		<i>Chest pain at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1018	0	0=No
891	1	1=Yes
462	9	9=Not Recorded
1		Missing

<i>PUL11F</i>		<i>Leg edema at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1218	0	0=No
289	1	1=Yes
865	9	9=Not Recorded

<i>PUL11G</i>		<i>Rescue bronchodilator at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
358	0	0=No
928	1	1=Yes
1086	9	9=Not Recorded

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<i>PUL11H</i>		<i>Dyspnea at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
496	0	0=No
1589	1	1=Yes
287	9	9=Not Recorded

<i>PUL11I</i>		<i>Dyspnea (at rest) at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
563	0	0=No
732	1	1=Yes
1077	9	9=Not Recorded

<i>PUL11J</i>		<i>Dyspnea (walking/on exertion) at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
394	0	0=No
1176	1	1=Yes
802	9	9=Not Recorded

<i>PUL11K</i>		<i>Woken up at night at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
418	0	0=No
92	1	1=Yes
1862	9	9=Not Recorded

<i>PUL11L</i>		<i>Fever at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1608	0	0=No
461	1	1=Yes
303	9	9=Not Recorded

<i>PUL11M</i>		<i>AMS at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2008	0	0=No
214	1	1=Yes
150	9	9=Not Recorded

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<i>PUL12</i>		<i>Reason COPD/chronic bronchitis/emphysema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2006	0	0=No/NR
366	1	1=Yes

<i>PUL13</i>		<i>Reason exacerbation of asthma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1825	0	0=No/NR
547	1	1=Yes

<i>PUL14</i>		<i>New onset of exacerbation prior</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1470	0	0=No/NR
902	1	1=Yes

<i>PUL15A</i>		<i>Asthma history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1072	0	0=No/NR
1300	1	1=Yes

<i>PUL15B</i>		<i>Chronic bronchitis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2302	0	0=No/NR
70	1	1=Yes

<i>PUL15C</i>		<i>Emphysema history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2277	0	0=No/NR
95	1	1=Yes

<i>PUL15D</i>		<i>COPD history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1692	0	0=No/NR
680	1	1=Yes

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<i>PUL15E</i>		<i>Pulmonary fibrosis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2336	0	0=No/NR
36	1	1=Yes

<i>PUL15F</i>		<i>Sarcoidosis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2370	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>
2327	0	0=No/NR
45	1	1=Yes

<i>PUL15H</i>		<i>Lung resection or lobectomy history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2361	0	0=No/NR
11	1	1=Yes

<i>PUL15I</i>		<i>Home oxygen history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2265	0	0=No/NR
107	1	1=Yes

<i>PUL15J</i>		<i>Pulmonary embolus history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2344	0	0=No/NR
28	1	1=Yes

<i>PUL15K</i>		<i>Pulmonary hypertension history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2309	0	0=No/NR
63	1	1=Yes

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<i>PUL15L</i>		<i>Cor pulmonale history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2196	0	0=No/NR
176	1	1=Yes

<i>PUL15M</i>		<i>OSA history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2075	0	0=No/NR
297	1	1=Yes

<i>PUL15N</i>		<i>Coronary artery disease history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1889	0	0=No/NR
483	1	1=Yes

<i>PUL15O</i>		<i>Heart failure history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2015	0	0=No/NR
357	1	1=Yes

<i>PUL15P</i>		<i>Atrial fibrillation/atrial flutter history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2248	0	0=No/NR
124	1	1=Yes

<i>PUL15Q</i>		<i>Diabetes history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1320	0	0=No/NR
1052	1	1=Yes

<i>PUL15R</i>		<i>Pulmonary Tuberculosis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2327	0	0=No/NR
45	1	1=Yes

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<i>PUL15S</i>		<i>Bronchiectasis history</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2340	0	0=No/NR
32	1	1=Yes

<i>PUL16A</i>		<i>Pre-bronchodilator FEV1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15	Range	19 - 99 (median=70 mean=62.7 std=26.7)
2357		Missing

<i>PUL16B</i>		<i>Post-bronchodilator FEV1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16	Range	18 - 150 (median=77 mean=76.2 std=37.0)
2356		Missing

<i>PUL17</i>		<i>FEV1/FVC ratio</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	53 - 89 (median=74.5 mean=73.33 std=10.58)
2354		Missing

<i>PUL17A</i>		<i>FEV1/FVC ratio Units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
0	1	1=proportion or decimal
19	2	2=percent (If percent, then assure not percent predicted for the ratio)
2353		Missing

<i>PUL18A</i>		<i>Active Upper Respiratory Infection</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1991	0	0=No/NR
381	1	1=Yes

<i>PUL18B</i>		<i>Active Pneumonia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2005	0	0=No/NR
367	1	1=Yes

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<i>PUL18C</i>		<i>Active Pulmonary embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2350	0	0=No/NR
22	1	1=Yes

<i>PUL18D</i>		<i>Active Myocardial infarction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2288	0	0=No/NR
84	1	1=Yes

<i>PUL18E</i>		<i>Active Heart failure exacerbation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2153	0	0=No/NR
219	1	1=Yes

<i>PUL18F</i>		<i>Active Atrial fibrillation/flutter</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2274	0	0=No/NR
98	1	1=Yes

<i>PUL18G</i>		<i>Active SVT/MAT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2350	0	0=No/NR
22	1	1=Yes

<i>PUL18H</i>		<i>Active CABG/Valvular Surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2337	0	0=No/NR
35	1	1=Yes

<i>PUL18I</i>		<i>Active Non-cardiac surgery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2095	0	0=No/NR
277	1	1=Yes

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<i>PUL19A</i>		<i>Heart rate arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2150	Range	0 - 199 (median=88.5 mean=90.41 std=19.82)
222		Missing

<i>PUL19B</i>		<i>Respiration rate arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2105	Range	0 - 51 (median=19 mean=19.8 std=3.9)
267		Missing

<i>PUL19C</i>		<i>Oxy sat arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1904	Range	6 - 115 (median=97 mean=96.1 std=4.8)
468		Missing

<i>PUL19C1</i>		<i>Oxy Sats on room air arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
143	0	0=No
920	1	1=Yes
953	9	9=NR
356		Missing

<i>PUL19C2</i>		<i>Level oxy arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
113	Range	2 - 180 (median=2 mean=21.0 std=38.3)
2153	.S	Legitimately skipped
106		Missing

<i>PUL19C3</i>		<i>Level oxy Units arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2153	.S	Legitimately skipped
91	1	1= Liters
22	2	2=Percent
106		Missing

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<i>PUL19D</i>		<i>Weight arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1334	Range	36.36 - 575 (median=85.71 mean=101.087 std=49.237)
1038		Missing

<i>PUL19D1</i>		<i>Weight units arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
224	1	1= Lbs
1099	2	2=Kg
1049		Missing

<i>PUL20A</i>		<i>Use of accessory muscles at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
587	0	0=No
82	1	1=Yes
1703	9	9=NR

<i>PUL20B</i>		<i>Cyanosis at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1183	0	0=No
17	1	1=Yes
1172	9	9=NR

<i>PUL20C</i>		<i>Clubbing at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
890	0	0=No
8	1	1=Yes
1474	9	9=NR

<i>PUL20D</i>		<i>JVD at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1283	0	0=No
69	1	1=Yes
1020	9	9=NR

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<i>PUL20E</i>		<i>Crackles/rales at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1375	0	0=No
449	1	1=Yes
548	9	9=NR

<i>PUL20F</i>		<i>Wheezing or rhonchi at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1107	0	0=No
1029	1	1=Yes
236	9	9=NR

<i>PUL20G</i>		<i>Dec unilateral breath at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
925	0	0=No
33	1	1=Yes
1414	9	9=NR

<i>PUL20H</i>		<i>Dec bilateral breath at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
456	0	0=No
534	1	1=Yes
1382	9	9=NR

<i>PUL20I</i>		<i>Prolonged expiratory time at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
214	0	0=No
79	1	1=Yes
2079	9	9=NR

<i>PUL20J</i>		<i>Egophony at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
170	0	0=No
10	1	1=Yes
2192	9	9=NR

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<i>PUL20K</i>		<i>Lower extremity edema at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1554	0	0=No
357	1	1=Yes
461	9	9=NR

<i>PUL21</i>		<i>Chest X-ray performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	0	0=No/NR
1842	1	1=Yes

<i>PUL22A</i>		<i>Hyperinflation on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1766	0	0=No/NR
76	1	1=Yes

<i>PUL22B</i>		<i>Flattened diaphragms on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1818	0	0=No/NR
24	1	1=Yes

<i>PUL22C</i>		<i>Consolidation or infiltrate on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1388	0	0=No/NR
454	1	1=Yes

<i>PUL22D</i>		<i>Scarring on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1788	0	0=No/NR
54	1	1=Yes

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<i>PUL22E</i>		<i>Nodules on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1838	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>
530	.S	Legitimately skipped
1838	0	0=No/NR
4	1	1=Yes

<i>PUL22G</i>		<i>Pulmonary edema on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1475	0	0=No/NR
367	1	1=Yes

<i>PUL22H</i>		<i>Bilateral pleural effusion on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1681	0	0=No/NR
161	1	1=Yes

<i>PUL22I</i>		<i>Unilateral pleural effusion on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1716	0	0=No/NR
126	1	1=Yes

<i>PUL22J</i>		<i>Emphysema on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1808	0	0=No/NR
34	1	1=Yes

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<i>PUL22K</i>		<i>Cardiomegaly on X-ray</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
530	.S	Legitimately skipped
1453	0	0=No/NR
389	1	1=Yes

<i>PUL23</i>		<i>Chest/lung CT scan or CTA performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1819	0	0=No/NR
552	1	1=Yes
1		Missing

<i>PUL24A</i>		<i>Emphysema on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
475	0	0=No/NR
77	1	1=Yes

<i>PUL24B</i>		<i>Nodules on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
532	0	0=No/NR
20	1	1=Yes

<i>PUL24C</i>		<i>Masses on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
540	0	0=No/NR
12	1	1=Yes

<i>PUL24D</i>		<i>Lymphadenopathy on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
488	0	0=No/NR
64	1	1=Yes

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<i>PUL24E</i>		<i>Ground glass changes on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
462	0	0=No/NR
90	1	1=Yes

<i>PUL24F</i>		<i>Pneumonia on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
489	0	0=No/NR
63	1	1=Yes

<i>PUL24G</i>		<i>Fibrosis or honeycombing on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
529	0	0=No/NR
23	1	1=Yes

<i>PUL24H</i>		<i>Filling defect-vascular (PE) on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
546	0	0=No/NR
6	1	1=Yes

<i>PUL24I</i>		<i>Filling defect-mucus plug on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
541	0	0=No/NR
11	1	1=Yes

<i>PUL24J</i>		<i>Cysts or blebs on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
542	0	0=No/NR
10	1	1=Yes

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<i>PUL24K</i>		<i>Atelectasis on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
372	0	0=No/NR
180	1	1=Yes

<i>PUL24L</i>		<i>Calcifications on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
477	0	0=No/NR
75	1	1=Yes

<i>PUL24M</i>		<i>Pulmonary embolus on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
537	0	0=No/NR
15	1	1=Yes

<i>PUL24N</i>		<i>Enlarged pulmonary artery on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
528	0	0=No/NR
24	1	1=Yes

<i>PUL24O</i>		<i>Bronchiectasis on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
513	0	0=No/NR
39	1	1=Yes

<i>PUL24P</i>		<i>Pulmonary edema on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
488	0	0=No/NR
64	1	1=Yes

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<i>PUL24Q</i>		<i>Cardiomegaly on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
472	0	0=No/NR
80	1	1=Yes

<i>PUL24R</i>		<i>Bilateral pleural effusion on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
458	0	0=No/NR
94	1	1=Yes

<i>PUL24S</i>		<i>Unilateral pleural effusion on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
506	0	0=No/NR
46	1	1=Yes

<i>PUL24T</i>		<i>Airway wall thickening on CT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1820	.S	Legitimately skipped
510	0	0=No/NR
42	1	1=Yes

<i>PUL25</i>		<i>Spirometry performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2349	0	0=No/NR
23	1	1=Yes

<i>PUL25A1</i>		<i>FEV1 spirometry</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
21	Range	0.66 - 3.07 (median=1.73 mean=1.810 std=0.675)
2349	.S	Legitimately skipped
2		Missing

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PUL25A2		FEV1 Percent Predicted spirometry
N	Value	Description
21	Range	19 - 118 (median=71 mean=69.3 std=23.5)
2349	.S	Legitimately skipped
2		Missing

PUL25B1		FVC spirometry
N	Value	Description
20	Range	0.66 - 4.91 (median=2.445 mean=2.5915 std=1.0821)
2349	.S	Legitimately skipped
3		Missing

PUL25B2		FVC Percent Predicted spirometry
N	Value	Description
20	Range	15 - 121 (median=75 mean=76.3 std=26.1)
2349	.S	Legitimately skipped
3		Missing

PUL25C1		FEV1/FVC ratio spirometry
N	Value	Description
20	Range	0.93 - 88 (median=71.5 mean=69.27 std=18.41)
2349	.S	Legitimately skipped
3		Missing

PUL25C2		Units for FEV1/FVC ratio spirometry
N	Value	Description
2349	.S	Legitimately skipped
1	1	1=proportion or decimal
17	2	2=percent (If percent, then assure not percent predicted for the ratio)
5		Missing

PUL26		Post-bronchodilator spirometry performed
N	Value	Description
2360	0	0=No/NR
12	1	1=Yes

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PUL26A1		FEV1 post
N	Value	Description
12	Range	0.63 - 3.28 (median=2.365 mean=2.2067 std=0.7789)
2360	.S	Legitimately skipped

PUL26A2		FEV1 Percent Predicted post
N	Value	Description
8	Range	18 - 81 (median=73 mean=63.9 std=21.4)
2360	.S	Legitimately skipped
4		Missing

PUL26B1		FVC post
N	Value	Description
8	Range	0.63 - 3.76 (median=2.885 mean=2.5763 std=1.1175)
2360	.S	Legitimately skipped
4		Missing

PUL26B2		FVC Percent Predicted post
N	Value	Description
8	Range	15 - 94 (median=78 mean=67.6 std=26.0)
2360	.S	Legitimately skipped
4		Missing

PUL26C1		FEV1/FVC ratio post
N	Value	Description
8	Range	64 - 89 (median=72.69 mean=74.548 std=9.248)
2360	.S	Legitimately skipped
4		Missing

PUL26C2		Units for FEV1/FVC ratio post
N	Value	Description
2360	.S	Legitimately skipped
0	1	1=proportion or decimal
8	2	2=percent (If percent, then assure not percent predicted for the ratio)
4		Missing

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<i>PUL27</i>		<i>PEFR or PEF obtained at event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2170	0	0=No/NR
201	1	1=Yes
1		Missing

<i>PUL27B</i>		<i>First PEF recording</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
195	Range	50 - 450 (median=200 mean=211.9 std=77.8)
2171	.S	Legitimately skipped
6		Missing

<i>PUL27C</i>		<i>Worst recording</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
193	Range	50 - 450 (median=200 mean=208.4 std=76.8)
2171	.S	Legitimately skipped
8		Missing

<i>PUL28</i>		<i>PERF or PEF obtained at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2218	0	0=No/NR
154	1	1=Yes

<i>PUL28B</i>		<i>Last PEF recording</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
154	Range	120 - 510 (median=300 mean=302.3 std=84.9)
2218	.S	Legitimately skipped

<i>PUL29</i>		<i>VQ Scan done</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2321	0	0=No/NR
51	1	1=Yes

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<i>PUL29A</i>		<i>VQ Scan results</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2321	.S	Legitimately skipped
2	1	1=High probability
1	2	2=Intermediate probability
21	3	3=Low probability
24	4	4=No evidence of Pulmonary Embolus
1	5	5=Indeterminate
2		Missing

<i>PUL30</i>		<i>TTE or TEE performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1862	0	0=No/NR
510	1	1=Yes

<i>PUL30A</i>		<i>TTE/TEE Ejection fraction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
463	Range	5 - 85 (median=56 mean=53.3 std=15.5)
1862	.S	Legitimately skipped
47		Missing

<i>PUL30B</i>		<i>TTE/TEE RVSP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
273	Range	5 - 85 (median=36 mean=37.6 std=14.7)
1862	.S	Legitimately skipped
237		Missing

<i>PUL30C</i>		<i>Right Ventricular Hypertrophy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1862	.S	Legitimately skipped
121	0	0=None
6	1	1=Present
1	2	2=Mild
1	3	3=Mod
0	4	4=Severe
376	9	9=NR
5		Missing

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<i>PUL30D</i>		<i>Impaired RV systolic function</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1862	.S	Legitimately skipped
286	0	0=None
10	1	1=Present
12	2	2=Mild
11	3	3=Mod
5	4	4=Severe
181	9	9=NR
5		Missing

<i>PUL30E</i>		<i>Pulmonary hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1862	.S	Legitimately skipped
70	0	0=None
10	1	1=Present
28	2	2=Mild
29	3	3=Mod
19	4	4=Severe
350	9	9=NR
4		Missing

<i>PUL30F</i>		<i>Tricuspid Regurgitation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1862	.S	Legitimately skipped
134	0	0=None
7	1	1=Present
177	2	2=Mild
48	3	3=Mod
16	4	4=Severe
123	9	9=NR
5		Missing

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<i>PUL30G</i>		<i>Diastolic dysfunction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1862	.S	Legitimately skipped
319	0	0=No/Unknown/NR
186	1	1=Yes
5		Missing

<i>PUL31</i>		<i>White Blood Cell Count</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1861	Range	1.7 - 174 (median=9.9 mean=11.12 std=6.69)
511		Missing

<i>PUL32A</i>		<i>First Hemoglobin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1864	Range	3.8 - 18.7 (median=12.1 mean=11.85 std=2.27)
508		Missing

<i>PUL33A</i>		<i>First Hematocrit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1834	Range	11.7 - 55 (median=36.7 mean=35.92 std=6.45)
538		Missing

<i>PUL34A</i>		<i>First Sodium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1834	Range	101 - 164 (median=138 mean=137.1 std=4.3)
538		Missing

<i>PUL35A</i>		<i>First Serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1850	Range	0.2 - 37 (median=0.9 mean=1.43 std=1.93)
522		Missing

<i>PUL36A</i>		<i>First BUN</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1835	Range	1.1 - 162 (median=17 mean=23.4 std=19.7)
537		Missing

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<i>PUL37A</i>		<i>First Bicarbonate</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1786	Range	4 - 91 (median=27 mean=27.1 std=4.8)
586		Missing

<i>PUL38A</i>		<i>First BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
275	Range	4 - 19300 (median=129 mean=779.3 std=2052.7)
2097		Missing

<i>PUL38B</i>		<i>Upper limit normal BNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
189	Range	23 - 125 (median=100 mean=99.6 std=5.9)
2183		Missing

<i>PUL39A</i>		<i>First ProBNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
314	Range	4 - 130000 (median=472 mean=6105.5 std=14822.0)
2058		Missing

<i>PUL39B</i>		<i>Upper limit normal ProBNP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
244	Range	99 - 70000 (median=298 mean=672.7 std=4470.9)
2128		Missing

<i>PUL40</i>		<i>ABGs obtained</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1829	0	0=No/NR
529	1	1=Yes
14		Missing

<i>PUL40A1</i>		<i>First blood gas pH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
507	Range	6.59 - 7.69 (median=7.4 mean=7.37 std=0.12)
1839	.S	Legitimately skipped
26		Missing

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PUL40B1		Last blood gas pH
N	Value	Description
247	Range	6.92 - 7.58 (median=7.405 mean=7.3857 std=0.0970)
1839	.S	Legitimately skipped
286		Missing

PUL40A2		First blood gas PaCO2
N	Value	Description
504	Range	10.6 - 138 (median=40.05 mean=43.886 std=16.113)
1839	.S	Legitimately skipped
29		Missing

PUL40B2		Last blood gas PaCO2
N	Value	Description
245	Range	21 - 111 (median=41 mean=43.8 std=14.2)
1839	.S	Legitimately skipped
288		Missing

PUL40A3		First blood gas PaO2
N	Value	Description
502	Range	19.1 - 623.3 (median=80.05 mean=110.399 std=90.610)
1839	.S	Legitimately skipped
31		Missing

PUL40B3		Last blood gas PaO2
N	Value	Description
243	Range	19.1 - 534 (median=88.1 mean=104.19 std=58.78)
1839	.S	Legitimately skipped
290		Missing

PUL40A4		First blood gas O2 Sat
N	Value	Description
421	Range	18 - 100 (median=96 mean=90.9 std=14.9)
1839	.S	Legitimately skipped
112		Missing

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<i>PUL40B4</i>		<i>Last blood gas O2 Sat</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
211	Range	31 - 100 (median=96.8 mean=94.25 std=9.80)
1839	.S	Legitimately skipped
322		Missing

<i>PUL40C</i>		<i>Blood gas on room air</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1839	.S	Legitimately skipped
213	0	0=No
145	1	1=Yes
168	9	9=Not Recorded
7		Missing

<i>PUL40C1</i>		<i>Level oxy blood gas</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
204	Range	0.6 - 100 (median=40 mean=50.1 std=33.9)
2157	.S	Legitimately skipped
11		Missing

<i>PUL40C2</i>		<i>Level oxy Units blood gas</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2157	.S	Legitimately skipped
33	1	1=Liters
173	2	2=Percent
9		Missing

<i>PUL41</i>		<i>Sputum culture done</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2104	0	0=No/NR
267	1	1=Yes
1		Missing

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PUL41A		Sputum Culture Results
N	Value	Description
2105	.S	Legitimately skipped
107	0	0=Neg
80	1	1=Pos
80	9	9=Not Recorded

PUL41B1		Haemophilus Influenzae sputum
N	Value	Description
2292	.S	Legitimately skipped
78	0	0=No
1	1	1=Yes
1		Missing

PUL41B2		Moraxella Catarrhalis sputum
N	Value	Description
2292	.S	Legitimately skipped
79	0	0=No
0	1	1=Yes
1		Missing

PUL41B3		Streptococcus pneumoniae sputum
N	Value	Description
2292	.S	Legitimately skipped
77	0	0=No
2	1	1=Yes
1		Missing

PUL41B4		Methicillin-resistant Staph sputum
N	Value	Description
2292	.S	Legitimately skipped
65	0	0=No
14	1	1=Yes
1		Missing

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PUL41B5		<i>Staphylococcus aureus sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2292	.S	Legitimately skipped
75	0	0=No
4	1	1=Yes
1		Missing

PUL41B6		<i>Mycoplasma pneumoniae sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2292	.S	Legitimately skipped
78	0	0=No
1	1	1=Yes
1		Missing

PUL41B7		<i>Pseudomonas Aureginosa sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2292	.S	Legitimately skipped
71	0	0=No
8	1	1=Yes
1		Missing

PUL41B8		<i>Chlamydophila pneumoniae sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2292	.S	Legitimately skipped
79	0	0=No
0	1	1=Yes
1		Missing

PUL41B9		<i>Oropharyngeal flora sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2292	.S	Legitimately skipped
63	0	0=No
16	1	1=Yes
1		Missing

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<i>PUL41B10</i>		<i>Other sputum</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
51	Present	Text suppressed
2292	~	Legitimately skipped
29		Missing

<i>PUL42</i>		<i>Blood culture done</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1828	0	0=No/NR
544	1	1=Yes

<i>PUL42A</i>		<i>Culture Results</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1828	.S	Legitimately skipped
320	0	0=Neg
50	1	1=Pos
173	9	9=Not Recorded
1		Missing

<i>PUL42B1</i>		<i>Haemophilus Influenzae blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2322	.S	Legitimately skipped
50	0	0=No
0	1	1=Yes

<i>PUL42B2</i>		<i>Moraxella Catarrhalis blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2322	.S	Legitimately skipped
50	0	0=No
0	1	1=Yes

<i>PUL42B3</i>		<i>Streptococcus pneumoniae blood</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2322	.S	Legitimately skipped
47	0	0=No
3	1	1=Yes

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PUL42B4		Methicillin-resistant Staph blood
N	Value	Description
2322	.S	Legitimately skipped
43	0	0=No
7	1	1=Yes

PUL42B5		Staphylococcus aureus blood
N	Value	Description
2322	.S	Legitimately skipped
47	0	0=No
3	1	1=Yes

PUL42B6		Other blood
N	Value	Description
42	Present	Text suppressed
2322	~	Legitimately skipped
8		Missing

PUL43		Influenza swab
N	Value	Description
249	0	0=Neg
49	1	1=Pos
2053	9	9=Not Recorded
21		Missing

PUL44		CPAP or BiPap
N	Value	Description
2149	0	0=No/NR
223	1	1=Yes

PUL45		Mechanical Ventilation
N	Value	Description
2166	0	0=No/NR
206	1	1=Yes

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<i>PUL46</i>		<i>Inhaled short-acting beta-agonists</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1197	0	0=No/NR
1175	1	1=Yes

<i>PUL47</i>		<i>Inhaled short-acting anticholinergic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1396	0	0=No/NR
976	1	1=Yes

<i>PUL48</i>		<i>Use of continuous nebulizers</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1247	0	0=No/NR
1125	1	1=Yes

<i>PUL49</i>		<i>Magnesium injections in ED</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2240	0	0=No/NR
132	1	1=Yes

<i>PUL50</i>		<i>Oxygen</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1441	0	0=No/NR
931	1	1=Yes

<i>PUL51</i>		<i>IV Antibiotics</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1582	0	0=No/NR
790	1	1=Yes

<i>PUL52</i>		<i>IV corticosteroid</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1519	0	0=No/NR
853	1	1=Yes

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<i>PUL53</i>		<i>IV lasix or furosemide</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2085	0	0=No/NR
287	1	1=Yes

<i>PUL54A</i>		<i>Antibiotics-oral onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1898	0	0=No/NR
474	1	1=Yes

<i>PUL54B</i>		<i>Antibiotics-oral discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1720	0	0=No/NR
652	1	1=Yes

<i>PUL55A</i>		<i>Systemic corticosteroid onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1680	0	0=No/NR
692	1	1=Yes

<i>PUL55B</i>		<i>Systemic corticosteroid discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1747	0	0=No/NR
625	1	1=Yes

<i>PUL56A</i>		<i>short acting beta-agonists onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1045	0	0=No/NR
1327	1	1=Yes

<i>PUL56B</i>		<i>short acting beta-agonists discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1424	0	0=No/NR
948	1	1=Yes

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PUL57A		long-acting beta-agonist onset
N	Value	Description
2266	0	0=No/NR
106	1	1=Yes

PUL57B		long-acting beta-agonist discharge
N	Value	Description
2281	0	0=No/NR
91	1	1=Yes

PUL58A		short-acting anticholinergics onset
N	Value	Description
1533	0	0=No/NR
839	1	1=Yes

PUL58B		short-acting anticholinergics discharge
N	Value	Description
2134	0	0=No/NR
238	1	1=Yes

PUL59A		long-acting anticholinergics onset
N	Value	Description
2270	0	0=No/NR
102	1	1=Yes

PUL59B		long-acting anticholinergics discharge
N	Value	Description
2272	0	0=No/NR
100	1	1=Yes

PUL60A		Inhaled corticosteroids onset
N	Value	Description
1887	0	0=No/NR
485	1	1=Yes

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<i>PUL60B</i>		<i>Inhaled corticosteroids discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1988	0	0=No/NR
384	1	1=Yes

<i>PUL61A</i>		<i>Nebulized bronchodilators onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1387	0	0=No/NR
985	1	1=Yes

<i>PUL61B</i>		<i>Nebulized bronchodilators discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2072	0	0=No/NR
300	1	1=Yes

<i>PUL62A</i>		<i>Leukotriene antagonist onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2020	0	0=No/NR
352	1	1=Yes

<i>PUL62B</i>		<i>Leukotriene antagonist discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2087	0	0=No/NR
285	1	1=Yes

<i>PUL63A</i>		<i>Home oxygen onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2247	0	0=No/NR
125	1	1=Yes

<i>PUL63B</i>		<i>Home oxygen discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2257	0	0=No/NR
115	1	1=Yes

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<i>PUL0A_FUTIME</i>		<i>Futime from Randomization date for (PUL0A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2372	Range	22 - 5100 (median=2660 mean=2775.5 std=850.8)

<i>PUL0D_FUTIME</i>		<i>Futime from Randomization date for (PUL0D)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2372	Range	1 - 4206 (median=1576 mean=1676.8 std=1028.0)

<i>PUL3A_FUTIME</i>		<i>Futime from Randomization date for (PUL3A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2357	Range	1 - 4206 (median=1577 mean=1679.4 std=1029.4)
15		Missing

<i>PUL3C_FUTIME</i>		<i>Futime from Randomization date for (PUL3C)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1789	Range	2 - 4207 (median=1654 mean=1736.2 std=1030.8)
583		Missing

<i>PUL4A_FUTIME</i>		<i>Futime from Randomization date for (PUL4A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2351	Range	4 - 4227 (median=1585 mean=1685.2 std=1024.1)
21		Missing

<i>PUL27A_FUTIME</i>		<i>Futime from Randomization date for (PUL27A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
193	Range	41 - 3874 (median=947 mean=1165.5 std=863.3)
2179		Missing

<i>PUL28A_FUTIME</i>		<i>Futime from Randomization date for (PUL28A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
147	Range	41 - 3874 (median=871 mean=1114.0 std=825.7)
2225		Missing

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Pulmonary Reviewer 1 Form: FORM CODE=PLD1, VERSION=A

ID		HCHS Participant ID
N	Value	Description
2333	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
1585	1	Occurrence 1
408	2	Occurrence 2
153	3	Occurrence 3
69	4	Occurrence 4
44	5	Occurrence 5
24	6	Occurrence 6
13	7	Occurrence 7
11	8	Occurrence 8
5	9	Occurrence 9
7	10	Occurrence 10
3	11	Occurrence 11
2	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

PLD11		Meet SOL criteria for CLRD
N	Value	Description
192	1	1= Definite
48	2	2= Highly Probable
1025	3	3= Probable
563	4	4= NO/unknown
505		Missing

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PLD12A		Asthma
N	Value	Description
564	.S	Legitimately skipped
68	1	1= Definite
130	2	2= Highly Probable
755	3	3= Probable
2	4	4= Definitely Not
310	5	5= Probably not/Unknown
504		Missing

PLD12B		COPD
N	Value	Description
564	.S	Legitimately skipped
12	1	1= Definite
23	2	2= Highly Probable
412	3	3= Probable
7	4	4= Definitely Not
811	5	5= Probably not/Unknown
504		Missing

PLD12C		Emphysema
N	Value	Description
564	.S	Legitimately skipped
44	1	1= Definite
1	2	2= Highly Probable
30	3	3= Probable
52	4	4= Definitely Not
1138	5	5= Probably not/Unknown
504		Missing

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<i>PLD12D</i>		<i>Chronic Bronchitis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564	.S	Legitimately skipped
0	1	1= Definite
0	2	2= Highly Probable
12	3	3= Probable
5	4	4= Definitely Not
1248	5	5= Probably not/Unknown
504		Missing

<i>PLD13</i>		<i>Predominant CLRD subtype</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564	.S	Legitimately skipped
843	1	1= Asthma
361	2	2= COPD
33	3	3= Emphysema
3	4	4= Chronic Bronchitis
25	5	5= Unsure
504		Missing

<i>PLD14</i>		<i>Evidence of other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
564	.S	Legitimately skipped
1160	0	0=No
105	1	1=Yes
504		Missing

<i>PLD14A</i>		<i>Specify other lung disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
105	Present	Text suppressed
1724	~	Legitimately skipped
504		Missing

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PLD15		Exacerbation of CLRD
N	Value	Description
564	.S	Legitimately skipped
201	1	1= Definite
349	2	2= Highly Probable
125	3	3= Probable
590	4	4= No/Unknown
504		Missing

PLD15A		Evidence for another diagnosis
N	Value	Description
1154	.S	Legitimately skipped
545	1	1= No, "LONE" CLRD exacerbation
130	2	2= Yes, "COMORBID" CLRD
504		Missing

PLD15B1		Pneumonia
N	Value	Description
1699	.S	Legitimately skipped
79	1	1=Yes
51	2	2=No/unknown
504		Missing

PLD15B2		Pulmonary embolus
N	Value	Description
1699	.S	Legitimately skipped
2	1	1=Yes
128	2	2=No/unknown
504		Missing

PLD15B3		Pneumothorax
N	Value	Description
1699	.S	Legitimately skipped
3	1	1=Yes
127	2	2=No/unknown
504		Missing

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PLD15B4		Acute myocardial infarction
N	Value	Description
1699	.S	Legitimately skipped
6	1	1=Yes
124	2	2=No/unknown
504		Missing

PLD15B5		Acute decompensated heart failure
N	Value	Description
1699	.S	Legitimately skipped
34	1	1=Yes
96	2	2=No/unknown
504		Missing

PLD15B5A		Type of ADHF
N	Value	Description
1795	.S	Legitimately skipped
21	1	1= Left-sided
4	2	2= Right-sided
9	3	3= Unknown
504		Missing

PLD15B6		Other condition
N	Value	Description
1699	.S	Legitimately skipped
25	1	1=Yes
105	2	2=No/unknown
504		Missing

PLD15B6A		Specify other condition
N	Value	Description
25	Present	Text suppressed
1804	~	Legitimately skipped
504		Missing

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PLD16		Comments
N	Value	Description
68	Present	Text suppressed
564	~	Legitimately skipped
1701		Missing

PLD10A_FUTIME		Futime from Randomization date for (PLD10A)
N	Value	Description
1828	Range	1289 - 5170 (median=3151.5 mean=3156.11 std=686.56)
505		Missing

PLD10D_FUTIME		Futime from Randomization date for (PLD10D)
N	Value	Description
2332	Range	1 - 4206 (median=1598.5 mean=1691.92 std=1024.57)
1		Missing

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Pulmonary Reviewer 2 Form: FORM CODE=PLD2, VERSION=A

ID		HCHS Participant ID
N	Value	Description
2329	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
1583	1	Occurrence 1
406	2	Occurrence 2
153	3	Occurrence 3
69	4	Occurrence 4
44	5	Occurrence 5
24	6	Occurrence 6
13	7	Occurrence 7
11	8	Occurrence 8
5	9	Occurrence 9
7	10	Occurrence 10
3	11	Occurrence 11
2	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

PLD21		Meet SOL criteria for CLRD
N	Value	Description
169	1	1= Definite
48	2	2= Highly Probable
1021	3	3= Probable
552	4	4= NO/unknown
539		Missing

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PLD22A		Asthma
N	Value	Description
552	.S	Legitimately skipped
60	1	1= Definite
124	2	2= Highly Probable
761	3	3= Probable
2	4	4= Definitely Not
291	5	5= Probably not/Unknown
539		Missing

PLD22B		COPD
N	Value	Description
552	.S	Legitimately skipped
8	1	1= Definite
13	2	2= Highly Probable
422	3	3= Probable
3	4	4= Definitely Not
792	5	5= Probably not/Unknown
539		Missing

PLD22C		Emphysema
N	Value	Description
552	.S	Legitimately skipped
37	1	1= Definite
3	2	2= Highly Probable
31	3	3= Probable
50	4	4= Definitely Not
1117	5	5= Probably not/Unknown
539		Missing

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PLD22D		Chronic Bronchitis
N	Value	Description
552	.S	Legitimately skipped
0	1	1= Definite
1	2	2= Highly Probable
9	3	3= Probable
4	4	4= Definitely Not
1224	5	5= Probably not/Unknown
539		Missing

PLD23		Predominant CLRD subtype
N	Value	Description
552	.S	Legitimately skipped
835	1	1= Asthma
350	2	2= COPD
31	3	3= Emphysema
5	4	4= Chronic Bronchitis
17	5	5= Unsure
539		Missing

PLD24		Evidence of other lung disease
N	Value	Description
552	.S	Legitimately skipped
1119	0	0=No
119	1	1=Yes
539		Missing

PLD24A		Specify other lung disease
N	Value	Description
119	Present	Text suppressed
1671	~	Legitimately skipped
539		Missing

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PLD25		Exacerbation of CLRD
N	Value	Description
552	.S	Legitimately skipped
189	1	1= Definite
365	2	2= Highly Probable
114	3	3= Probable
569	4	4= No/Unknown
540		Missing

PLD25A		Evidence for another diagnosis
N	Value	Description
1122	.S	Legitimately skipped
532	1	1= No, "LONE" CLRD exacerbation
136	2	2= Yes, "COMORBID" CLRD
539		Missing

PLD25B1		Pneumonia
N	Value	Description
1654	.S	Legitimately skipped
79	1	1=Yes
57	2	2=No/unknown
539		Missing

PLD25B2		Pulmonary embolus
N	Value	Description
1654	.S	Legitimately skipped
0	1	1=Yes
136	2	2=No/unknown
539		Missing

PLD25B3		Pneumothorax
N	Value	Description
1654	.S	Legitimately skipped
3	1	1=Yes
133	2	2=No/unknown
539		Missing

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PLD25B4		Acute myocardial infarction
N	Value	Description
1654	.S	Legitimately skipped
2	1	1=Yes
134	2	2=No/unknown
539		Missing

PLD25B5		Acute decompensated heart failure
N	Value	Description
1654	.S	Legitimately skipped
31	1	1=Yes
105	2	2=No/unknown
539		Missing

PLD25B5A		Type of ADHF
N	Value	Description
1759	.S	Legitimately skipped
16	1	1= Left-sided
8	2	2= Right-sided
7	3	3= Unknown
539		Missing

PLD25B6		Other condition
N	Value	Description
1654	.S	Legitimately skipped
40	1	1=Yes
96	2	2=No/unknown
539		Missing

PLD25B6A		Specify other condition
N	Value	Description
40	Present	Text suppressed
1750	~	Legitimately skipped
539		Missing

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PLD26		Comments
N	Value	Description
47	Present	Text suppressed
552	~	Legitimately skipped
1730		Missing

PLD20A_FUTIME		Futime from Randomization date for (PLD20A)
N	Value	Description
1790	Range	420 - 4817 (median=3123 mean=3101.4 std=662.4)
539		Missing

PLD20D_FUTIME		Futime from Randomization date for (PLD20D)
N	Value	Description
2328	Range	1 - 4206 (median=1599 mean=1693.6 std=1024.3)
1		Missing

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Pulmonary Adjudication Reviewer Form: FORM CODE=PLD3, VERSION=A

ID		HCHS Participant ID
N	Value	Description
468	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
311	1	Occurrence 1
99	2	Occurrence 2
31	3	Occurrence 3
10	4	Occurrence 4
8	5	Occurrence 5
4	6	Occurrence 6
3	8	Occurrence 8
1	9	Occurrence 9
1	11	Occurrence 11

PLD31		Meet SOL criteria for CLRD
N	Value	Description
93	1	1= Definite
29	2	2= Highly Probable
255	3	3= Probable
40	4	4= NO/unknown
51		Missing

PLD32A		Asthma
N	Value	Description
40	.S	Legitimately skipped
27	1	1= Definite
56	2	2= Highly Probable
203	3	3= Probable
0	4	4= Definitely Not
91	5	5= Probably not/Unknown
51		Missing

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PLD32B		COPD
N	Value	Description
40	.S	Legitimately skipped
5	1	1= Definite
13	2	2= Highly Probable
118	3	3= Probable
1	4	4= Definitely Not
240	5	5= Probably not/Unknown
51		Missing

PLD32C		Emphysema
N	Value	Description
40	.S	Legitimately skipped
18	1	1= Definite
1	2	2= Highly Probable
13	3	3= Probable
5	4	4= Definitely Not
340	5	5= Probably not/Unknown
51		Missing

PLD32D		Chronic Bronchitis
N	Value	Description
40	.S	Legitimately skipped
0	1	1= Definite
0	2	2= Highly Probable
6	3	3= Probable
3	4	4= Definitely Not
368	5	5= Probably not/Unknown
51		Missing

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PLD33		Predominant CLRD subtype
N	Value	Description
40	.S	Legitimately skipped
252	1	1= Asthma
104	2	2= COPD
12	3	3= Emphysema
2	4	4= Chronic Bronchitis
7	5	5= Unsure
51		Missing

PLD34		Evidence of other lung disease
N	Value	Description
40	.S	Legitimately skipped
346	0	0=No
31	1	1=Yes
51		Missing

PLD34A		Specify other lung disease
N	Value	Description
31	Present	Text suppressed
386	~	Legitimately skipped
51		Missing

PLD35		Exacerbation of CLRD
N	Value	Description
40	.S	Legitimately skipped
86	1	1= Definite
115	2	2= Highly Probable
56	3	3= Probable
120	4	4= No/Unknown
51		Missing

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PLD35A		Evidence for another diagnosis
N	Value	Description
160	.S	Legitimately skipped
186	1	1= No, "LONE" CLRD exacerbation
71	2	2= Yes, "COMORBID" CLRD
51		Missing

PLD35B1		Pneumonia
N	Value	Description
346	.S	Legitimately skipped
38	1	1=Yes
33	2	2=No/unknown
51		Missing

PLD35B2		Pulmonary embolus
N	Value	Description
346	.S	Legitimately skipped
1	1	1=Yes
70	2	2=No/unknown
51		Missing

PLD35B3		Pneumothorax
N	Value	Description
346	.S	Legitimately skipped
2	1	1=Yes
69	2	2=No/unknown
51		Missing

PLD35B4		Acute myocardial infarction
N	Value	Description
346	.S	Legitimately skipped
3	1	1=Yes
68	2	2=No/unknown
51		Missing

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PLD35B5		Acute decompensated heart failure
N	Value	Description
346	.S	Legitimately skipped
18	1	1=Yes
53	2	2=No/unknown
51		Missing

PLD35B5A		Type of ADHF
N	Value	Description
399	.S	Legitimately skipped
13	1	1= Left-sided
2	2	2= Right-sided
3	3	3= Unknown
51		Missing

PLD35B6		Other condition
N	Value	Description
346	.S	Legitimately skipped
18	1	1=Yes
53	2	2=No/unknown
51		Missing

PLD35B6A		Specify other condition
N	Value	Description
18	Present	Text suppressed
399	~	Legitimately skipped
51		Missing

PLD36		Comments
N	Value	Description
49	Present	Text suppressed
40	~	Legitimately skipped
379		Missing

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PLD30A_FUTIME		Futime from Randomization date for (PLD30A)
N	Value	Description
417	Range	328 - 5185 (median=3536 mean=3496.0 std=682.1)
51		Missing

PLD30D_FUTIME		Futime from Randomization date for (PLD30D)
N	Value	Description
468	Range	1 - 3168 (median=1114.5 mean=1230.25 std=776.99)

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Stroke Abstraction Form: FORM CODE=STR, VERSION=1

ID		HCHS Participant ID
N	Value	Description
452	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
333	1	Occurrence 1
79	2	Occurrence 2
22	3	Occurrence 3
6	4	Occurrence 4
8	5	Occurrence 5
2	6	Occurrence 6
2	7	Occurrence 7

STR1A		Time of arrival
N	Value	Description
438	Range	0:00:00 - 23:59:00
14		Missing

STR1AHR		Hour of Time of arrival
N	Value	Description
438	Range	0 - 23 (median=13 mean=13.1 std=5.5)
14		Missing

STR1AMN		Minute of Time of arrival
N	Value	Description
438	Range	0 - 59 (median=30 mean=29.3 std=17.6)
14		Missing

STR1ASC		Second of Time of arrival
N	Value	Description
452		Missing

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STR2		Primary admitting code
N	Value	Description
393	Present	Text suppressed
59		Missing

STR4		Primary discharge code
N	Value	Description
405	Present	Text suppressed
47		Missing

STR5		Arrival mode
N	Value	Description
184	1	1=EMS
96	2	2=Private transport/walk-in
11	3	3=Transfer from another hospital
0	4	4=Mobile stroke unit
161	9	9=Unknown

STR6		Transferred from this hospital to another
N	Value	Description
438	0	0=No
11	1	1=Yes
3	9	9=Unknown

STR7		DNR/DNI present
N	Value	Description
135	0	0=No
36	1	1=Yes
281	9	9=Unknown

STR8		New neurological signs/symptoms
N	Value	Description
91	0	0=No
353	1	1=Yes
8	9	9=Unknown

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STR9		Stroke or TIA this hospitalization
N	Value	Description
426	0	0=No/NR
26	1	1=Yes

STR10		event in setting of a procedure
N	Value	Description
427	0	0=No/NR
25	1	1=Yes

STR10A1		cerebral angiogram
N	Value	Description
427	.S	Legitimately skipped
23	0	0=No/NR
2	1	1=Yes

STR10A2		coronary angiogram
N	Value	Description
427	.S	Legitimately skipped
20	0	0=No/NR
5	1	1=Yes

STR10A3		cardiac surgery or procedures
N	Value	Description
427	.S	Legitimately skipped
20	0	0=No/NR
5	1	1=Yes

STR10A4		carotid endarterectomy
N	Value	Description
427	.S	Legitimately skipped
25	0	0=No/NR
0	1	1=Yes

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STR10A5		carotid stenting
N	Value	Description
427	.S	Legitimately skipped
25	0	0=No/NR
0	1	1=Yes

STR10A6		other procedure performed
N	Value	Description
427	.S	Legitimately skipped
7	0	0=No/NR
18	1	1=Yes

STR10A6A		other procedure
N	Value	Description
18	Present	Text suppressed
434	~	Legitimately skipped

STR11A		symptom onset date known
N	Value	Description
66	0	0=No
325	1	1=Yes
61		Missing

STR11C		symptom onset time known
N	Value	Description
127	.S	Legitimately skipped
192	0	0=No
132	1	1=Yes
1		Missing

STR11D		symptom onset time
N	Value	Description
132	Range	0:33:00 - 23:40:00
320	.S	Legitimately skipped

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STR11DHR		Hour of symptom onset time
N	Value	Description
132	Range	0 - 23 (median=13 mean=12.7 std=6.0)
320		Missing

STR11DMN		Minute of symptom onset time
N	Value	Description
132	Range	0 - 59 (median=0 mean=12.0 std=17.2)
320		Missing

STR11DSC		Second of symptom onset time
N	Value	Description
452		Missing

STR12A		free of deficits date known
N	Value	Description
67	0	0=No
324	1	1=Yes
61		Missing

STR12C		free of deficits time known
N	Value	Description
128	.S	Legitimately skipped
184	0	0=No
139	1	1=Yes
1		Missing

STR12D		free of deficits time
N	Value	Description
139	Range	0:33:00 - 23:40:00
313	.S	Legitimately skipped

STR12DHR		Hour of free of deficits time
N	Value	Description
139	Range	0 - 23 (median=13 mean=13.1 std=6.1)
313		Missing

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STR12DMN		Minute of free of deficits time
N	Value	Description
139	Range	0 - 59 (median=0 mean=11.8 std=16.9)
313		Missing

STR12DSC		Second of free of deficits time
N	Value	Description
452		Missing

STR13		history of stroke
N	Value	Description
213	0	0=No
196	1	1=Yes
43	9	9=Unknown

STR13A		previous stroke date known
N	Value	Description
256	.S	Legitimately skipped
87	0	0=No
109	1	1=Yes

STR13C		Previous stroke type
N	Value	Description
256	.S	Legitimately skipped
156	1	1=Ischemic
8	2	2=Intracranial hemorrhage
7	3	3=Subarachnoid hemorrhage
25	9	9=Unknown

STR13C1		history of hematoma evacuation
N	Value	Description
444	.S	Legitimately skipped
0	0	0=No
1	1	1=Yes
7	9	9=Unknown

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STR13C2		history of intra-hematoma tPA
N	Value	Description
444	.S	Legitimately skipped
0	0	0=No
0	1	1=Yes
8	9	9=Unknown

STR13C3		history of aneurysm
N	Value	Description
445	.S	Legitimately skipped
1	0	0=No
3	1	1=Yes
3	9	9=Unknown

STR13C4		history of aneurysm coiling
N	Value	Description
445	.S	Legitimately skipped
1	0	0=No
0	1	1=Yes
6	9	9=Unknown

STR13C5		history of aneurysm clipping
N	Value	Description
445	.S	Legitimately skipped
0	0	0=No
2	1	1=Yes
5	9	9=Unknown

STR14		history of TIA
N	Value	Description
257	0	0=No
45	1	1=Yes
150	9	9=Unknown

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STR14A		previous TIA date known
N	Value	Description
407	.S	Legitimately skipped
35	0	0=No
10	1	1=Yes

STR15		history of CAD
N	Value	Description
279	0	0=No
131	1	1=Yes
42	9	9=Unknown

STR16		history of PVD
N	Value	Description
373	0	0=No
27	1	1=Yes
52	9	9=Unknown

STR17		history of carotid disease
N	Value	Description
360	0	0=No
35	1	1=Yes
57	9	9=Unknown

STR18		history of ESRD
N	Value	Description
377	0	0=No
24	1	1=Yes
51	9	9=Unknown

STR19		history of diabetes
N	Value	Description
171	0	0=No
249	1	1=Yes
32	9	9=Unknown

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STR20		history of HTN
N	Value	Description
56	0	0=No
372	1	1=Yes
24	9	9=Unknown

STR21		history of dyslipidemia
N	Value	Description
129	0	0=No
253	1	1=Yes
70	9	9=Unknown

STR22		history of smoking
N	Value	Description
239	0	0=No
156	1	1=Yes
57	9	9=Unknown

STR23		history of alcohol use disorder
N	Value	Description
317	0	0=No
58	1	1=Yes
77	9	9=Unknown

STR24		history of illicit stimulant use four weeks prior
N	Value	Description
307	0	0=No
25	1	1=Yes
120	9	9=Unknown

STR25A		myocardial infarction
N	Value	Description
439	0	0=No/NR
13	1	1=Yes

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STR25B		<i>intracardiac thrombus or intracardiac tumor</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
450	0	0=No/NR
2	1	1=Yes

STR25C		<i>atrial fibrillation/atrial flutter</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	0	0=No/NR
50	1	1=Yes

STR25C1		<i>24 hours of cardiac monitoring</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
222	0	0=No/NR
230	1	1=Yes

STR25D		<i>left-sided atrial enlargement on echocardiogram</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
364	0	0=No/NR
88	1	1=Yes

STR25E		<i>rheumatic heart disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
447	0	0=No/NR
5	1	1=Yes

STR25F		<i>systemic embolus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
450	0	0=No/NR
2	1	1=Yes

STR25G		<i>hematologic abnormality (hypercoagulable state)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
446	0	0=No/NR
6	1	1=Yes

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STR25H		hematologic abnormality (hemorrhagic)
N	Value	Description
446	0	0=No/NR
6	1	1=Yes

STR25I		migraine headaches
N	Value	Description
423	0	0=No/NR
29	1	1=Yes

STR25J		left-sided bioprosthetic valve
N	Value	Description
443	0	0=No/NR
9	1	1=Yes

STR25K		left-sided mechanical valve
N	Value	Description
445	0	0=No/NR
7	1	1=Yes

STR26		time between onset and resolution of new neuro symptoms
N	Value	Description
66	1	1=Less than 24 hours
68	2	2=24-48 hours
35	3	3=Greater than 48 hours
181	4	4=Unknown
102	5	5=Not applicable

STR27		NIHSS recorded
N	Value	Description
352	0	0=No
99	1	1=Yes
1		Missing

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STR27A		NIHSS score
N	Value	Description
99	Range	0 - 27 (median=3 mean=5.4 std=6.4)
353	.S	Legitimately skipped

STR27B1		LOCa
N	Value	Description
100	.S	Legitimately skipped
313	0	0
15	1	1
8	2	2
16	3	3

STR27B2		LOCb
N	Value	Description
100	.S	Legitimately skipped
280	0	0
43	1	1
29	2	2

STR27B3		LOCc
N	Value	Description
100	.S	Legitimately skipped
304	0	0
19	1	1
29	2	2

STR27B4		Best Gaze
N	Value	Description
100	.S	Legitimately skipped
324	0	0
24	1	1
4	2	2

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STR27B5		Visual Fields
N	Value	Description
100	.S	Legitimately skipped
327	0	0
12	1	1
9	2	2
4	3	3

STR27B6		Facial Palsy
N	Value	Description
100	.S	Legitimately skipped
286	0	0
62	1	1
2	2	2
2	3	3

STR27B7A		Motor Left Arm
N	Value	Description
100	.S	Legitimately skipped
278	0	0
47	1	1
5	2	2
2	3	3
20	4	4

STR27B7B		Motor Right Arm
N	Value	Description
100	.S	Legitimately skipped
279	0	0
51	1	1
6	2	2
3	3	3
13	4	4

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STR27B8A		Motor Left Leg
N	Value	Description
100	.S	Legitimately skipped
283	0	0
40	1	1
7	2	2
5	3	3
17	4	4

STR27B8B		Motor Right Leg
N	Value	Description
100	.S	Legitimately skipped
286	0	0
46	1	1
5	2	2
3	3	3
12	4	4

STR27B9		Limb Ataxia
N	Value	Description
100	.S	Legitimately skipped
330	0	0
18	1	1
4	2	2

STR27B10		Sensory
N	Value	Description
100	.S	Legitimately skipped
288	0	0
61	1	1
3	2	2

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STR27B11		Best Language
N	Value	Description
100	.S	Legitimately skipped
302	0	0
31	1	1
9	2	2
10	3	3

STR27B12		Dysarthria
N	Value	Description
100	.S	Legitimately skipped
284	0	0
63	1	1
5	2	2

STR27B13		Extinction and Inattention
N	Value	Description
100	.S	Legitimately skipped
342	0	0
3	1	1
7	2	2

STR27B14		Total Score
N	Value	Description
352	Range	0 - 38 (median=2 mean=3.4 std=5.0)
100	.S	Legitimately skipped

STR28		asleep at time of event
N	Value	Description
192	0	0=No
65	1	1=Yes
195	9	9=Unknown

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STR29		severe headache at onset or admission
N	Value	Description
204	0	0=No
48	1	1=Yes
200	9	9=Unknown

STR30		Vomiting
N	Value	Description
198	0	0=No
74	1	1=Yes
180	9	9=Unknown

STR31		Blurry Vision
N	Value	Description
132	0	0=No
42	1	1=Yes
278	9	9=Unknown

STR32		Diplopia
N	Value	Description
122	0	0=No
15	1	1=Yes
315	9	9=Unknown

STR33		Vertigo
N	Value	Description
90	0	0=No
25	1	1=Yes
337	9	9=Unknown

STR34		Dizziness
N	Value	Description
25	.S	Legitimately skipped
86	0	0=No
94	1	1=Yes
247	9	9=Unknown

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STR35		Seizure
N	Value	Description
99	0	0=No
24	1	1=Yes
329	9	9=Unknown

STR35A		Seizure as first neurologic symptom
N	Value	Description
428	.S	Legitimately skipped
5	0	0=No
10	1	1=Yes
9	9	9=Unknown

STR36		Level of Consciousness
N	Value	Description
353	.S	Legitimately skipped
62	0	0=Normal
33	1	1=Abnormal
3	2	2=Unconscious
1	9	9=Unknown

STR37		Aphasia
N	Value	Description
353	.S	Legitimately skipped
56	0	0=No
25	1	1=Yes
18	9	9=Unknown

STR38		Hemianopsia
N	Value	Description
353	.S	Legitimately skipped
46	0	0=No
13	1	1=Yes
40	9	9=Unknown

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STR39		<i>abnormality of cranial nerves</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
353	.S	Legitimately skipped
52	0	0=No
10	1	1=Yes
37	9	9=Unknown

STR39A		<i>location of abnormality</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
442	.S	Legitimately skipped
4	1	1=Left
5	2	2=Right
1	3	3=Both
0	9	9=NR

STR40		<i>Horizontal gaze palsy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
353	.S	Legitimately skipped
52	0	0=No
9	1	1=Yes
38	9	9=Unknown

STR40A		<i>Location of Gaze Deviation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
443	.S	Legitimately skipped
3	1	1=to Left
5	2	2=to Right
0	3	3=Both
1	9	9=NR

STR41		<i>Dysphagia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
353	.S	Legitimately skipped
44	0	0=No
9	1	1=Yes
46	9	9=Unknown

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STR42		Dysarthria
N	Value	Description
353	.S	Legitimately skipped
50	0	0=No
39	1	1=Yes
10	9	9=Unknown

STR43		weakness
N	Value	Description
353	.S	Legitimately skipped
23	0	0=No
72	1	1=Yes
4	9	9=Unknown

STR43A		face weakness
N	Value	Description
380	.S	Legitimately skipped
21	1	1=Left
16	2	2=Right
3	3	3=Both
12	4	4=none
20	9	9=NR

STR43B		arm weakness
N	Value	Description
380	.S	Legitimately skipped
30	1	1=Left
22	2	2=Right
11	3	3=Both
4	4	4=none
5	9	9=NR

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STR43C		leg weakness
N	Value	Description
380	.S	Legitimately skipped
27	1	1=Left
17	2	2=Right
9	3	3=Both
11	4	4=none
8	9	9=NR

STR44		sensory deficit
N	Value	Description
353	.S	Legitimately skipped
52	0	0=No
37	1	1=Yes
10	9	9=Unknown

STR44A		sensory deficit face
N	Value	Description
415	.S	Legitimately skipped
13	1	1=Left
5	2	2=Right
2	3	3=Both
7	4	4=none
10	9	9=NR

STR44B		sensory deficit arm
N	Value	Description
415	.S	Legitimately skipped
17	1	1=Left
10	2	2=Right
1	3	3=Both
4	4	4=none
5	9	9=NR

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STR44C		sensory deficit leg
N	Value	Description
415	.S	Legitimately skipped
16	1	1=Left
9	2	2=Right
3	3	3=Both
4	4	4=none
5	9	9=NR

STR45		visual fields abnormality
N	Value	Description
353	.S	Legitimately skipped
55	0	0=No
17	1	1=Yes
27	9	9=Unknown

STR46		ataxic gait
N	Value	Description
353	.S	Legitimately skipped
43	0	0=No
8	1	1=Yes
48	9	9=Unknown

STR47		hemiataxia
N	Value	Description
353	.S	Legitimately skipped
39	0	0=No
7	1	1=Yes
53	9	9=Unknown

STR48A1		first systolic blood pressure
N	Value	Description
391	Range	64 - 270 (median=148 mean=151.4 std=32.3)
61		Missing

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STR48A2		first diastolic blood pressure
N	Value	Description
389	Range	31 - 160 (median=79 mean=81.3 std=18.4)
63		Missing

STR49		first glucose level
N	Value	Description
389	Range	49 - 1221 (median=127 mean=161.7 std=112.0)
63		Missing

STR50		lumbar puncture performed
N	Value	Description
439	0	0=No/NR
13	1	1=Yes

STR50A		lumbar puncture diagnosis
N	Value	Description
439	.S	Legitimately skipped
4	0	0=Normal
2	1	1=Abnormal
7	9	9=Unknown

STR51		CT scan of head performed
N	Value	Description
109	0	0=No
231	1	1=Yes, 1 CT
108	2	2=Yes, 2 or more CTs
4	9	9=NR

STR51B		First CT acute or subacute stroke diagnosed
N	Value	Description
113	.S	Legitimately skipped
222	0	0=No
106	1	1=Yes
11	9	9=Unknown/NR

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STR51D		Last CT acute or subacute stroke or previously unseen stroke diagnosed
N	Value	Description
344	.S	Legitimately skipped
44	0	0=No
55	1	1=Yes
9	9	9=Unknown/NR

STR52		MRI including head performed
N	Value	Description
196	0	0=No
227	1	1=Yes
29	9	9=Unknown/NR

STR52A		Acute or subacute stroke diagnosed on MRI
N	Value	Description
225	.S	Legitimately skipped
102	0	0=No
115	1	1=Yes
10	9	9=Unknown/NR

STR52B		DWI performed
N	Value	Description
225	.S	Legitimately skipped
16	0	0=No
155	1	1=Yes
56	9	9=Unknown/NR

STR52C		DWI diagnosis
N	Value	Description
297	.S	Legitimately skipped
66	0	0=Normal
80	1	1=Abnormal
9	9	9=Unknown/NR

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STR53		cerebrovascular angiography performed
N	Value	Description
164	0	0=No
263	1	1=Yes
25	9	9=NR

STR53B		CTA head
N	Value	Description
189	.S	Legitimately skipped
148	0	0=No
94	1	1=Yes
21	9	9=NR

STR53C		CTA neck
N	Value	Description
189	.S	Legitimately skipped
158	0	0=No
80	1	1=Yes
25	9	9=NR

STR53D		MRA head
N	Value	Description
189	.S	Legitimately skipped
123	0	0=No
116	1	1=Yes
24	9	9=NR

STR53E		MRA neck
N	Value	Description
189	.S	Legitimately skipped
183	0	0=No
48	1	1=Yes
32	9	9=NR

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STR53F		Carotid doppler
N	Value	Description
189	.S	Legitimately skipped
104	0	0=No
141	1	1=Yes
18	9	9=NR

STR53G		Transcranial Doppler
N	Value	Description
189	.S	Legitimately skipped
227	0	0=No
6	1	1=Yes
30	9	9=NR

STR53H		Catheter Angiogram
N	Value	Description
189	.S	Legitimately skipped
224	0	0=No
6	1	1=Yes
33	9	9=NR

STR53I		Stenosis - Right Internal Carotid Artery
N	Value	Description
189	.S	Legitimately skipped
160	0	0=No
33	1	1=Yes
70	9	9=NR

STR53J		Stenosis - Left Internal Carotid Artery
N	Value	Description
189	.S	Legitimately skipped
153	0	0=No
36	1	1=Yes
74	9	9=NR

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STR53K		Stenosis - Right Vertebral Artery
N	Value	Description
189	.S	Legitimately skipped
117	0	0=No
4	1	1=Yes
142	9	9=NR

STR53L		Stenosis - Left Vertebral Artery
N	Value	Description
189	.S	Legitimately skipped
119	0	0=No
2	1	1=Yes
142	9	9=NR

STR53M		Stenosis - Right Internal Carotid Artery
N	Value	Description
189	.S	Legitimately skipped
135	0	0=No
23	1	1=Yes
105	9	9=NR

STR53N		Stenosis - Left Internal Carotid Artery
N	Value	Description
189	.S	Legitimately skipped
136	0	0=No
24	1	1=Yes
103	9	9=NR

STR53O		Stenosis - Right Vertebral Artery
N	Value	Description
189	.S	Legitimately skipped
134	0	0=No
8	1	1=Yes
121	9	9=NR

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STR53P		Stenosis - Left Vertebral Artery
N	Value	Description
189	.S	Legitimately skipped
136	0	0=No
3	1	1=Yes
124	9	9=NR

STR53Q		Stenosis - Right Middle Cerebral Artery
N	Value	Description
189	.S	Legitimately skipped
129	0	0=No
10	1	1=Yes
124	9	9=NR

STR53R		Stenosis - Left Middle Cerebral Artery
N	Value	Description
189	.S	Legitimately skipped
130	0	0=No
13	1	1=Yes
120	9	9=NR

STR53S		Stenosis - Right Anterior Cerebral Artery
N	Value	Description
189	.S	Legitimately skipped
136	0	0=No
2	1	1=Yes
125	9	9=NR

STR53T		Stenosis - Left Anterior Cerebral Artery
N	Value	Description
189	.S	Legitimately skipped
139	0	0=No
3	1	1=Yes
121	9	9=NR

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STR53U		Stenosis - Right Posterior Cerebral Artery
N	Value	Description
189	.S	Legitimately skipped
131	0	0=No
9	1	1=Yes
123	9	9=NR

STR53V		Stenosis - Left Posterior Cerebral Artery
N	Value	Description
189	.S	Legitimately skipped
132	0	0=No
8	1	1=Yes
123	9	9=NR

STR53W		Stenosis - Basilar Artery
N	Value	Description
189	.S	Legitimately skipped
130	0	0=No
6	1	1=Yes
127	9	9=NR

STR54		transthoracic echocardiography study performed
N	Value	Description
228	0	0=No/NR
224	1	1=Yes

STR54A		transthoracic ejection fraction
N	Value	Description
201	Range	20 - 84 (median=58 mean=56.9 std=12.8)
228	.S	Legitimately skipped
23		Missing

STR55		transesophageal echocardiography study performed
N	Value	Description
427	0	0=No/NR
25	1	1=Yes

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STR55A		<i>transesophageal ejection fraction</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14	Range	20 - 70 (median=55 mean=51.7 std=15.8)
427	.S	Legitimately skipped
11		Missing

STR56A		<i>intracardiac thrombus</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
219	.S	Legitimately skipped
71	0	0=No
2	1	1=Yes
160	9	9=NR

STR56B		<i>mitral stenosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
219	.S	Legitimately skipped
59	0	0=No
3	1	1=Yes
171	9	9=NR

STR56C		<i>mitral regurgitation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
219	.S	Legitimately skipped
70	0	0=No
107	1	1=Yes
56	9	9=NR

STR56D		<i>aortic stenosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
219	.S	Legitimately skipped
88	0	0=No
11	1	1=Yes
134	9	9=NR

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STR56E		aortic regurgitation
N	Value	Description
219	.S	Legitimately skipped
98	0	0=No
43	1	1=Yes
92	9	9=NR

STR56F		poor ventricular function
N	Value	Description
219	.S	Legitimately skipped
120	0	0=No
60	1	1=Yes
53	9	9=NR

STR56G		dilated left ventricle
N	Value	Description
219	.S	Legitimately skipped
145	0	0=No
22	1	1=Yes
66	9	9=NR

STR56H		ascending aortic arch atheroma
N	Value	Description
219	.S	Legitimately skipped
46	0	0=No
7	1	1=Yes
180	9	9=NR

STR56H1		ascending aortic arch atheroma diagnosis
N	Value	Description
445	.S	Legitimately skipped
4	1	1=Mild
0	2	2=Moderate
0	3	3=Severe
3	9	9=NR

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STR56I		atrioseptal aneurysm
N	Value	Description
219	.S	Legitimately skipped
15	0	0=No
3	1	1=Yes
215	9	9=NR

STR56J		patent foramen ovale (PFO)
N	Value	Description
219	.S	Legitimately skipped
26	0	0=No
6	1	1=Yes
201	9	9=NR

STR56K		valve vegetations
N	Value	Description
219	.S	Legitimately skipped
58	0	0=No
5	1	1=Yes
170	9	9=NR

STR56L		artificial valve
N	Value	Description
219	.S	Legitimately skipped
48	0	0=No
7	1	1=Yes
178	9	9=NR

STR56M		left atrial enlargement
N	Value	Description
219	.S	Legitimately skipped
87	0	0=No
84	1	1=Yes
62	9	9=NR

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STR57		first sodium
N	Value	Description
398	Range	117 - 148 (median=139 mean=138.7 std=4.0)
54		Missing

STR58		first creatinine
N	Value	Description
402	Range	0.3 - 19 (median=0.9 mean=1.27 std=1.65)
50		Missing

STR59		first BUN
N	Value	Description
397	Range	5 - 160 (median=17 mean=21.8 std=18.2)
55		Missing

STR60		first Hgb
N	Value	Description
409	Range	3.9 - 19 (median=13.1 mean=12.86 std=2.02)
43		Missing

STR61		first Hct
N	Value	Description
392	Range	19 - 53.3 (median=39.3 mean=38.66 std=5.50)
60		Missing

STR62		first INR
N	Value	Description
282	Range	0.8 - 10.2 (median=1 mean=1.2 std=0.9)
170		Missing

STR63		first PTT
N	Value	Description
251	Range	10 - 81 (median=27 mean=29.4 std=7.6)
201		Missing

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STR63A		Activated PTT
N	Value	Description
201	.S	Legitimately skipped
72	0	0=No
135	1	1=Yes
44	9	9=NR

STR64		first platelet count
N	Value	Description
396	Range	7 - 833 (median=236 mean=247.3 std=95.3)
56		Missing

STR65		Total Cholesterol
N	Value	Description
148	Range	49 - 329 (median=169 mean=176.0 std=45.5)
304		Missing

STR66		LDL
N	Value	Description
148	Range	8 - 214 (median=96 mean=100.4 std=39.1)
304		Missing

STR67		HbA1c
N	Value	Description
124	Range	4.6 - 38 (median=6.95 mean=7.704 std=3.350)
328		Missing

STR68		receive thrombolytic treatment
N	Value	Description
395	0	0=No
14	1	1=Yes
43	9	9=NR

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STR68A		<i>symptomatic ICH post tPA</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
438	.S	Legitimately skipped
10	0	0=No
0	1	1=Yes
4	9	9=NR

STR69		<i>endovascular therapy for ischemic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
412	0	0=No
9	1	1=Yes
31	9	9=NR

STR69A		<i>symptomatic ICH post EVT?</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
443	.S	Legitimately skipped
5	0	0=No
0	1	1=Yes
4	9	9=NR

STR70		<i>osmotic/hypertonic treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
411	0	0=No
8	1	1=Yes
33	9	9=NR

STR71		<i>decompressive hemicraniectomy needed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
418	0	0=No
9	1	1=Yes
25	9	9=NR

STR72		<i>alive at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
31	0	0=No
421	1	1=Yes

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STR72A		<i>autopsy performed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
421	.S	Legitimately skipped
24	0	0=No
2	1	1=Yes
5	9	9=NR

STR72A1		<i>autopsy diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
450	.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
2		Missing

STR73		<i>ambulate independently before hospitalization</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
31	.S	Legitimately skipped
17	0	0=No
380	1	1=Yes
24	9	9=Unknown/NR

STR74		<i>ambulate independently at discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
31	.S	Legitimately skipped
59	0	0=No
279	1	1=Yes
83	9	9=Unknown/NR

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STR75		where discharged to
N	Value	Description
31	.S	Legitimately skipped
298	1	1=Home
43	2	2=Acute Rehabilitation
49	3	3=Skilled Nursing Facility
3	4	4=Assisted Living
4	5	5=Hospice
24	6	6=Other/Unknown

STR76		ADLs required
N	Value	Description
31	.S	Legitimately skipped
190	0	0=No
109	1	1=Yes
122	9	9=Unknown/NR

STR77		modified rankin scale done at discharge
N	Value	Description
31	.S	Legitimately skipped
410	0	0=No
4	1	1=Yes
7	9	9=Unknown/NR

STR77A		modified rankin scale score
N	Value	Description
448	.S	Legitimately skipped
4	0	0
0	1	1
0	2	2
0	3	3
0	4	4
0	5	5
0	6	6

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STR77B		discharged ADLs/walk capabilities
N	Value	Description
35	.S	Legitimately skipped
147	0	0=Back to Normal
54	1	1=Exhibiting S/S related to stroke but otherwise without disability
33	2	2=Able to ambulate independently with some disability
16	3	3=Dependent on others for some ADLs, but able to walk
22	4	4=Moderately severe disability
9	5	5=Bedridden (requires 24/7 care)
136	9	9=Unknown/NR

STR78A		prior to admission anticoagulants
N	Value	Description
432	0	0=No/NR
20	1	1=Yes

STR78B		prior to admission beta blockers
N	Value	Description
298	0	0=No/NR
154	1	1=Yes

STR78C		prior to admission CCB
N	Value	Description
356	0	0=No/NR
96	1	1=Yes

STR78D		prior to admission ACE or ARB
N	Value	Description
229	0	0=No/NR
223	1	1=Yes

STR78E		prior to admission scheduled aspirin
N	Value	Description
251	0	0=No/NR
201	1	1=Yes

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STR78F		prior to admission heparin
N	Value	Description
443	0	0=No/NR
9	1	1=Yes

STR78G		prior to admission Coumadin
N	Value	Description
425	0	0=No/NR
27	1	1=Yes

STR78H		prior to admission anti-platelet agent (non-aspirin)
N	Value	Description
383	0	0=No/NR
69	1	1=Yes

STR78I		prior to admission statin
N	Value	Description
240	0	0=No/NR
212	1	1=Yes

STR79A		during/after admission anticoagulants
N	Value	Description
429	0	0=No/NR
23	1	1=Yes

STR79B		during/after admission beta blockers
N	Value	Description
269	0	0=No/NR
183	1	1=Yes

STR79C		during/after admission CCB
N	Value	Description
296	0	0=No/NR
156	1	1=Yes

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STR79D		during/after admission ACE or ARB
N	Value	Description
242	0	0=No/NR
210	1	1=Yes

STR79E		during/after admission scheduled aspirin
N	Value	Description
187	0	0=No/NR
265	1	1=Yes

STR79F		during/after admission heparin
N	Value	Description
331	0	0=No/NR
121	1	1=Yes

STR79G		during/after admission Coumadin
N	Value	Description
413	0	0=No/NR
39	1	1=Yes

STR79H		during/after admission anti-platelet agent (non-aspirin)
N	Value	Description
344	0	0=No/NR
108	1	1=Yes

STR79I		during/after admission statin
N	Value	Description
179	0	0=No/NR
273	1	1=Yes

STR0A_FUTIME		Futime from Randomization date for (STR0A)
N	Value	Description
452	Range	3330 - 5109 (median=4031.5 mean=4017.32 std=358.94)

STR0D_FUTIME		Futime from Randomization date for (STR0D)
N	Value	Description
452	Range	14 - 3943 (median=1586.5 mean=1716.40 std=1063.56)

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STR1_FUTIME		Futime from Randomization date for (STR1)
N	Value	Description
452	Range	14 - 3943 (median=1586 mean=1715.5 std=1063.2)

STR3_FUTIME		Futime from Randomization date for (STR3)
N	Value	Description
451	Range	21 - 4366 (median=1604 mean=1733.9 std=1073.5)
1		Missing

STR11B_FUTIME		Futime from Randomization date for (STR11B)
N	Value	Description
324	Range	25 - 3939 (median=1559 mean=1700.9 std=1098.6)
128		Missing

STR12B_FUTIME		Futime from Randomization date for (STR12B)
N	Value	Description
323	Range	25 - 3939 (median=1565 mean=1707.4 std=1098.2)
129		Missing

STR13B_FUTIME		Futime from Randomization date for (STR13B)
N	Value	Description
30	Range	-539 - 2629 (median=986 mean=894.6 std=621.8)
422		Missing

STR14B_FUTIME		Futime from Randomization date for (STR14B)
N	Value	Description
3	Range	644 - 3480 (median=1185 mean=1769.7 std=1505.7)
449		Missing

STR51A_FUTIME		Futime from Randomization date for (STR51A)
N	Value	Description
327	Range	36 - 3939 (median=1597 mean=1728.1 std=1080.5)
125		Missing

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STR51C_FUTIME		Futime from Randomization date for (STR51C)
N	Value	Description
99	Range	52 - 3861 (median=1325 mean=1562.0 std=1042.1)
353		Missing

STR53A_FUTIME		Futime from Randomization date for (STR53A)
N	Value	Description
240	Range	52 - 3943 (median=1668 mean=1767.5 std=1094.9)
212		Missing

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Stroke Reviewer 1 Form: FORM CODE=STD1, VERSION=1

ID		HCHS Participant ID
N	Value	Description
450	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
331	1	Occurrence 1
79	2	Occurrence 2
22	3	Occurrence 3
6	4	Occurrence 4
8	5	Occurrence 5
2	6	Occurrence 6
2	7	Occurrence 7

STD11		primary diagnosis
N	Value	Description
206	1	1 = Not a TIA or Stroke
34	2	2 = TIA
188	3	3 = Stroke
22		Missing

STD12		stroke type
N	Value	Description
240	.S	Legitimately skipped
137	1	1 = Brain Infarction
18	2	2 = Subarachnoid Hemorrhage
25	3	3 = Intracerebral Hemorrhage
4	4	4 = Other Stroke Type
4	5	5 = Unknown Stroke Type
22		Missing

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STD12A		other stroke type specify
N	Value	Description
4	Present	Text suppressed
424	~	Legitimately skipped
22		Missing

STD12B		unknown stroke type specify
N	Value	Description
4	Present	Text suppressed
424	~	Legitimately skipped
22		Missing

STD13		brain infarct subtype
N	Value	Description
287	.S	Legitimately skipped
3	0	0 = N/A (not applicable)
7	1	1 = Large vessel extracranial atheroembolic
5	2	2 = Large vessel intracranial atheroembolic
19	3	3 = Cardioembolic
34	4	4 = Lacunar Infarction
8	5	5 = Acute ischemic stroke of other known etiology
18	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)
38	8	8 = Ischemic stroke of unknown cause (workup is incomplete)
22		Missing

STD13A		acute ischemic other known specify
N	Value	Description
8	Present	Text suppressed
420	~	Legitimately skipped
22		Missing

STD13B		ischemic unknown cause
N	Value	Description
9	Present	Text suppressed
419	~	Legitimately skipped
22		Missing

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STD14		Was the Brain Infarction Procedure-Related?
N	Value	Description
290	.S	Legitimately skipped
129	0	0=No
7	1	1=Yes
2	9	9=Unknown
22		Missing

STD14A		specify procedure related infarction
N	Value	Description
7	Present	Text suppressed
421	~	Legitimately skipped
22		Missing

STD15		subarachnoid hemorrhage subtype
N	Value	Description
406	.S	Legitimately skipped
3	0	0=N/A (not applicable)
9	1	1=Aneurysmal
3	2	2=Sulcal
2	3	3=Perimesencephalic
3	4	4=Other
2	9	9=Unknown
22		Missing

STD15A		subarachnoid subtype other specify
N	Value	Description
3	Present	Text suppressed
425	~	Legitimately skipped
22		Missing

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STD16		subarachnoid intraventricular extension
N	Value	Description
409	.S	Legitimately skipped
13	0	0=No
5	1	1=Yes
1	9	9=Unknown
22		Missing

STD17		subarachnoid intraparenchymal extension
N	Value	Description
409	.S	Legitimately skipped
14	0	0=No
5	1	1=Yes
0	9	9=Unknown
22		Missing

STD18		Intracerebral Hemorrhage Subtype
N	Value	Description
399	.S	Legitimately skipped
2	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
2	6	6=Primary Intraventricular Hemorrhage
5	7	7=Other
4	9	9=Unknown
22		Missing

STD18A		intracerebral Hemorrhage Subtype other specify
N	Value	Description
5	Present	Text suppressed
423	~	Legitimately skipped
22		Missing

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STD19		<i>intracerebral intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
401	.S	Legitimately skipped
19	0	0=No
7	1	1=Yes
1	9	9=Unknown
22		Missing

STD110		<i>core of hemorrhage description</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
401	.S	Legitimately skipped
7	1	1=Lobar
16	2	2=Deep
2	3	3=Cerebellar
2	9	9=Unknown
22		Missing

STD111		<i>hemorrhagic procedure-related</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
381	.S	Legitimately skipped
45	0	0=No or N/A
1	1	1=Yes
1	9	9=Unknown
22		Missing

STD111A		<i>hemorrhagic procedure-related specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Present	Text suppressed
427	~	Legitimately skipped
22		Missing

STD112		<i>overall reviewer comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
258	Present	Text suppressed
192		Missing

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STD10A_FUTIME		Futime from Randomization date for (STD10A)
N	Value	Description
428	Range	3434 - 5011 (median=4208 mean=4190.1 std=336.5)
22		Missing

STD10D_FUTIME		Futime from Randomization date for (STD10D)
N	Value	Description
450	Range	14 - 3943 (median=1586.5 mean=1712.91 std=1062.81)

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Section E: Stroke Ascertainment

Stroke Reviewer 2 Form: FORM CODE=STD2, VERSION=1

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

<i>OCCURRENCE</i>		<i>Occurrence</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
331	1	Occurrence 1
79	2	Occurrence 2
22	3	Occurrence 3
6	4	Occurrence 4
8	5	Occurrence 5
2	6	Occurrence 6
2	7	Occurrence 7

<i>STD21</i>		<i>primary diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
205	1	1 = Not a TIA or Stroke
41	2	2 = TIA
182	3	3 = Stroke
22		Missing

<i>STD22</i>		<i>stroke type</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
246	.S	Legitimately skipped
129	1	1 = Brain Infarction
20	2	2 = Subarachnoid Hemorrhage
26	3	3 = Intracerebral Hemorrhage
3	4	4 = Other Stroke Type
4	5	5 = Unknown Stroke Type
22		Missing

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STD22A		other stroke type specify
N	Value	Description
3	Present	Text suppressed
425	~	Legitimately skipped
22		Missing

STD22B		unknown stroke type specify
N	Value	Description
4	Present	Text suppressed
424	~	Legitimately skipped
22		Missing

STD23		brain infarct subtype
N	Value	Description
296	.S	Legitimately skipped
1	0	0 = N/A (not applicable)
5	1	1 = Large vessel extracranial atheroembolic
9	2	2 = Large vessel intracranial atheroembolic
18	3	3 = Cardioembolic
36	4	4 = Lacunar Infarction
5	5	5 = Acute ischemic stroke of other known etiology
20	6	6 = Ischemic stroke of unknown cause (no probable etiology despite complete workup)
5	7	7 = Ischemic stroke of unknown cause (more than one etiology)
33	8	8 = Ischemic stroke of unknown cause (workup is incomplete)
22		Missing

STD23A		acute ischemic other known specify
N	Value	Description
5	Present	Text suppressed
423	~	Legitimately skipped
22		Missing

STD23B		ischemic unknown cause
N	Value	Description
5	Present	Text suppressed
423	~	Legitimately skipped
22		Missing

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STD24		Was the Brain Infarction Procedure-Related?
N	Value	Description
297	.S	Legitimately skipped
122	0	0=No
7	1	1=Yes
2	9	9=Unknown
22		Missing

STD24A		specify procedure related infarction
N	Value	Description
7	Present	Text suppressed
421	~	Legitimately skipped
22		Missing

STD25		subarachnoid hemorrhage subtype
N	Value	Description
405	.S	Legitimately skipped
3	0	0=N/A (not applicable)
9	1	1=Aneurysmal
4	2	2=Sulcal
2	3	3=Perimesencephalic
3	4	4=Other
2	9	9=Unknown
22		Missing

STD25A		subarachnoid subtype other specify
N	Value	Description
3	Present	Text suppressed
425	~	Legitimately skipped
22		Missing

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Section E: Stroke Ascertainment

STD26		subarachnoid intraventricular extension
N	Value	Description
408	.S	Legitimately skipped
11	0	0=No
5	1	1=Yes
4	9	9=Unknown
22		Missing

STD27		subarachnoid intraparenchymal extension
N	Value	Description
408	.S	Legitimately skipped
13	0	0=No
4	1	1=Yes
3	9	9=Unknown
22		Missing

STD28		Intracerebral Hemorrhage Subtype
N	Value	Description
399	.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
5	9	9=Unknown
22		Missing

STD28A		intracerebral Hemorrhage Subtype other specify
N	Value	Description
2	Present	Text suppressed
426	~	Legitimately skipped
22		Missing

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STD29		<i>intracerebral intraventricular extension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
17	0	0=No
9	1	1=Yes
0	9	9=Unknown
22		Missing

STD210		<i>core of hemorrhage description</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
402	.S	Legitimately skipped
9	1	1=Lobar
14	2	2=Deep
3	3	3=Cerebellar
0	9	9=Unknown
22		Missing

STD211		<i>hemorrhagic procedure-related</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
379	.S	Legitimately skipped
47	0	0=No or N/A
1	1	1=Yes
1	9	9=Unknown
22		Missing

STD211A		<i>hemorrhagic procedure-related specify</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Present	Text suppressed
427	~	Legitimately skipped
22		Missing

STD212		<i>overall reviewer comments</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
230	Present	Text suppressed
220		Missing

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STD20A_FUTIME		Futime from Randomization date for (STD20A)
N	Value	Description
428	Range	3425 - 5098 (median=4200.5 mean=4173.88 std=342.56)
22		Missing

STD20D_FUTIME		Futime from Randomization date for (STD20D)
N	Value	Description
450	Range	14 - 3943 (median=1586.5 mean=1712.91 std=1062.81)

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Stroke Adjudication Reviewer Form: FORM CODE=STD3, VERSION=1

ID		HCHS Participant ID
N	Value	Description
66	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
40	1	Occurrence 1
21	2	Occurrence 2
5	3	Occurrence 3

STD31		primary diagnosis
N	Value	Description
34	1	1 = Not a TIA or Stroke
11	2	2 = TIA
16	3	3 = Stroke
5		Missing

STD32		stroke type
N	Value	Description
45	.S	Legitimately skipped
9	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
5		Missing

STD32A		other stroke type specify
N	Value	Description
2	Present	Text suppressed
59	~	Legitimately skipped
5		Missing

Chapter 1: Endpoint Ascertainment

Section E: Stroke Ascertainment

STD32B		unknown stroke type specify
N	Value	Description
1	Present	Text suppressed
60	~	Legitimately skipped
5		Missing

STD33		brain infarct subtype
N	Value	Description
50	.S	Legitimately skipped
2	0	0 = N/A (not applicable)
1	1	1 = Large vessel extracranial atheroembolic
0	2	2 = Large vessel intracranial atheroembolic
1	3	3 = Cardioembolic
3	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)
0	7	7 = Ischemic stroke of unknown cause (more than one etiology)
3	8	8 = Ischemic stroke of unknown cause (workup is incomplete)
5		Missing

STD33A		acute ischemic other known specify
N	Value	Description
61	~	Legitimately skipped
5		Missing

STD33B		ischemic unknown cause
N	Value	Description
61	~	Legitimately skipped
5		Missing

STD34		Was the Brain Infarction Procedure-Related?
N	Value	Description
52	.S	Legitimately skipped
8	0	0=No
0	1	1=Yes
1	9	9=Unknown
5		Missing

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Section E: Stroke Ascertainment

STD34A		specify procedure related infarction
N	Value	Description
61	~	Legitimately skipped
5		Missing

STD35		subarachnoid hemorrhage subtype
N	Value	Description
57	.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
5		Missing

STD35A		subarachnoid subtype other specify
N	Value	Description
61	~	Legitimately skipped
5		Missing

STD36		subarachnoid intraventricular extension
N	Value	Description
58	.S	Legitimately skipped
3	0	0=No
0	1	1=Yes
0	9	9=Unknown
5		Missing

STD37		subarachnoid intraparenchymal extension
N	Value	Description
58	.S	Legitimately skipped
1	0	0=No
2	1	1=Yes
0	9	9=Unknown
5		Missing

Chapter 1: Endpoint Ascertainment

Section E: Stroke Ascertainment

STD38		Intracerebral Hemorrhage Subtype
N	Value	Description
57	.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 Cavernous
0	5	5=Endocarditis
0	6	6=Primary Intraventricular Hemorrhage
1	7	7=Other
1	9	9=Unknown
5		Missing

STD38A		intracerebral Hemorrhage Subtype other specify
N	Value	Description
1	Present	Text suppressed
60	~	Legitimately skipped
5		Missing

STD39		intracerebral intraventricular extension
N	Value	Description
58	.S	Legitimately skipped
2	0	0=No
1	1	1=Yes
0	9	9=Unknown
5		Missing

STD310		core of hemorrhage description
N	Value	Description
58	.S	Legitimately skipped
2	1	1=Lobar
0	2	2=Deep
1	3	3=Cerebellar
0	9	9=Unknown
5		Missing

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Section E: Stroke Ascertainment

STD311		hemorrhagic procedure-related
N	Value	Description
55	.S	Legitimately skipped
5	0	0=No or N/A
0	1	1=Yes
1	9	9=Unknown
5		Missing

STD311A		hemorrhagic procedure-related specify
N	Value	Description
61	~	Legitimately skipped
5		Missing

STD312		overall reviewer comments
N	Value	Description
46	Present	Text suppressed
20		Missing

STD30A_FUTIME		Futime from Randomization date for (STD30A)
N	Value	Description
61	Range	3759 - 5037 (median=4405 mean=4391.4 std=357.9)
5		Missing

STD30D_FUTIME		Futime from Randomization date for (STD30D)
N	Value	Description
66	Range	81 - 3876 (median=1298 mean=1454.6 std=932.3)

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Section F: Pregnancy Complication Ascertainment

Pregnancy Complication Abstraction Form: FORM CODE=PRC, VERSION=1

ID		HCHS Participant ID
N	Value	Description
127	Present	Text suppressed

OCCURRENCE		Occurrence
N	Value	Description
109	1	Occurrence 1
14	2	Occurrence 2
3	3	Occurrence 3
1	4	Occurrence 4

PRC1		presenting complaint
N	Value	Description
126	Range	1 - 6 (median=1 mean=1.7 std=1.6)
1		Missing

PRC2		24 hour stay
N	Value	Description
127	Range	0 - 0 (median=0 mean=0.0 std=0.0)

PRC3A		Time of arrival
N	Value	Description
110	Range	0:37:00 - 23:31:00
17		Missing

PRC3AHR		Hour of Time of arrival
N	Value	Description
110	Range	0 - 23 (median=11 mean=11.5 std=5.7)
17		Missing

PRC3AMN		Minute of Time of arrival
N	Value	Description
110	Range	0 - 57 (median=28.5 mean=26.29 std=16.62)
17		Missing

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<i>PRC3ASC</i>		<i>Second of Time of arrival</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 0 (median=0 mean=0.0 std=0.0)
52		Missing

<i>PRC4A</i>		<i>Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	1:00:00 - 23:59:00
55		Missing

<i>PRC4AHR</i>		<i>Hour of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	1 - 23 (median=15 mean=15.0 std=4.1)
55		Missing

<i>PRC4AMN</i>		<i>Minute of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 59 (median=30 mean=27.2 std=18.4)
55		Missing

<i>PRC4ASC</i>		<i>Second of Time of discharge</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
50	Range	0 - 0 (median=0 mean=0.0 std=0.0)
77		Missing

<i>PRC5</i>		<i>Admitting diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
102	Present	Text suppressed
25		Missing

<i>PRC6</i>		<i>Discharge diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
109	Present	Text suppressed
18		Missing

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Section F: Pregnancy Complication Ascertainment

<i>PRC7</i>		<i>patient referred</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	0 - 1 (median=0 mean=0.2 std=0.4)

<i>PRC8</i>		<i>patient transferred</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	0 - 1 (median=0 mean=0.0 std=0.1)

<i>PRC9</i>		<i>patient alive</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	1 - 1 (median=1 mean=1.0 std=0.0)

<i>PRC11</i>		<i>EDC method</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Range	1 - 9 (median=9 mean=6.4 std=3.2)

<i>PRC12A</i>		<i>headache</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

<i>PRC12B</i>		<i>blurry vision</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC12C</i>		<i>right upper quadrant pain or epigastric pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC12D</i>		<i>nausea, vomiting</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.1)
1		Missing

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<i>PRC12E</i>		<i>vaginal bleeding</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC12F</i>		<i>seizure</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 0 (median=0 mean=0.0 std=0.0)
1		Missing

<i>PRC13A</i>		<i>diabetes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

<i>PRC13B</i>		<i>gestational diabetes in prior pregnancies</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

<i>PRC13C</i>		<i>gestational diabetes in current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=1 mean=0.5 std=0.5)
1		Missing

<i>PRC13D</i>		<i>hypertension prior to pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.2 std=0.4)
1		Missing

<i>PRC13E</i>		<i>gestational hypertension in prior pregnancies</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

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Section F: Pregnancy Complication Ascertainment

<i>PRC13F</i>		<i>gestational hypertension in current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.3 std=0.5)
1		Missing

<i>PRC13G</i>		<i>preeclampsia in prior pregnancies</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC13H</i>		<i>preeclampsia in current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.2 std=0.4)
1		Missing

<i>PRC13I</i>		<i>thyroid disease</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

<i>PRC13J</i>		<i>smoking prior to current pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC13K</i>		<i>smoking during pregnancy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC13L</i>		<i>known seizure disorder</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.1)
1		Missing

Chapter 1: Endpoint Ascertainment

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<i>PRC14</i>		<i>qualifying BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.3 std=0.5)
1		Missing

<i>PRC14A2</i>		<i>Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24	Range	1:02:00 - 20:53:00
85	.S	Legitimately skipped
18		Missing

<i>PRC14A2HR</i>		<i>Hour of Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24	Range	1 - 20 (median=15 mean=13.3 std=5.2)
103		Missing

<i>PRC14A2MN</i>		<i>Minute of Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24	Range	0 - 53 (median=30.5 mean=27.79 std=18.97)
103		Missing

<i>PRC14A2SC</i>		<i>Second of Initial BP time</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15	Range	0 - 0 (median=0 mean=0.0 std=0.0)
112		Missing

<i>PRC14A3</i>		<i>initial systolic BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	120 - 200 (median=150 mean=155.0 std=16.6)
85	.S	Legitimately skipped

<i>PRC14A4</i>		<i>initial diastolic BP</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
34	Range	72 - 116 (median=93 mean=92.8 std=11.0)
85	.S	Legitimately skipped
8		Missing

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PRC14B2		confirming BP time
N	Value	Description
10	Range	1:51:00 - 21:55:00
85	.S	Legitimately skipped
32		Missing

PRC14B2HR		Hour of confirming BP time
N	Value	Description
10	Range	1 - 21 (median=15.5 mean=13.70 std=7.73)
117		Missing

PRC14B2MN		Minute of confirming BP time
N	Value	Description
10	Range	0 - 55 (median=22.5 mean=23.00 std=19.84)
117		Missing

PRC14B2SC		Second of confirming BP time
N	Value	Description
7	Range	0 - 0 (median=0 mean=0.0 std=0.0)
120		Missing

PRC14B3		confirming systolic BP
N	Value	Description
18	Range	140 - 186 (median=149.5 mean=153.72 std=12.02)
85	.S	Legitimately skipped
24		Missing

PRC14B4		confirming diastolic BP
N	Value	Description
16	Range	58 - 120 (median=92 mean=92.6 std=14.9)
85	.S	Legitimately skipped
26		Missing

PRC15		anti-hypertensive meds
N	Value	Description
126	Range	0 - 9 (median=9 mean=7.1 std=3.6)
1		Missing

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<i>PRC16</i>		<i>vaginal exam</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 9 (median=1 mean=3.9 std=3.9)
1		Missing

<i>PRC16A</i>		<i>dilation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 10 (median=3 mean=3.2 std=2.9)
53	.S	Legitimately skipped
2		Missing

<i>PRC16B</i>		<i>effacement</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
62	Range	0 - 100 (median=65 mean=56.7 std=36.8)
53	.S	Legitimately skipped
12		Missing

<i>PRC17</i>		<i>prenatal records</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=1 mean=0.6 std=0.5)
1		Missing

<i>PRC17A</i>		<i>baseline urine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 1 (median=0 mean=0.1 std=0.3)
52	.S	Legitimately skipped

<i>PRC17A2</i>		<i>total urine volume</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	2375 - 2375 (median=2375 mean=2375.0 std=)
121	.S	Legitimately skipped
5		Missing

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<i>PRC17A3</i>		<i>total protein</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5	Range	168 - 432 (median=349 mean=325.4 std=97.7)
121	.S	Legitimately skipped
1		Missing

<i>PRC17A3A</i>		<i>protein units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	1 - 2 (median=1 mean=1.3 std=0.5)
121	.S	Legitimately skipped
2		Missing

<i>PRC17B</i>		<i>glucose loading test</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 1 (median=0 mean=0.3 std=0.5)
52	.S	Legitimately skipped

<i>PRC17B1</i>		<i>glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
19	Range	50 - 220 (median=148 mean=143.8 std=46.4)
103	.S	Legitimately skipped
5		Missing

<i>PRC17C</i>		<i>type of OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 3 (median=0 mean=0.8 std=1.3)
52	.S	Legitimately skipped

<i>PRC17C1A</i>		<i>OGTT discrete values</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
22	Range	0 - 1 (median=1 mean=0.7 std=0.5)
105	.S	Legitimately skipped

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<i>PRC17C1B</i>		<i>fasting glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	84 - 150 (median=95 mean=96.8 std=16.7)
111	.S	Legitimately skipped
3		Missing

<i>PRC17C1C</i>		<i>1 hr glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16	Range	103 - 270 (median=183.5 mean=177.94 std=43.04)
111	.S	Legitimately skipped

<i>PRC17C1E</i>		<i>3 hr glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14	Range	83 - 176 (median=128 mean=124.3 std=29.8)
111	.S	Legitimately skipped
2		Missing

<i>PRC17C2</i>		<i>OGTT number of values reported</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6	Range	0 - 0 (median=0 mean=0.0 std=0.0)
121	.S	Legitimately skipped

<i>PRC17C2A</i>		<i>number of abnormal values</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	.S	Legitimately skipped

<i>PRC17C3</i>		<i>OGTT documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6	Range	1 - 2 (median=1 mean=1.3 std=0.5)
121	.S	Legitimately skipped

<i>PRC17D</i>		<i>prenatal Hgb A1c</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 1 (median=0 mean=0.2 std=0.4)
52	.S	Legitimately skipped

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PRC17D1A		Hgb A1c value less than or equal to 20
N	Value	Description
7	Range	5.2 - 9.3 (median=5.8 mean=6.41 std=1.41)
114	.S	Legitimately skipped
6		Missing

PRC17D2A		Hgb A1c greater than 20 value
N	Value	Description
9	Range	5 - 8.4 (median=5.5 mean=5.71 std=1.03)
114	.S	Legitimately skipped
4		Missing

PRC17E		outpatient pressure
N	Value	Description
75	Range	0 - 1 (median=0 mean=0.4 std=0.5)
52	.S	Legitimately skipped

PRC17E1A		highest systolic less than or equal to 20 value
N	Value	Description
8	Range	100 - 143 (median=135 mean=126.1 std=17.9)
95	.S	Legitimately skipped
24		Missing

PRC17E2A		highest systolic greater than 20 value
N	Value	Description
28	Range	108 - 160 (median=128 mean=130.7 std=13.9)
95	.S	Legitimately skipped
4		Missing

PRC17E3A		highest diastolic less than or equal to 20 date
N	Value	Description
8	Range	70 - 90 (median=76.5 mean=77.50 std=6.95)
95	.S	Legitimately skipped
24		Missing

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PRC17E4A		highest diastolic greater than 20 date
N	Value	Description
28	Range	54 - 100 (median=77.5 mean=78.46 std=10.19)
95	.S	Legitimately skipped
4		Missing

PRC17F		weight gain available
N	Value	Description
75	Range	0 - 1 (median=0 mean=0.4 std=0.5)
52	.S	Legitimately skipped

PRC17F1		pre weight
N	Value	Description
11	Range	72.7 - 230 (median=96.3 mean=120.01 std=51.27)
94	.S	Legitimately skipped
22		Missing

PRC17F1A		pre weight units
N	Value	Description
17	Range	1 - 9 (median=2 mean=4.1 std=3.8)
94	.S	Legitimately skipped
16		Missing

PRC17F2		first weight
N	Value	Description
5	Range	63.1 - 173.5 (median=151 mean=135.5 std=43.3)
94	.S	Legitimately skipped
28		Missing

PRC17F2A		first weight units
N	Value	Description
10	Range	1 - 9 (median=5.5 mean=5.40 std=3.81)
94	.S	Legitimately skipped
23		Missing

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PRC17F3		<i>last weight</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	62.6 - 237 (median=120.6 mean=135.90 std=54.40)
94	.S	Legitimately skipped
8		Missing

PRC17F3A		<i>last weight units</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	1 - 2 (median=1 mean=1.4 std=0.5)
94	.S	Legitimately skipped
8		Missing

PRC17G		<i>outpatient diabetes tx</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 1 (median=1 mean=0.6 std=0.5)
52	.S	Legitimately skipped

PRC17G1		<i>outpatient insulin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	0 - 1 (median=0 mean=0.4 std=0.5)
85	.S	Legitimately skipped

PRC17G2		<i>outpatient glyburide</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	0 - 1 (median=0 mean=0.4 std=0.5)
85	.S	Legitimately skipped

PRC17G3		<i>outpatient metformin</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	0 - 1 (median=0 mean=0.1 std=0.4)
85	.S	Legitimately skipped

PRC17G4		<i>outpatient other med</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	0 - 1 (median=0 mean=0.2 std=0.4)
85	.S	Legitimately skipped

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PRC17G4A		outpatient med name
N	Value	Description
7	Present	Text suppressed
120	~	Legitimately skipped

PRC18A		12 or 24 hour urine test
N	Value	Description
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

PRC18A2		urine total volume
N	Value	Description
1	Range	2375 - 2375 (median=2375 mean=2375.0 std=)
115	.S	Legitimately skipped
11		Missing

PRC18A3		urine total protein
N	Value	Description
10	Range	0 - 2142 (median=356 mean=599.0 std=622.0)
115	.S	Legitimately skipped
2		Missing

PRC18A4		24 hour creatinine
N	Value	Description
1	Range	2.1 - 2.1 (median=2.1 mean=2.10 std=)
115	.S	Legitimately skipped
11		Missing

PRC18A5		urine duration
N	Value	Description
10	Range	2 - 2 (median=2 mean=2.0 std=0.0)
115	.S	Legitimately skipped
2		Missing

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<i>PRC18B</i>		<i>last urine protein</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.2)
1		Missing

<i>PRC18B2</i>		<i>last urine protein value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	Range	0.43 - 1211 (median=0.715 mean=200.7238 std=430.3331)
119	.S	Legitimately skipped

<i>PRC19A</i>		<i>platelet count</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=1 mean=0.7 std=0.5)
1		Missing

<i>PRC19A1</i>		<i>lowest platelet value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
85	Range	104 - 493 (median=198 mean=214.1 std=67.9)
42	.S	Legitimately skipped

<i>PRC19B</i>		<i>AST</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.3 std=0.5)
1		Missing

<i>PRC19B1</i>		<i>highest AST value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	14 - 408 (median=23.5 mean=47.60 std=80.82)
85	.S	Legitimately skipped

<i>PRC19B2</i>		<i>lab limit SGOT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	0 - 1 (median=1 mean=0.8 std=0.4)
85	.S	Legitimately skipped

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<i>PRC19B2A</i>		<i>Lab limit SGOT value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
32	Range	32 - 60 (median=41 mean=41.3 std=6.7)
95	.S	Legitimately skipped

<i>PRC19C</i>		<i>ALT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.3 std=0.5)
1		Missing

<i>PRC19C1</i>		<i>highest ALT value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	7 - 402 (median=19.5 mean=40.71 std=71.39)
85	.S	Legitimately skipped

<i>PRC19C2</i>		<i>lab limit ALT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	0 - 1 (median=1 mean=0.8 std=0.4)
85	.S	Legitimately skipped

<i>PRC19C2A</i>		<i>lab limit ALT value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
32	Range	24 - 78 (median=52 mean=51.0 std=10.2)
95	.S	Legitimately skipped

<i>PRC19D</i>		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.3 std=0.5)
1		Missing

<i>PRC19D1</i>		<i>highest serum creatinine value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
43	Range	0.39 - 2.13 (median=0.56 mean=0.618 std=0.271)
84	.S	Legitimately skipped

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<i>PRC19E</i>		<i>uric acid</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.2 std=0.4)
1		Missing

<i>PRC19E1</i>		<i>highest acid value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	2.9 - 10.6 (median=4.9 mean=5.13 std=1.51)
98	.S	Legitimately skipped

<i>PRC19F</i>		<i>LDH</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.2 std=0.4)
1		Missing

<i>PRC19F1</i>		<i>highest LDH value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
22	Range	147 - 792 (median=212.5 mean=279.50 std=167.45)
105	.S	Legitimately skipped

<i>PRC19G</i>		<i>admission hematocrit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=1 mean=0.6 std=0.5)
1		Missing

<i>PRC19G1</i>		<i>admission hematocrit value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
77	Range	25.2 - 44.8 (median=35.8 mean=35.11 std=3.83)
50	.S	Legitimately skipped

<i>PRC19H</i>		<i>post delivery hematocrit</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=1 mean=0.6 std=0.5)
1		Missing

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<i>PRC19H1</i>		<i>lowest hematocrit value</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
80	Range	19.8 - 39.6 (median=30.9 mean=30.71 std=4.19)
47	.S	Legitimately skipped

<i>PRC19I</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 0 (median=0 mean=0.0 std=0.0)
1		Missing

<i>PRC20A</i>		<i>magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	1 - 9 (median=9 mean=7.9 std=2.5)
1		Missing

<i>PRC20B</i>		<i>insulin (subcutaneous)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.2 std=0.4)
1		Missing

<i>PRC20C</i>		<i>insulin (IV or drip)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.1)
1		Missing

<i>PRC20D</i>		<i>antihypertensives IV</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC20E</i>		<i>antihypertensives PO</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

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<i>PRC20F</i>		<i>betamethason/dexamethasone IM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.1 std=0.3)
1		Missing

<i>PRC20F1</i>		<i>prematurity</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16	Range	0 - 1 (median=1 mean=0.8 std=0.4)
111	.S	Legitimately skipped

<i>PRC20G</i>		<i>tocolysis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=0 mean=0.0 std=0.2)
1		Missing

<i>PRC21</i>		<i>patient deliver</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
126	Range	0 - 1 (median=1 mean=0.9 std=0.3)
1		Missing

<i>PRC21B</i>		<i>Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0:02:00 - 23:48:00
10	.S	Legitimately skipped
42		Missing

<i>PRC21BHR</i>		<i>Hour of Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 23 (median=11 mean=11.4 std=6.7)
52		Missing

<i>PRC21BMN</i>		<i>Minute of Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
75	Range	0 - 59 (median=28 mean=28.4 std=19.0)
52		Missing

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<i>PRC21BSC</i>		<i>Second of Time of delivery</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
50	Range	0 - 0 (median=0 mean=0.0 std=0.0)
77		Missing

<i>PRC22A</i>		<i>gestional weeks</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
114	Range	27 - 41 (median=39 mean=37.8 std=2.5)
10	.S	Legitimately skipped
3		Missing

<i>PRC22B</i>		<i>gestional day</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
112	Range	0 - 7 (median=3 mean=2.7 std=2.2)
10	.S	Legitimately skipped
5		Missing

<i>PRC23</i>		<i>type of labor</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
116	Range	1 - 9 (median=2 mean=2.5 std=1.5)
10	.S	Legitimately skipped
1		Missing

<i>PRC24</i>		<i>scheduled delivery reason</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
77	Range	1 - 17 (median=9 mean=10.6 std=4.2)
50	.S	Legitimately skipped

<i>PRC24A</i>		<i>scheduled delivery other</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	Present	Text suppressed
119	~	Legitimately skipped

<i>PRC25</i>		<i>c-section indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	2 - 17 (median=17 mean=13.7 std=4.9)
10	.S	Legitimately skipped

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<i>PRC25A</i>		<i>c-section specify reason</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Present	Text suppressed
123	~	Legitimately skipped

<i>PRC26</i>		<i>skin incision</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
56	Range	1 - 9 (median=2 mean=4.9 std=3.5)
71	.S	Legitimately skipped

<i>PRC27</i>		<i>uterine incision</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
56	Range	2 - 9 (median=3 mean=4.0 std=2.3)
71	.S	Legitimately skipped

<i>PRC28A</i>		<i>placental abruption</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

<i>PRC28B</i>		<i>postpartum hemorrhage</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.2)
10	.S	Legitimately skipped

<i>PRC28C</i>		<i>third degree laceration</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

<i>PRC28D</i>		<i>fourth degree laceration</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 0 (median=0 mean=0.0 std=0.0)
10	.S	Legitimately skipped

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<i>PRC28E</i>		<i>injury to viscera</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 0 (median=0 mean=0.0 std=0.0)
10	.S	Legitimately skipped

<i>PRC28F</i>		<i>intubation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 0 (median=0 mean=0.0 std=0.0)
10	.S	Legitimately skipped

<i>PRC28G</i>		<i>ICU admission</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

<i>PRC28H</i>		<i>transfusion</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.2)
10	.S	Legitimately skipped

<i>PRC28H1</i>		<i>platelets</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

<i>PRC28H2</i>		<i>cryoprecipitate</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 0 (median=0 mean=0.0 std=0.0)
10	.S	Legitimately skipped

<i>PRC28H3</i>		<i>frozen plasma</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

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<i>PRC28I</i>		<i>tubal ligation</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.2 std=0.4)
10	.S	Legitimately skipped

<i>PRC28J</i>		<i>dilation and curettage</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 0 (median=0 mean=0.0 std=0.0)
10	.S	Legitimately skipped

<i>PRC28K</i>		<i>hysterectomy</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

<i>PRC28L</i>		<i>intrapartum IV/IM narcotic</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

<i>PRC28M</i>		<i>intrapartum regional epidural</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.4 std=0.5)
10	.S	Legitimately skipped

<i>PRC28N</i>		<i>intrapartum regional spinal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.4 std=0.5)
10	.S	Legitimately skipped

<i>PRC28O</i>		<i>intrapartum general anesthesia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	0 - 1 (median=0 mean=0.0 std=0.1)
10	.S	Legitimately skipped

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<i>PRC29</i>		<i>infants born</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	1 - 9 (median=1 mean=1.2 std=1.1)
10	.S	Legitimately skipped

<i>PRC30A1</i>		<i>grams 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
73	Range	810 - 4730 (median=3255 mean=3161.7 std=664.1)
12	.S	Legitimately skipped
42		Missing

<i>PRC30A2</i>		<i>pounds 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
22	Range	3 - 10 (median=7 mean=6.7 std=1.8)
12	.S	Legitimately skipped
93		Missing

<i>PRC30A3</i>		<i>ounces 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
20	Range	1 - 15 (median=11 mean=10.5 std=3.8)
12	.S	Legitimately skipped
95		Missing

<i>PRC30B</i>		<i>one min apgar 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
92	Range	0 - 9 (median=9 mean=8.1 std=1.6)
12	.S	Legitimately skipped
23		Missing

<i>PRC30C</i>		<i>five min apgar 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
92	Range	0 - 9 (median=9 mean=8.8 std=1.0)
12	.S	Legitimately skipped
23		Missing

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<i>PRC30D</i>		<i>shoulder dystocia 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
112	Range	0 - 9 (median=9 mean=7.2 std=3.6)
12	.S	Legitimately skipped
3		Missing

<i>PRC30E</i>		<i>birth trauma 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
114	Range	0 - 9 (median=9 mean=8.0 std=2.9)
12	.S	Legitimately skipped
1		Missing

<i>PRC30F</i>		<i>stillbirth 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
114	Range	0 - 9 (median=0 mean=0.2 std=1.2)
12	.S	Legitimately skipped
1		Missing

<i>PRC30G</i>		<i>type of delivery 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
114	Range	1 - 9 (median=5 mean=3.3 std=2.1)
12	.S	Legitimately skipped
1		Missing

<i>PRC30H</i>		<i>infant sex 1</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
114	Range	1 - 9 (median=2 mean=3.1 std=3.0)
12	.S	Legitimately skipped
1		Missing

<i>PRC31A1</i>		<i>grams 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	1320 - 1320 (median=1320 mean=1320.0 std=)
124	.S	Legitimately skipped
2		Missing

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PRC31A2		pounds 2
N	Value	Description
124	.S	Legitimately skipped
3		Missing

PRC31A3		ounces 2
N	Value	Description
124	.S	Legitimately skipped
3		Missing

PRC31B		one min apgar 2
N	Value	Description
2	Range	5 - 6 (median=5.5 mean=5.50 std=0.71)
124	.S	Legitimately skipped
1		Missing

PRC31C		five min apgar 2
N	Value	Description
2	Range	8 - 8 (median=8 mean=8.0 std=0.0)
124	.S	Legitimately skipped
1		Missing

PRC31D		shoulder dystocia 2
N	Value	Description
3	Range	0 - 9 (median=9 mean=6.0 std=5.2)
124	.S	Legitimately skipped

PRC31E		birth trauma 2
N	Value	Description
3	Range	9 - 9 (median=9 mean=9.0 std=0.0)
124	.S	Legitimately skipped

PRC31F		stillbirth 2
N	Value	Description
3	Range	0 - 0 (median=0 mean=0.0 std=0.0)
124	.S	Legitimately skipped

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PRC31G		type of delivery 2
N	Value	Description
3	Range	1 - 5 (median=5 mean=3.7 std=2.3)
124	.S	Legitimately skipped

PRC31H		infant sex 2
N	Value	Description
3	Range	1 - 2 (median=1 mean=1.3 std=0.6)
124	.S	Legitimately skipped

PRC32A1		grams 3
N	Value	Description
127	.S	Legitimately skipped

PRC32A2		pounds 3
N	Value	Description
127	.S	Legitimately skipped

PRC32A3		ounces 3
N	Value	Description
127	.S	Legitimately skipped

PRC32B		one min apgar 3
N	Value	Description
127	.S	Legitimately skipped

PRC32C		five min apgar 3
N	Value	Description
127	.S	Legitimately skipped

PRC32D		shoulder dystocia 3
N	Value	Description
127	.S	Legitimately skipped

PRC32E		birth trauma 3
N	Value	Description
127	.S	Legitimately skipped

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PRC32F		stillbirth
N	Value	Description
127	.S	Legitimately skipped

PRC32G		type of delivery 3
N	Value	Description
127	.S	Legitimately skipped

PRC32H		infant sex 3
N	Value	Description
127	.S	Legitimately skipped

PRC33A1		grams 4
N	Value	Description
127	.S	Legitimately skipped

PRC33A2		pounds 4
N	Value	Description
127	.S	Legitimately skipped

PRC33A3		ounces 4
N	Value	Description
127	.S	Legitimately skipped

PRC33B		one min apgar 4
N	Value	Description
127	.S	Legitimately skipped

PRC33C		five min apgar 4
N	Value	Description
127	.S	Legitimately skipped

PRC33D		shoulder dystocia 4
N	Value	Description
127	.S	Legitimately skipped

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PRC33E		birth trauma 4
N	Value	Description
127	.S	Legitimately skipped

PRC33F		stillbirth 4
N	Value	Description
127	.S	Legitimately skipped

PRC33G		type of delivery 4
N	Value	Description
127	.S	Legitimately skipped

PRC33H		infant sex 4
N	Value	Description
127	.S	Legitimately skipped

PRC0A_FUTIME		Futime from Randomization date for (PRC0A)
N	Value	Description
127	Range	2093 - 3703 (median=2756 mean=2708.0 std=372.0)

PRC0D_FUTIME		Futime from Randomization date for (PRC0D)
N	Value	Description
127	Range	212 - 3405 (median=1333 mean=1433.4 std=803.4)

PRC3_FUTIME		Futime from Randomization date for (PRC3)
N	Value	Description
126	Range	212 - 3405 (median=1330 mean=1421.1 std=806.4)
1		Missing

PRC3B_FUTIME		Futime from Randomization date for (PRC3B)
N	Value	Description
127	Range	212 - 3405 (median=1333 mean=1433.4 std=803.4)

PRC4_FUTIME		Futime from Randomization date for (PRC4)
N	Value	Description
126	Range	242 - 3427 (median=1350 mean=1445.5 std=806.8)
1		Missing

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<i>PRC10_FUTIME</i>		<i>Futime from Randomization date for (PRC10)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
101	Range	252 - 3493 (median=1489 mean=1501.5 std=844.5)
26		Missing

<i>PRC14A1_FUTIME</i>		<i>Futime from Randomization date for (PRC14A1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
41	Range	240 - 3405 (median=1674 mean=1555.3 std=911.6)
86		Missing

<i>PRC14B1_FUTIME</i>		<i>Futime from Randomization date for (PRC14B1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17	Range	240 - 3405 (median=1771 mean=1652.8 std=860.1)
110		Missing

<i>PRC17A1_FUTIME</i>		<i>Futime from Randomization date for (PRC17A1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	273 - 1174 (median=958.5 mean=841.00 std=404.61)
123		Missing

<i>PRC17B2_FUTIME</i>		<i>Futime from Randomization date for (PRC17B2)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	99 - 2631 (median=903 mean=1116.6 std=1038.9)
120		Missing

<i>PRC17C1_FUTIME</i>		<i>Futime from Randomization date for (PRC17C1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
10	Range	129 - 2640 (median=1188.5 mean=1193.90 std=954.45)
117		Missing

<i>PRC17D1B_FUTIME</i>		<i>Futime from Randomization date for (PRC17D1B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	1546 - 2631 (median=2045 mean=2066.8 std=450.4)
123		Missing

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<i>PRC17D2B_FUTIME</i>		<i>Futime from Randomization date for (PRC17D2B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5	Range	185 - 2210 (median=1532 mean=1377.2 std=775.7)
122		Missing

<i>PRC17E1B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E1B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	338 - 1893 (median=1222 mean=1168.8 std=725.7)
123		Missing

<i>PRC17E2B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E2B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	279 - 3350 (median=1602 mean=1617.3 std=787.4)
102		Missing

<i>PRC17E3B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E3B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	338 - 1886 (median=1255 mean=1183.5 std=738.5)
123		Missing

<i>PRC17E4B_FUTIME</i>		<i>Futime from Randomization date for (PRC17E4B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
26	Range	279 - 3350 (median=1608.5 mean=1627.15 std=771.02)
101		Missing

<i>PRC17F2B_FUTIME</i>		<i>Futime from Randomization date for (PRC17F2B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
4	Range	134 - 1880 (median=552.5 mean=779.75 std=780.99)
123		Missing

<i>PRC17F3B_FUTIME</i>		<i>Futime from Randomization date for (PRC17F3B)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
23	Range	276 - 2404 (median=1583 mean=1374.1 std=687.1)
104		Missing

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<i>PRC18A1_FUTIME</i>		<i>Futime from Randomization date for (PRC18A1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	273 - 1675 (median=1081 mean=939.4 std=495.8)
120		Missing

<i>PRC18B1_FUTIME</i>		<i>Futime from Randomization date for (PRC18B1)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	Range	1944 - 3405 (median=2250 mean=2470.5 std=562.7)
119		Missing

<i>PRC21A_FUTIME</i>		<i>Futime from Randomization date for (PRC21A)</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
109	Range	240 - 3417 (median=1338 mean=1451.5 std=807.2)
18		Missing

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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

OCCURRENCE		Occurrence
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
127	Range	1 - 3 (median=2 mean=1.7 std=0.7)

PCD12		prior hypertension
N	Value	Description
29	Range	0 - 1 (median=0 mean=0.4 std=0.5)
17	.S	Legitimately skipped
81		Missing

PCD12A1		SBP Prior
N	Value	Description
12	Range	0 - 9 (median=0.5 mean=1.17 std=2.52)
34	.S	Legitimately skipped
81		Missing

PCD12A2		self-report hypertension
N	Value	Description
12	Range	0 - 9 (median=0.5 mean=1.17 std=2.52)
34	.S	Legitimately skipped
81		Missing

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PCD12A3		documentation of hypertensions
N	Value	Description
12	Range	0 - 1 (median=1 mean=0.8 std=0.5)
34	.S	Legitimately skipped
81		Missing

PCD13		SBP less than 20 weeks
N	Value	Description
27	Range	0 - 9 (median=9 mean=7.4 std=3.4)
17	.S	Legitimately skipped
83		Missing

PCD13A		documentation of treatment
N	Value	Description
26	Range	0 - 9 (median=9 mean=7.3 std=3.5)
17	.S	Legitimately skipped
84		Missing

PCD14		hypertensive blood pressures 4 hours apart
N	Value	Description
27	Range	0 - 9 (median=9 mean=4.8 std=4.4)
17	.S	Legitimately skipped
83		Missing

PCD15		proteinuria documented
N	Value	Description
28	Range	1 - 9 (median=5.5 mean=4.93 std=2.77)
17	.S	Legitimately skipped
82		Missing

PCD16A		two occasions
N	Value	Description
29	Range	0 - 9 (median=1 mean=3.6 std=4.3)
17	.S	Legitimately skipped
81		Missing

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PCD16A1		<i>more than 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6	Range	0 - 9 (median=1 mean=3.5 std=4.3)
40	.S	Legitimately skipped
81		Missing

PCD16B		<i>anti-hypertensive medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=1 mean=3.3 std=4.2)
17	.S	Legitimately skipped
81		Missing

PCD16C		<i>elevated liver enzymes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=0 mean=2.5 std=4.1)
17	.S	Legitimately skipped
81		Missing

PCD16D		<i>persistent pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=0 mean=2.5 std=4.1)
17	.S	Legitimately skipped
81		Missing

PCD16E		<i>thrombocytopenia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=0 mean=0.9 std=2.8)
17	.S	Legitimately skipped
81		Missing

PCD16F		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=0 mean=2.8 std=4.2)
17	.S	Legitimately skipped
81		Missing

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<i>PCD16G</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=9 mean=7.5 std=3.4)
17	.S	Legitimately skipped
81		Missing

<i>PCD16H</i>		<i>new onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=0 mean=2.6 std=4.0)
17	.S	Legitimately skipped
81		Missing

<i>PCD17</i>		<i>intrapartum magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	0 - 9 (median=9 mean=5.8 std=4.2)
17	.S	Legitimately skipped
81		Missing

<i>PCD17A</i>		<i>magnesium indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
5	Range	0 - 1 (median=1 mean=0.8 std=0.4)
41	.S	Legitimately skipped
81		Missing

<i>PCD18</i>		<i>grand mal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	4 - 9 (median=4 mean=4.2 std=0.9)
17	.S	Legitimately skipped
81		Missing

<i>PCD19</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	1 - 5 (median=4 mean=3.4 std=1.1)
17	.S	Legitimately skipped
81		Missing

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<i>PCD110</i>		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	1 - 5 (median=4 mean=3.1 std=1.2)
17	.S	Legitimately skipped
81		Missing

<i>PCD111</i>		<i>severe features</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	1 - 4 (median=4 mean=3.7 std=0.8)
17	.S	Legitimately skipped
81		Missing

<i>PCD112</i>		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
29	Range	4 - 4 (median=4 mean=4.0 std=0.0)
17	.S	Legitimately skipped
81		Missing

<i>PCD113</i>		<i>pregestational DM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24	Range	0 - 1 (median=0 mean=0.3 std=0.5)
21	.S	Legitimately skipped
82		Missing

<i>PCD114A</i>		<i>fasting glucose ?126</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	Range	0 - 1 (median=0 mean=0.4 std=0.5)
37	.S	Legitimately skipped
82		Missing

<i>PCD114B</i>		<i>2h glucose ? 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	Range	0 - 9 (median=5 mean=4.8 std=4.6)
37	.S	Legitimately skipped
82		Missing

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<i>PCD114C</i>		<i>prior HbA1c ? 6.5</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	Range	0 - 9 (median=1 mean=1.6 std=3.0)
37	.S	Legitimately skipped
82		Missing

<i>PCD114D</i>		<i>pregestational DM documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	0 - 1 (median=0 mean=0.4 std=0.5)
37	.S	Legitimately skipped
83		Missing

<i>PCD114E</i>		<i>DM self-report</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
8	Range	0 - 9 (median=0.5 mean=1.50 std=3.07)
37	.S	Legitimately skipped
82		Missing

<i>PCD115</i>		<i>glucose intolerance present</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
24	Range	0 - 9 (median=9 mean=5.9 std=4.1)
21	.S	Legitimately skipped
82		Missing

<i>PCD115A1</i>		<i>3 hour OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	1 - 9 (median=9 mean=5.6 std=4.3)
38	.S	Legitimately skipped
82		Missing

<i>PCD115A2</i>		<i>2h 75g OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	9 - 9 (median=9 mean=9.0 std=0.0)
38	.S	Legitimately skipped
82		Missing

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<i>PCD115A3</i>		<i>50g GLT > 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	0 - 9 (median=9 mean=6.4 std=4.4)
38	.S	Legitimately skipped
82		Missing

<i>PCD115A4</i>		<i>two or more glucose > 125</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	1 - 9 (median=9 mean=7.9 std=3.0)
38	.S	Legitimately skipped
82		Missing

<i>PCD115A5</i>		<i>abnormal or elevated OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	0 - 9 (median=9 mean=6.6 std=4.2)
38	.S	Legitimately skipped
82		Missing

<i>PCD115B</i>		<i>Non-diagnostic glucose intolerance</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17	Range	0 - 9 (median=9 mean=7.5 std=3.4)
28	.S	Legitimately skipped
82		Missing

<i>PCD115B1</i>		<i>elevated 1 hour GLT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	9 - 9 (median=9 mean=9.0 std=)
44	.S	Legitimately skipped
82		Missing

<i>PCD115B2</i>		<i>evidence of abnormal glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	1 - 1 (median=1 mean=1.0 std=)
44	.S	Legitimately skipped
82		Missing

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PCD116		treatment with insulin or oral
N	Value	Description
24	Range	0 - 9 (median=1 mean=1.6 std=2.3)
21	.S	Legitimately skipped
82		Missing

PCD117		GDM diagnosis
N	Value	Description
24	Range	0 - 1 (median=1 mean=0.8 std=0.4)
21	.S	Legitimately skipped
82		Missing

PCD117A		physician mention GDM
N	Value	Description
24	Range	0 - 1 (median=1 mean=0.8 std=0.4)
21	.S	Legitimately skipped
82		Missing

PCD118		Gestational Diabetes classification
N	Value	Description
24	Range	1 - 6 (median=1 mean=2.6 std=2.1)
21	.S	Legitimately skipped
82		Missing

PCD119		Comments
N	Value	Description
22	Present	Text suppressed
105		Missing

PCD10A_FUTIME		Futime from Randomization date for (PCD10A)
N	Value	Description
45	Range	2563 - 3896 (median=3173 mean=3217.3 std=331.3)
82		Missing

PCD10D_FUTIME		Futime from Randomization date for (PCD10D)
N	Value	Description
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

OCCURRENCE		Occurrence
N	Value	Description
109	1	Occurrence 1
14	2	Occurrence 2
3	3	Occurrence 3
1	4	Occurrence 4

PCD219		Comments
N	Value	Description
34	Present	Text suppressed
93		Missing

PCD21		PCD reviews required
N	Value	Description
127	Range	1 - 3 (median=2 mean=1.7 std=0.7)

PCD22		prior hypertension
N	Value	Description
59	Range	0 - 9 (median=0 mean=0.7 std=2.0)
59	.S	Legitimately skipped
9		Missing

PCD22A1		SBP Prior
N	Value	Description
18	Range	0 - 9 (median=0 mean=1.7 std=3.4)
101	.S	Legitimately skipped
8		Missing

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<i>PCD22A2</i>		<i>self-report hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	0 - 9 (median=1 mean=3.2 std=4.2)
101	.S	Legitimately skipped
8		Missing

<i>PCD22A3</i>		<i>documentation of hypertensions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	0 - 9 (median=1 mean=1.7 std=2.7)
101	.S	Legitimately skipped
8		Missing

<i>PCD23</i>		<i>SBP less than 20 weeks</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=9 mean=5.7 std=4.4)
59	.S	Legitimately skipped
8		Missing

<i>PCD23A</i>		<i>documentation of treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
58	Range	0 - 9 (median=9 mean=5.0 std=4.5)
59	.S	Legitimately skipped
10		Missing

<i>PCD24</i>		<i>hypertensive blood pressures 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
57	Range	0 - 9 (median=9 mean=5.1 std=4.3)
59	.S	Legitimately skipped
11		Missing

<i>PCD25</i>		<i>proteinuria documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
59	Range	1 - 9 (median=6 mean=6.2 std=2.8)
59	.S	Legitimately skipped
9		Missing

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PCD26A		<i>two occasions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=1 mean=4.3 std=4.4)
59	.S	Legitimately skipped
8		Missing

PCD26A1		<i>more than 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6	Range	0 - 1 (median=1 mean=0.7 std=0.5)
113	.S	Legitimately skipped
8		Missing

PCD26B		<i>anti-hypertensive medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=0.5 mean=3.30 std=4.23)
59	.S	Legitimately skipped
8		Missing

PCD26C		<i>elevated liver enzymes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=0 mean=2.6 std=4.1)
59	.S	Legitimately skipped
8		Missing

PCD26D		<i>persistent pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=0 mean=2.4 std=4.0)
59	.S	Legitimately skipped
8		Missing

PCD26E		<i>thrombocytopenia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=0 mean=2.6 std=4.1)
59	.S	Legitimately skipped
8		Missing

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<i>PCD26F</i>		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=0 mean=2.7 std=4.2)
59	.S	Legitimately skipped
8		Missing

<i>PCD26G</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=0 mean=3.2 std=4.3)
59	.S	Legitimately skipped
8		Missing

<i>PCD26H</i>		<i>new onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
60	Range	0 - 9 (median=0 mean=1.8 std=3.5)
59	.S	Legitimately skipped
8		Missing

<i>PCD27</i>		<i>intrapartum magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
59	Range	0 - 9 (median=1 mean=4.6 std=4.3)
59	.S	Legitimately skipped
9		Missing

<i>PCD27A</i>		<i>magnesium indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 9 (median=9 mean=5.8 std=4.3)
113	.S	Legitimately skipped
1		Missing

<i>PCD28</i>		<i>grand mal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
59	Range	4 - 9 (median=4 mean=4.3 std=1.1)
59	.S	Legitimately skipped
9		Missing

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<i>PCD29</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
59	Range	1 - 5 (median=4 mean=3.3 std=1.2)
59	.S	Legitimately skipped
9		Missing

<i>PCD210</i>		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
59	Range	1 - 5 (median=3 mean=3.1 std=1.2)
59	.S	Legitimately skipped
9		Missing

<i>PCD211</i>		<i>severe features</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
59	Range	1 - 5 (median=4 mean=3.8 std=0.8)
59	.S	Legitimately skipped
9		Missing

<i>PCD212</i>		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
59	Range	3 - 5 (median=4 mean=4.0 std=0.2)
59	.S	Legitimately skipped
9		Missing

<i>PCD213</i>		<i>pregestational DM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
66	Range	0 - 9 (median=0 mean=0.7 std=2.2)
53	.S	Legitimately skipped
8		Missing

<i>PCD214A</i>		<i>fasting glucose ?126</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
12	Range	0 - 9 (median=5 mean=4.8 std=4.5)
107	.S	Legitimately skipped
8		Missing

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PCD214B		2h glucose ? 200
N	Value	Description
12	Range	0 - 9 (median=9 mean=6.2 std=4.2)
107	.S	Legitimately skipped
8		Missing

PCD214C		prior HbA1c ? 6.5
N	Value	Description
12	Range	0 - 9 (median=1 mean=4.1 std=4.4)
107	.S	Legitimately skipped
8		Missing

PCD214D		pregestational DM documented
N	Value	Description
11	Range	1 - 9 (median=1 mean=2.5 std=3.2)
107	.S	Legitimately skipped
9		Missing

PCD214E		DM self-report
N	Value	Description
12	Range	0 - 9 (median=1 mean=4.1 std=4.4)
107	.S	Legitimately skipped
8		Missing

PCD215		glucose intolerance present
N	Value	Description
65	Range	0 - 9 (median=9 mean=5.4 std=4.1)
53	.S	Legitimately skipped
9		Missing

PCD215A1		3 hour OGTT
N	Value	Description
25	Range	0 - 9 (median=9 mean=5.3 std=4.2)
94	.S	Legitimately skipped
8		Missing

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<i>PCD215A2</i>		<i>2h 75g OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	0 - 9 (median=9 mean=6.3 std=4.0)
94	.S	Legitimately skipped
8		Missing

<i>PCD215A3</i>		<i>50g GLT > 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	0 - 9 (median=9 mean=5.8 std=4.4)
94	.S	Legitimately skipped
8		Missing

<i>PCD215A4</i>		<i>two or more glucose > 125</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	0 - 9 (median=9 mean=7.6 std=3.4)
94	.S	Legitimately skipped
8		Missing

<i>PCD215A5</i>		<i>abnormal or elevated OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
25	Range	0 - 9 (median=9 mean=6.1 std=4.3)
94	.S	Legitimately skipped
8		Missing

<i>PCD215B</i>		<i>Non-diagnostic glucose intolerance</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
42	Range	0 - 9 (median=9 mean=6.4 std=4.0)
77	.S	Legitimately skipped
8		Missing

<i>PCD215B1</i>		<i>elevated 1 hour GLT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	1 - 9 (median=9 mean=7.9 std=3.0)
112	.S	Legitimately skipped
8		Missing

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<i>PCD215B2</i>		<i>evidence of abnormal glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	1 - 9 (median=9 mean=5.6 std=4.3)
112	.S	Legitimately skipped
8		Missing

<i>PCD216</i>		<i>treatment with insulin or oral</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
65	Range	0 - 9 (median=1 mean=2.7 std=3.5)
53	.S	Legitimately skipped
9		Missing

<i>PCD217</i>		<i>GDM diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
65	Range	0 - 9 (median=1 mean=1.3 std=1.7)
53	.S	Legitimately skipped
9		Missing

<i>PCD217A</i>		<i>physician mention GDM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
64	Range	0 - 9 (median=1 mean=1.3 std=1.8)
53	.S	Legitimately skipped
10		Missing

<i>PCD218</i>		<i>Gestational Diabetes classification</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
66	Range	1 - 6 (median=2 mean=2.5 std=1.7)
53	.S	Legitimately skipped
8		Missing

<i>FORMSTATUS_PCD 2</i>		<i>Form Status</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Present	Text suppressed

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<i>FORMLOCKED_PCD2</i>		<i>Form Locked</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	No	

<i>FORMSIGNED_PCD2</i>		<i>Form Signed</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	No	

<i>LOGINNAME_PCD2</i>		<i>Login Name</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Present	Text suppressed

<i>LOCATIONNAME_PCD2</i>		<i>Location Name</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
33	B	
44	C	
19	M	
31	.S	Legitimately skipped

<i>DATESTAMPINITIAL_PCD2</i>		<i>Date Stamp Initial</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Present	Text suppressed

<i>TIMESTAMPINITIAL_PCD2</i>		<i>Time Stamp Initial</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Present	Text suppressed

<i>DATESTAMPLASTMOD_PCD2</i>		<i>Date Stamp Last Modified</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
127	Present	Text suppressed

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TIMESTAMPLASTMOD_PCD2		Time Stamp Last Modified
N	Value	Description
127	Present	Text suppressed

PCD20A_FUTIME		Futime from Randomization date for (PCD20A)
N	Value	Description
101	Range	2585 - 3868 (median=3147 mean=3192.2 std=293.5)
26		Missing

PCD20D_FUTIME		Futime from Randomization date for (PCD20D)
N	Value	Description
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

OCCURRENCE		Occurrence
N	Value	Description
107	1	Occurrence 1
13	2	Occurrence 2
3	3	Occurrence 3
1	4	Occurrence 4

PCD319		Comments
N	Value	Description
71	Present	Text suppressed
53		Missing

PCD31		PCD reviews required
N	Value	Description
124	Range	1 - 3 (median=2 mean=1.8 std=0.7)

PCD32		prior hypertension
N	Value	Description
72	Range	0 - 9 (median=0 mean=0.5 std=1.1)
52	.S	Legitimately skipped

PCD32A1		SBP Prior
N	Value	Description
26	Range	0 - 1 (median=0 mean=0.3 std=0.5)
97	.S	Legitimately skipped
1		Missing

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<i>PCD32A2</i>		<i>self-report hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
26	Range	0 - 1 (median=0 mean=0.3 std=0.5)
97	.S	Legitimately skipped
1		Missing

<i>PCD32A3</i>		<i>documentation of hypertensions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
27	Range	0 - 1 (median=1 mean=0.8 std=0.4)
97	.S	Legitimately skipped

<i>PCD33</i>		<i>SBP less than 20 weeks</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
70	Range	0 - 9 (median=9 mean=8.2 std=2.5)
52	.S	Legitimately skipped
2		Missing

<i>PCD33A</i>		<i>documentation of treatment</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
63	Range	0 - 9 (median=9 mean=8.6 std=1.9)
52	.S	Legitimately skipped
9		Missing

<i>PCD34</i>		<i>hypertensive blood pressures 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=9 mean=5.0 std=4.2)
52	.S	Legitimately skipped

<i>PCD35</i>		<i>proteinuria documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
71	Range	1 - 9 (median=6 mean=5.6 std=2.9)
52	.S	Legitimately skipped
1		Missing

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<i>PCD36A</i>		<i>two occasions</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=9 mean=4.9 std=4.4)
52	.S	Legitimately skipped

<i>PCD36A1</i>		<i>more than 4 hours apart</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	0 - 9 (median=1 mean=3.0 std=4.1)
116	.S	Legitimately skipped
1		Missing

<i>PCD36B</i>		<i>anti-hypertensive medication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=9 mean=4.7 std=4.4)
52	.S	Legitimately skipped

<i>PCD36C</i>		<i>elevated liver enzymes</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=0 mean=3.4 std=4.4)
52	.S	Legitimately skipped

<i>PCD36D</i>		<i>persistent pain</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=9 mean=4.9 std=4.5)
52	.S	Legitimately skipped

<i>PCD36E</i>		<i>thrombocytopenia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=0 mean=3.0 std=4.3)
52	.S	Legitimately skipped

<i>PCD36F</i>		<i>serum creatinine</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=0 mean=3.7 std=4.4)
52	.S	Legitimately skipped

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<i>PCD36G</i>		<i>pulmonary edema</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=9 mean=8.3 std=2.5)
52	.S	Legitimately skipped

<i>PCD36H</i>		<i>new onset</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=1 mean=4.4 std=4.4)
52	.S	Legitimately skipped

<i>PCD37</i>		<i>intrapartum magnesium</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	0 - 9 (median=9 mean=4.9 std=4.3)
52	.S	Legitimately skipped

<i>PCD37A</i>		<i>magnesium indication</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
15	Range	1 - 1 (median=1 mean=1.0 std=0.0)
107	.S	Legitimately skipped
2		Missing

<i>PCD38</i>		<i>grand mal</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	4 - 4 (median=4 mean=4.0 std=0.0)
52	.S	Legitimately skipped

<i>PCD39</i>		<i>gestational hypertension</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	1 - 5 (median=4 mean=3.5 std=0.9)
52	.S	Legitimately skipped

<i>PCD310</i>		<i>preeclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	1 - 5 (median=4 mean=3.1 std=1.4)
52	.S	Legitimately skipped

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<i>PCD311</i>		<i>severe features</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	1 - 5 (median=4 mean=3.7 std=1.1)
52	.S	Legitimately skipped

<i>PCD312</i>		<i>eclampsia</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
72	Range	4 - 4 (median=4 mean=4.0 std=0.0)
52	.S	Legitimately skipped

<i>PCD313</i>		<i>pregestatal DM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	Range	0 - 1 (median=0 mean=0.2 std=0.4)
48	.S	Legitimately skipped

<i>PCD314A</i>		<i>fasting glucose ?126</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	0 - 1 (median=0 mean=0.4 std=0.5)
106	.S	Legitimately skipped

<i>PCD314B</i>		<i>2h glucose ? 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	0 - 9 (median=1 mean=3.3 std=4.1)
106	.S	Legitimately skipped

<i>PCD314C</i>		<i>prior HbA1c ? 6.5</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	0 - 1 (median=0.5 mean=0.50 std=0.51)
106	.S	Legitimately skipped

<i>PCD314D</i>		<i>pregestational DM documented</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	0 - 9 (median=1 mean=1.1 std=2.0)
106	.S	Legitimately skipped

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<i>PCD314E</i>		<i>DM self-report</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
18	Range	0 - 9 (median=0 mean=0.8 std=2.1)
106	.S	Legitimately skipped

<i>PCD315</i>		<i>glucose intolerance present</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	Range	0 - 9 (median=9 mean=6.7 std=3.8)
48	.S	Legitimately skipped

<i>PCD315A1</i>		<i>3 hour OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 9 (median=1 mean=1.2 std=2.4)
110	.S	Legitimately skipped
1		Missing

<i>PCD315A2</i>		<i>2h 75g OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 9 (median=0 mean=1.4 std=3.4)
110	.S	Legitimately skipped
1		Missing

<i>PCD315A3</i>		<i>50g GLT > 200</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 9 (median=0 mean=0.9 std=2.5)
110	.S	Legitimately skipped
1		Missing

<i>PCD315A4</i>		<i>two or more glucose > 125</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
13	Range	0 - 9 (median=0 mean=1.4 std=3.4)
110	.S	Legitimately skipped
1		Missing

Chapter 1: Endpoint Ascertainment

Section F: Pregnancy Complication Ascertainment

<i>PCD315A5</i>		<i>abnormal or elevated OGTT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
14	Range	0 - 9 (median=0 mean=1.6 std=3.2)
110	.S	Legitimately skipped

<i>PCD315B</i>		<i>Non-diagnostic glucose intolerance</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
61	Range	0 - 9 (median=9 mean=7.6 std=3.1)
62	.S	Legitimately skipped
1		Missing

<i>PCD315B1</i>		<i>elevated 1 hour GLT</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	1 - 9 (median=1 mean=3.3 std=3.9)
116	.S	Legitimately skipped
1		Missing

<i>PCD315B2</i>		<i>evidence of abnormal glucose</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
7	Range	0 - 9 (median=0 mean=1.6 std=3.3)
116	.S	Legitimately skipped
1		Missing

<i>PCD316</i>		<i>treatment with insulin or oral</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	Range	0 - 9 (median=1 mean=1.6 std=2.8)
48	.S	Legitimately skipped

<i>PCD317</i>		<i>GDM diagnosis</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	Range	0 - 1 (median=1 mean=0.8 std=0.4)
48	.S	Legitimately skipped

<i>PCD317A</i>		<i>physician mention GDM</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
76	Range	0 - 1 (median=1 mean=0.7 std=0.4)
48	.S	Legitimately skipped

Chapter 1: Endpoint Ascertainment

Section F: Pregnancy Complication Ascertainment

PCD318		Gestational Diabetes classification
N	Value	Description
76	Range	1 - 6 (median=2 mean=2.9 std=2.0)
48	.S	Legitimately skipped

FORMSTATUS_PCD3		Form Status
N	Value	Description
124	Present	Text suppressed

FORMLOCKED_PCD3		Form Locked
N	Value	Description
124	No	

FORMSIGNED_PCD3		Form Signed
N	Value	Description
124	No	

LOGINNAME_PCD3		Login Name
N	Value	Description
124	Present	Text suppressed

LOCATIONNAME_PCD3		Location Name
N	Value	Description
31	B	
43	C	
19	M	
31	.S	Legitimately skipped

DATESTAMPINITIAL_PCD3		Date Stamp Initial
N	Value	Description
124	Present	Text suppressed

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Section F: Pregnancy Complication Ascertainment

TIMESTAMPINITIAL_PCD3		Time Stamp Initial
N	Value	Description
124	Present	Text suppressed

DATESTAMPLASTM OD_PCD3		Date Stamp Last Modified
N	Value	Description
124	Present	Text suppressed

TIMESTAMPLASTMO D_PCD3		Time Stamp Last Modified
N	Value	Description
124	Present	Text suppressed

PCD30A_FUTIME		Futime from Randomization date for (PCD30A)
N	Value	Description
124	Range	2855 - 3981 (median=3261 mean=3317.1 std=293.0)

PCD30D_FUTIME		Futime from Randomization date for (PCD30D)
N	Value	Description
124	Range	212 - 3405 (median=1346.5 mean=1449.21 std=782.42)

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

HCHS incident MI Event Derived Variable file 2008-2019: FORM CODE=INCIDENT_MI_EVNT, VERSION=1

INCIDENT_MI_2019		Incident Myocardial Infarction (MI) Event by 2019
N	Value	Description
0	0	No
138	1	Yes

RECUR_MI_2019_1		MI as 1st Recur MI 2019
N	Value	Description
0	0	No
15	1	Yes
123		Missing

RECUR_MI_2019_2		MI as 2nd Recur MI 2019
N	Value	Description
0	0	No
5	1	Yes
133		Missing

FUTIME_MI		Follow up time of MI event, censor or the end of follow-up by 2019
N	Value	Description
138	Range	119 - 3949 (median=1915.5 mean=1909.67 std=1025.93)

FUTIME_MI_1		Follow up time 1st Recur MI
N	Value	Description
15	Range	336 - 3794 (median=2548 mean=2433.7 std=1033.5)
123		Missing

FUTIME_MI_2		Follow up time 2nd Recur MI
N	Value	Description
5	Range	1986 - 3750 (median=2403 mean=2620.8 std=681.8)
133		Missing

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

HCHS incident HF Event Derived Variable file 2008-2019: FORM CODE=INCIDENT_HF_EVNT, VERSION=1

INCIDENT_HF_2019 Incident Heart Failure (HF) Event by 2019		
N	Value	Description
0	0	No
184	1	Yes
16		Missing

RECUR_HF_2019_1 HF as 1st Recur HF 2019		
N	Value	Description
0	0	No
62	1	Yes
138		Missing

RECUR_HF_2019_2 HF as 2nd Recur HF 2019		
N	Value	Description
0	0	No
31	1	Yes
169		Missing

RECUR_HF_2019_3 HF as 3rd Recur HF 2019		
N	Value	Description
0	0	No
15	1	Yes
185		Missing

RECUR_HF_2019_4 HF as 4th Recur HF 2019		
N	Value	Description
0	0	No
6	1	Yes
194		Missing

RECUR_HF_2019_5 HF as 5th Recur HF 2019		
N	Value	Description
0	0	No
3	1	Yes
197		Missing

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

RECUR_HF_2019_6		HF as 6th Recur HF 2019
N	Value	Description
0	0	No
2	1	Yes
198		Missing

RECUR_HF_2019_7		HF as 7th Recur HF 2019
N	Value	Description
0	0	No
1	1	Yes
199		Missing

RECUR_HF_2019_8		HF as 8th Recur HF 2019
N	Value	Description
0	0	No
1	1	Yes
199		Missing

FUTIME_HF		Follow up time of HF event, censor or the end of follow-up by 2019
N	Value	Description
200	Range	12 - 4016 (median=1789.5 mean=1900.20 std=1042.70)

FUTIME_HF_1		Follow up time 1st Recur HF
N	Value	Description
62	Range	142 - 3895 (median=1989.5 mean=2035.97 std=908.90)
138		Missing

FUTIME_HF_2		Follow up time 2nd Recur HF
N	Value	Description
31	Range	934 - 3979 (median=2405 mean=2279.2 std=874.4)
169		Missing

FUTIME_HF_3		Follow up time 3rd Recur HF
N	Value	Description
15	Range	1006 - 4130 (median=2242 mean=2392.1 std=987.7)
185		Missing

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

<i>FUTIME_HF_4</i>		<i>Follow up time 4th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
6	Range	1378 - 3480 (median=2414.5 mean=2460.17 std=725.17)
194		Missing

<i>FUTIME_HF_5</i>		<i>Follow up time 5th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
3	Range	2259 - 3494 (median=2596 mean=2783.0 std=638.4)
197		Missing

<i>FUTIME_HF_6</i>		<i>Follow up time 6th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
2	Range	2604 - 3543 (median=3073.5 mean=3073.50 std=663.97)
198		Missing

<i>FUTIME_HF_7</i>		<i>Follow up time 7th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	2615 - 2615 (median=2615 mean=2615.0 std=)
199		Missing

<i>FUTIME_HF_8</i>		<i>Follow up time 8th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
1	Range	2683 - 2683 (median=2683 mean=2683.0 std=)
199		Missing

<i>HFPEF</i>		<i>HFpEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
69	0	No
105	1	Yes
26		Missing

<i>HFrEF</i>		<i>HFrEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
65	0	No
109	1	Yes
26		Missing

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

HCHS incident PUL Event Derived Variable file 2008-2019: FORM CODE=INCIDENT_PUL_EVNT, VERSION=1

ASTHMA_PLD		PLD define or highly probable Asthma by 2019
N	Value	Description
14	0	No
116	1	Yes

COPD_PLD		PLD define or highly probable COPD by 2019
N	Value	Description
113	0	No
17	1	Yes

CLRD_2019		Chronic Lower Respiratory Disease by 2019
N	Value	Description
0	0	No
130	1	Yes

CLRD_ASTHMA_2019		CLRD due to Asthma by 2019
N	Value	Description
13	0	No
117	1	Yes

CLRD_COPD_2019		CLRD due to COPD by 2019
N	Value	Description
113	0	No
17	1	Yes

EXACERBATION_CLRD		Exacerbation of CLRD by 2019
N	Value	Description
33	0	No
97	1	Yes

FUTIME_CLRD		Follow up time of CLRD event, censor or the end of follow-up by 2019
N	Value	Description
130	Range	74 - 2602 (median=739.5 mean=944.50 std=640.27)

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

<i>FUTURE_CLRD_AST HMA</i>		<i>Follow up time of ASTHMA event, censor or the end of follow-up by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
117	Range	74 - 2602 (median=717 mean=908.1 std=641.6)
13		Missing

<i>FUTURE_CLRD_COP D</i>		<i>Follow up time of COPD event, censor or the end of follow-up by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
17	Range	426 - 2403 (median=1415 mean=1393.7 std=661.5)
113		Missing

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

HCHS incident STR Event Derived Variable file 2008-2019: FORM CODE=INCIDENT_STR_EVNT, VERSION=1

INCIDENT_STR_2019		Incident Stroke (STR) Event by 2019
N	Value	Description
0	0	No
139	1	Yes

RECUR_STR_2019_1		STR as Recur STR 2019 1
N	Value	Description
0	0	No
19	1	Yes
120		Missing

RECUR_STR_2019_2		STR as Recur STR 2019 2
N	Value	Description
0	0	No
2	1	Yes
137		Missing

FUTIME_STR		Follow up time of STR event, censor or the end of follow-up by 2019
N	Value	Description
139	Range	42 - 3734 (median=1537 mean=1617.5 std=997.0)

FUTIME_STR_1		Recur FUTIME STR 1
N	Value	Description
19	Range	545 - 3323 (median=1320 mean=1580.1 std=904.4)
120		Missing

FUTIME_STR_2		Recur FUTIME STR 2
N	Value	Description
2	Range	2675 - 2730 (median=2702.5 mean=2702.50 std=38.89)
137		Missing

Chapter 2: Endpoint Derived Variable File

Section A: Incident Event Derived Variable File

HCHS incident PRC Event Derived Variable file all years: FORM CODE=INCIDENT_PRC_EVNT, VERSION=1

INCIDENT_GHTN		Pregnancy Complications Diagnosis of GHTN (Preeclampsia to Eclampsia) Event
N	Value	Description
0	0	No
23	1	Yes
104		Missing

INCIDENT_GDM		Pregnancy Complications Diagnosis of GDM Event
N	Value	Description
0	0	No
45	1	Yes
82		Missing

INCIDENT_PCD		Pregnancy Complications Diagnosis (PCD) Event
N	Value	Description
0	0	No
104	1	Yes
23		Missing

Chapter 2: Endpoint Derived Variable File

Section B: Combined Event Derived Variable File for All Incident Events
(N=16,415)

HCHS Combined Incident Event Derived Variable file 2008-2019: FORM CODE=EVENT_PART_DERV, VERSION=1

ASTHMA_PLD		PLD define or highly probable Asthma by 2019
N	Value	Description
14	0	No
116	1	Yes
16285		Missing

COPD_PLD		PLD define or highly probable COPD by 2019
N	Value	Description
113	0	No
17	1	Yes
16285		Missing

CLRD_2019		Chronic Lower Respiratory Disease by 2019
N	Value	Description
16285	0	No
130	1	Yes

CLRD_ASTHMA_2019		CLRD due to Asthma by 2019
N	Value	Description
16298	0	No
117	1	Yes

CLRD_COPD_2019		CLRD due to COPD by 2019
N	Value	Description
16398	0	No
17	1	Yes

EXACERBATION_CLRD		Exacerbation of CLRD by 2019
N	Value	Description
33	0	No
97	1	Yes
16285		Missing

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Section B: Combined Event Derived Variable File for All Incident Events
(N=16,415)

<i>FUTURE_CLRD</i>		<i>Follow up time of CLRD event, censor or the end of follow-up by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3554 mean=3502.9 std=554.2)

<i>FUTURE_CLRD_AST HMA</i>		<i>Follow up time of ASTHMA event, censor or the end of follow-up by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3554 mean=3504.3 std=551.5)

<i>FUTURE_CLRD_COP D</i>		<i>Follow up time of COPD event, censor or the end of follow-up by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3559 mean=3521.0 std=508.2)

<i>INCIDENT_MI_2019</i>		<i>Incident Myocardial Infarction (MI) Event by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16277	0	No
138	1	Yes

<i>RECUR_MI_2019_1</i>		<i>MI as 1st Recur MI 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16400	0	No
15	1	Yes

<i>RECUR_MI_2019_2</i>		<i>MI as 2nd Recur MI 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16410	0	No
5	1	Yes

<i>FUTURE_MI</i>		<i>Follow up time of MI event, censor or the end of follow-up by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3555 mean=3510.4 std=529.6)

<i>FUTURE_MI_1</i>		<i>Follow up time 1st Recur MI</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3559 mean=3522.1 std=505.7)

Chapter 2: Endpoint Derived Variable File

Section B: Combined Event Derived Variable File for All Incident Events
(N=16,415)

FUTIME_MI_2		Follow up time 2nd Recur MI
N	Value	Description
16415	Range	25 - 4319 (median=3559 mean=3522.6 std=504.5)

INCIDENT_HF_2019		Incident Heart Failure (HF) Event by 2019
N	Value	Description
16215	0	No
184	1	Yes
16		Missing

RECUR_HF_2019_1		HF as 1st Recur HF 2019
N	Value	Description
16353	0	No
62	1	Yes

RECUR_HF_2019_2		HF as 2nd Recur HF 2019
N	Value	Description
16384	0	No
31	1	Yes

RECUR_HF_2019_3		HF as 3rd Recur HF 2019
N	Value	Description
16400	0	No
15	1	Yes

RECUR_HF_2019_4		HF as 4th Recur HF 2019
N	Value	Description
16409	0	No
6	1	Yes

RECUR_HF_2019_5		HF as 5th Recur HF 2019
N	Value	Description
16412	0	No
3	1	Yes

Chapter 2: Endpoint Derived Variable File

Section B: Combined Event Derived Variable File for All Incident Events
(N=16,415)

RECUR_HF_2019_6		HF as 6th Recur HF 2019
N	Value	Description
16413	0	No
2	1	Yes

RECUR_HF_2019_7		HF as 7th Recur HF 2019
N	Value	Description
16414	0	No
1	1	Yes

RECUR_HF_2019_8		HF as 8th Recur HF 2019
N	Value	Description
16414	0	No
1	1	Yes

FUTIME_HF		Follow up time of HF event, censor or the end of follow-up by 2019
N	Value	Description
16415	Range	12 - 4319 (median=3554 mean=3507.2 std=537.1)

FUTIME_HF_1		Follow up time 1st Recur HF
N	Value	Description
16415	Range	25 - 4319 (median=3556 mean=3518.4 std=513.2)

FUTIME_HF_2		Follow up time 2nd Recur HF
N	Value	Description
16415	Range	25 - 4319 (median=3559 mean=3521.1 std=507.4)

FUTIME_HF_3		Follow up time 3rd Recur HF
N	Value	Description
16415	Range	25 - 4319 (median=3559 mean=3522.0 std=505.7)

FUTIME_HF_4		Follow up time 4th Recur HF
N	Value	Description
16415	Range	25 - 4319 (median=3559 mean=3522.7 std=504.5)

Chapter 2: Endpoint Derived Variable File

Section B: Combined Event Derived Variable File for All Incident Events
(N=16,415)

<i>FUTURE_HF_5</i>		<i>Follow up time 5th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3559 mean=3522.7 std=504.4)

<i>FUTURE_HF_6</i>		<i>Follow up time 6th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3559 mean=3522.8 std=504.3)

<i>FUTURE_HF_7</i>		<i>Follow up time 7th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3560 mean=3522.8 std=504.3)

<i>FUTURE_HF_8</i>		<i>Follow up time 8th Recur HF</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3560 mean=3522.8 std=504.3)

<i>HFPEF</i>		<i>HFpEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
69	0	No
105	1	Yes
16241		Missing

<i>HFrEF</i>		<i>HFrEF simplified version</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
65	0	No
109	1	Yes
16241		Missing

<i>INCIDENT_GHTN</i>		<i>Pregnancy Complications Diagnosis of GHTN (Preeclampsia to Eclampsia) Event</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
0	0	No
22	1	Yes
16393		Missing

Chapter 2: Endpoint Derived Variable File

Section B: Combined Event Derived Variable File for All Incident Events
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INCIDENT_GDM		Pregnancy Complications Diagnosis of GDM Event
N	Value	Description
0	0	No
39	1	Yes
16376		Missing

INCIDENT_PCD		Pregnancy Complications Diagnosis (PCD) Event
N	Value	Description
0	0	No
93	1	Yes
16322		Missing

INCIDENT_STR_2019		Incident Stroke (STR) Event by 2019
N	Value	Description
16276	0	No
139	1	Yes

RECUR_STR_2019_1		STR as Recur STR 2019 1
N	Value	Description
16396	0	No
19	1	Yes

RECUR_STR_2019_2		STR as Recur STR 2019 2
N	Value	Description
16413	0	No
2	1	Yes

FUTIME_STR		Follow up time of STR event, censor or the end of follow-up by 2019
N	Value	Description
16415	Range	25 - 4319 (median=3554 mean=3508.9 std=534.7)

FUTIME_STR_1		Recur FUTIME STR 1
N	Value	Description
16415	Range	25 - 4319 (median=3559 mean=3521.1 std=508.3)

Chapter 2: Endpoint Derived Variable File

Section B: Combined Event Derived Variable File for All Incident Events
(N=16,415)

<i>FUTIME_STR_2</i>		<i>Recur FUTIME STR 2</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	25 - 4319 (median=3560 mean=3522.8 std=504.3)

<i>INCIDENT_CVD_2019</i>		<i>Incident CVD (MI, HF or Stroke) Event by 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16019	0	No
396	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 2019</i>
<i>N</i>	<i>Value</i>	<i>Description</i>
16415	Range	12 - 4319 (median=3549 mean=3485.2 std=580.1)