

Hidden Burden of Electronic Health Record-Identified Familial Hypercholesterolemia: Clinical Outcomes and Cost of Medical Care

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Background—Familial hypercholesterolemia (FH), is a historically underdiagnosed, undertreated, high-risk condition that is associated with a high burden of cardiovascular morbidity and mortality. In this study, we use a population-based approach using electronic health record (EHR)-based algorithms to identify FH. We report the major adverse cardiovascular events, mortality, and cost of medical care associated with this diagnosis.

Methods and Results—In our 1.18 million EHR-eligible cohort, *International Classification of Diseases, Ninth Revision (ICD-9)* codedefined hyperlipidemia was categorized into FH and non-FH groups using an EHR algorithm designed using the modified Dutch Lipid Clinic Network criteria. Major adverse cardiovascular events, mortality, and cost of medical care were analyzed. A priori associated variables/confounders were used for multivariate analyses using binary logistic regression and linear regression with propensity score—based weighted methods as appropriate. EHR FH was identified in 32 613 individuals, which was 2.7% of the 1.18 million EHR cohort and 13.7% of 237 903 patients with hyperlipidemia. FH had higher rates of myocardial infarction (14.77% versus 8.33%; *P*<0.0001), heart failure (11.82% versus 10.50%; *P*<0.0001), and, after adjusting for traditional risk factors, significantly correlated to a composite major adverse cardiovascular events variable (odds ratio, 4.02; 95% Cl, 3.88–4.16; *P*<0.0001), mortality (odds ratio, 1.20; Cl, 1.15–1.26; *P*<0.0001), and higher total revenue per-year (incidence rate ratio, 1.30; 95% Cl, 1.28–1.33; *P*<0.0001).

Conclusions—EHR-based algorithms discovered a disproportionately high prevalence of FH in our medical cohort, which was associated with worse outcomes and higher costs of medical care. This data-driven approach allows for a more precise method to identify traditionally high-risk groups within large populations allowing for targeted prevention and therapeutic strategies. (*J Am Heart Assoc.* 2019;8:e011822. DOI: 10.1161/JAHA.118.011822.)

Key Words: familial hypercholesterolemia • major adverse cardiovascular events • mortality • subclinical atherosclerosis risk factor • subclinical familial hypercholesterolemia

T he precision medicine model proposes customization of health care to individual patients, the success of which is largely dependent on early and accurate diagnosis. Familial hypercholesterolemia (FH)—a genetic disorder characterized

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by elevated low-density lipoprotein cholesterol (LDL-C) levels and early cardiovascular disease, is 1 of the 3 Centers for Disease Control and Prevention (CDC) designated Tier 1 public health genomic conditions, based on available evidence-based guidelines.¹ The major clinical manifestation of FH, premature atherosclerosis, is thought to result from the prolonged exposure of the vasculature to high levels of LDL-C. Clinical cardiovascular heart disease occurs at a higher frequency and at an earlier age in patients with FH than in patients without FH or patients with polygenetic causes of elevated LDL-C.² Traditional estimates of FH in the general population vary significantly, from 1:500 to 1:137,³ with the added challenge of lack of specific International Classification of Diseases (ICD) coding until recently. Despite advances in our understanding of the pathophysiology of FH, significant numbers remain undiagnosed and undertreated in relation to LDL-C targets.⁴ While the cost effectiveness of universal screening, early identification, and treatment is still evolving,⁵ availability of electronic health records (EHRs) and validated

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Clinical Perspective

What Is New?

- Familial hypercholesterolemia is characterized by lifelong elevation of circulating low-density lipoproteins and is associated with high cardiovascular mortality and morbidity; however, because of the variability in its presentation and lack of awareness, the condition is often underdiagnosed.
- Within our integrated health system, we evaluated novel ways of identifying and examining the sequelae of familial hypercholesterolemia in a population-based manner.
- This was done by using an electronic health record-based algorithm for detection and risk stratification of familial hypercholesterolemia and its associated comorbidities and cardiovascular outcomes and by using the total cost of care to examine the higher cost per year associated with the various categories of the familial hypercholesterolemia phenotype.

What Are the Clinical Implications?

 This methodology can allow health systems to study the drivers of worse clinical outcomes and higher costs of care to identify areas of opportunity to target limited resources.

clinical criteria may offer a rapid and efficient approach to population-based screening to identify high-risk individuals for targeted interventions. Additionally, evaluation of interventions in FH is complicated by the paucity of relevant economic data.⁶ The goal of this study is to use EHR-based algorithms to implement a population-based screening approach to identify the hidden burden of FH and study the trends of major adverse cardiovascular events (MACE), mortality and cost of care associated with this diagnosis.

Methods

The data that support the findings of this study are available from the corresponding author upon reasonable request.

Data Extraction

We queried our EHR database of over 1.5 million Geisinger Health System (GHS) patients (January 2000 to August 2016) with *ICD, Ninth Revision (ICD-9)* codes for hyperlipidemia (272.0, 272.1, 272.2, 272.3, 272.4). This project was approved as an expedited study on the basis of use of deidentified data by the Institutional Review Board at Geisinger Medical Center. We excluded patients aged <18 years or \geq 85 years, leaving an eligible study population of 1.18 million patients. An internally validated algorithm (see below) developed at Geisinger using modified Dutch Lipid Clinic Network Criteria (DLCN) (Table S1) was applied to these patients to categorize them into definite, probable, possible, and unlikely FH categories.⁷ The DLCN criteria were modified (Tables S2 and S3) to suit the EHR-derived data set. The first 3 of these discrete groups was defined to represent the hidden burden of FH in our population and were analyzed together as the "FH cohort."

The EHR-based algorithm based on the modified DLCN criteria for FH using a structured data set was validated internally by chart reviewing 250 randomly chosen case and control patients (definite, 2; probable, 13; possible, 111; unlikely, 124). A total of 125 of 126 patients were found to be accurately assigned to the positive group by the algorithm, for a positive predictive value of 99.21%. A total of 118 of 124 were found to be true-negative patients (unlikely FH), for a negative predictive value of 95.16% (for details of diagnostic accuracy see Tables S4 and S5). Six patients moved from the unlikely category to possible, for a sensitivity score of 95.42%. One patient went from possible to unlikely, for a specificity score of 99.16% (Table S6). Ten patients had errors found but stayed unlikely, and 1 patient remained in the definite category. Six of the 10 algorithm definitions were found to have 100% accuracy after chart reviews. Our EHR-based algorithm for identifying relevant family history in first-degree relatives performed poorly. Ten additional patients were found to have a family history of heart disease during the chart review of clinic notes, and 2 patients were deemed to have a family history of hyperlipidemia.

Outcomes

Primary outcomes of interest were MACE, mortality, and cost of care.

Mortality

Mortality was defined as all-cause mortality identified within the EHR. Our database interfaces with the Social Security Death Index on a biweekly basis to reflect updated information on vital status.

Major Adverse Cardiovascular Events

MACE was defined as a categorical variable denoting occurrence of the first instance of any of the following: all-cause death, myocardial infarction, percutaneous coronary intervention, or coronary artery bypass. Each of these clinical events were identified by using *ICD-9* codes (Table S7) in patients' EHR to obtain information relevant to MACE. Validation of the EHR-derived MACE and clinical variables was done by manual chart review in 100 randomly chosen subjects and showed good diagnostic accuracy (95% myocardial infarction, 95% heart failure, 100% ischemic stroke, 100% percutaneous coronary intervention, 100% bypass grafting, and 100% *ICD*).

Cost of Care

To identify the hidden economic burden of FH on healthcare system, we sought to determine differences in medical care expenditures between FH and non-FH hyperlipidemia cohorts on the basis of revenue data from medical care service use within our health system. Because of stepwise integration of several satellite sites within the system, financial data were not available for early years and some patients. We dealt with this by analyzing the clinical characteristics of patients with missing data to those with available data and using complete case analysis only. Inflation adjusted financial data for each year were calculated in terms of the 2015 data, according to the latest consumer price index data.⁸ We then compared adjusted median revenue between the 2 groups for the years 2005 through 2015. Because the revenue received by the healthcare system may vary depending on the payer mix, to calculate the absolute magnitude of difference in expenditure between the 2 groups we used a variable "Med Net Revenue," which adjusts all payer revenues to Medicare rates for GHS. These data were available only for the years 2014 and 2015.

Data Analyses

Statistical Analysis System (SAS) Version 9.4 (SAS Institute Inc., Cary, NC) was used for statistical analyses. Descriptive variables were expressed as mean with standard deviation in cases of normal distribution, and median with interguartile range (IQR) in case of nonnormal distribution. Normality was assessed by the Kolmogorov-Smirnoff test. Categorical variables are expressed as counts with percentages. Comparisons between continuous variables were conducted with nonparametric Mann-Whitney test for mean ranks for those variables with nonnormal distribution, whereas the chi-square test was performed to test the independence of categorical variables. To account for biases attributable to the observed confounders from baseline characteristics, the stabilized inverse probability of treatment weights was implemented by weighting the responses on the basis of their propensity scores in the regression model. The propensity scores were estimated on the basis of observed confounders that were significantly different between FH and non-FH cohorts including age, sex, smoking history, comorbid conditions, and the lost to follow-up indicator. Lost to follow-up was defined as no visit in EHR 2 years before the end of the study. A priori associated variables/confounders were included in the logistic regression, after adjusting for age, sex, smoking status, diabetes mellitus, hypertension, and LDL-C (maximum) values to evaluate if EHR FH significantly predicted outcomes.⁹ Odds ratios with corresponding 95% CIs were presented. By including the total follow-up time (defined as the years between first and last encounter over 2005–2015) as the offset variable to normalize the total adjusted revenue to a per-year basis, a negative binomial regression model using inverse probability of treatment weights was performed to evaluate the impact of FH on the total revenue after adjusting for a priori associated variables/confounders. Incidence rate ratios with corresponding 95% CIs were presented. A *P* value of <0.05 was considered significant.

Results

In our 1.18 million EHR-eligible cohort, 237 903 patients had hyperlipidemia (49.5% female, 96.6% white). Median age of the entire cohort was 63 years (IQR, 20 years), with an overall available median EHR follow-up of 10 years (IQR, 11 years). FH phenotype was identified using the EHR-based algorithm in 32 613 individuals, which was 2.76% of the entire EHR cohort (definite, 0.03%; probable, 0.16%; possible, 2.55%) and constituted 13.7% of patients with hyperlipidemia (Figure 1).

In univariate analysis, the FH cohort was noted to have a higher percentage of females (56.52% versus 48.51%; P<0.0001) and history of smoking (54.11% versus 50.73%; P < 0.0001). Baseline demographic characteristics as well as use of medications and diagnostics tests in 2 groups are shown in Table 1. Deceased patients in the FH cohort were younger at the time of death compared with those in the non-FH cohort (median, 70 years versus 75 years; P<0.0001). Cumulatively, a MACE event was observed at a younger age in the FH cohort (54 years versus 67 years; P<0.0001). Patients in the FH cohort experienced higher rate of specific MACE events, including myocardial infarction (14.77% versus 8.33%; *P*<0.0001) and heart failure (11.82% versus 10.50%; P<0.0001) (Table 2). Furthermore, total cholesterol, triglycerides, LDL-C, and high-density lipoprotein cholesterol levels were all higher in the FH cohort (Table 1). The FH cohort had a significantly higher median value of maximum LDL-C levels (highest LDL-C documented in EHR per patient) compared with that in non-FH cohort (202 mg/dL versus 137 mg/dL; P<0.0001). Analysis of medication data revealed higher use of statins (79.10% versus 57.82%; P<0.0001), high-potency statins (42.15% versus 19.49%; P<0.0001), and proprotein convertase subtilisin/kexin type 9 inhibitors (0.18% versus 0.02%; P<0.0001) in FH cohort compared with their non-FH counterparts (Table 1). Overall, there were a total of 11.3% lost to follow-up for MACE, and 12.5% lost to follow-up for mortality. In the logistic regression model, by weighting the responses on the basis of their propensity scores (inverse probability of treatment weights), EHR FH correlated with



Figure 1. FH categories within hyperlipidemia cohort of 237 903 individuals. FH indicates familial hypercholesterolemia.

MACE (incidence rate ratio, 4.02; 95% Cl, 3.88–4.16; P<0.0001) and mortality (incidence rate ratio, 1.20; 95% Cl, 1.15–1.26; P<0.0001) after adjusting for descriptive characteristics that differed between cohorts such as age, sex, smoking status, diabetes mellitus, hypertension, and maximum LDL (Figure 2).

Cost Analysis

Using a complete case analysis methodology, inflationadjusted median annual revenue for the FH cohort was higher for each year from 2005 through 2015 (Table 3). The total follow-up time for each patient during the study period for analysis (2005–2015) was used as an offset variable to normalize the total amount to a per-year basis. The median follow-up time was 11 years (IQR, 3 years) for the FH cohort and 10 years (IQR, 6 years) for the non-FH cohort.

In the negative binomial regression model, by including the total follow-up time (defined as the years between first and last encounter over 2005–2015) as the offset variable to normalize the total adjusted revenue to a per-year basis, the

incidence rate ratio for the total revenue per year in the FH cohort was 1.30 (95% Cl, 1.28–1.33; P<0.0001), after adjusting for age, sex, smoking, diabetes mellitus, hypertension, and maximum LDL-C. There were no significant differences in baseline characteristics of patients with missing financial data compared with those for whom data were available.

Medicare rate-adjusted median revenue for the FH cohort was 19% higher than the non-FH cohort for the year 2014 (\$1026 versus \$860; P<0.0001) and 28% higher in 2015 (\$1089 versus \$850; P<0.0001). These trends were maintained when the analysis was repeated with exclusion of possible FH (Tables S8 and S9).

Discussion

Our study is one of few based on integrated health system data with population coverage of >3 million residents, including several areas designated as rural, with longitudinal follow-up of >10 years. Using the modified DLCN criteria, an EHR-based algorithm in our population was able to identify an FH cohort within an already high-risk diagnosis of

Table 1. Univariate Comparison of Baseline Characteristics Between FH and Non-FH Hyperlipidemia Cohort

	FH Cohort		Non-FH Cohort			
	n	%	n	%	P Value	
Total	32 613	13.71	205 290	86.29		
Sex (male)	14 179	43.48	105 696	51.49	<0.0001	
Race (white, inclusive of Hispanic ethnicity)	31 860	97.69	198 135	96.51	<0.0001	
Smoking history (yes)	17 647	54.11	104 132	50.73	<0.0001	
Medications	·					
Statins	25 796	79.10	118 688	57.82	<0.0001	
High-potency statins*	13 747	42.15	40 008	19.49	<0.0001	
PCSK-9 inhibitors	59	0.18	51	0.02	<0.0001	
β-blockers	13 288	40.74	68 062	33.15	<0.0001	
Calcium channel blockers	7490	22.97	44 186	21.52	<0.0001	
ACE inhibitors	14 034	43.03	79 250	38.60	<0.0001	
Loop diuretics	6349	19.47	35 288	17.19	<0.0001	
Antiplatelets	5837	17.90	22 431	10.93	<0.0001	
Anticoagulants	4712	14.45	28 585	13.92	0.01	
Diagnostic tests			1	1		
ECG	23 740	72.79	132 775	64.69	<0.0001	
Echocardiogram	15 439	47.34	77 316	37.66	<0.0001	
	Median	IQR	Median	IQR	<i>P</i> Value	
Age	61	17	63	19	<0.0001	
Lipid profiles						
Total cholesterol						
Maximum [†]	264	40	194	57	<0.0001	
Average [‡]	200	42	161	45	<0.0001	
Delta [§]	131	68	58	70	<0.0001	
LDL cholesterol						
Maximum [†]	202	27	137	49	<0.0001	
Average [‡]	148	35	110	38	<0.0001	
Delta [§]	111	61	42	59	<0.0001	
HDL cholesterol						
Maximum [†]	57	21	54	22	<0.0001	
Average [‡]	49	17	48	19	<0.0001	
Delta [§]	18	15	12	16	<0.0001	
Triglycerides						
Maximum [†]	237	163	181	144	<0.0001	
Average [‡]	171	105	143	100	<0.0001	
Delta [§]	133	142	75	122	<0.0001	

ACE indicates angiotensin-converting enzyme; FH, familial hypercholesterolemia; HDL, high-density lipoprotein; IQR, interquartile range; LDL, low-density lipoprotein; PCSK-9, proprotein convertase subtilisin/kexin type 9.

*High potency is rosuvastatin \geq 20 mg, atorvastatin \geq 40 mg.

[†]Maximum indicates highest value for a patient for the cholesterol type.

 $^{\ddagger}\text{Average indicates (max+min)/2 for a patient for the cholesterol type.}$

[§]Delta indicates difference between maximum and minimum values for a patient for the cholesterol type.

Table 2. Univariate Comparison of Comorbidities BetweenFH and Non-FH Cohort

	FH Cohort		Non-FH Col			
	n	%	n	%	P Value	
Total	32 613	13.71	205 290	86.29		
Hypertension	21 318	65.37	133 137	64.85	0.07	
Renal disease	5659	17.35	30 958	15.08	<0.0001	
Cancer	5938	18.21	37 488	18.26	0.82	
Angina	2446	7.50	7354	3.58	<0.0001	
Myocardial infarction	4817	14.77	17 092	8.33	<0.0001	
Heart failure	3854	11.82	21 557	10.50	< 0.0001	
MACE*	9202	28.22 6.49	40 832	19.89 6.35	<0.0001 0.33	
Ischemic stroke	2118		13 044			
Ventricular arrhythmias	1701	5.22	9710	4.73	0.0001	
Percutaneous coronary intervention	3086	9.46	8150	3.97	<0.0001	
Coronary artery bypass grafting	2488	7.63	10 427	5.08	<0.0001	
Implantable cardioverter defibrillator	577	1.77	2857	1.39	<0.0001	

FH indicates familial hypercholesterolemia; MACE, major adverse cardiovascular events. *MACE is a composite of death, myocardial infarction, percutaneous coronary interventions, and coronary artery bypass grafting.

hyperlipidemia. Use of the modified DLCN criteria in this population-based manner is methodology not unique to our study.^{10,11} The designation of FH in the overall hyperlipidemia group had a prevalence of 13.7% and 2.7% of our entire EHR cohort. The prevalence of FH is variable^{4,10,12,13} depending on the population studied and case definitions used, ranging from 1:137 (0.7%)¹⁰ to 1:250 (0.4%).³ The SEARCH (Screening Employees and Residents in the Community for Hypercholesterolemia) study by Safarova et al¹¹ was based on 131 000 individuals seen in primary care practice and used a unique phenotyping algorithm using structured data and natural language processing for family history and presence of FH stigmata on physical examination for identification of FH in EHR and identified a prevalence of FH of 0.32%. This study comes the closest in methodology to our examination, but there are important differences. First, we used data from the entire spectrum of healthcare delivery systems, including outpatient as well as inpatient data, irrespective of the type of practice. Another notable distinction is our inclusion of "possible FH" in the FH cohort definition. Although this may lead to a degree of misclassification bias for FH (as evidenced by a total prevalence of 2.7% in the entire cohort), our approach was to be inclusive, as this strategy helped identify higher-risk individuals within the larger EHR cohort. We felt that this was prudent, as the ability of EHR data alone to accurately differentiate between possible and probable FH may be limited. Furthermore, we believe that being inclusive better suits the objective to preemptively screen patients to identify those likely to be at higher risk who would be candidates for more intensive clinical phenotyping to confirm FH status. A combination of only definite and probable FH groups in our study would yield a prevalence of 0.19% in our entire EHR population and 0.98% in the hyperlipidemia cohort. We report a higher proportion of female subjects in our FH cohort as has been reported previously in a meta-analysis of 6 large population-based studies that included 37 889 patients.14

The EHR-based FH designation in our study was associated with higher cholesterol levels and traditional comorbidities and correlated with MACE and all-cause mortality. Benn et al¹⁰ examined the prevalence of FH and the risk of cardiovascular disease in a population of 69 016 individuals from the Danish general population and reported an odds ratio of 13.2 (95% Cl, 10.0-17.4) for coronary artery disease for patients with FH not receiving lipid-modifying therapies and 10.3 (95% CI, 7.8-13.8) for patients with FH receiving lipid-modifying therapies, compared with patients without FH and not receiving lipid-modifying therapies. Perak et al¹⁵ reported that the FH phenotype was associated with substantially elevated 30-year coronary heart disease risk, with hazard ratios up to 5.0 (95% Cl, 1.1-21.7). Similar patterns of results were found for total atherosclerotic cardiovascular disease risk, with hazard ratios up to 4.1 (95% Cl, 1.2-13.4). The variable phenotypic expression of heterozygous FH is modulated in part by the underlying traditional cardiovascular risk factors,¹⁶ including obesity, diabetes mellitus, smoking, hypertension, male sex, and age, in addition to the risk associated with increased LDL-C. Using an ICD-based approach, we identified clinical outcomes that would represent the typical burden of atherosclerotic vascular disorder and combined them to define MACE for our study. To assess the utility of our EHR-based FH assignment in correlating to outcomes, we decided a priori to adjust for traditional risk factors, namely, age, sex, smoking status, diabetes mellitus, hypertension, and maximum LDL-C in our multivariate model. We observed odds ratios of 4.02 and 1.20 for MACE and mortality, respectively. These slightly lower odds ratios with regard to the studies cited above are expected, as our cohort is not derived from an FH diseasespecific database. More importantly, our comparison population also has hyperlipidemia, which constitutes a difference in methodology from the quoted studies and would lower the reported odds ratios.



Figure 2. Multivariate models showing outcomes of MACE, mortality, and cost of care for FH diagnosis in the hyperlipidemia cohort. *Adjusted for age, sex, smoking, diabetes mellitus, hypertension and LDL (max). EHR indicates electronic health record; FH, familial hypercholesterolemia; IRR, incidence rate ratio; LDL, low-density lipoprotein; MACE, major adverse cardiovascular events.

Our purpose with this investigation was to show the viability of a large population-based methodology to identify diverse FH phenotypes within an already high-risk group of hyperlipidemias and show that such a methodology has consistency in identifying phenotypes associated with adverse outcomes and increased costs of care. We report on differences in death, ischemic heart disease, congestive heart failure, and implantable cardiac defibrillator use. While worse cardiac outcomes such as myocardial infarction have been reported in FH patients previously,¹⁷ the data on ischemic stroke have been variable.¹⁸ This more expansive assessment of comorbidities allowed us to attempt to explain the higher costs of care.

Only 79% of our EHR FH cohort was on statin treatment and only 42% on high-potency statins. Undertreatment of FH as seen in our analysis has been widely reported previously.^{4,19–21} These data again underline the role preemptive identification can play in early and adequate treatment of these individuals.

In our study, using revenue data, we determined that the FH designation was associated with consistently higher median costs of care across 12 years of longitudinal data. FH

individuals can incur significant costs to the healthcare system over their lifetime attributable to premature atherosclerosis, and more so if not identified and treated.⁴ Prior studies exploring the economics of FH have been evaluations of screening strategies²² or therapeutic interventions,²³ but less is known about the overall economic burden associated with FH. This is a needed area of research to support FH-related economic evaluations. Our study provides evidence of higher total median costs (reflected in revenue) for FH patients almost twice that of their non-FH hyperlipidemia counterparts. This differential is maintained across time and appears to be expanding. Using an inverse probability of treatment weights multivariable model, the EHR FH flag was associated with higher median total revenue per year (incidence rate ratio, 1.30; P<0.0001). These results add to current knowledge about higher morbidity and associated cost of care for this high-risk population and can be helpful in studying the cost effectiveness of novel therapies. We believe that our approach is a novel way of assessing the hidden economic impact of FH in a hospital cohort and can be extrapolated to other health systems.

Table 3. Cost of Care: Comparison of Yearly Revenue for theYears 2005–2015 (US\$)

	Adjusted F	Adjusted Revenue 2005–2015								
	FH		Non-FH							
	Median	IQR	Median	IQR	P Value					
2005	810	1914	687	1586	<0.0001					
2006	852	2048	724	1697	< 0.0001					
2007	902	2186	752	1818	< 0.0001					
2008	920	2210	790	1868	<0.0001					
2009	983 2461		847	2070	< 0.0001					
2010	1043	2680	874	2184	<0.0001					
2011	1044	2627	867	2153 2439 2737	< 0.0001					
2012	1063	2989	907		<0.0001					
2013	1166	3370	974		< 0.0001					
2014	1294	4028	1093	3116	< 0.0001					
2015	1307	3818	1005	2846	< 0.0001					
Total adjusted revenue (2005–2015)	al adjusted 17 071 venue 005–2015)		43 024 11 178		<0.0001					
Med net revenu	Med net revenue									
2014	1026	2687	860	2152	<0.0001					
2015	1089	2788	850	2123	< 0.0001					

FH indicates familial hypercholesterolemia; IQR, interquartile range.

Cost-effectiveness analyses of a targeted genetic (cascade) screening program for relatives of patients with FH have been extensively published previously.^{24,25} However, the yield of an EHR-based phenotyping algorithm to identify FH within the larger hyperlipidemia cohort as performed in our study is less common and could represent an important missed opportunity to use already available clinical data, which could then be advanced to the next level of targeted screening in the most cost-effective manner. This potentially promising approach of using EHR-based phenotyping to identify FH patients and then screen family members to efficiently achieve targeted population-based FH screening is in need of additional evidence of effectiveness and costs for economic evaluation. Our approach is similar to that of the FH Foundation that recently launched the Find FH program, a machine-learning algorithm used to identify individuals with probable FH using EHR data, laboratory results, and claims databases.²⁶

Limitations

This is a retrospective observational study based on a population-based cohort in a single healthcare system that is geographically limited and ethnically homogenous. Therefore, generalizability of our results to other, more diverse populations may be limited. However, generalizability of our methodology and approach to identify these high-risk subjects is possible and desirable. Secondary to the retrospective nature of our study that used a preexisting general EHR, our FH phenotyping was likely affected by incomplete information regarding FH-specific physical examination findings (in particular tendon xanthomas and arcus) and family history. These missing data may lead to an underestimate of the true prevalence of probable and definite FH in our population. Similarly, the use of EHR data to determine outcomes has its limitations regarding the potential for missing data, but manual review of the records validated that this approach was effective in our system. GHS may represent an ideal case in this regard given the high number of patients receiving most or all of their care at GHS. Finally, estimating the absolute increase in cardiovascular risk resulting from FH is complicated, as case ascertainment using healthcare systembased EHR data is likely to be biased toward patients experiencing symptoms and cardiovascular events.²⁷ Nonetheless, this is the nature of the big-data approach and is likely balanced by the real-life pragmatic information being gathered by this methodology. Cost data were available for medical care utilization only within the GHS, and omitted medical care from other providers; however, it is unknown whether and the extent to which these additional services would change the relative cost outcomes.

Conclusion

Our study demonstrates a novel and pragmatic approach relying on standardized clinical criteria for identification of FH in a previously available EHR database. We were able to demonstrate the clinical significance of this approach by showing a statistically significant association between patients identified as having FH and adverse cardiovascular outcomes and higher costs of care, compared with those with hyperlipidemia not meeting the criteria for FH. The everincreasing use of EHR systems may broaden the appeal for such a population-based approach to identify high-risk patient groups, initiate guideline-based interventions, and improve clinical outcomes.

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Disclosures

None.

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SUPPLEMENTAL MATERIAL

 Table S1. Dutch Lipid Clinic Network diagnostic criteria for familial hypercholesterolemia*.

Sup	pplementary Table S1 : Dutch Lipid Clinic Network diagnostic criteria for familial hypercholesterolemia*					
Criteria		Points				
Family history	First-degree relative with known premature (men: <55 years; women: <60 years) coronary and vascular disease, or First-degree relative with known LDLC† above the 95th percentile					
	First-degree relative with tendinous xanthomata and/or arcus cornealis, or Children aged less than 18 years with LDLC above the 95th percentile	2				
Clinical history	Patient with premature (men: <55 years; women: <60 years) coronary artery disease	2				
Cirlical filstory	Patient with premature (men: <55 years; women: <60 years) cerebral or peripheral vascular disease	1				
Physical examination	Tendinous xanthomata	6				
T Thysical examination	Arcus cornealis prior to age 45 years	4				
	LDLC, ≥8.5	8				
Cholesterol levels	LDLC, 6.5–8.4	5				
(mmol/liter)	LDLC, 5.0–6.4					
	LDLC, 4.0–4.9					
DNA analysis	Functional mutation in the LDLR gene	8				
Diagnosis	(diagnosis is based on the total number of points obtained)					
A "definite"	FH† diagnosis requires more than 8 points					
A "probable	e" FH diagnosis requires 6–8 points					
A "possible	e" FH diagnosis requires 3–5 points					
* World He	alth Organization. Familial hypercholesterolemia—report of a second WHO Consultation. Geneva, Switzerland: World	Health				
Organizatio	on, 1999. (WHO publication no. WHO/HGN/FH/CONS/99.2). (15).					
† LDLC, lo	w density lipoprotein cholesterol; FH, familial hypercholesterolemia.					

Table S2. EHR based algorithm based on Dutch Lipid Clinic Network criteria to identify familial hypercholesterolemia.

Dutch Lipid Clinic Network Criteria	Implementation
First-degree relative with premature coronary and/or vascular disease (men \leq 55 years, women \leq 60 years)	 Searched for 'Heart Disease' in family history (not limited to first-degree relatives as this information was not always available) Searched for 'Family history of premature heart disease' in
	problem list
First-degree relative with known LDL-C \geq 95th percentile for age and sex	- Searched for 'Family history of hyperlipidemia in problem list
First-degree relative with tendon xanthomata and/or arcus cornealis,	- Data unavailable
Children aged \leq 18 years with known LDL-C \geq 95th percentile for age and sex	Used known mother/child links (available since 2010) to search for children with LDL \geq 95th percentile (LOINC: 13457-7, 18262-6, 2089-1, 55440-2 and lab result value between 230 and 90000.
Clinical History	
Patient with premature coronary artery disease (men \leq 55 years, women \leq 60 years)	Used electronic phenotyping to identify patients with premature coronary artery disease. Used P004 to pull these patients. A previously validated algorithm.
Patient with premature cerebral or peripheral vascular disease (men \leq 55 years, women \leq 60 years)	Searched for ICD9 diagnosis codes in encounters and problem list (PVD: 249.7, 249.70, 249.71, 250.70, 250.71, 250.73, 440.20, 440.21, 440.22, 440.23, 440.24, 443, 443.0, 443.1, 443.2, 443.21, 443.22, 443.23, 443.24, 443.29, 443.8, 443.81, 443.82, 443.89, 443.9, V12.59 CVD: 199.1, 436, 437, 437.0, 437.1, 437.8, 437.9, 438, 438.0, 438.1, 438.10, 438.11, 438.12, 438.13, 438.14, 438.19, 438.2, 438.20, 438.21, 438.22, 438.3, 438.30, 438.31, 438.32, 438.4, 438.40, 438.41, 438.42, 438.5, 438.50, 438.51, 438.52, 438.53, 438.6, 438.7, 438.8, 438.81, 438.82, 438.83, 438.84, 438.85, 438.89, 438.9, 674.0, 674.00, 674.02, V12.59) Patients had to have two or more diagnoses in their EHR.
Physical Examination	
Tendon xanthomata	

	Searched for ICD9 diagnosis codes in encounters and problem list (272.7). Patients had to have two or more diagnoses in their EHR.
Arcus cornealis at age \leq 45 years	Searched for ICD9 diagnosis codes in encounters and problem list (371.41)
LDL-C (mg/dl)	
	- Used maximum lifetime outpatient LDL-C level
LDL-C < 155	
$155 \ge LDL-C < 189$	
$190 \ge \text{LDL-C} < 249$	
$250 \ge \text{LDL-C} < 329$	
$LDL-C \ge 330$	
DNA Analysis – functional variant in LDLR, APOB or PCSK9 gene	
LDLR	
APOB	
PCSK9	

 Table S3. Count of Patients Meeting Each Definition.

	D • 4	Meeting criteria in	Not meeting criteria			
Dutch Lipid Clinic Network Criteria	Points	EHR, N (%)	in EHR, N (%)	Implemented using Modified criteria (below)		
				Searched for 'Heart Disease' in family history		
First-degree relative with premature				(not limited to first-degree relatives as this		
coronary and/or vascular disease (men ≤ 55	1	108,742 (45.70%)	129,161 (54.29%)	information was not always available)		
years, women ≤ 60 years)				Searched for 'Family history of premature heart		
				disease' in problem list		
First-degree relative with known LDL-C \geq				Searched for 'Family history of hyperlipidemia in		
95th percentile for age and sex	entile for age and sex 1 15 (0.009		237,888 (99.99%)	problem list		
First-degree relative with tendon xanthomata						
and/or arcus cornealis	2	N/A	N/A	Data unavailable		
				Used known mother/child links (available since		
				2010) to search for children with $LDL \ge 95$ th		
Children aged ≤ 18 years with known LDL-	2	0 (0.00%)*	237,903 (100.00%)	percentile (LOINC: 13457-7, 18262-6, 2089-1,		
$C \ge 95$ th percentile for age and sex				55440-2 where lab value between 230 AND		
				90000)		

Clinical History				
Patient with premature coronary artery disease (men \leq 55 years, women \leq 60 years)	2	9,809 (4.12%)	228,094 (95.87%)	Used electronic phenotyping to identify patients with premature coronary artery disease. Used P004 to pull these patients
Patient with premature cerebral or peripheral vascular disease (men ≤ 55 years, women ≤ 60 years)	1	2,722 (1.14%)	235,181 (98.85%)	Searched for ICD9 diagnosis codes in encounters and problem list (PVD: 249.7, 249.70, 249.71, 250.70, 250.71, 250.73, 440.20, 440.21, 440.22, 440.23, 440.24, 443, 443.0, 443.1, 443.2, 443.21, 443.22, 443.23, 443.24, 443.29, 443.8, 443.81, 443.82, 443.89, 443.9, V12.59 CVD: 199.1, 436, 437, 437.0, 437.1, 437.8, 437.9, 438, 438.0, 438.1, 438.10, 438.11, 438.12, 438.13, 438.14, 438.19, 438.2, 438.20, 438.21, 438.22, 438.3, 438.30, 438.31, 438.32, 438.4, 438.40, 438.41, 438.42, 438.5, 438.50, 438.51, 438.52, 438.53, 438.6, 438.7, 438.8, 438.81, 438.82, 438.83, 438.84, 438.85, 438.89, 438.9, 674.0, 674.00, 674.02, V12.59) Patients had to have two or more diagnoses in their EHR.

Physical Examination				
Tendon xanthomata	6	27 (0.00%)*	237,876 (99.99%)	Searched for ICD9 diagnosis codes in encounters and problem list (272.7). Patients had to have two or more diagnoses in their EHR.
Arcus cornealis at age \leq 45 years	4	1 (0.00%)*	237,902 (99.99%)	Searched for ICD9 diagnosis codes in encounters and problem list (371.41). Patients had to have two or more diagnoses in their EHR.
LDL-C (mg/dl)				
None recorded	-	9,573 (4.02%)	-	Used maximum lifetime outpatient LDL-C level (LOINC 13457-7, 18262-6, 2089-1, 55440-2)
LDL-C < 155	0	138,351 (58.12%)	-	
155 ≥ LDL-C < 189	1	61,955(26.04%)	-	
$190 \ge \text{LDL-C} < 249$	3	25,494 (10.72%)	-	
250 ≥ LDL-C < 329	5	2,178 (0.92%)	-	

$LDL-C \ge 330$	8	352 (0.15%)	-						
DNA Analysis – functional variant in LDLR, APOB or PCSK9 gene (variant data was not available for the entire cohort so %s are not presented).									
LDLR	8	85	-						
APOB	8	76	-						
PCSK9	8	19	-						

Definition 1	Family his	story of hea	rt diseas	se	Sensitivity 92.37			
	Def1+	Def1-			Specificity	98.32%		
Def1+	12	21	2		PPV	98.37%		
Def1-	1	10	117		NPV	92.13%		
	13	31	119	250	Accuracy	95.20%		
Definition 2*	First-degr LDL-C ≥ 95t	ee relative w h percentile sex	ith know for age	wn and	Sensitivity	0.00%		
	Def2+	Def2-			Specificity	100.00%		
Def2+		0	0		PPV	N/A		
Def2-		2	248		NPV	99.20%		
		2	248	250	Accuracy	99.20%		
Definition 5	Patient wit	th premature ease (men ≤ men ≤ 60 ve	Sensitivity	93 94%				
Dominion o	Def5+	Def5-	uloj		Specificity	99 54%		
Def5+	3	31	1		PPV	96.88%		
Def5-		2	216		NPV	99.08%		
	3	33	217	250	Accuracy	98.80%		
					·			
	Patient with peripheral v	lor ∋n≤						
Definition 6	55 years	55 years, women ≤ 60 years)			Sensitivity	92.86%		
	Def6+	Def6-			Specificity	100.00%		
Def6+	1	13	0		PPV	100.00%		
Def6-		1	236		NPV	99.58%		
	1	4	236	250	Accuracy	99.60%		

Table S4. PPV, NPV, Sensitivity and Specificity to Definitions.

*analysis affected by unavailable data in the EHR for this field so modified criteria (Definition 1) was used

Broken Down by Groups and Over All Sensitivity 95.16% Unlikely+ Unlikely-Specificity 100.00% 0 PPV Unlikely+ 118 100.00% 126 Unlikely-6 NPV 95.45% 124 126 250 97.60% Accuracy Sensitivity 100.00% Possible+ Possible-Specificity 99.29% Possible+ 110 1 PPV 99.10% 139 NPV Possible-0 100.00% 110 140 250 99.60% Accuracy Sensitivity 100.00% Probable+ Probable-100.00% Specificity Probable+ 13 0 PPV 100.00% 0 237 NPV Probable-100.00% 13 237 250 Accuracy 100.00% 100.00% Sensitivity Definitive+ **Definitive-**Specificity 100.00% 2 0 PPV Definitive+ 100.00% Definitive-0 248 NPV 100.00% 248 250 100.00% 2 Accuracy Sensitivity 95.42% 99.16% FH All+ FH All-Specificity FH All+ 125 1 PPV 99.21% FH All-118 NPV 6 95.16% 131 119 250 97.20% Accuracy

Table S5. PPV, NPV, Sensitivity and Specificity by FH Category.

			EHR_F		CHART											
		EHR_FH	Н	CHART	FH_SC											
Ptid	PosNeg	CAT	SCORE	FH_CAT	R	DEF1	DEF2	DEF3	DEF4	DEF5	DEF6	DEF7	DEF8	DEF9	DEF10	ValidYN
1	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
2	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
3	0	UNLIKELY	2	POSSIBLE	3	0	0	0	0	1	0	0	0	0	0	0
4	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
5	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
6	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
7	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
8	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
9	1	POSSIBLE	3			1	0	0	0	0	1	0	0	1	0	1
10	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
11	1	POSSIBLE	3			1	0	0	0	1	0	0	0	0	0	1
12	0	UNLIKELY	2	POSSIBLE	3	0	0	0	0	1	0	0	0	0	0	0
13	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
14	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
15	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
16	0	UNLIKELY	2	POSSIBLE	3	0	0	0	0	1	0	0	0	0	0	0
17	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
18	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
19	1	POSSIBLE	5			0	0	0	0	1	0	0	0	1	0	1
20	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
21	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
22	0	UNLIKELY	2	POSSIBLE	3	0	0	0	0	0	1	0	0	1	0	0
23	0	UNLIKELY	2	POSSIBLE	3	0	0	0	0	1	0	0	0	0	0	0
24	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
25	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
26	0	UNLIKELY	2			0	0	0	0	1	0	0	0	0	0	1
27	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
28	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
29	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
30	0	UNLIKELY	0		Ī	0	0	0	0	0	0	0	0	0	0	1

 Table S6. Validation of DLCN-derived EHR FH algorithm by manual chart review.

31	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
32	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
33	0	UNLIKELY	1	POSSIBLE	3	1	0	0	0	0	0	0	0	0	0	0
34	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
35	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
36	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
37	1	PROBABLE	6			1	0	0	0	0	0	0	0	1	0	1
38	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
39	0	UNLIKELY	2			0	0	0	0	1	0	0	0	0	0	1
40	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
41	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
42	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
43	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
44	1	PROBABLE	6			1	0	0	0	1	1	0	0	1	0	1
45	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
46	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
47	0	UNLIKELY	1			0	0	0	0	0	0	0	0	1	0	1
48	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
49	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
50	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
51	1	PROBABLE	6			1	0	0	0	1	0	0	0	1	0	1
52	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
53	0	UNLIKELY	2			0	0	0	0	1	0	0	0	0	0	1
54	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
55	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
56	0	UNLIKELY	1			0	0	0	0	0	0	0	0	1	0	1
57	1	POSSIBLE	4			1	0	0	0	1	1	0	0	1	0	1
58	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
59	0	UNLIKELY	1			0	0	0	0	0	0	0	0	1	0	1
60	0	UNLIKELY	0	UNLIKELY	1	0	0	0	0	0	0	0	0	0	0	0
61	0	UNLIKELY	1			0	0	0	0	0	0	0	0	1	0	1
62	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
63	1	POSSIBLE	3			1	0	0	0	0	1	0	0	1	0	1
64	0	UNLIKELY	1			0	0	0	0	0	0	0	0	1	0	1
65	1	POSSIBLE	5			0	0	0	0	0	0	0	0	1	0	1

		DEFINITIV		DEFINITIV												
66	1	E	11	Е	9	1	0	0	0	1	0	0	0	1	0	0
67	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
68	0	UNLIKELY	1	UNLIKELY	1	0	0	0	0	0	0	0	0	1	0	0
69	0	UNLIKELY	0	UNLIKELY	1	0	0	0	0	0	0	0	0	0	0	0
70	1	POSSIBLE	3			1	0	0	0	1	0	0	0	0	0	1
71	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
72	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
73	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
74	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
75	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
76	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
77	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
78	1	POSSIBLE	5			1	0	0	0	0	1	0	0	1	0	1
79	1	POSSIBLE	3			1	0	0	0	1	0	0	0	0	0	1
80	0	UNLIKELY	1			0	0	0	0	0	0	0	0	1	0	1
81	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
82	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
83	0	UNLIKELY	1	UNLIKELY	2	1	0	0	0	0	0	0	0	0	0	0
84	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
85	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
86	1	POSSIBLE	4			1	0	0	0	1	0	0	0	1	0	1
87	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
88	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
89	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
90	0	UNLIKELY	1			0	0	0	0	0	0	0	0	1	0	1
91	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
92	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
93	0	UNLIKELY	2			1	0	0	0	0	1	0	0	0	0	1
94	0	UNLIKELY	2			0	0	0	0	1	0	0	0	0	0	1
95	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
96	1	POSSIBLE	3			1	0	0	0	1	0	0	0	0	0	1
97	0	UNLIKELY	0	UNLIKELY	1	0	0	0	0	0	0	0	0	0	0	0
98	1	PROBABLE	6			1	0	0	0	1	0	0	0	1	0	1
99	0	UNLIKELY	1	UNLIKELY	0	1	0	0	0	0	0	0	0	0	0	0

100	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
101	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
102	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
103	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
104	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
105	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
106	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
107	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
108	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
109	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
110	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
111	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
112	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
113	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
114	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
115	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
116	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
117	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
118	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
119	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
120	0	UNLIKELY	2			0	0	0	0	1	0	0	0	0	0	1
121	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
122	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
123	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
124	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
125	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
126	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
127	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
128	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
129	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
130	0	UNLIKELY	0	UNLIKELY	1	0	0	0	0	0	0	0	0	0	0	0
131	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
132	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
133	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
134	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1

135	0	UNLIKELY	1	UNLIKELY	2	0	0	0	0	0	1	0	0	0	0	0
136	0	UNLIKELY	0	UNLIKELY	1	0	0	0	0	0	0	0	0	0	0	0
137	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
138	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
139	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
140	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
141	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
142	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
143	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
144	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
145	0	UNLIKELY	2			1	0	0	0	0	0	0	0	1	0	1
146	0	UNLIKELY	2			0	0	0	0	1	0	0	0	0	0	1
147	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
148	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
149	0	UNLIKELY	0			0	0	0	0	0	0	0	0	0	0	1
150	0	UNLIKELY	1			1	0	0	0	0	0	0	0	0	0	1
151	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
152	1	PROBABLE	6			1	0	0	0	0	0	0	0	1	0	1
153	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
154	1	POSSIBLE	3	UNLIKELY	2	1	0	0	0	0	1	0	0	1	0	0
155	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
156	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
157	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
158	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
159	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
160	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
161	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
162	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
163	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1
164	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
165	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
166	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
167	1	PROBABLE	6			1	0	0	0	0	0	0	0	1	0	1
168	1	POSSIBLE	4			1	0	0	0	0	0	0	0	1	0	1
169	1	POSSIBLE	3			0	0	0	0	0	0	0	0	1	0	1

170	1	PROBABLE	6		1	0	0	0	0	0	0	0	1	0	1
171	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
172	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
173	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
174	1	POSSIBLE	5		0	0	0	0	0	0	0	0	1	0	1
175	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
176	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
177	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
178	1	POSSIBLE	3		1	0	0	0	1	0	0	0	0	0	1
179	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
180	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
181	1	POSSIBLE	3		1	0	0	0	1	0	0	0	0	0	1
182	1	POSSIBLE	3		1	0	0	0	1	1	0	0	0	0	1
183	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
184	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
185	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
186	1	POSSIBLE	3		1	0	0	0	1	0	0	0	0	0	1
187	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
188	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
189	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
190	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
191	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
192	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
193	1	POSSIBLE	3		1	0	0	0	1	0	0	0	0	0	1
194	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
195	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
196	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
197	1	POSSIBLE	3		1	0	0	0	1	1	0	0	0	0	1
198	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
199	1	PROBABLE	6		1	0	0	0	1	0	0	0	1	0	1
200	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
201	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
202	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
203	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
204	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1

205	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
206	1	PROBABLE	6		1	0	0	0	0	0	0	0	1	0	1
207	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
208	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
209	1	PROBABLE	7		1	0	0	0	0	1	0	0	1	0	1
210	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
211	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
212	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
213	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
214	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
215	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
216	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
217	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
218	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
219	1	POSSIBLE	5		1	0	0	0	0	1	0	0	1	0	1
220	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
221	1	PROBABLE	8		0	0	0	0	0	0	0	0	1	0	1
222	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
223	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
224	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
225	1	PROBABLE	6		1	0	0	0	0	0	0	0	1	0	1
226	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
227	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
		DEFINITIV													
228	1	E	9		1	0	0	0	0	0	0	0	1	0	1
229	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
230	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
231	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
232	1	POSSIBLE	3		0	0	0	0	1	0	0	0	1	0	1
233	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
234	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
235	1	POSSIBLE	3		1	0	0	0	1	0	0	0	0	0	1
236	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
237	1	POSSIBLE	3		1	0	0	0	1	0	0	0	0	0	1
238	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1

239	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
240	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
241	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
242	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
243	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
244	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
245	1	PROBABLE	8		0	0	0	0	0	0	0	0	1	0	1
246	1	POSSIBLE	4		1	0	0	0	0	0	0	0	1	0	1
247	1	POSSIBLE	5		0	0	0	0	0	0	0	0	1	0	1
248	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1
249	1	POSSIBLE	4		1	0	0	0	1	0	0	0	1	0	1
250	1	POSSIBLE	3		0	0	0	0	0	0	0	0	1	0	1

 Table S7. International Classification of Disease-9 codes used to identify the outcomes.

Diagnosis	ICD 9 Codes
Myocardial Infarction	410.00, 410.0, 410.01, 410.02, 410.1, 410.10, 410.11, 410.12, 410.20, 410.2, 410.21, 410.22, 410.30, 410.31, 410.32, 410.40, 410.4, 410.41, 410.42, 410.50, 410.51, 410.52, 410.60, 410.61, 410.62, 410.7, 410.70, 410.71, 410.72, 410.8, 410.80, 410.81, 410.82, 410.9, 410.90, 410.91, 410.92, 411.0, 412, 429.7, 429.71, 429.79
Heart Failure	398.91, 428.0, 428, 428.1, 428.20, 428.2, 428.21, 428.22, 428.23, 428.3, 428.30, 428.31, 428.32, 428.33, 428.40, 428.4, 428.41, 428.42, 428.43, 428.9
Ischemic Stroke	433.01, 433.11, 433.21, 433.31, 433.81, 433.91, 434.01, 434.11, 434.91
PCI	00.66, 17.55, 36.02, 36.03, 36.04, 36.05, 36.06, 36.07, 36.09, 36.01, C9600, C9601, C9602, C9604, C9605, C9606, C9607, G0290
CABG	36.10, 36.11, 36.12, 36.13, 36.14, 36.15, 36.16, 36.19
ICD	89.49, 37.94, 37.95, 37.96, 37.97, 37.98, C1721, C1722, C1777, C1882, C1895, C1896

		FH	1	Non-FH	P-value
	Median	IQR	Median	IQR	
Total adjusted	16425	41595	11153	30844	< 0.0001
Revenue (2005-2015)					
		FH	1	Non-FH	
Med Net Revenue	Median	IQR	Median	IQR	p-value
2014	1007	2586	860	2152	< 0.0001
2015	1065	2709	850	2123	< 0.0001

Table S8. Cost Analysis -DEFINITIVE and PROBABLE categories combined as FH and compared to UNLIKELY (POSSIBLE excluded).

		FH	N	Non-FH	P-value
	Median	IQR	Median	IQR	
Total adjusted	16425	41595	11252	31198	< 0.0001
Revenue (2005-2015)					
		FH	1	Non-FH	
Med Net Revenue	Median	IQR	Median	IQR	p-value
2014	1007	2586	864	2171	< 0.0001
2015	1065	2709	854	2141	< 0.0001

Table S9. Cost Analysis - DEFINITIVE and PROBABLE combined as FH and compared to UNLIKELY and POSSIBLE (combined as Non-FH).