

## Prediction of Pediatric Death in the Year after Hospitalization: A Population-Level Retrospective Cohort Study

Chris Feudtner, M.D., Ph.D., M.P.H.,<sup>1,5</sup> Kari R. Hexem, M.P.H.,<sup>1</sup> Mayadah Shabbout, M.S.,<sup>3</sup> James A. Feinstein, M.D.,<sup>1</sup> Julie Sochalski, Ph.D., R.N.,<sup>4,5</sup> and Jeffery H. Silber, M.D., Ph.D.<sup>2,5</sup>

### Abstract

**Background:** The study of how the quality of pediatric end-of-life care varies across systems of health care delivery and financing is hampered by lack of methods to adjust for the probability of death in populations of ill children.

**Objective:** To develop a prognostication models using administratively available data to predict the probability of in-hospital and 1-year postdischarge death.

**Methods:** Retrospective cohort study of 0–21 year old patients admitted to Pennsylvania hospitals from 1994–2001 and followed for 1-year postdischarge mortality, assessing logistic regression models ability to predict in-hospital and 1-year postdischarge deaths.

**Results:** Among 678,365 subjects there were 2,202 deaths that occurred during the hospitalization (0.32% of cohort) and 860 deaths that occurred 365 days or less after hospital discharge (0.13% of cohort). The model predicting hospitalization deaths exhibited a C statistic of 0.91, with sensitivity of 65.9% and specificity of 92.9% at the 99th percentile cutpoint; while the model predicting 1-year postdischarge deaths exhibited a C statistic of 0.92, with sensitivity of 56.1% and specificity of 98.4% at the 99th percentile cutpoint.

**Conclusions:** Population-level mortality prognostication of hospitalized children using administratively available data is feasible, assisting the comparison of health care services delivered to children with the highest probability of dying during and after a hospital admission.

### Introduction

THE PROVISION OF PEDIATRIC PALLIATIVE CARE in the United States occurs in a context in which the death of children due to a medical condition is an infrequent event.<sup>1</sup> Because of this rarity, the study of pediatric palliative and end-of-life care benefits when conducted from the perspective of large populations, both providing an adequate sample size and also enabling the comparison, among hospitals and across health care systems, of processes of care and of outcomes. In order for studies of this regional or national scale to be feasible, economically and logistically, pediatric researchers use preexisting data, often involving hospital administrative data.<sup>2–6</sup>

An analytic linchpin in the conduct of such population-level health services research studies is the ability to adjust

for individual patient characteristics so that comparisons of hospitals and health care systems in fact compare differences in processes of care, and not merely detect confounded associations arising from differences in the clinical characteristics of patients cared for by these hospitals or systems.<sup>7,8</sup> In the particular case of pediatric palliative care health services research, a key concern is the ability to predict the probability that a given type patient will die based on the available demographic and clinical information. Sufficiently accurate calculation of the probability of mortality would enable comparison of the processes of care, such as receipt of palliative or hospice services, or of specific medical interventions, or location of death at home or in a hospital, among patients with similar likelihood of dying.

Importantly, for this form of analytic comparison, emphasis is placed on the phrase “sufficiently accurate,” which

<sup>1</sup>Pediatric Advanced Care Team, <sup>2</sup>Center for Outcomes Research, The Children’s Hospital of Philadelphia, Philadelphia, Pennsylvania.

<sup>3</sup>GlaxoSmithKline, Philadelphia, Pennsylvania.

<sup>4</sup>School of Nursing, <sup>5</sup>The Leonard Davis Institute, The University of Pennsylvania, Philadelphia, Pennsylvania.

implies some degree of tolerance of individual-level predictive error, requiring instead that the group-level predictions of mortality are well-calibrated.<sup>9</sup> This tolerance contrasts with the need for virtually perfect accuracy for any scheme of mortality prediction used to guide the care of individual patients, a goal that has eluded various mortality risk scoring systems in the setting of hospital intensive care units.<sup>10-15</sup>

Guided by this line of motivation, we sought to develop, evaluate, and validate a practical pediatric mortality prediction model, based on administrative hospitalization data drawn from a retrospective cohort of children admitted to all Pennsylvania hospitals during a 6-year period and who were assessed for mortality during the year after discharge from the hospital. We used two models, one to predict in-hospital deaths and another to predict 1-year postdischarge deaths (Fig. 1). We also aimed to compare the parsimonious model, the components of which were based on our prior population-level research regarding pediatric mortality and end-of-life care, with similar models that also contained severity of illness and risk of mortality scores that were based on more detailed laboratory and physiologic data about patients at the time of admission to the hospital and on the full complement of diagnoses at the time of discharge.

**Methods**

**Study design**

We conducted a retrospective cohort study of persons less than 21 years of age who were hospitalized in the Commonwealth of Pennsylvania between 1995 and 2000, with 1 year of prehospitalization observation (or less, for subjects

under a year of age) to assess for previous hospitalizations, and 1-year of posthospitalization discharge observation to assess 1-year postdischarge mortality. This study was approved by The Children’s Hospital of Philadelphia Committee for the Protection of Human Subjects.

**Data source**

We used data specially prepared by the Pennsylvania Healthcare Cost Containment Council (PHC4). PHC4 matched the administrative hospital record data to death certificate data. We excluded records of normal newborns, patients discharged from psychiatric facilities, and patients who died due to trauma, homicides, suicides, and sudden infant death syndrome.

**Selection of index hospitalization and study time**

If a patient had more than one hospitalization between 1995 and 2000, we randomly selected a single hospitalization as the index hospitalization. The study time frame for each subject extended from the 365 days prior to the day of admission to the index hospitalization, to 365 days after the index hospitalization discharge.

**Outcome of interest**

Patients who were observed to have died 365 days or less after the index hospitalization discharge had the outcome of interest, which were subclassified as having either occurred during the index hospitalization (“hospitalization death”) or during the ensuing 365 days (“1-year postdischarge death”).

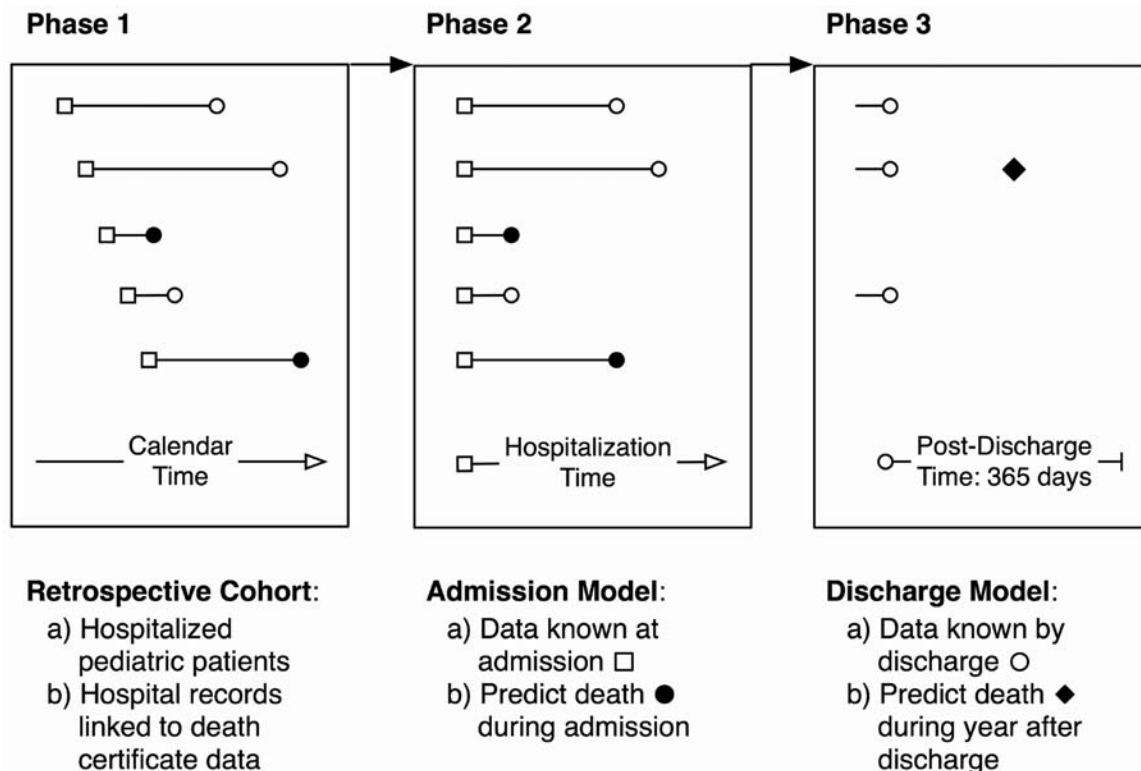


FIG. 1. Retrospective cohort study design for both models.

### Predictor variables

Predictor variables from the index hospitalization were conceptualized as either known at the time of admission to the hospital or at the time of discharge from the hospitalization. Data known at the time of admission included demographic information, certain historical clinical information (including any diagnoses during a prior admission of a Complex Chronic Condition [CCC] based on the *International Classification of Disease-9 Clinical Modification* codes<sup>16-19</sup>), and the facility type for the index hospitalization (classified as pediatric hospital or other). Data known by the end of the index hospitalization included that hospitalization's length of stay, CCC diagnoses, the All Patient Refined-Diagnosis Related Group (APR-DRG), and the APR-DRG-specific risk of mortality (ROM) index rating. The APR-DRG major diagnostic codes were included as 24 indicator variables, and interacted with the APR-DRG specific risk of mortality score, which were ranked ordinally from 0 to 4.

### Primary statistical analysis

Two "base" logistic regression models were built to estimate the likelihood of having the outcome of in-hospital and 1-year postdischarge deaths 365 days or less after the index hospitalization discharge. Covariates in both models included information that could be known at the time of hospital admission, as well as indicator variables for 10 of the largest Pennsylvania hospitals as fixed effects. The "admission model" did not contain length of stay or the APR-DRG and APR-ROM measures, since the information upon which these are based is known only at the time of discharge. The "discharge model" did include these variables. These models were then compared to models that included the physiologic MediQual Severity Index.

To evaluate and compare the fit for each of the models, we evaluated each model's C statistic, Brier score<sup>20</sup> and Hosmer-Lemeshow test statistic.<sup>21</sup> To demonstrate the performance of the chosen model, we also provide a classification of the observed and predicted readmission rates and the sensitivity and specificity at the 95th and 99th percentile probability of death cutpoints.

The models were developed using a random selection of cases comprising half of the original sample, with subsequent assessment of the models on the other "validation" half of the sample. Since the model-fit assessments from the validation subsample were almost identical to the development subsample (for the hospitalization model, C statistics of 0.907 for the development subsample and 0.908 for the validation subsample; for the 1-year postdischarge model, 0.928 for development subsample, and 0.940 for the validation subsample), we report the results using the entire sample.

### Additional statistical analyses

To assess for potential bias arising from missing data, we performed multiple imputation (using the Stata "ice" command) to compare C statistic results of the models in datasets with imputed missing data. To assess whether the performance of the final prediction model depended upon the particular composition of our dataset, we performed internal validation using the regular bootstrap with 1,000 replications using resampling with replacement.<sup>22</sup>

All analyses were performed using SAS version 9.1 (SAS Institute Inc, Cary, NC) and Stata SE version 10.0 (Stata Corp, College Station, TX). Due to large sample size, *p* values less than 0.001 were considered statistically significant.

## Results

### Characteristics of subjects in the cohort

Among the 678,861 hospitalization records of patients between the ages of 0 and 21 years who were hospitalized in Pennsylvania between 1995 and 2000, there were 2,202 patients (0.32%) who died during the index hospitalization, and 860 (0.13%) patients who died 1 year or less postdischarge from their index hospitalization discharge (Table 1).

### Predictive performance of the hospitalization death models

The C statistic for the hospitalization death model without the severity index was 0.912, and with the severity index was 0.937 (Table 2), both high values that suggest "outstanding" discriminative predictive performance.<sup>21</sup> Brier scores (with the lowest possible score of 0 connoting perfect prediction) were identical (0.0031) for both models (Table 2). At the 95th and 99th percentiles of the probabilities of death predicted by both models, the specificity of these cutoff values exceeded 92.9%, with sensitivity ranging from 21.6% to 30.7% at the 99th percentile cutoff and from 65.9% to 71.6% at the 95th percentile, for the models without and with the severity score, respectively (Table 2).

To assess the calibration of the hospitalization deaths models, we plotted the degree of correspondence between the observed proportions of patients who actually died across the range of the predicted probabilities of death (Fig. 2, upper panel), demonstrating a high degree of accuracy of the predictions for both models across the range of predicted probabilities; the model without the severity index generated a small range of predicted mortality values, 0.00006 to 0.3060, whereas the model with the severity index generated values of predicted mortality extending to 0.7888. We also divided the sample into 10 equally sized groups, based on the predicted probability, tabulated the observed number of deaths and the predicted number of deaths in each group (Table 3). This analysis revealed that the model without the severity index was more accurately calibrated than the model with the severity index. The Hosmer-Lemeshow test detected a statistically significant degree of miscalibration in both models, due to the extremely large sample size of the models, as the differences between the observed and expected values within each group are relatively small.

### Predictive performance of the one-year postdischarge models

The C statistic for the 1-year postdischarge death model without the severity index was 0.920, and the C statistic for the model with the severity index was 0.924. Brier scores for both models were very similar, 0.0012 without the severity index and 0.0013 with the severity index. At the 95th and 99th percentile and of 56.1% of the predicted probability for both models, the specificity at these cutoff values exceeded 98.3%, with sensitivity of 32.2% to 33.4% at the 99th percentile and of 56.1% to 57.8% at the 95th percentile cutoff,

TABLE 1. DEMOGRAPHIC CHARACTERISTICS OF PENNSYLVANIA PEDIATRIC SUBJECTS WHO DIED THREE HUNDRED SIXTY-FIVE DAYS OR LESS AFTER HOSPITAL DISCHARGE

	n =	Total 678,635 (100%)	Death in hospital		Death after discharge	
			2202 (0.32%)	p-value <sup>b</sup>	860 (0.13%)	p-value <sup>b</sup>
Age	<1 month	59,351 (8.8%)	1,496 (2.5%)	< 0.0001	12 (0.02%)	< 0.0001
	1–11 months	99,983 (14.7%)	330 (0.3%)		131 (0.13%)	
	1–9 y	199,734 (29.4%)	177 (0.1%)		333 (0.17%)	
	10–18 y	218,912 (32.3%)	137 (0.1%)		272 (0.12%)	
	19–20 y	100,655 (14.8%)	62 (0.1%)		112 (0.11%)	
Gender	Female	354,945 (52.3%)	962 (0.3%)	< 0.0001	403 (0.11%)	0.001
	Male	323,613 (47.7%)	1,240 (0.4%)		457 (0.14%)	
	Missing	77 (0.01%)				
Race	White	449,250 (66.2%)	1,365 (0.3%)	< 0.0001	639 (0.14%)	0.007
	Black	122,010 (18.0%)	356 (0.3%)		135 (0.11%)	
	Other	33,587 (5.0%)	145 (0.4%)		34 (0.10%)	
	Missing	73,788 (10.9%)				
Year admitted <sup>a</sup>	1995	127,756 (18.8%)	430 (0.3%)	0.279	149 (0.12%)	0.586
	1996	116,297 (17.1%)	336 (0.3%)		160 (0.14%)	
	1997	111,050 (16.7%)	368 (0.3%)		132 (0.12%)	
	1998	103,806 (15.3%)	337 (0.3%)		129 (0.12%)	
	1999	110,718 (16.3%)	336 (0.3%)		158 (0.14%)	
	2000	109,008 (16.1%)	395 (0.4%)		132 (0.12%)	
Payer	None	22,600 (3.3%)	97 (0.4%)		11 (0.05%)	
	Medicaid	178,092 (26.2%)	512 (0.3%)		247 (0.14%)	
	Blue Cross	226,415 (33.4%)	760 (0.3%)		279 (0.12%)	
	HMO	138,102 (20.4%)	435 (0.3%)		155 (0.11%)	
	Other <sup>a</sup>	110,663 (16.3%)	397 (0.4%)		166 (0.15%)	
	Missing	2,763 (0.41%)				
Median Income for zip code	≤\$24,000	57,761 (8.5%)	167 (0.3%)		38 (0.07%)	< 0.0001
	\$24,001–\$29,000	102,243 (15.1%)	310 (0.3%)		112 (0.11%)	
	\$29,001–\$35,000	131,076 (19.3%)	415 (0.3%)		177 (0.14%)	
	\$35,001–\$46,000	156,071 (23.0%)	540 (0.4%)		214 (0.14%)	
	\$46,001–\$60,000	83,692 (12.3%)	327 (0.4%)		119 (0.14%)	
	>\$60,001	60,578 (8.9%)	225 (0.4%)		58 (0.10%)	
	Missing	87,214 (12.9%)				
Complex chronic conditions	None	642,563 (94.7%)	1,688 (0.3%)		637 (0.10%)	< 0.0001 <sup>c</sup>
	Neuromuscular	5,545 (0.8%)	45 (0.8%)		35 (0.64%)	
	Cardiovascular	10,154 (1.5%)	254 (2.5%)		39 (0.39%)	
	Respiratory	2,117 (0.31%)	31 (1.5%)		8 (0.38%)	
	Metabolic	749 (0.1%)	17 (2.3%)		3 (0.41%)	
	Congenital or Genetic	1,808 (0.3%)	76 (4.2%)		6 (0.35%)	
	Malignancy	6,611 (1.0%)	68 (1.0%)		116 (1.77%)	
	Other	7,100 (1.1%)	14 (0.2%)		11 (0.16%)	
	Missing	1,988 (0.3%)				
Number of prior hospital admissions	0	609,788 (89.9%)	2,032 (0.3%)		303 (0.05%)	< 0.0001
	1	46,205 (6.8%)	68 (0.2%)		158 (0.34%)	
	2	12,037 (1.8%)	23 (0.2%)		107 (0.89%)	
	3+	10,605 (1.6%)	79 (0.7%)		292 (2.78%)	
Previous complex chronic condition	No	672,894 (99.2%)	2,136 (0.3%)		599 (0.09%)	< 0.0001
	Yes	5,741 (0.9%)	66 (1.2%)		261 (4.60%)	
Length of stay (days)	<2 days	206,256 (30.4%)	894 (0.4%)		156 (0.08%)	
	2–7 days	407,152 (60.0%)	560 (0.1%)		452 (0.11%)	
	8–14 days	37,346 (5.5%)	234 (0.6%)		132 (0.36%)	
	15+ days	27,974 (4.1%)	514 (1.8%)		120 (0.44%)	
Facility type	Pediatric	175,673 (25.9%)	945 (0.5%)		387 (0.22%)	< 0.0001
	General	502,962 (74.1%)	1,257 (0.3%)		473 (0.09%)	
Severity score	0	303,259 (44.7%)	139 (0.05%)		139 (0.05%)	< 0.0001
	1	162,785 (24.0%)	263 (0.2%)		309 (0.19%)	
	2	23,284 (3.4%)	311 (1.3%)		154 (0.67%)	
	3	6,736 (1.0%)	389 (5.8%)		81 (1.28%)	
	4	1,030 (0.2%)	203 (19.7%)		9 (1.09%)	
	Missing	181,541 (26.8%)				

<sup>a</sup>The category “other insurance” included self-insured individuals.

<sup>b</sup>Univariable logistic regression.

<sup>c</sup>Metabolic and congenital were grouped together for the  $\chi^2$  test.

TABLE 2. PERFORMANCE STATISTICS FOR MODELS

Model	C statistic	Brier score	Occurrence of death at 95th and 99th percentiles of predicted probability		
				95%	99%
Hospitalization death					
A. Without severity score	0.9122	0.0031	Sensitivity	65.9%	21.6%
			Specificity	92.9%	99.2%
B. With severity score	0.9364	0.0031	Sensitivity	71.6%	30.7%
			Specificity	94.5%	99.2%
1-year postdischarge death					
A. Without severity score	0.9204	0.0012	Sensitivity	56.1%	32.2%
			Specificity	98.4%	99.7%
B. With severity score	0.9238	0.0013	Sensitivity	57.8%	33.4%
			Specificity	98.3%	99.7%

for the models without and with the severity score, respectively (Table 2).

To assess the calibration of the 1-year postdischarge death models, we again plotted the degree of correspondence between the observed proportions of patients who died across the range of the predicted probabilities of death (Fig. 2, lower panel), demonstrating a high-degree of accuracy of the pre-

dictions for both models across the range of predicted probabilities extending from 0 to 0.6552 for the model without the severity index, and extending to 0.7630 for the model with the severity index. We divided the sample into 10 equally sized groups (Table 3); this analysis again revealed that the model without the severity index was more accurately calibrated than the model with the severity index (and

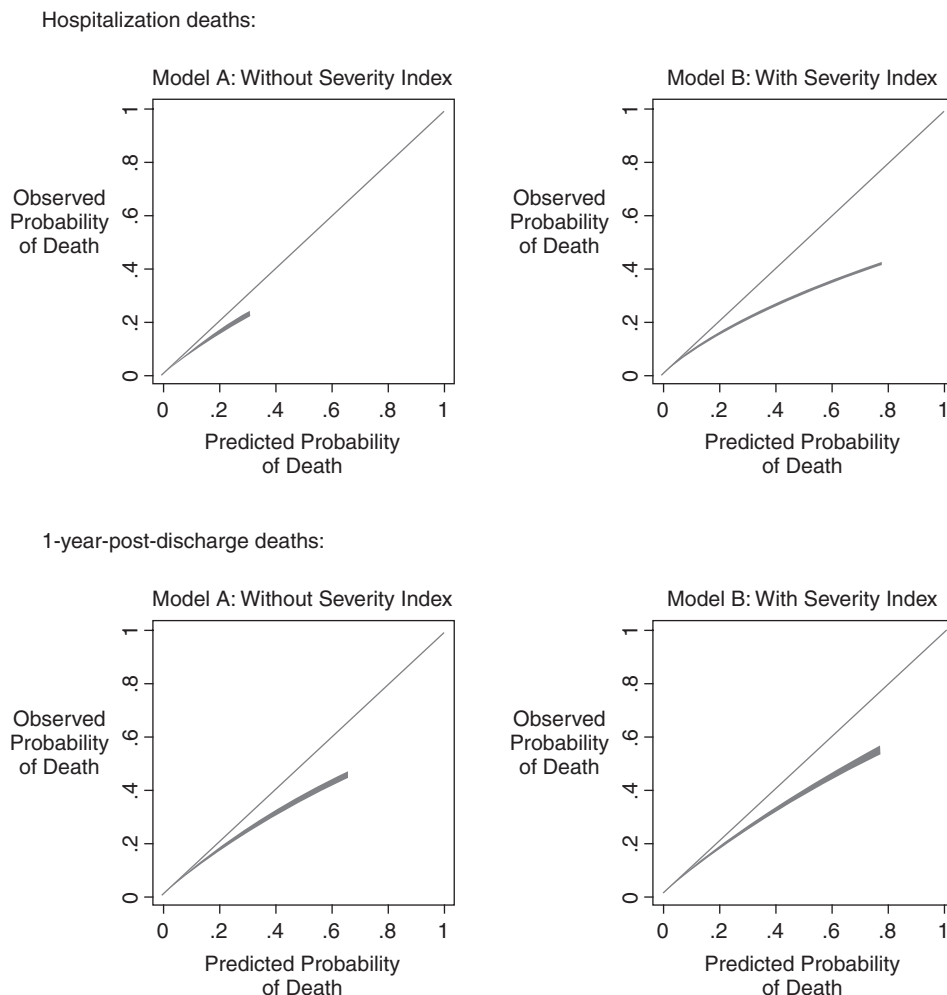


FIG. 2. Calibration of the mortality prediction models with 95% confidence intervals, which appear as very narrow light gray bands surrounding the linear estimate.

TABLE 3. CALIBRATION OF MODELS

Group	1	2	3	4	5	6	7	8	9	10
Hospitalization deaths										
A: Without Severity Index										
Probability	0.0002	0.0003	0.0003	0.0003	0.0006	0.0009	0.0012	0.0019	0.0076	0.3060
Observed	8	11	11	16	15	52	67	70	293	1659
Expected	14.4	16.6	18.4	20.8	26.7	50.6	68.2	94.4	261.3	1630.5
Hosmer-Lemeshow $\chi^2 = 24.69$ , $p$ value = 0.0018										
B: With Severity Index										
Probability	0.0002	0.0002	0.0003	0.0003	0.0005	0.0007	0.0010	0.0019	0.0058	0.7788
Observed	9	3	6	3	11	27	37	51	241	1814
Expected	11.8	14.8	16.8	20.0	26.8	41.3	56.9	87.2	222.9	1703.4
Hosmer-Lemeshow $\chi^2 = 76.67$ , $p$ value < 0.0001										
One-year postdischarge deaths										
A: Without Severity Index										
Probability	0.0000	0.0001	0.0001	0.0002	0.0003	0.0004	0.0005	0.0007	0.0015	0.6555
Observed	5	5	4	5	11	13	17	28	50	573
Expected	1.1	2.4	5.0	8.6	13.1	17.6	23.4	32.9	55.0	551.9
Hosmer-Lemeshow $\chi^2 = 24.41$ , $p$ value = 0.0020										
B: With Severity Index										
Probability	0.0000	0.0001	0.0001	0.0002	0.0003	0.0003	0.0005	0.0007	0.0015	0.7632
Observed	5	5	3	3	14	16	10	25	52	578
Expected	1.1	2.3	4.7	8.3	12.6	16.6	21.8	30.9	52.4	560.3
Hosmer-Lemeshow $\chi^2 = 30.18$ , $p$ value = 0.0020										

again detected a statistically significant degree of miscalibration due to the exceedingly large sample size, with relatively small observed-versus-expected differences).

**Sensitivity analyses of the hospitalization and postdischarge models**

We conducted two sensitivity analyses to assess the whether the discriminative accuracy of the models (as measured by the C statistic) was influenced by either the small amount of missing data in the original sample, or by the precise configuration of the original sample. Multiple imputation techniques were used to address the potential bias of

missing data, and bootstrap re-sampling to address the possibility of model over-fitting (Table 4), we observed essentially identical C statistics and thus no evidence to suggest either type of bias in any of the models.

**Adjusted associations between patient characteristics and odds of death during index hospitalization**

The adjusted odds of death during the index hospitalization (Table 5) was most strongly associated with patient age, with infants less than 1 month of age at the time of hospital admission having 30 times the likelihood of mortality than patients older than 1 year of age at the time of hospital ad-

TABLE 4. SENSITIVITY ANALYSES OF DISCRIMINATIVE ACCURACY OF MODELS COMPARING RESULTS FROM ORIGINAL SAMPLE WITH IMPUTED SAMPLES AND BOOTSTRAP SAMPLES

	<i>C statistic</i>	
	Original	Imputed
Multiple Imputation Samples		
1. Hospitalization death model with no severity score	0.9122	0.9104
2. Hospitalization death model with severity score	0.9364	0.9512
3. 1-year postdischarge death model with no severity score	0.9204	0.9208
4. 1-year postdischarge death model with severity score	0.9238	0.9246
Bootstrap Samples		
1. Hospitalization death model with no severity score	0.9122	0.9100
2. Hospitalization death model with severity score	0.9364	0.9364
3. 1-year postdischarge death model with no severity score	0.9204	0.9206
4. 1-year postdischarge death model with severity score	0.9238	0.9241

mission (infants 1–11 months of age have a 5-fold increase in the odds of mortality). The size of the age association was comparable to the size of the association with the five levels of the severity score, which demonstrated a 36-fold increase in the odds of death for those patients who had the highest score, compared to those with the lowest score. Examining the association with specific types of complex chronic conditions, patients with malignancies had the highest adjusted

odds ratio for the model that did not include the severity score (odds ratio [OR] 5.3, 95% confidence interval [CI]: 4.1–6.9), but a considerably lower score for the model that did include the severity score (OR 1.6, 95%CI: 1.2–2.1), suggesting that the risk of mortality for pediatric oncology patients was strongly associated with the components of the physiologic severity score that were gathered at the time of hospital admission. This contrasts with patients with meta-

TABLE 5. ADJUSTED ASSOCIATIONS OF PATIENT CHARACTERISTICS AND ODDS OF DEATH IN THE HOSPITALIZATION DEATHS MODELS

		Base model n = 678558		With severity index n = 678558	
		OR (95% CI)	p-value	OR (95% CI)	p-value
Age	<1 month	38.0 (29.0–49.9)	< 0.0001	31.2 (23.7–41.0)	< 0.0001
	1–11 months	4.7 (3.5–6.2)		4.8 (3.6–6.4)	
	1–9 y	1.1 (0.8–1.5)		1.2 (0.9–1.6)	
	10–18 y	0.9 (0.7–1.2)		0.9 (0.7–1.3)	
	19–20 y	Ref		Ref	
Gender	Male	1.1 (1.0–1.2)	0.057	1.1 (1.0–1.2)	0.163
	Female	Ref		Ref	
Race	White	Ref	0.3547	Ref	0.5227
	Black	1.0 (0.8–1.1)		0.9 (0.8–1.1)	
	Other	1.1 (0.9–1.3)		1.1 (0.9–1.3)	
Year admitted	1995	1.1 (0.9–1.2)	0.0013	0.7 (0.6–0.8)	< 0.0001
	1996	0.8 (0.7–1.0)		0.6 (0.5–0.7)	
	1997	0.9 (0.8–1.1)		0.6 (0.5–0.8)	
	1998	0.8 (0.7–1.0)		0.6 (0.5–0.7)	
	1999	0.8 (0.7–1.0)		0.6 (0.5–0.7)	
	2000	Ref		Ref	
Payer	None	1.6 (1.3–2.0)	0.0002	1.6 (1.3–2.1)	0.0003
	Medicaid	1.0 (0.9–1.2)		1.0 (0.9–1.2)	
	HMO	1.0 (0.8–1.1)		1.0 (0.9–1.2)	
	Other	0.9 (0.8–1.1)		1.0 (0.8–1.1)	
	Commercial	Ref		Ref	
Median income for subject's zip code	≤\$24,000	Ref	0.2540	Ref	0.3570
	\$24,001–\$29,000	0.9 (0.8–1.1)		0.9 (0.8–1.1)	
	\$29,001–\$35,000	0.9 (0.7–1.0)		0.9 (0.8–1.0)	
	\$35,001–\$46,000	0.9 (0.8–1.1)		1.0 (0.9–1.1)	
	\$46,001–\$60,000	1.0 (0.9–1.2)		1.1 (0.9–1.2)	
	>\$60,001	1.0 (0.8–1.1)		1.0 (0.9–1.2)	
Complex chronic condition	None	Ref	< 0.0001	Ref	< 0.0001
	Neuromuscular	1.8 (1.3–2.4)		1.3 (0.9–1.7)	
	Cardiovascular	2.8 (2.4–3.2)		1.1 (0.9–1.3)	
	Respiratory	2.0 (1.4–2.9)		1.8 (1.2–2.6)	
	Metabolic	4.4 (2.7–7.3)		4.0 (2.4–6.6)	
	Congenital	4.2 (3.2–5.3)		2.3 (1.7–3.0)	
	Malignancy	5.3 (4.1–6.9)		1.6 (1.2–2.1)	
	Other	0.3 (0.2–0.6)		0.3 (0.2–0.4)	
Previous chronic complex condition	No	Ref	< 0.0001	Ref	< 0.0001
	Yes	2.1 (1.5–3.0)		2.4 (1.7–3.5)	
Prior Admission	0	Ref	< 0.0001	Ref	< 0.0001
	1	1.5 (1.1–1.9)		1.6 (1.2–2.0)	
	2	2.0 (1.3–3.2)		2.1 (1.3–3.2)	
	+3	6.7 (4.9–9.2)		6.3 (4.6–8.6)	
Facility	Pediatric hospital	0.3 (0.2–0.4)	< 0.0001	0.3 (0.2–0.4)	< 0.0001
	Other	Ref		Ref	
Severity score	0			Ref	< 0.0001
	1			0.7 (0.6–0.8)	
	2			3.4 (3.0–3.9)	
	3			10.0 (8.7–11.5)	
	4			36.1 (29.7–44.0)	

The *p* values for each variable represent the significance test for the entire block of values of that variable. OR, odds ratio; CI, confidence interval.

bolic disorders, who had the second highest adjusted OR in the model that did not include the severity score (OR 4.4, 95% CI: 2.7–7.3) and the highest and essentially unchanged adjust OR in the model that included the severity score (OR 4.0, 95% CI: 2.4–6.6), suggesting that for these patients, the

severity score did not denote a higher risk of mortality. Similarly, the association between the history of 3 or more prior admission and the odds of mortality was essentially the same for the models without (OR 6.7, 95% CI: 4.9–9.2) and with (OR 6.3, 95% CI: 4.6–8.6) the severity index.

TABLE 6. ADJUSTED ASSOCIATIONS OF PATIENT CHARACTERISTICS AND ODDS OF DEATH IN THE ONE-YEAR POSTDISCHARGE DEATHS MODELS

		<i>Base model</i> n = 549,035		<i>With severity index</i> n = 549,035	
		OR (95% CI)	p-value	OR (95% CI)	p-value
Age	<1 month				
	1–11 months	0.9 (0.6–1.2)	0.52	0.9 (0.6–1.2)	0.48
	1–9 y	0.8 (0.6–1.0)		0.8 (0.6–1.0)	
	10–18 y	0.8 (0.6–1.0)		0.8 (0.6–1.0)	
	19–20 y	Ref		Ref	
Gender	Male	0.9 (0.8–1.1)	0.207	0.9 (0.8–1.1)	0.187
	Female	Ref		Ref	
Race	White	Ref	0.08	Ref	0.1308
	Black	0.8 (0.6–1.0)		0.8 (0.6–1.0)	
	Other	0.8 (0.5–1.2)		0.8 (0.5–1.2)	
Year	1995		0.0153		0.0025
	1996	1.5 (1.2–2.0)		1.7 (1.3–2.2)	
	1997	1.3 (1.0–1.7)		1.3 (1.0–1.7)	
	1998	1.3 (1.0–1.7)		1.3 (1.0–1.7)	
	1999	1.4 (1.1–1.8)		1.4 (1.1–1.8)	
	2000	Ref		Ref	
Payer	None	0.7 (0.4–1.4)	0.2134	0.7 (0.4–1.4)	0.2258
	Medicaid	0.8 (0.7–1.0)		0.8 (0.7–1.0)	
	HMO	0.9 (0.7–1.2)		1.0 (0.8–1.3)	
	Other	1.1 (0.8–1.4)		1.1 (0.8–1.4)	
	Commercial	Ref		Ref	
Median income for subject's zip code	≤\$24,000	Ref	0.0542	Ref	0.0644
	\$24,001–\$29,000	1.3 (1.0–1.7)		1.3 (1.0–1.7)	
	\$29,001–\$35,000	1.3 (1.0–1.7)		1.3 (1.0–1.7)	
	\$35,001–\$46,000	1.2 (0.9–1.5)		1.2 (0.9–1.6)	
	\$46,001–\$60,000	1.4 (1.1–1.9)		1.4 (1.1–1.9)	
	>\$60,001	1.0 (0.7–1.4)		0.9 (0.6–1.4)	
Chronic complex condition	None	Ref	0.0023	Ref	0.0003
	Neuromuscular	1.9 (1.2–3.0)		2.2 (1.4–3.4)	
	Cardiovascular	1.0 (0.6–1.7)		1.1 (0.6–1.9)	
	Respiratory	0.7 (0.3–1.8)		0.9 (0.4–2.3)	
	Metabolic and congenital	0.5 (0.1–2.5)		0.5 (0.1–2.4)	
	Malignancy	3.9 (1.5–10.1)		4.0 (1.6–10.4)	
	Other	0.6 (0.3–1.2)		0.6 (0.3–1.2)	
Previous complex chronic condition	No	Ref	< 0.0001	Ref	< 0.0001
	Yes	1.5 (1.2–1.9)			
Prior Admission	0	Ref	< 0.0001	Ref	
	1	6.9 (5.5–8.7)		7.3 (5.7–9.2)	
	2	14.6 (11.2–19.1)		15.3 (11.7–20.0)	
	+3	20.2 (15.9–25.7)		21.9 (17.3–27.9)	
Facility	Pediatric Hospital	2.5 (0.6–10.5)	0.196	2.6 (0.6–11.1)	0.182
	Other	Ref		Ref	
Length of stay	<2 days	Ref	< 0.0001	Ref	0.001
	2–7 days	1.2 (1.0–1.5)		1.2 (0.9–1.4)	
	8–14 days	1.9 (1.5–2.6)		1.7 (1.3–2.2)	
	15+ days	1.9 (1.4–2.6)		1.6 (1.2–2.2)	
Severity score	0			Ref	
	1			1.3 (1.1–1.6)	
	2			2.9 (2.2–3.7)	
	3			4.1 (2.9–5.7)	
	4			3.6 (1.6–7.9)	

The p values for each variable represent the significance test for the entire block of values of that variable. OR, odds ratio; CI, confidence interval.



The adjusted odds of mortality during the 1-year postdischarge from the index hospitalization (Table 6) was most strongly associated with the number of prior admissions that had occurring during the year before the index hospitalization, with a 20-fold increase (95% CI: 15.9–25.7) in the odds of death among those subjects who had been previously admitted 3 or more times compared to patients who had not been previously admitted. The admission physiologic severity index continued to be significantly associated with postdischarge mortality, but to an attenuated degree compared to the association observed during the index hospitalization, with only a 4-fold (95% CI: 1.6–7.9) increase in the likelihood of death among subjects in the higher index risk levels compared to the lowest level. The types of medical condition most associated with postdischarge death were malignancies (4-fold increase, 95% CI: 1.5–10.1) and neuromuscular conditions (2-fold increase, 95% CI: 1.2–3.0), and having had a diagnosis of any complex chronic condition during a prior hospital admission (1.5-fold increase, 95% CI: 1.2–1.9); none of these associations was substantially different in the models without or with the severity index. Subjects who had been admitted to a pediatric hospital for the index admission had a 2.5-fold increase (95% CI: 0.6–10.5) in the odds of death during the year after discharge, and the likelihood of mortality after discharge was not associated with subjects' age.

#### Association of predicted probability of death after discharge and location of death

To assess the possible utility of the predicted probability of death, we examined the association between the 1-year postdischarge mortality probability prediction and the proportion of deaths that occurred in either the hospital or the home setting. Among all subjects (Fig. 3, upper panel), as the predicted probability of death increased, the proportion of subjects who died either in the hospital or at home increased, with a slight acceleration in the proportion who died at home as the predicted probability of death increased beyond 0.4. Among only those subjects who had oncologic diagnoses (Fig. 3, lower panel), the same pattern, but more accentuated, was evident. These patterns would be consistent with a pattern of care whereby the planning and provision of home-based end-of-life care services occurs to a greater degree once a certain threshold of 1-year postdischarge mortality risk is exceeded.

#### Discussion

Using readily available administrative data from hospital discharge records, we were able in this large cohort of hospitalized children in Pennsylvania to predict both in-hospital mortality and mortality during the year after discharge with an extremely high level of accuracy. The addition of a "physiologic severity index" that was based upon specially collected clinical and laboratory data for the subjects improved the accuracy of the in-hospital mortality model to a modest degree, but had essentially no impact on the 1-year postdischarge model. Furthermore, these predictions of mortality were associated with demonstrable differences in outcome, specifically regarding location of death among those patients who survived the initial hospitalization, attesting both to the validity of the construct and highlighting how these predictions could be used to assess differences in systems of health care.

This study has several strengths that should be kept in mind. First, applying predictive modeling of mortality probability to populations of patients as an analytic aid in the assessment of end-of-life care and outcomes is innovative, opening new avenues and approaches to palliative care health services research. Second, the population-basis of the cohort minimizes the risk of sampling bias. Third, careful selection of variables contained in the hospitalization model excluded information that was likely unknown on the initial day of the index hospitalization, such as the ensuing length of stay or the final diagnoses as conveyed by the APR-DRG classification. Fourth, the stability of the accuracy of the predictive models was affirmed both by the initial split-sample assessment and by further bootstrap resampling validation, as well as assessing that potential bias due to missing data was minimal through the multiple imputation of missing values sensitivity analysis.

Turning to consider weaknesses of the study, first, we cannot evaluate the completeness or accuracy of the match of hospital records to any corresponding death certificate information. We do note, however, that the occurrence of either failed matches (whereby subjects who died after discharge from the hospital were not identified as having died) or incorrect matches (whereby subjects were classified as having died when in fact they did not) would most likely have resulted in nondifferential misclassification biasing our

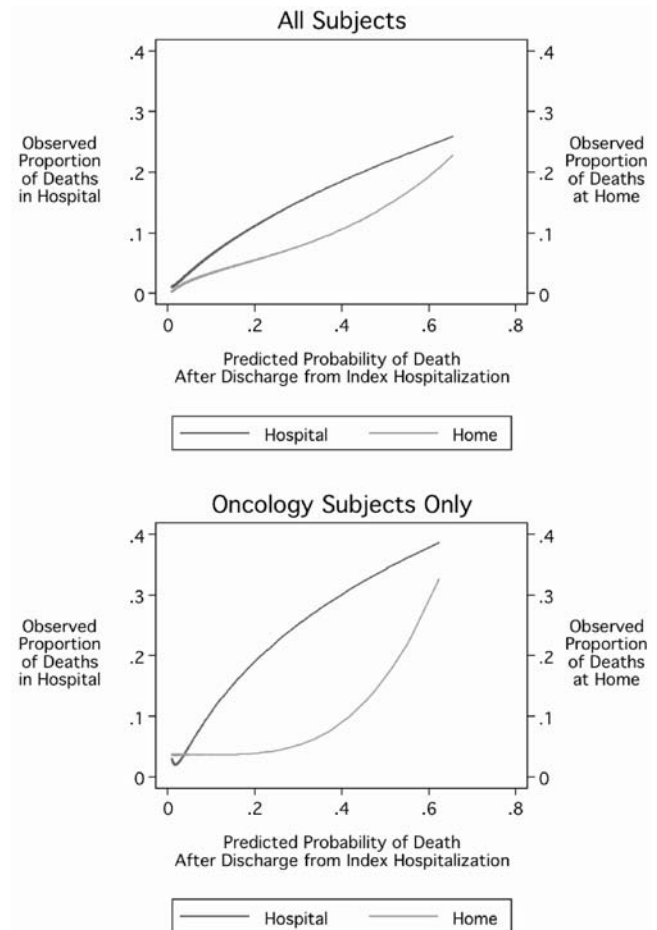


FIG. 3. One-year postdischarge probability of death and location of death.

results toward the null. Second, our results cannot be assuredly extrapolated outside of the cohort, to either other locations or other time periods. Finally, reemphasize a crucial point, these models do not demonstrate sufficient accuracy to be used for individual-level predictions.

We conclude by underscoring three important implications of this study. First, the methodology developed in this study enables population-level comparisons across different systems of health care, adjusting for differences between these systems in terms of patient "case mix" and the associated differences mortality risks, thereby enabling better comparisons of how differences in the processes of care between the systems affect outcomes. For example, with appropriate adjustment for subjects' risk of mortality after discharge, one could analyze how different methods of managing the process of introducing hospice services to patients or families affected the likelihood of deaths occurring back in the hospital or at home. Second, the ability to predict with substantial accuracy the likelihood of death during and then after hospitalization, using only routine administrative hospitalization data, suggests that such comparative studies can be performed widely at relatively low cost. Finally, the strong association between the history of prior hospitalizations and the likelihood of death highlights the value of datasets that enable the longitudinal linkage of hospitalization records for subjects.

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### Author Disclosure Statement

No competing financial interests exist.

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Address reprint requests to:

Chris Feudtner, M.D.

General Pediatrics

3535 Market Street, Room 1523

The Children's Hospital of Philadelphia

34<sup>th</sup> and Civic Center Boulevard

Philadelphia, PA 19104

E-mail: feudtner@email.chop.edu