

A Model to Predict Risk of Hyperkalemia in Patients with Chronic Kidney Disease Using a Large Administrative Claims Database

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Background: Chronic kidney disease (CKD) is responsible for substantial clinical and economic burden. Drugs that inhibit the renin-angiotensin-aldosterone system inhibitors (RAASi) slow CKD progression in many common clinical scenarios. Guideline-directed medical therapy requires maximal recommended doses of RAASi, which clinicians are often reluctant to prescribe because of the associated risk of hyperkalemia (HK).

Objective: This study aims to develop and validate a model to identify individuals with CKD at elevated risk for developing HK over a 12-month period on the basis of lab, medical, and pharmacy claims.

Methods: Using claims from a large US healthcare payer, we developed a model to predict the probability of individuals identified with CKD but not HK in 2016 (baseline year [BY]) who developed HK in 2017 (prediction year [PY]). The study population was comprised of members continuously enrolled with medical and pharmacy benefits and CKD (BY). Members were excluded from the analysis if they had HK (by lab results or diagnosis code) or dialysis (BY). Prediction model performance measures included area under the receiver operating characteristic curve (AUROC), calibration, and gain and lift charts.

Results: Of 435,512 members identified with CKD but not HK (BY), 6235 (1.43%) showed incident HK (PY). Compared with individuals without incident HK (PY), these members had a higher comorbidity burden, use of RAASi, and healthcare utilization. The AUROC and calibration analyses showed good predictive accuracy (area under the curve [AUC]=0.843 and calibration). The top 2 HK-prediction deciles identified 75.94% of members who went on to develop HK (PY).

Conclusion: Guideline-recommended doses of RAASi therapy can be limited by the risk of HK. Novel potassium binders may permit more patients at risk to benefit from these maximal RAASi doses. This predictive model successfully identified the risk of developing HK up to 1 year in advance.

Keywords: chronic kidney disease, hyperkalemia, RAAS inhibitors, potassium binder

Introduction

Chronic kidney disease (CKD) is responsible for substantial clinical, economic, and humanistic burden. The United States (US) adult prevalence of CKD in 2013–2016 was 14.8% for all stages and 6.9% for moderate-to-severe (stages 3–5) CKD.¹

As kidney function declines, the association of CKD with comorbidities, including hypertension, diabetes, and cardiovascular disease,² drives rising healthcare expenditures.^{3,4} For example, a 2016 analysis using clinical data of patients

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prescribed renin-angiotensin-aldosterone system (RAAS) inhibitors (RAASi) found that annual costs (2016 dollars, excluding dialysis) rose exponentially from \$7537 in patients without CKD to \$76,969 in patients with stage 4–5 CKD and to over \$121,000 in patients with end-stage renal disease (ESRD).⁵

Earlier diagnosis and evidence-based management of CKD are key to slowing progression and preventing the complications of advanced CKD. The RAAS blockade plays a major role in delaying or, in some patients perhaps halting, kidney failure in most types of CKD.^{6–10} In the RENAAL study, the investigators estimated that the RAAS blockade, with the optimal dose of losartan, delayed the development of ESRD by 2 years in patients with type 2 diabetes with nephropathy.⁶

RAASi drugs currently marketed in the US are classified by their mode of action: angiotensin-converting enzyme inhibitors (ACEi), angiotensin-receptor blockers (ARB), mineralocorticoid receptor antagonists (MRA; eg, aldosterone), and direct renin inhibitors (DRI). Importantly, those at greatest risk of CKD progression (advanced CKD and diabetes and/or heart failure [HF] with reduced ejection fraction) stand to benefit most.^{11,12}

Achieving the full clinical benefits of RAASi medications requires the use of maximum or optimal doses used in clinical trials, but these drugs are frequently not prescribed or are underprescribed because of the inherent risk of hyperkalemia (HK). A large clinical study found that only 26.8% of patients with CKD, HF, or both received optimal doses; other studies have confirmed RAASi underprescribing for patients with guideline-recommended cardiorenal indications.^{13–16}

Other factors associated with HK include reduced kidney function, age, male gender, diabetes, HF, adrenal insufficiency, and certain medications, such as nonsteroidal anti-inflammatory drugs and RAASi.¹⁷ The likelihood of developing HK increases as kidney function declines. A large clinical data analysis found the 5-year HK prevalence to be 23.5% in patients with HF, 29.5% in patients with CKD stages 3–4, and 47.6% in patients with both.¹⁸

The presence of HK further adds to the cost of CKD. A claims study of patients with CKD and/or HF found that after multivariable adjustment, those with HK cost \$25,156 more annually than matched patients without HK.¹⁹ In addition, a 2016 clinical data study found that in CKD stage 4–5, HK expenditures were 38% and 22% higher for commercial and Medicare patients, respectively.⁵

The clinical and potential economic value of optimal-dose RAASi in slowing CKD progression highlights the importance of predicting the risk of developing HK, especially because interventions to prevent HK would enable most patients to remain optimally treated.^{17,20} With administrative claims and electronic medical records, predictive analytics can be applied to combine the contribution of multiple risk factors into a single probabilistic estimate to guide healthcare.²¹ The objective of this study was to develop and validate a predictive model to identify patients with CKD likely to develop HK within a large national health plan, based on a broad set of demographic, clinical, and healthcare-resource utilization (HCRU) characteristics found in administrative claims data.

Data and Methods

Data

The model was developed to predict the occurrence of HK in the prediction year (PY, 2017) of individuals identified with CKD in the baseline year (BY, 2016). Individuals were excluded if they had HK or underwent dialysis in the BY or were enrolled in an end-of-life care program or hospice in either the BY or PY.

Data included medical and pharmacy claims and lab-test results of deidentified individuals continuously enrolled from January 2016 through December 2017 in a fully insured commercial or Medicare Advantage plan operated by a large national health plan. Membership in the study population required continuous medical and pharmacy coverage throughout both BY and PY; diagnosis in BY of CKD (at least 1 claim with a principal or secondary diagnosis of CKD or from estimated glomerular filtration rate [eGFR]—see [Table S1](#)); and being at least 18 years old on January 1, 2016.

Disease comorbidities were identified based on medical and pharmacy claims and lab results. The burden of chronic diseases was estimated from the original conditions and weights of the Charlson-Deyo Comorbidity Index using the International Statistical Classification of Disease and Related Health Problems, 10th revision (ICD-10) codes.^{22,23}

HK was identified if an individual met at least 1 of the following criteria: (1) ≥ 2 serum potassium >5.0 mmol/L on different dates (Logical Observation Identifiers Names and Codes [LOINC]: 2823–3); (2) ≥ 2 claims with principal or secondary diagnosis of HK (International

Classification of Diseases, 10th Revision, Clinical Modification [ICD-10-CM]: E87.5); (3) 1 claim with HK and 1 potassium >5.0 mmol/L; (4) ≥ 1 dispensed prescription for sodium polystyrene sulfonate or patiomer (the potassium binders approved by the FDA prior to 2017).

RAASi medications (see [Table S2](#)) included ACEi, ARB, MRA, and DRI. Patients were categorized as receiving optimal dosing if they were dispensed the maximum package-labeled dose in the BY, whereas patients who were dispensed any dose below that were categorized as receiving suboptimal dosing.²⁴ RAASi adherence was estimated by proportion of days covered (PDC) as defined by the Centers for Medicare & Medicaid Services (CMS) performance measure D12, renin-angiotensin system (RAS) antagonists.²⁵

Descriptive Statistical Analysis

All baseline characteristics were analyzed descriptively. Means (\pm SD) and medians (IQR) were reported for continuous variables, and frequencies (%) were reported for categorical variables. Statistical significance was assessed with the Student's *t*-test or Wilcoxon rank-sum test or Kruskal–Wallis test for continuous variables and χ^2 test for categorical variables.

Development and Validation of the Prediction Model

Data on study group members were randomly divided into 2 equal sets for model development and validation. The development phase focused on selection of PY HK predictor variables from a variety of candidates, including demographics, CKD stage, comorbidities, drugs, lab results, and HCRU.

The first step in the development phase excluded variables lacking significant ($P \leq 0.05$) univariate association with HK. This step yielded 46 candidate predictor variables, which, in the next step, were subjected to stepwise logistic regression to identify those that were independently correlated with HK. The probability for entry of a variable was set at 0.01 and for removal at 0.1. Adjusted odds ratios (ORs) were calculated, along with their associated 95% confidence intervals (CIs). Interactions between predictive variables in the model were checked using a pooled interaction test.²⁶ Multicollinearity was checked using Pearson's correlation coefficient. The model's goodness-of-fit was tested using deviance, Hosmer–Lemeshow, and log-likelihood.²⁷

The model's performance was validated on the validation dataset. The area under a receiver operating characteristic curve (AUROC) was calculated to determine the model's ability to discriminate between those who developed HK and those who did not during the PY follow-up period. AUROC was computed by a nonparametric method.²⁸ An AUROC of 1 implies perfect discrimination, whereas an AUROC of 0.5 implies the model performs no better than chance.²⁹

Calibration was assessed by plotting observed versus predicted probability of HK in the PY among patients grouped by deciles of predicted probability as estimated by the model (decile 1: lowest risk; decile 10: highest risk). Risk accuracy was estimated as the mean absolute difference (error) between the predicted and observed (actual) probability of HK across the 10 risk strata. Risk stratification performance was assessed by (1) the mean actual risk (observed probability) increases between strata, and (2) the risk ratio between the observed probability in the highest-risk strata to that in the lowest-risk strata.

A gains chart was used to compare predictive accuracy with the prediction model versus without the prediction model. The data were sorted in descending order of the probability of HK estimated by the model, binned by deciles, and the cumulative rate of HK at each decile was evaluated, from the highest-risk decile to the lowest-risk decile. The greater the area between the gain curve and the baseline (random selection without a prediction model), the better the model's performance. The model's goodness-of-fit was further tested using deviance, Hosmer–Lemeshow, and log-likelihood.

All data management, statistical analyses, and predictive modelling were performed using SAS software, version 9.4 (SAS Institute, Cary, NC, US). The study was approved by an independent institutional review board, Sterling IRB, before proceeding.

Results

Baseline Characteristics

Of the 435,512 individuals who met the study criteria, 57% were female, 42% lived in the South region, and 54% were enrolled in a Medicare Advantage plan ([Figure 1](#), [Table 1](#)). [Table 1](#) compares BY characteristics of study group members who went on to exhibit (6235; 1.44%) or not exhibit (429,277; 98.57%) evidence of HK in the PY. Members who were first identified with

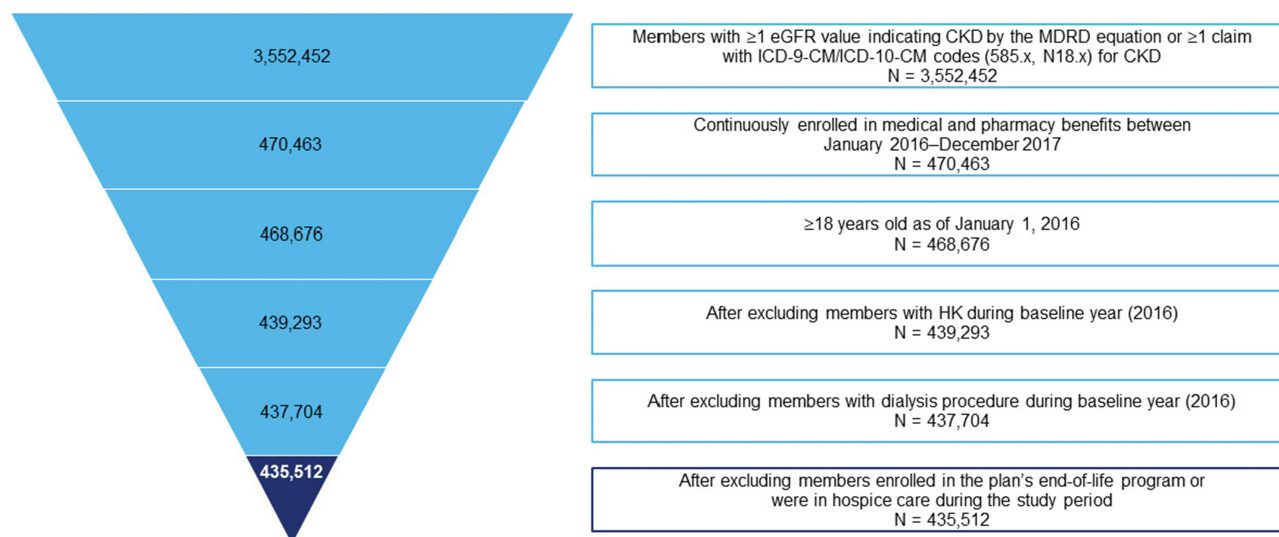


Figure 1 Study population flowchart.

Abbreviations: CKD, chronic kidney disease; eGFR, estimated glomerular filtration rate; ICD-9-CM, International Classification of Diseases, 9th Revision, Clinical Modification; ICD-10-CM, International Classification of Diseases, 10th Revision, Clinical Modification; HK, hyperkalemia; MDRD, modification of diet in renal disease.

HK in the PY tended to be older, male, of Southern residence, urban, Medicare-insured, with a higher Charlson-Deyo Comorbidity Index (CCI). They were more likely to have comorbid hypertension, hyperlipidemia, diabetes or congestive HF, and more advanced CKD (Table 2).

In addition, members of the HK subgroup were 52% likelier to use RAASi in the BY (70% versus 46.1% for the non-HK group; Table 3; Table S3). Submaximal RAASi dosing was widespread, but overall, adherence as measured by PDC $\geq 80\%$ was similar in the HK and non-HK groups. Serum potassium and urine protein excretion were greater in the HK group (Table 2).

Variables in the Logistic Regression Model

Forty-six variables with significant univariate association to HK first appearing in the PY (see Table S4) were reduced to 21 by stepwise logistic regression. Table 4 shows that CKD stage, higher BY potassium, use of ACEi, MRA, and calcineurin inhibitors; and certain comorbidities, including diabetes, independently predicted HK. Suboptimal RAASi dose and adherence were equally predictive of HK, but dose and adherence together were not more predictive.

Validation of the Logistic Prediction Model

Nine predictive models were developed and compared by their AUROC during validation (Figure 2). Among the

models based solely on single-feature variables, laboratory-only (M5) performed best (AUROC: 0.784), followed by CKD stage and comorbidities (M2, AUROC: 0.741), whereas models based solely on drugs (M3) or HCRU (M4) or demographics (M1) yielded poor AUROCs. Fitting the model using all predictive variables (full model) yielded a logistic regression model with 27 nonzero coefficients, plus an intercept. The full model's AUROC was 0.843, demonstrating good discriminative performance.

Figure 3 shows the prediction model's risk estimation of HK and stratification performance. The model's estimated average HK risk across the population closely matched the average actual risk (1.43% versus 1.51%, respectively). The mean absolute error was 1.48%, suggesting good accuracy calibration. There was, on average, a 1% increase in risk with each higher stratum, and those in the highest-risk stratum had 240 times the probability of developing HK in the PY than those in the lowest-risk stratum (9.15% versus 0.04%, respectively). The second-highest risk strata's actual risk was 2.9%, significantly higher than the overall mean of 1.43% for the cohort. Model calibration showed that the top 2 prediction deciles (20% of the study population) identified 75.94% of individuals first identified with hyperkalemia in the PY.

The model somewhat overpredicted deciles 1–8 and was very close for deciles 9 and 10 in people who actually developed HK. Model gains are shown in Figure 4. For

Table 1 Baseline Patient Demographics

Characteristics	Study Population	BY HK	BY No HK	P value
	(N=435,512)	(N=6235)	(N=429,277)	
Age				
Mean (SD)	61.3 (16.3)	69.3 (11.8)	61.1 (16.3)	<0.0001
Median (IQR)	64 (50–73)	71 (62–77)	63 (50–73)	<0.0001
Gender, n (%)				
Male	186,261 (42.8)	3281 (52.6)	182,980 (42.6)	<0.0001
Female	249,228 (57.2)	2954 (47.4)	246,274 (57.4)	
Geography, n (%)				
Midwest	77,548 (17.8)	853 (13.7)	76,695 (17.9)	<0.0001
Northeast	132,504 (30.4)	1703 (27.3)	130,801 (30.5)	
South	185,599 (42.6)	3302 (53.0)	182,297 (42.5)	
West	39,613 (9.1)	377 (6.1)	39,236 (9.2)	
Urban/rural, n (%)				
Urban	162,723 (37.4)	2825 (45.3)	159,898 (37.3)	<0.0001
Suburban	125,973 (28.94)	1647 (26.4)	124,326 (29.00)	
Rural	146,558 (33.7)	1763 (28.3)	144,795 (33.8)	
Insurance, n (%)				
Commercial	199,155 (45.7)	1293 (20.7)	197,862 (46.1)	<0.0001
Medicare Advantage	236,357 (54.3)	4942 (79.3)	231,415 (53.9)	

Abbreviations: BY, baseline year; HK, hyperkalemia; IQR, interquartile range; SD, standard deviation.

Table 2 Clinical Characteristics and Comorbidities

Characteristics	Study Population	BY HK	BY No HK	P value
	(N=435,512)	(N=6235)	(N=429,277)	
Charlson-Deyo Comorbidity Index				
Mean (SD)	1.5 (2.0)	2.52 (2.4)	1.5 (2.0)	<0.0001
Median (IQR)	1 (0–2)	2 (1–4)	1 (0–2)	
CKD stage, n (%)				
Stage 1	104,319 (24.0)	376 (6.0)	103,943 (24.2)	<0.0001
Stage 2	210,353 (48.3)	2309 (37.0)	208,044 (48.5)	
Stage 3	108,583 (24.9)	2814 (45.1)	105,769 (24.6)	
Stage 4	9891 (2.3)	621 (10.0)	9270 (2.2)	
Stage 5	2366 (0.5)	115 (1.8)	2251 (0.5)	
Key comorbidities, n (%)				
Hyperlipidemia	276,289 (63.4)	5365 (86.0)	270,917 (63.1)	<0.0001
Hypertension	278,336 (63.9)	5270 (84.5)	273,063 (63.6)	<0.0001
Diabetes	110,881 (25.5)	3139 (50.4)	109,294 (25.5)	<0.0001
Obesity	89,324 (20.5)	1708 (27.4)	87,615 (20.4)	<0.0001
Congestive HF	40,067 (9.2)	1327 (21.3)	38,764 (9.0)	<0.0001
Selected baseline biometrics				
Serum K ⁺ , mmol/L	4.43	4.71	4.33	<0.0001
Hemoglobin, g/dL	13.69	13.23	13.70	<0.0001
Urine protein, mg/24 h	177.91	259	173.00	<0.0001
Proteinuria, %	2.1	4.0	2.0	<0.0001

Abbreviations: BY, baseline year; HF, heart failure, HK, hyperkalemia, IQR, interquartile range; K⁺, potassium; SD, standard deviation.

Table 3 Baseline Medications

Characteristics	Study Population	BY HK	BY No HK	P value
	(N=435,512)	(N=6235)	(N=429,277)	
RAASi use, n (%)				
ACEi	108,269 (24.9)	2532 (40.6)	105,737 (24.6)	<0.0001
ARB	83,325 (19.1)	1536 (24.6)	81,789 (19.1)	<0.0001
MRA	10,339 (2.4)	281 (4.5)	10,058 (2.3)	<0.0001
Other RAASi	528 (0.1)	19 (0.3)	509 (0.1)	0.0003
RAASi PDC ≥80%, n (%)				
Stage 1	14,196 (53.8)	101 (58.1)	14,095 (53.8)	0.2856
Stage 2	55,918 (63.2)	791 (63.2)	55,127 (63.2)	0.9776
Stage 3	44,915 (66.1)	1354 (66.1)	43,561 (66.1)	0.9904
Stage 4	4053 (61.8)	306 (65.0)	3747 (61.5)	0.1401
Stage 5	692 (54.7)	44 (61.1)	648 (54.3)	0.2746
RAASi at optimal dose, n (%)				
Stage 1	6612 (25.1)	33 (19.0)	6579 (25.1)	0.0653
Stage 2	26,311 (28.7)	291 (23.3)	26,020 (29.8)	<0.0001
Stage 3	23,243 (34.2)	631 (30.8)	22,612 (34.3)	0.0008
Stage 4	2329 (35.5)	156 (33.1)	2173 (35.7)	0.2719
Stage 5	456 (36.0)	27 (37.5)	429 (35.9)	0.8012
Baseline K⁺ affecting non-RAASi use, n (%)				
NSAIDs	90,124 (20.7)	1335 (21.4)	88,789 (20.7)	0.1611
Calcineurin inhibitors	917 (0.2)	48 (0.8)	869 (0.2)	<0.0001
Beta-blockers	98,485 (22.6)	2058 (33.0)	96,427 (22.5)	<0.0001

Abbreviations: ACEi, angiotensin-converting enzyme inhibitors; ARB, angiotensin-receptor blocker; BY, baseline year; K⁺, potassium; MRA, mineralocorticoid-receptor antagonist; NSAID, nonsteroidal anti-inflammatory drug; PDC, proportion of days covered; RAASi, renin-angiotensin-aldosterone system inhibitors.

a given decile, gain is the cumulative number of individuals correctly identified with HK up to that point, divided by the total number of individuals with HK. For example, for the second-highest decile, gain = 1572/2070 = 75.94%.

Discussion

Early identification and appropriate management with pharmacologic inhibition of the RAAS has been shown to mitigate the consequences of CKD and slow the progression to ESRD. Yet more than one-third of new ESRD patients receive little or no pre-ESRD nephrology care,³⁰ despite dissemination of kidney disease and diabetes practice guidelines.^{31–33} To accelerate improving the quality and outcomes of kidney care, the US Department of Health and Human Services recently launched the Advancing American Kidney Health initiative, targeting early CKD identification and incentivizing evidence-based management of advanced CKD and ESRD.^{34,35}

Using a large national administrative claims database, we identified members with CKD but without HK or dialysis. We developed and internally validated a model

to predict the occurrence of HK in the following year. The performance characteristics of the logistic regression model were comparable to those of a neural network and decision tree model (data not shown in this article). We therefore selected the logistic model because of its general applicability in a clinical or care-management setting.

The model's performance was promising for potential use and further development in health plan-based care management: AUROC was 0.843; the numbers of predicted versus observed individuals with HK were well matched in the highest probability deciles; and a cutoff of the top 20% scores captured more than three-quarters of those who exhibited HK.

There is much evidence supporting maximal recommended RAASi dosing, if tolerated.^{6,7} An individualized approach using predictive analytics to identify patients at high risk of HK may allow clinicians to treat to guideline-recommended RAASi dosing.

Until recently, options for reducing chronic HK have been limited, but the recent introduction of novel

Table 4 List of Predictive Variables in the Final Logistic Model

Parameter	Regression Coefficient (SE)	Odds Ratio (95% CI)	P value
Intercept	-12.16 (0.34)		<0.0001
CKD stage 2 versus I	0.50 (0.09)	1.65 (1.38–1.98)	<0.0001
CKD stage 3 versus I	1.21 (0.09)	3.34 (2.77–4.02)	<0.0001
CKD stage 4 versus I	2.01 (0.11)	7.47 (6.02–9.28)	<0.0001
CKD stage 5 versus I	2.45 (0.19)	11.62 (7.90–17.07)	<0.0001
Potassium, per 0.1mmol/L higher	2.15 (0.05)	8.59 (7.85–9.40)	<0.0001
Calcineurin inhibitors use	1.49 (0.21)	4.43 (2.98–6.97)	<0.0001
Diabetes mellitus	0.41 (0.04)	1.51 (1.39–1.63)	<0.0001
Northeast versus Midwest	0.13 (0.06)	1.14 (1.01–1.29)	0.0170
South versus Midwest	0.45 (0.06)	1.57 (1.40–1.77)	<0.0001
West versus Midwest	0.28 (0.09)	1.32 (1.09–1.58)	0.0102
Suburban versus Rural	0.05 (0.04)	1.05 (0.96–1.17)	0.0568
Urban versus Rural	0.29 (0.05)	1.33 (1.22–1.46)	<0.0001
Hyperlipidemia	0.39 (0.07)	1.47 (1.28–1.67)	<0.0001
Osteoporosis	0.24 (0.06)	1.27 (1.12–1.45)	<0.0001
MRA use	0.23 (0.07)	1.26 (1.08–1.46)	<0.0001
ACE inhibitors use	0.22 (0.04)	1.25 (1.16–1.35)	<0.0001
Peripheral artery disease	0.17 (0.05)	1.19 (1.09–1.31)	0.0002
Malignant neoplasms	0.15 (0.05)	1.16 (1.04–1.29)	0.0107
Chronic obstructive pulmonary disease	0.10 (0.04)	1.11 (1.00–1.23)	0.0498
Number of comorbid conditions	0.07 (0.01)	1.07 (1.05–1.08)	<0.0001
Hemoglobin, per 1g/dL higher	-0.09 (0.01)	0.91 (0.89–0.94)	0.0028
Optimal RAASi dose	-0.09 (0.01)	0.91 (0.83–0.99)	0.0097
Inpatient admission – all cause	-0.14 (0.05)	0.87 (0.78–0.97)	0.0072
Commercial versus medicare plan	-0.16 (0.06)	0.85 (0.75–0.95)	0.0035
Primary care visit	-0.38 (0.04)	0.68 (0.62–0.74)	<0.0001
Female versus Male	-0.46 (0.05)	0.63 (0.57–0.69)	<0.0001
Potassium sparing diuretics use	-0.63 (0.07)	0.53 (0.37–0.74)	0.0003

Abbreviations: ACE, angiotensin-converting enzyme; CI, confidence interval; CKD, chronic kidney disease; COPD, chronic obstructive pulmonary disease; K⁺, potassium; MRA, mineralocorticoid-receptor antagonist; ORs, odds ratios.

potassium binders patiromer and sodium zirconium cyclo-silicate may allow patients to safely achieve higher RAASi doses—underscoring the value of being able to quantitatively predict who will develop HK.³⁶

To our knowledge, no HK predictive model has been validated using data typically available from health plans. The few published predictive models are based on clinical data from integrated health systems. Johnson and colleagues developed a model to predict quintile risk of HK (serum potassium ≥ 5.5 mmol/L) in patients with eGFR

<60 mL/min/m² within 90 days of starting lisinopril without diuretics. The observed risk was 6.9% and 0.7% for the highest and lowest quintiles, respectively; predicted and observed risks agreed within 1%, but formal performance characteristics were not reported.³⁶ A model based on clinical data and validated in a different healthcare system found that 6 variables (male, diabetes, congestive heart failure, use of potassium-sparing diuretics, lower eGFR, and higher baseline serum potassium) yielded an AUROC of 0.845 and a decile calibration plot comparable to our health plan data model.³⁷

A potential advantage of administrative claims over clinical data is that it essentially covers an individual's entire continuum of care. Still, claims may under identify HK,³⁸ lab results may not be available, and false positive and negative identifications might arise from lags in claims adjudication. Health plans are familiar with addressing these limitations, and our findings of approximately 6% HK prevalence in both years (BY/PY) is reassuring in that regard. In total, 21 independent predictors were included in the final model. The model may be theoretical in clinical practice, and we usually transform the model's parameters into a score; but with artificial intelligence technology, it could be implemented in clinical decision-making systems to identify high risk or subpopulations likely to benefit from an early intervention.

A population health program must weigh the marginal cost of outreach against the probability and potential impact of a true identification to maximize the value of its resources. The clinical and economic importance of optimizing RAASi dosing,^{39–41} together with the emergence of novel potassium binders,^{24,39–44} make it high value to quantitatively predict the onset of HK in CKD. Health plans will need to adjust their already robust, informatics-driven identification, risk-stratification, and efficient outreach programs to address these concerns.

Conclusion

By slowing the progression of CKD, RAASi have transformed its management, but HK can threaten the optimal use and dosing of these drugs. To facilitate broader, safe use of RAASi, we demonstrated that a predictive model based on administrative claims and laboratory results provides an acceptable level of accuracy in identifying both very high and very low risk CKD patients

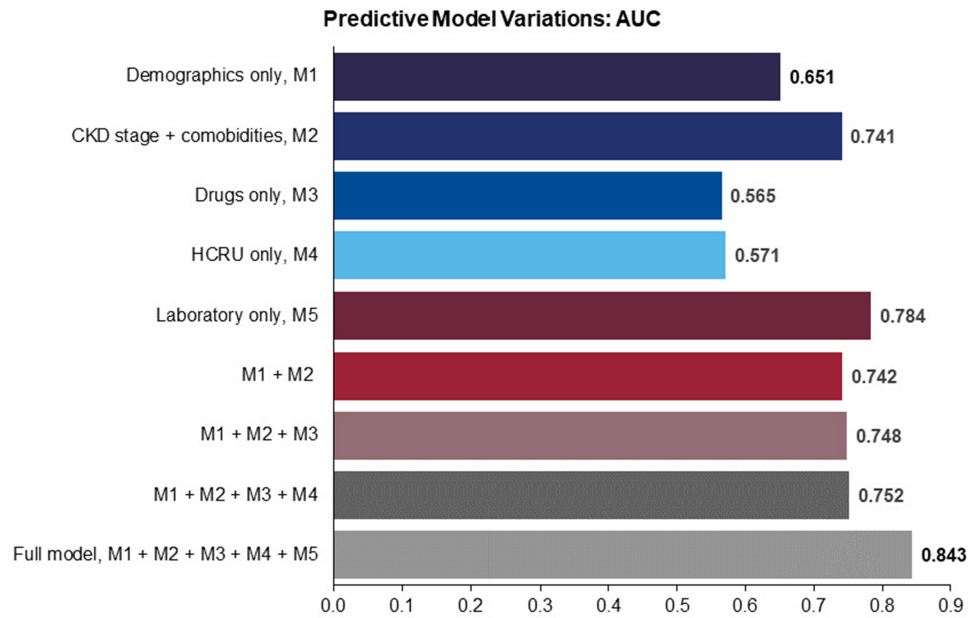


Figure 2 Nine candidate models based on single and combinations of variables.

Abbreviations: AUROC, area under the receiver operating characteristic curve; CKD, chronic kidney disease; HCRU, healthcare resource utilization.

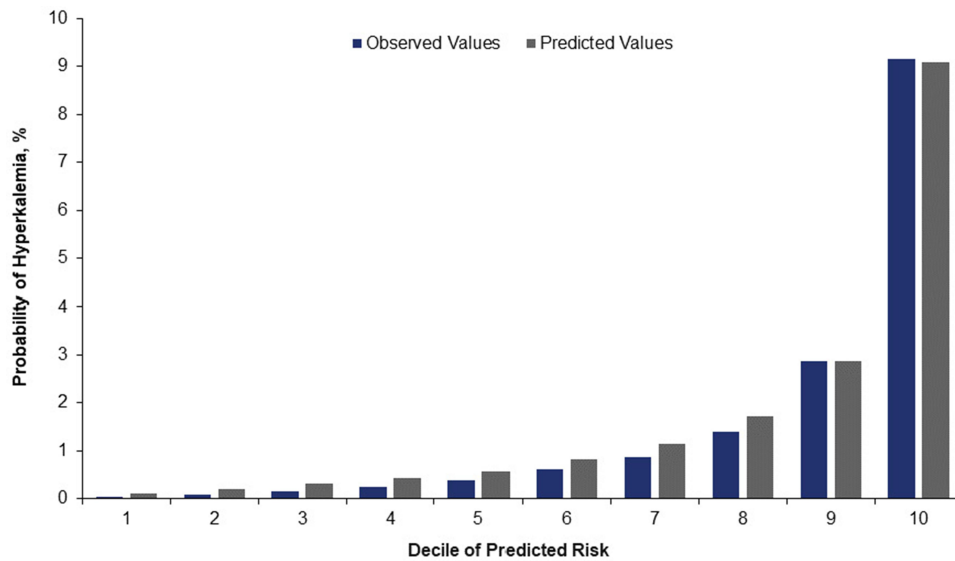


Figure 3 HK risk estimation and stratification performance. Calibration plot of observed versus predicted risk of HK during the follow-up period. The predicted risk estimated by the model stratifies the population and yields estimates of the average risk of HK (gray bar) within each decile (risk stratum). The estimates are compared to the actual (observed) risk in each decile (blue bar).

for HK in a large health plan. The model has been internally validated, but still needs external validation. When validated, this model could improve health

outcomes and lower associated costs in the CKD population at the point of care or in population health strategies.

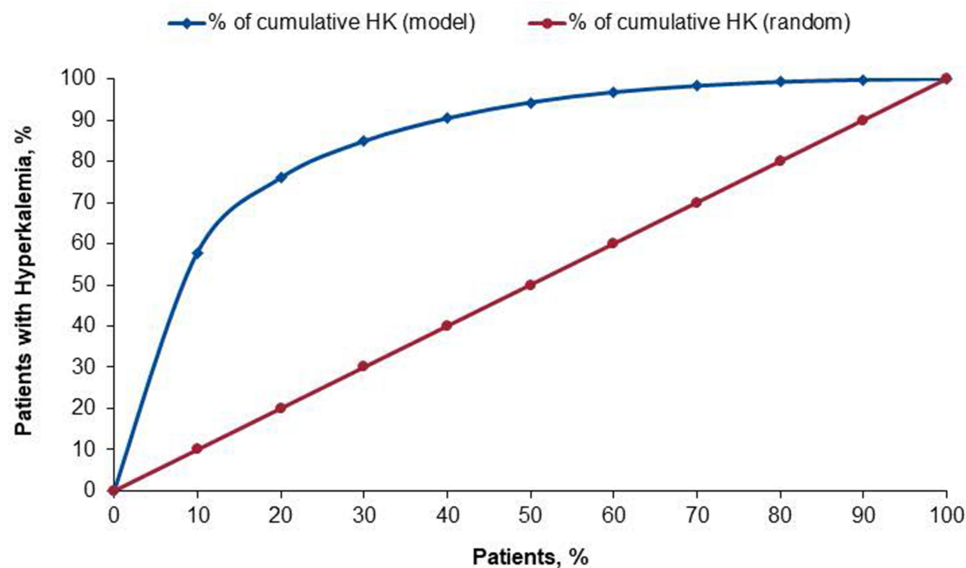


Figure 4 Model Validation: Gains chart. The predicted risk stratifies the population and evaluates the cumulative rate of actual HK at each decile (blue line) within each decile. Starting from the highest- toward the lowest-risk decile, the cumulative rate of HK is compared to the rate without predictive model (random selection) in each decile (red line).

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Author Contributions

All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; agreed to submit to the current journal; gave final approval of the version to be published; and agree to be accountable for all aspects of the work.

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