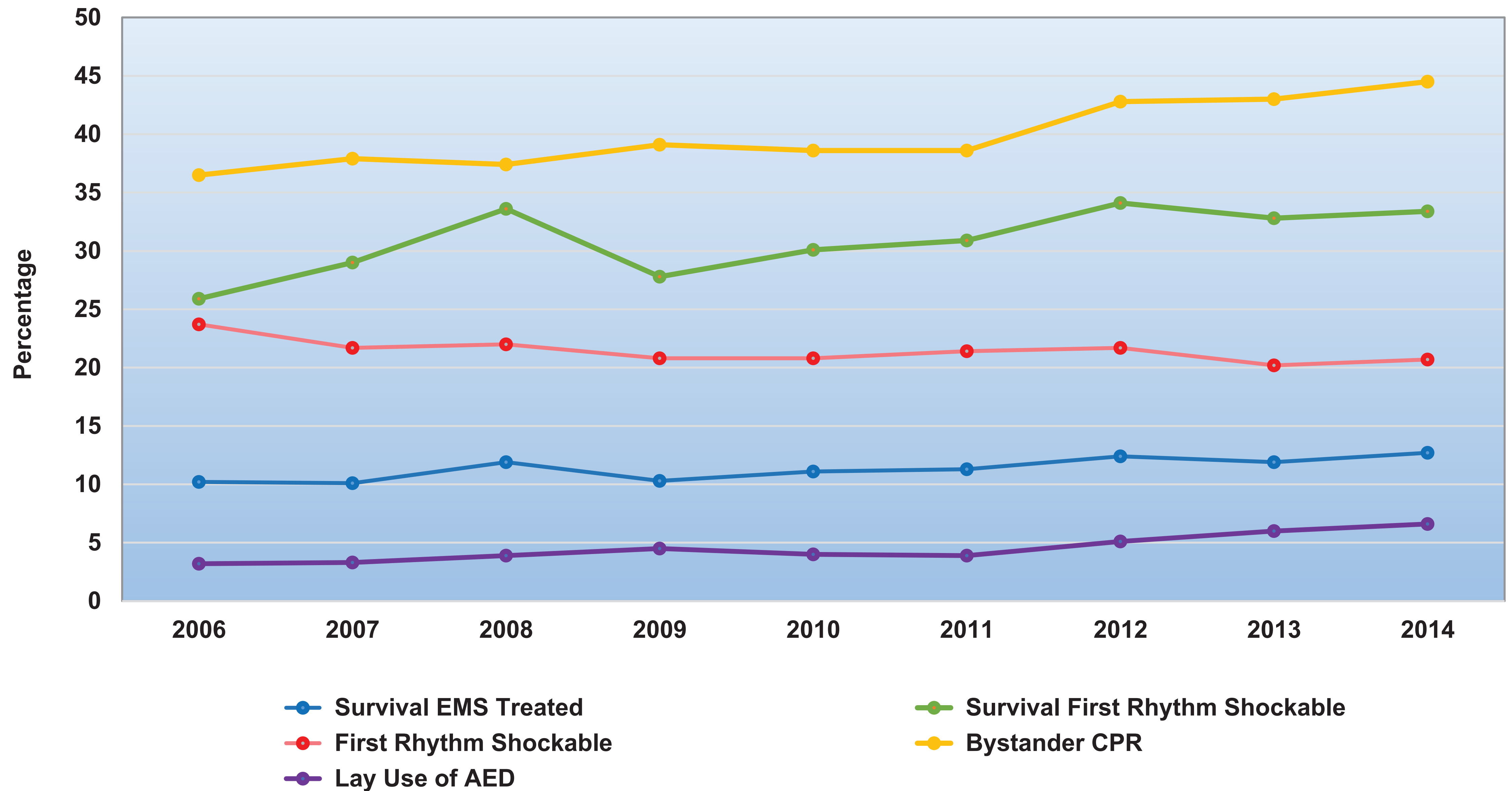


Predicting and explaining in-hospital mortality following cardiac arrest.

A machine learning approach



Clinical paper

APACHE II scoring to predict outcome in post-cardiac arrest

Michael W. Donnino^{a,b,*}, Justin D. Saliccioli^a, Andre Dejam^c, Tyler Giberson^a, Brandon Giberson^a, Cristal Cristia^a, Shiva Gautam^d, Michael N. Cocchi^{a,e}

Conclusions: APACHE II score is a poor predictor of outcome at time zero for out-of-hospital cardiac arrest (OHCA) post-arrest patients consistent with the original development of the score in the critically ill. For in-hospital cardiac arrest (IHCA) at time zero and for both IHCA and OHCA at 24 h and beyond, the APACHE II score was a modest indicator of illness severity and predictor of mortality/neurologic morbidity.

Clinical paper

Survival and outcome prediction using the Apache III and the out-of-hospital cardiac arrest (OHCA) score in patients treated in the intensive care unit (ICU) following out-of-hospital, in-hospital or ICU cardiac arrest[☆]

M.B. Skrifvars^{a,b,*}, B. Varghese^a, M.J. Parr^{a,c}

Conclusions: Latency to ROSC seems to be the most important determinant of survival in patients following ICU care after a cardiac arrest in this single center trial. The OHCA score and the Apache III score offer moderate predictive accuracy in ICU cardiac arrest patients but correlated weakly with each other. Illness severity adjustment for cardiac arrest patients in ICU should include features of both these scoring systems.

Effectiveness of SAPS III to predict hospital mortality for post-cardiac arrest patients[☆]

Magali Bisbal^{a,b,*}, Elisabeth Jouve^c, Laurent Papazian^{d,e}, Sophie de Bourmont^{a,b}, Gilles Perrin^b, Beatrice Eon^b, Marc Gainnier^{a,b}

Conclusions: The SAPS III did not predict mortality in patients admitted to ICU after CA. The amount of time before specialized CPR, the low-flow interval and the absence of an initial ventricular arrhythmia appeared to be independently associated with mortality and these factors should be used to predict mortality for these patients.

Key predictive variables

Summed from the literature.

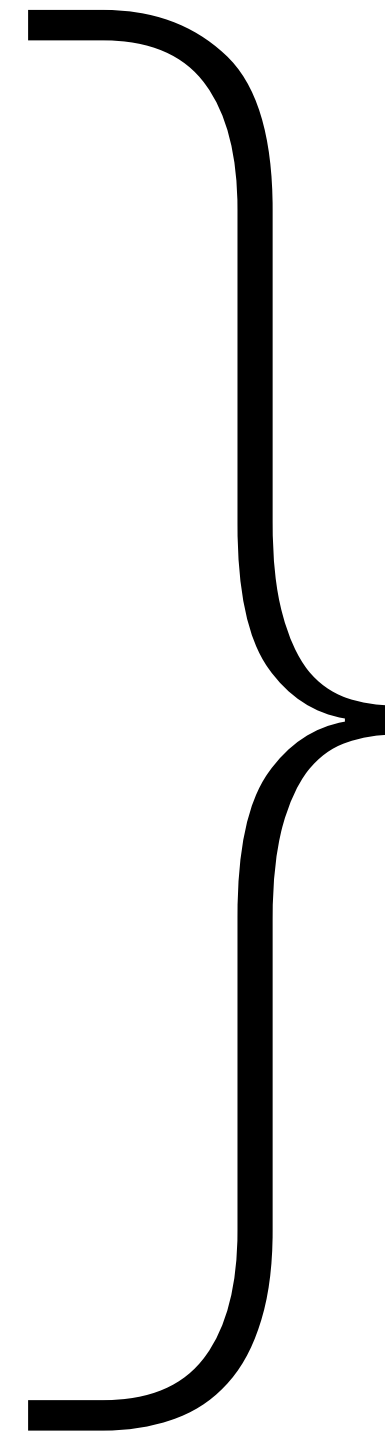
rhythm on admission

time to ROSC

bystander CPR

time to adrenaline

duration of CPR



simplicity of prognostication

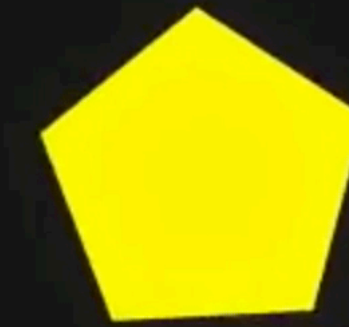
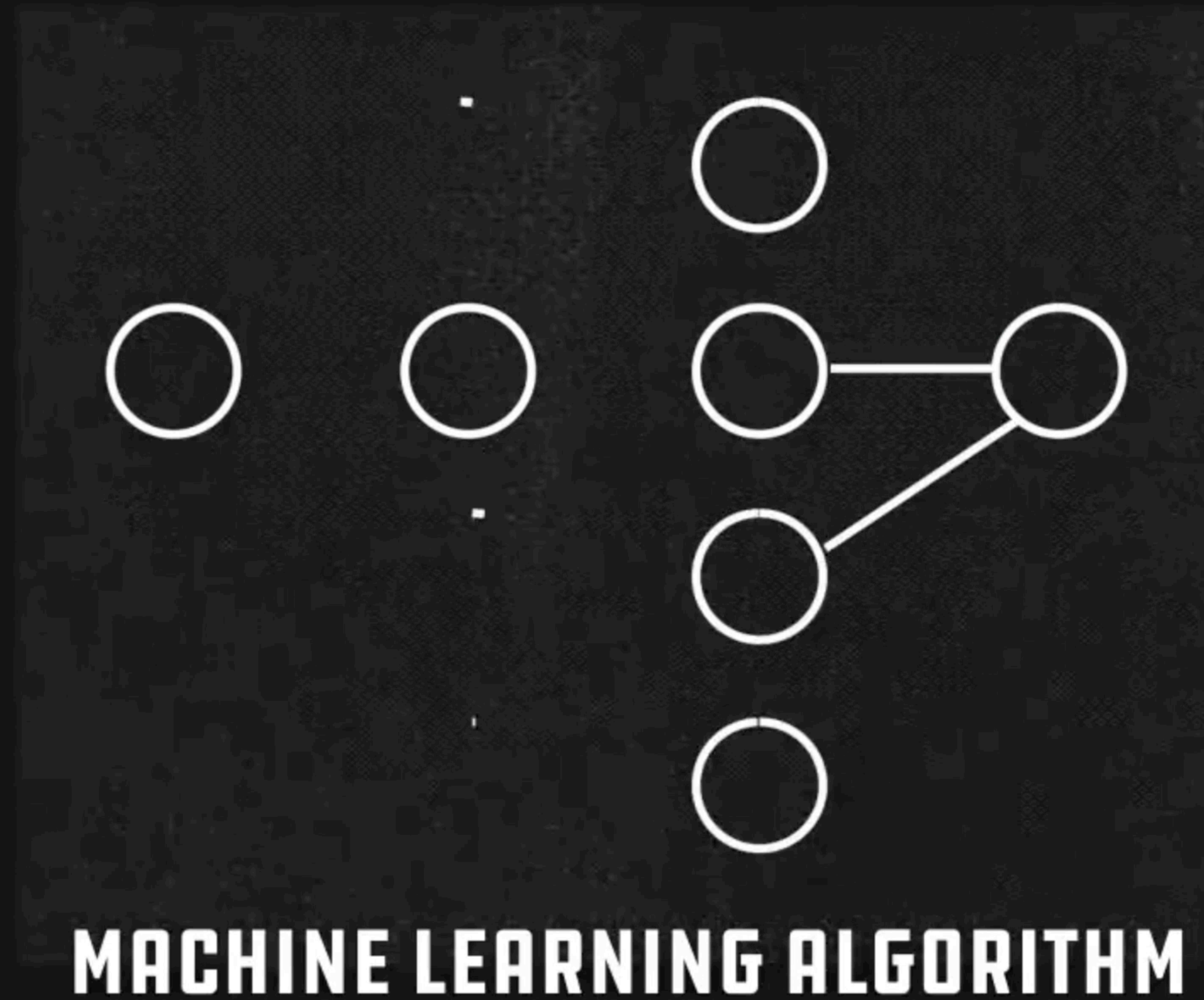
After hearing for several decades that computers will soon be able to assist with difficult diagnoses, the practicing physician may well wonder why the revolution has not occurred.

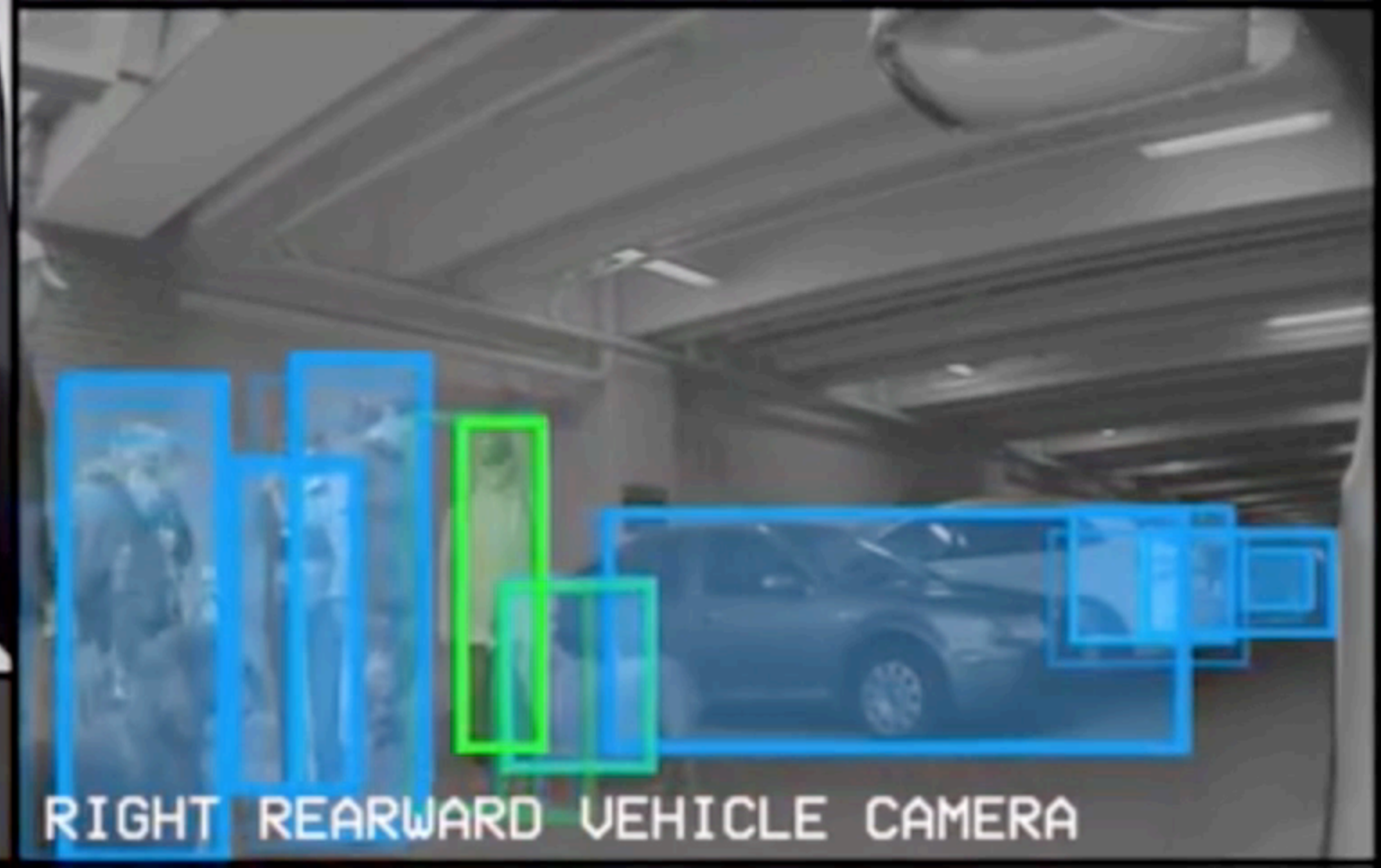
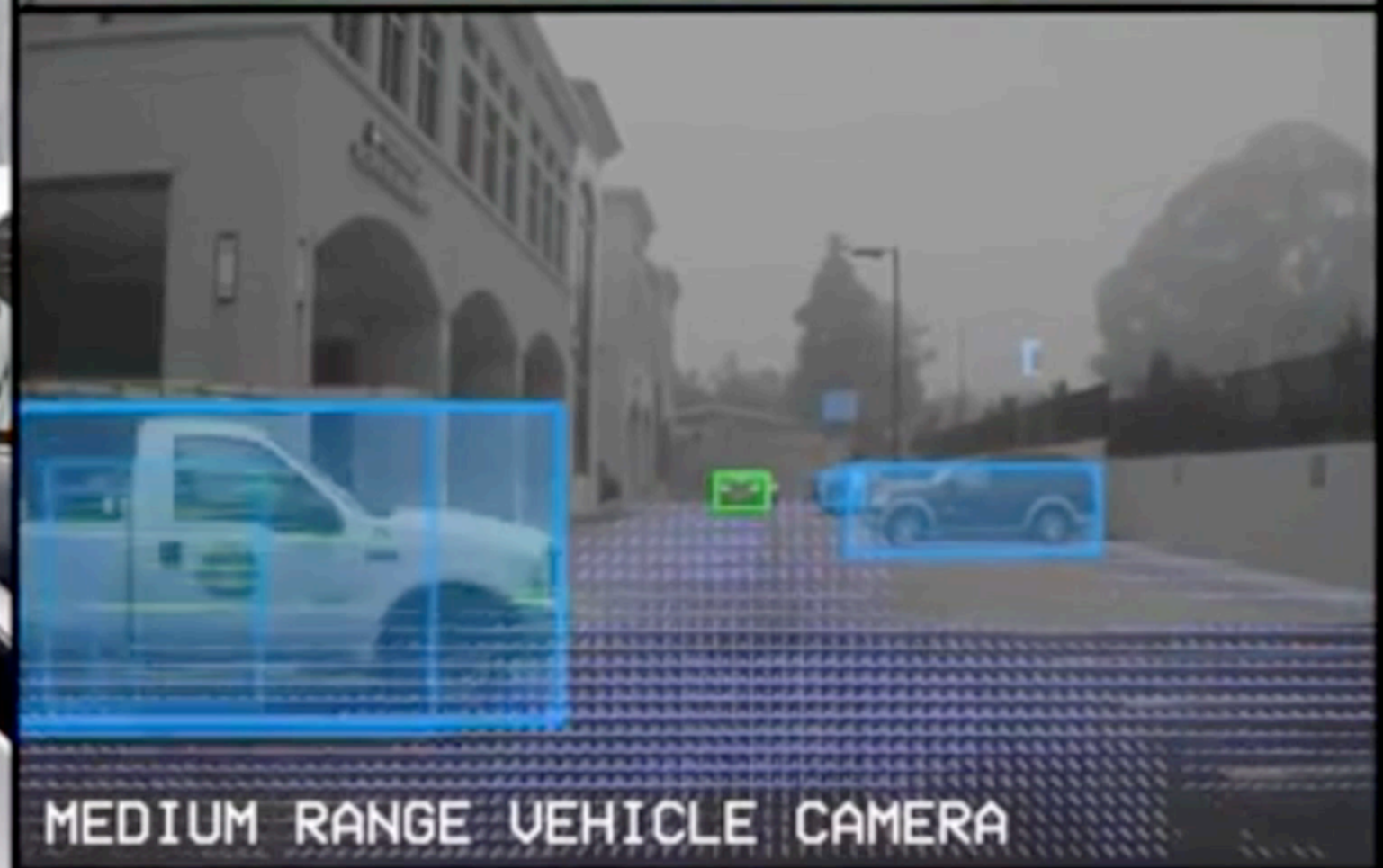
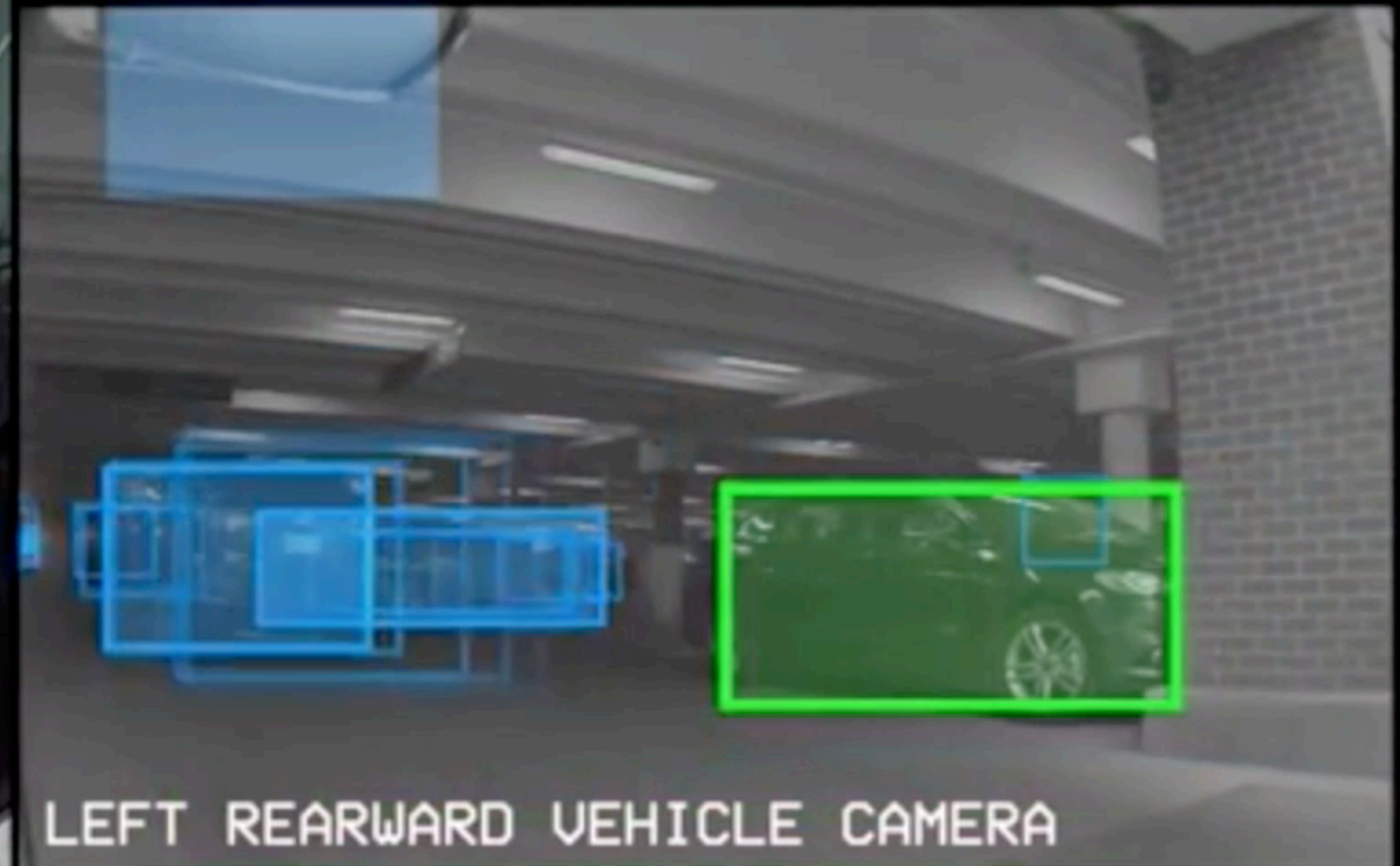
A P R I L 1 9 8 7



The NEW ENGLAND
JOURNAL of MEDICINE

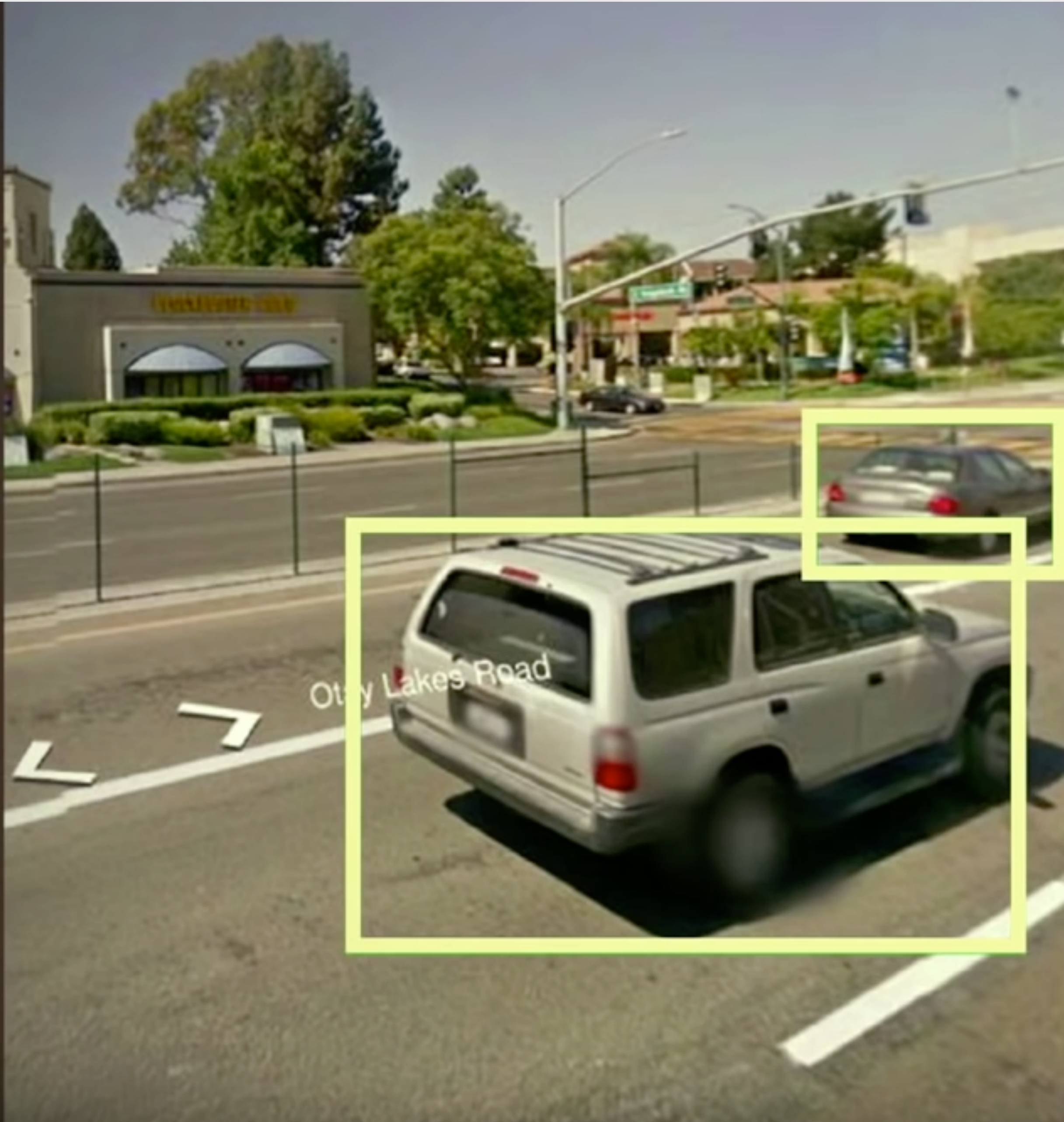
“Machine learning turns this around: in goes the data and the desired result and out comes the algorithm that turns one into the other.”





Legend for the main driving view:

- MOTION FLOW
- LANE LINES
- LANE LINES
- ROAD FLOW
- IN-PATH OBJECTS
- ROAD LIGHTS
- OBJECTS
- ROAD SIGNS

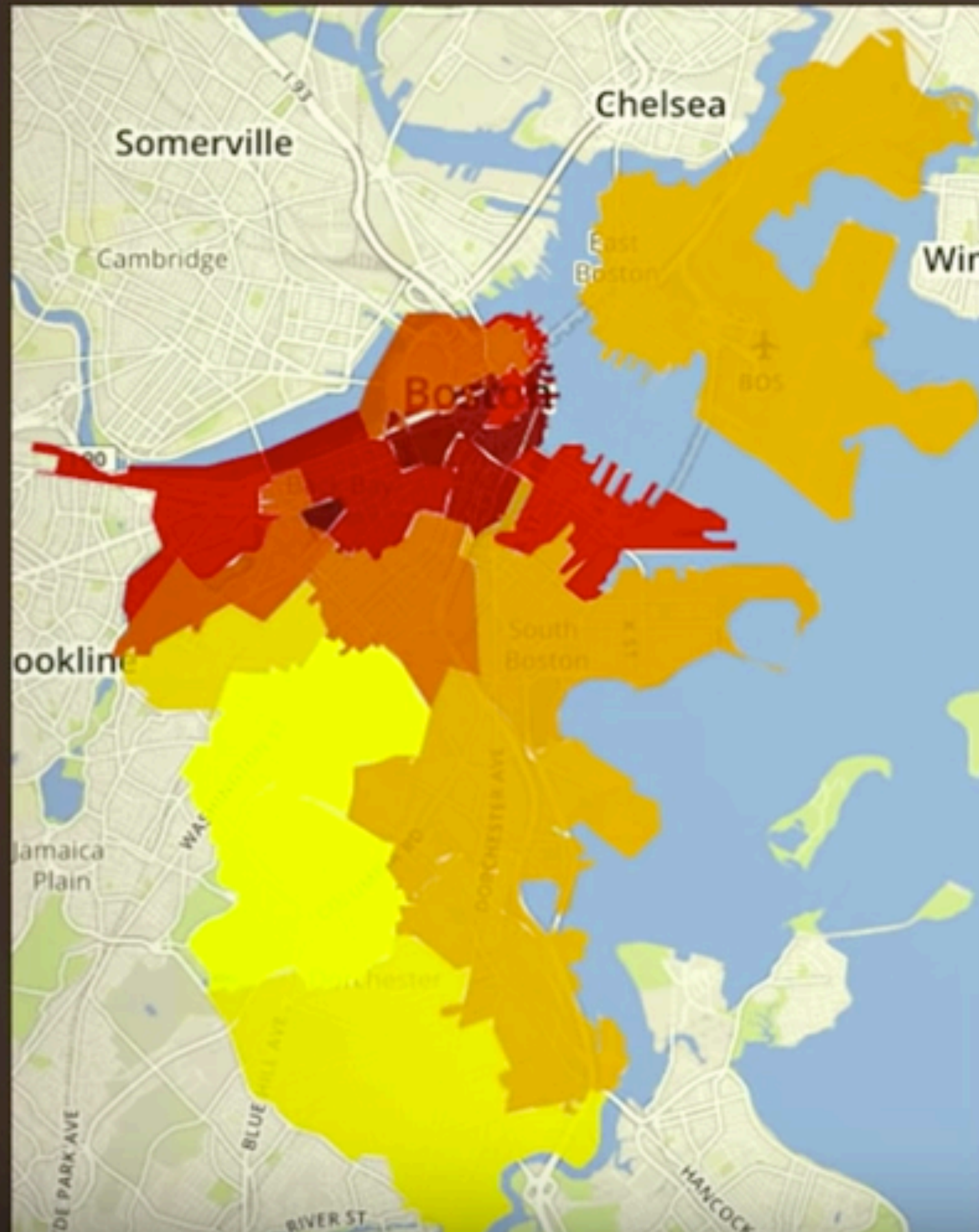


2005 Nissan
Sentra 1.8 S

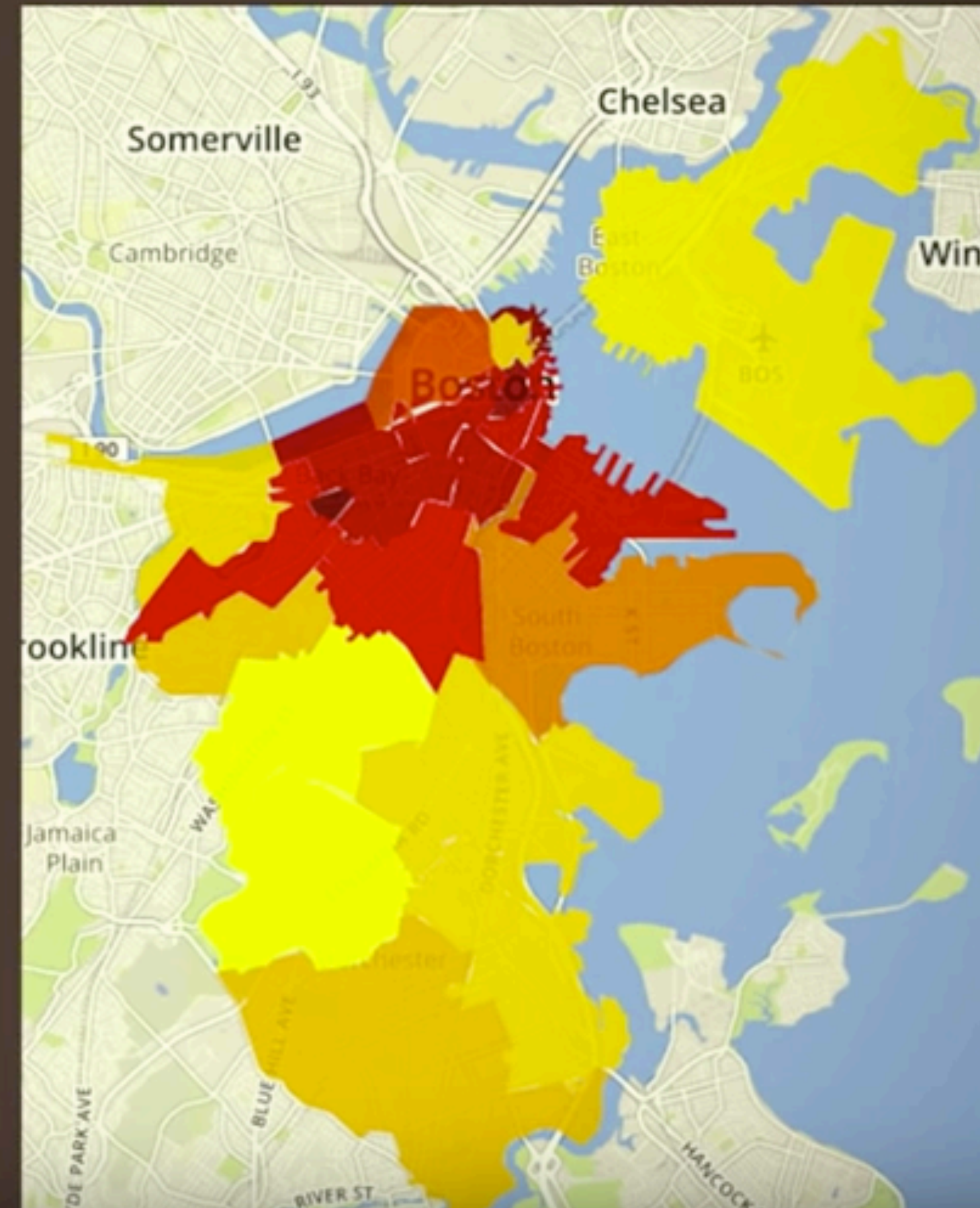
2000 Toyota
4Runner Limited

Income in Boston, MA

Actual

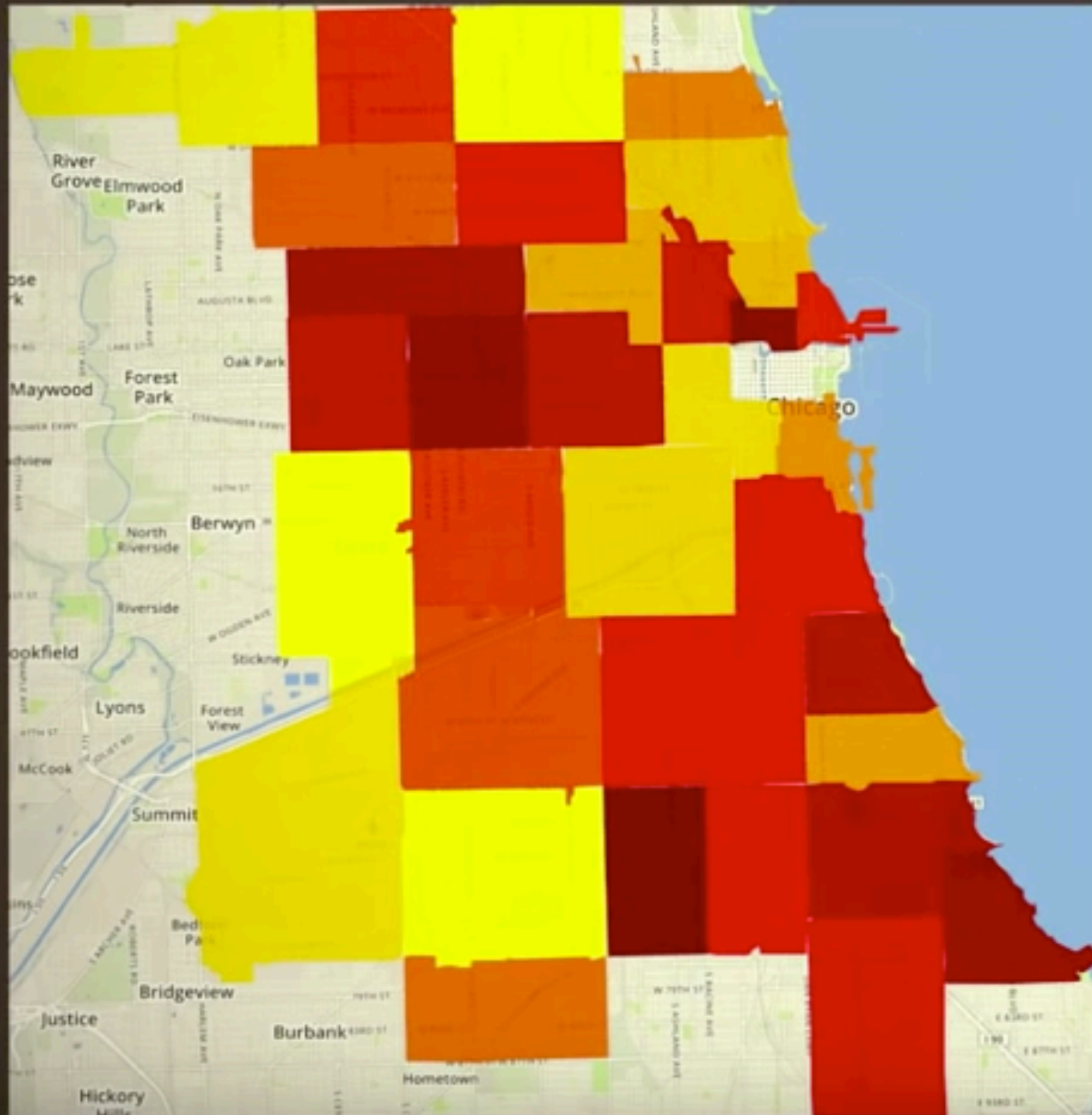


Predicted by Car Price

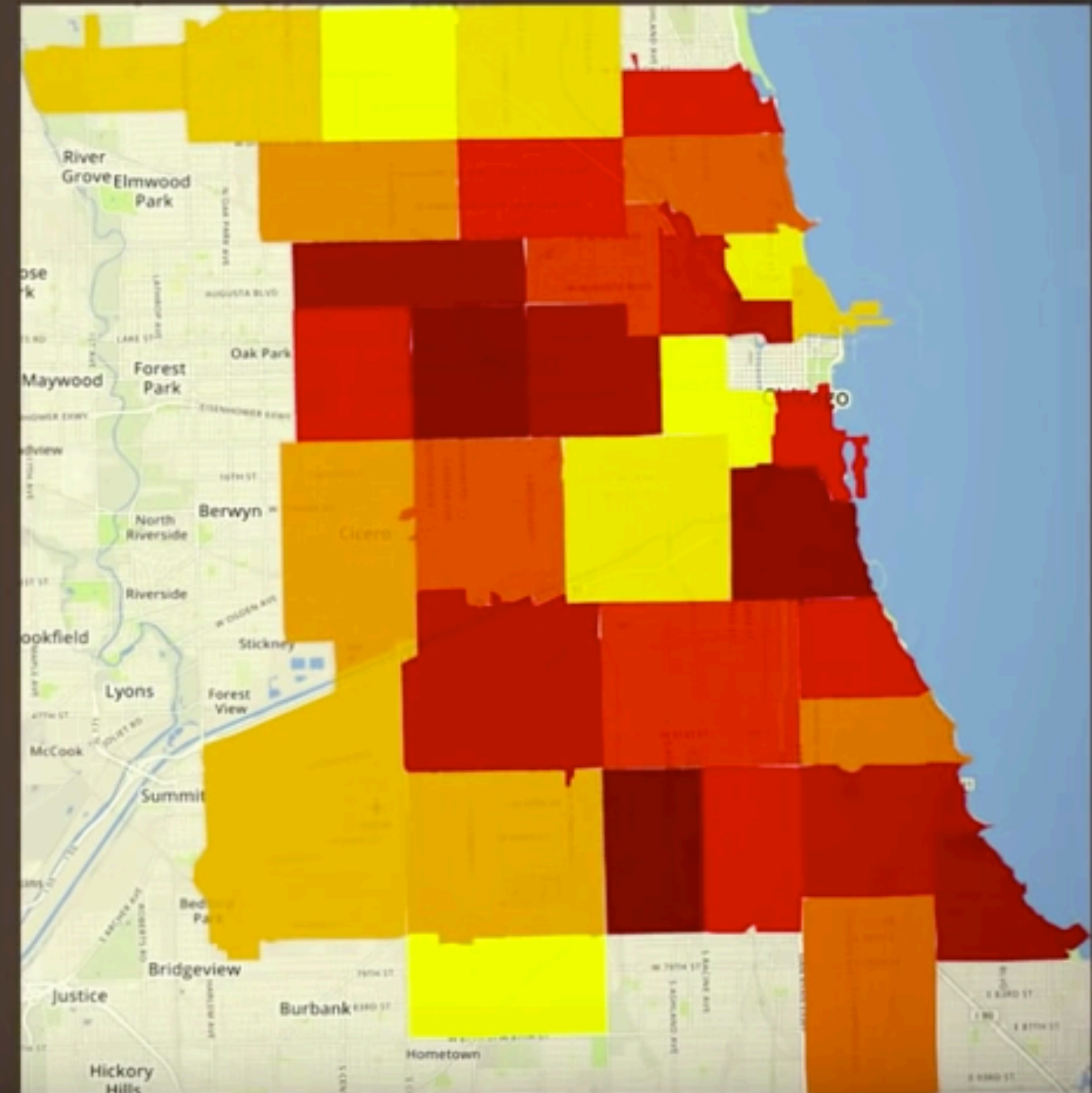


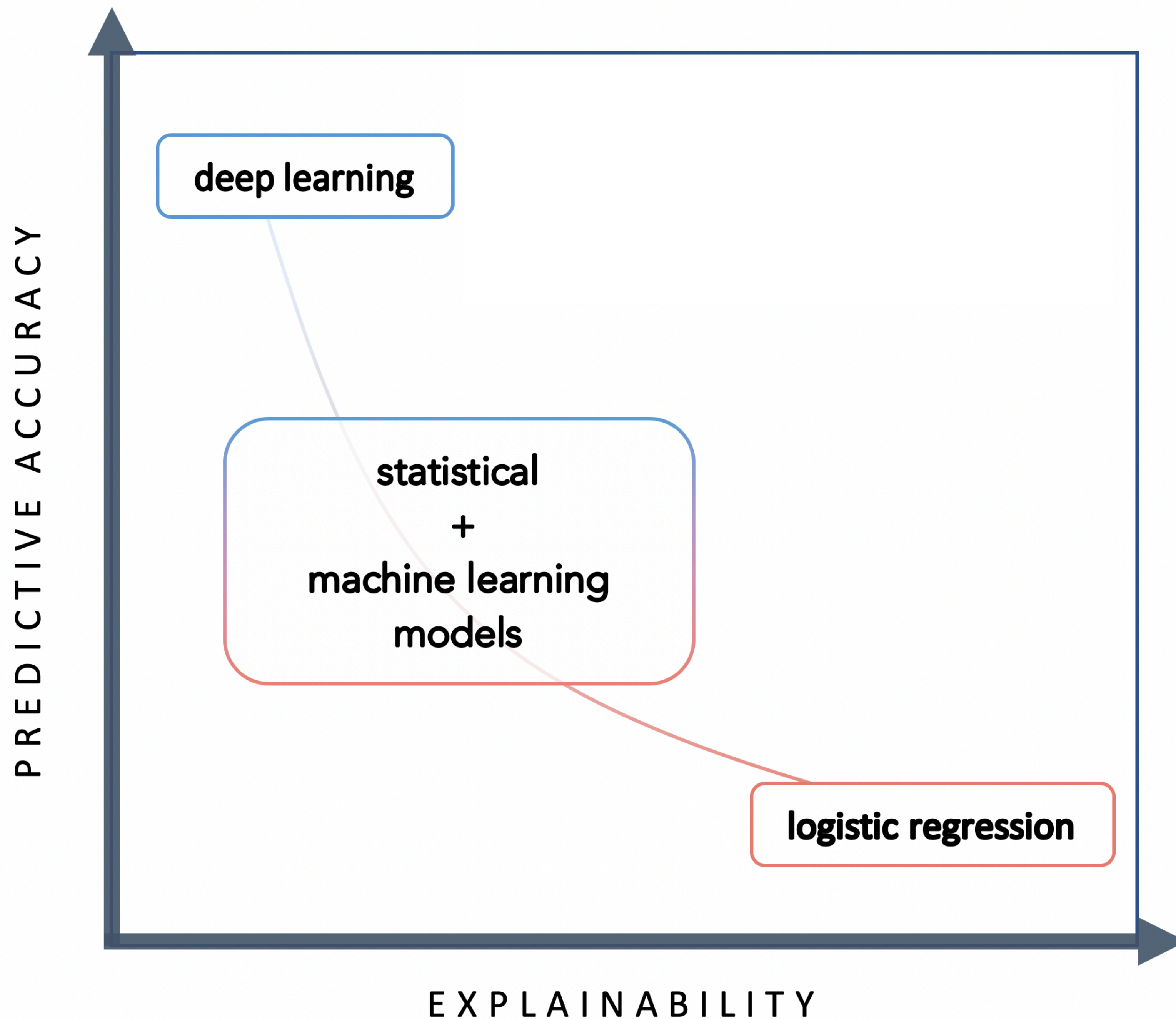
Crime in Chicago, IL

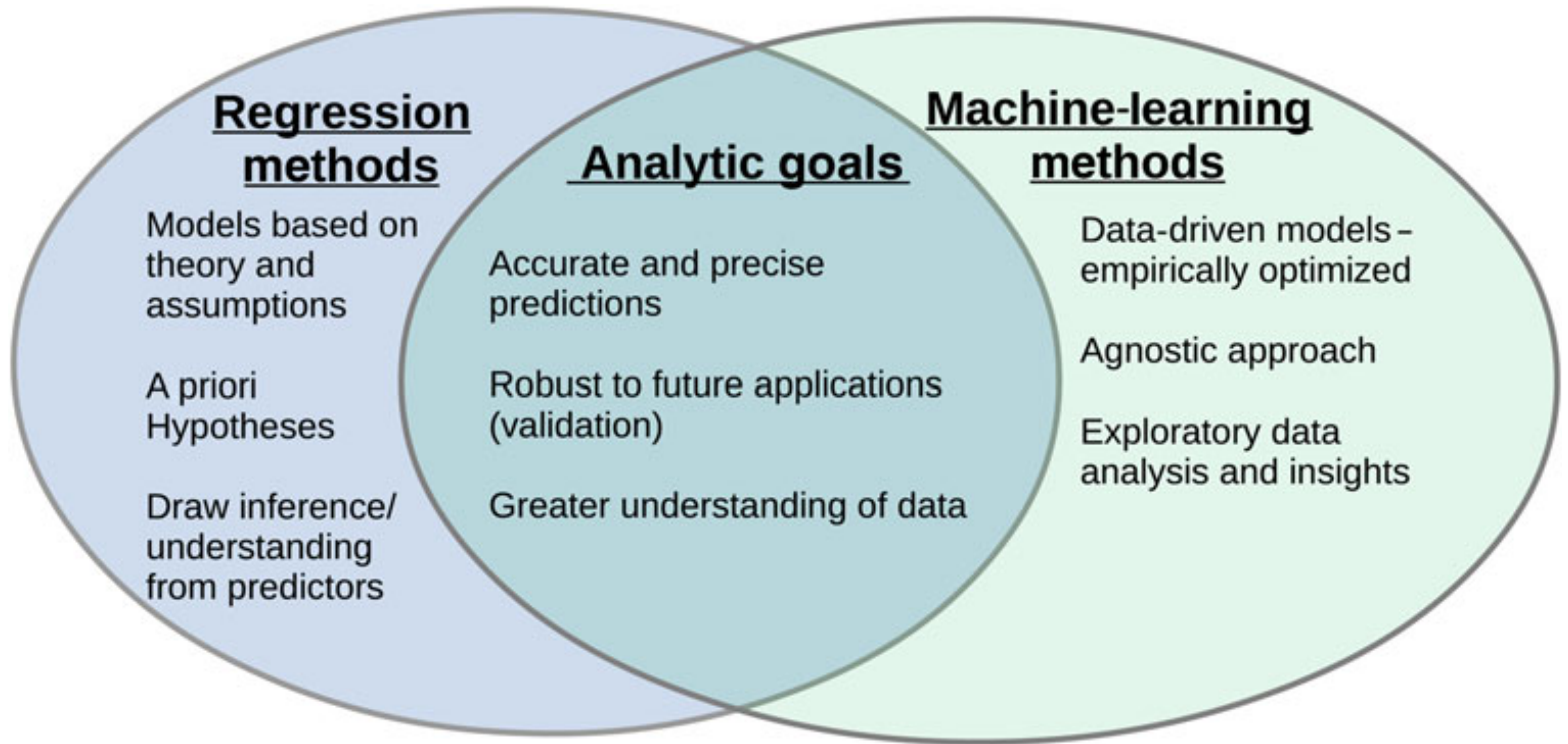
Actual



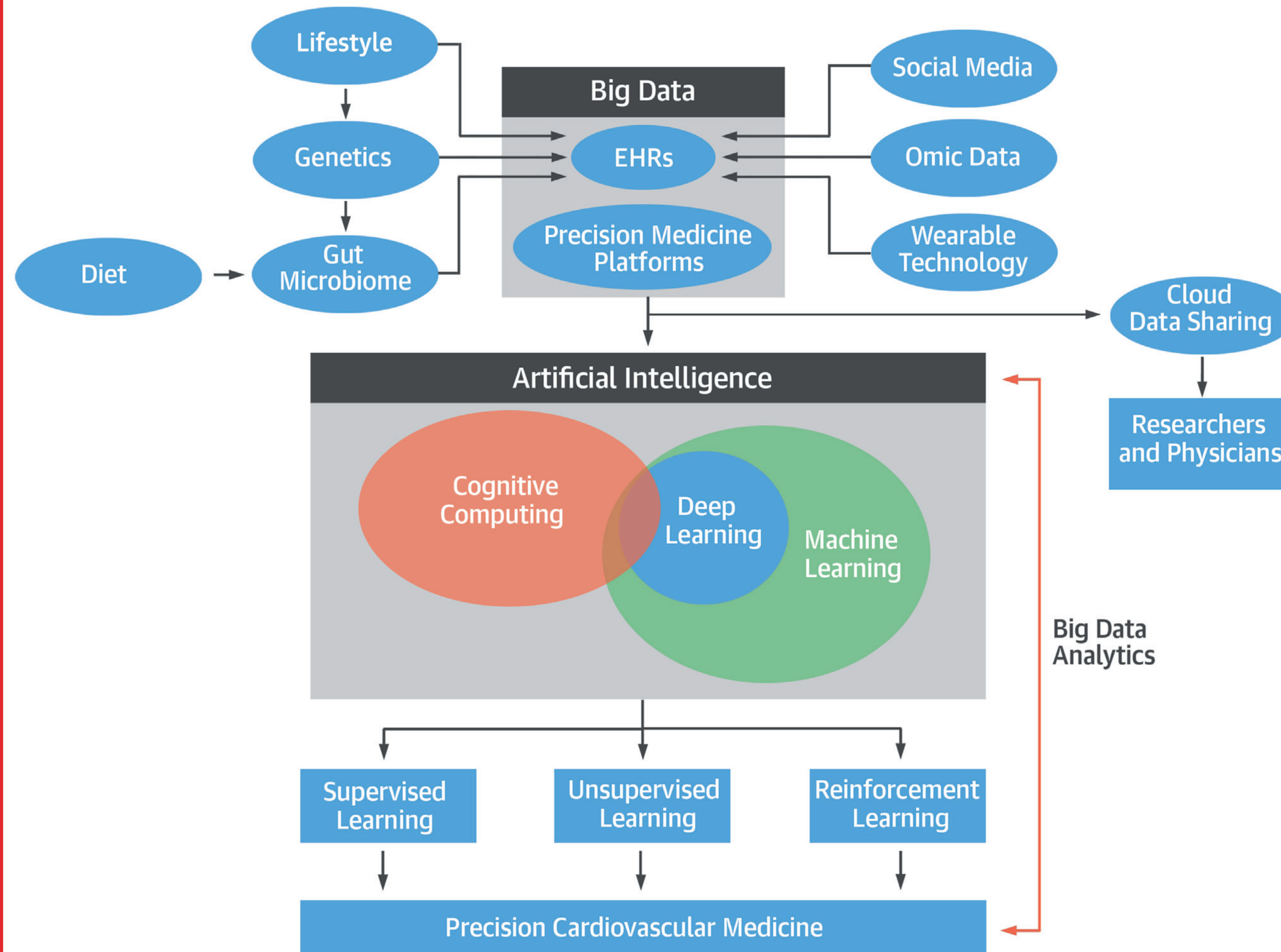
Predicted by Car Price







CENTRAL ILLUSTRATION Artificial Intelligence in Precision Cardiovascular Medicine



Krittanawong, C. et al. J Am Coll Cardiol. 2017;69(21):2657-64.

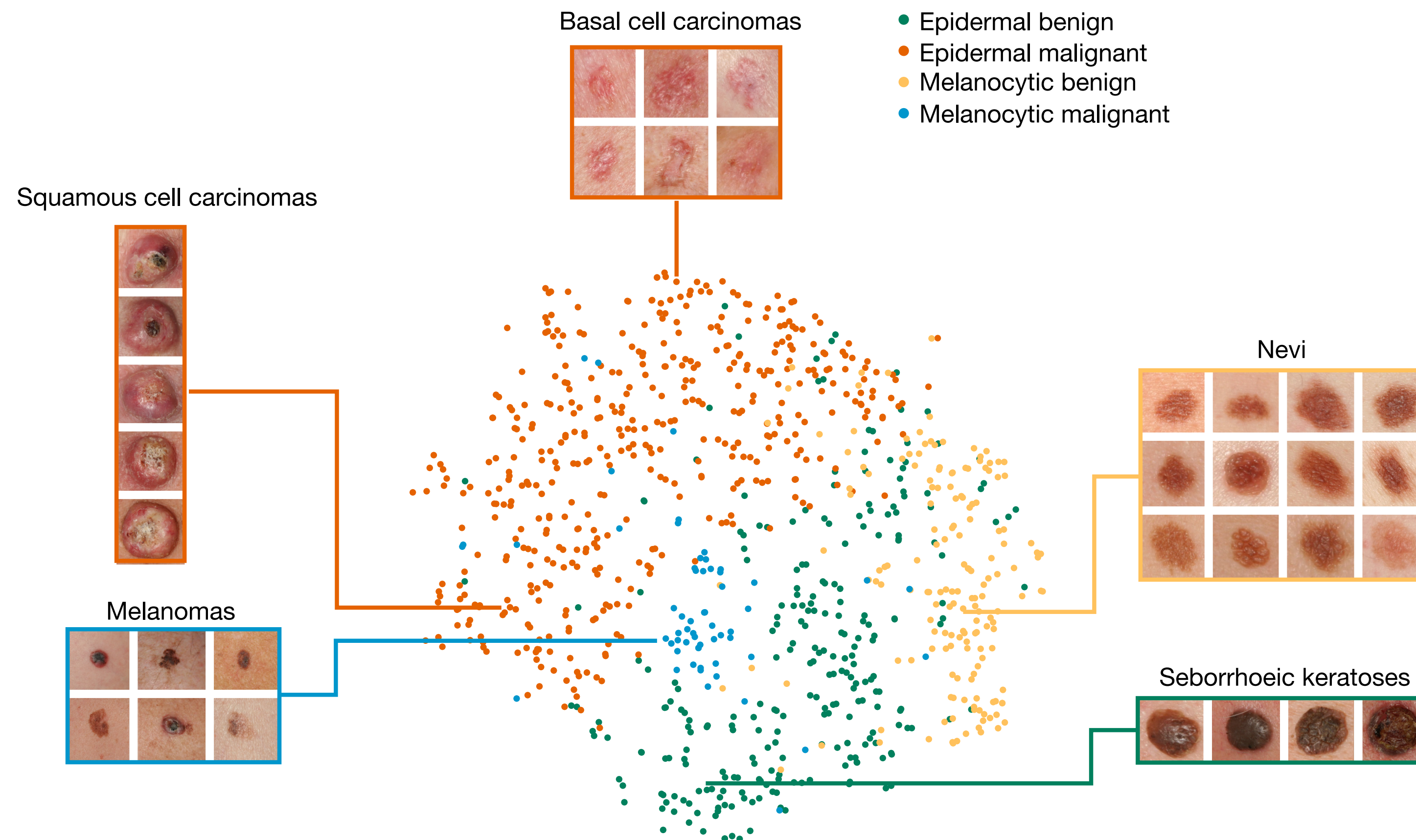
Big data (genetics, social media, environmental, and lifestyle-related factors, or "omic" data) can be stored through EHRs or precision medicine platforms, and can be shared for data analysis with other physicians or researchers through secure cloud systems. Big data analytics using artificial intelligence (machine learning, deep learning, or cognitive computing) and 3 main types of learning algorithms (supervised, unsupervised, and reinforcement learning) will enable precision cardiovascular medicine. EHR = electronic health record.

We can automate anything a person can do with less than one second of thinking.







Andrew Ng

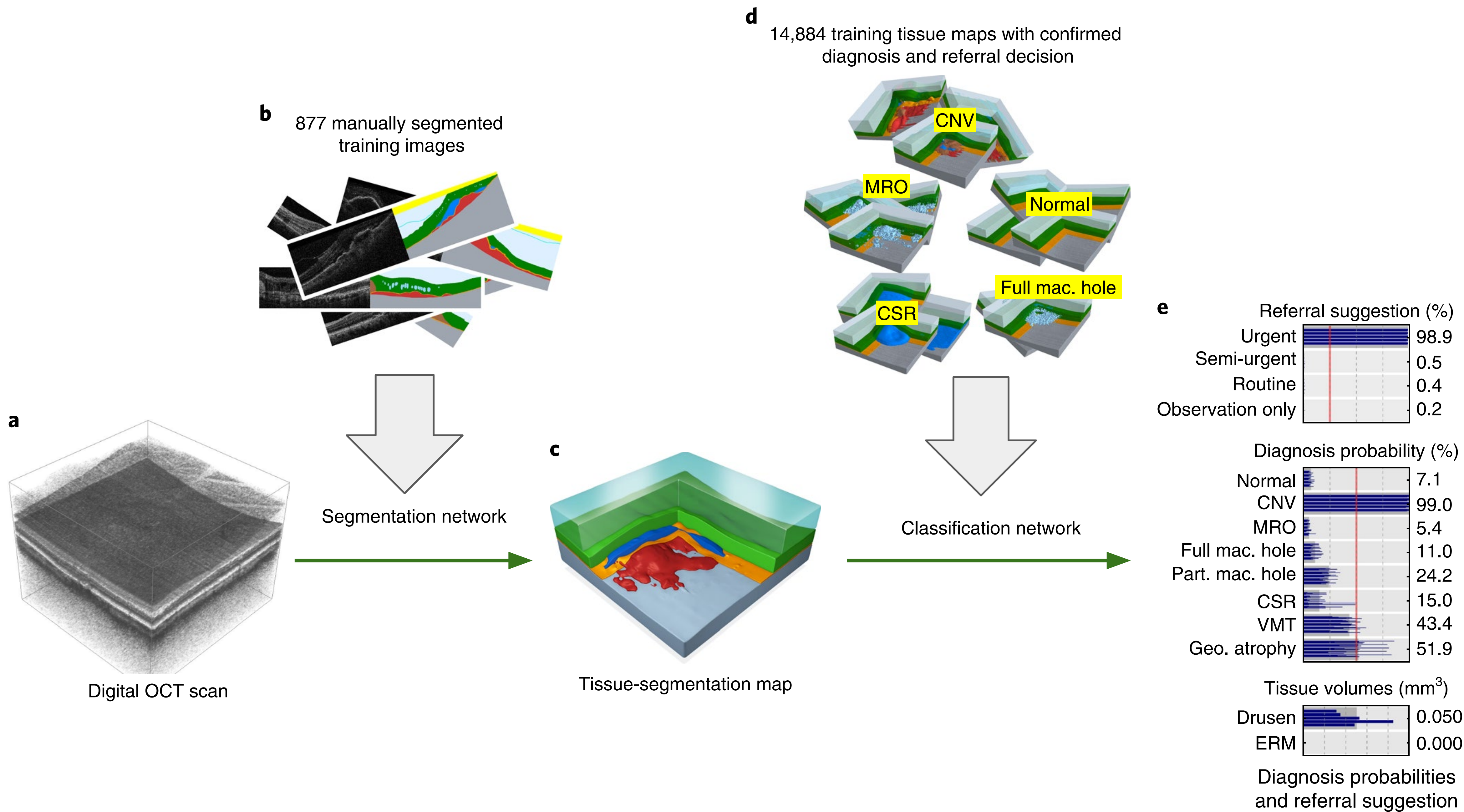
Dermatologist-level classification of skin cancer with deep neural networks

Andre Esteva^{1*}, Brett Kuprel^{1*}, Roberto A. Novoa^{2,3}, Justin Ko², Susan M. Swetter^{2,4}, Helen M. Blau⁵ & Sebastian Thrun⁶



Clinically applicable deep learning for diagnosis and referral in retinal disease

Jeffrey De Fauw¹, Joseph R. Ledsam¹, Bernardino Romera-Paredes¹, Stanislav Nikolov¹, Nenad Tomasev¹, Sam Blackwell¹, Harry Askham¹, Xavier Glorot¹, Brendan O'Donoghue¹, Daniel Visentin¹, George van den Driessche¹, Balaji Lakshminarayanan¹, Clemens Meyer¹, Faith Mackinder¹, Simon Bouton¹, Kareem Ayoub¹, Reena Chopra ², Dominic King¹, Alan Karthikesalingam¹, Cían O. Hughes ^{1,3}, Rosalind Raine³, Julian Hughes², Dawn A. Sim², Catherine Egan², Adnan Tufail², Hugh Montgomery ³, Demis Hassabis¹, Geraint Rees ³, Trevor Back¹, Peng T. Khaw², Mustafa Suleyman¹, Julien Cornebise^{1,3,4}, Pearse A. Keane ^{2,4}★ and Olaf Ronneberger ^{1,4}★



New Research Aims to Solve the Problem of AI Bias in “Black Box” Algorithms

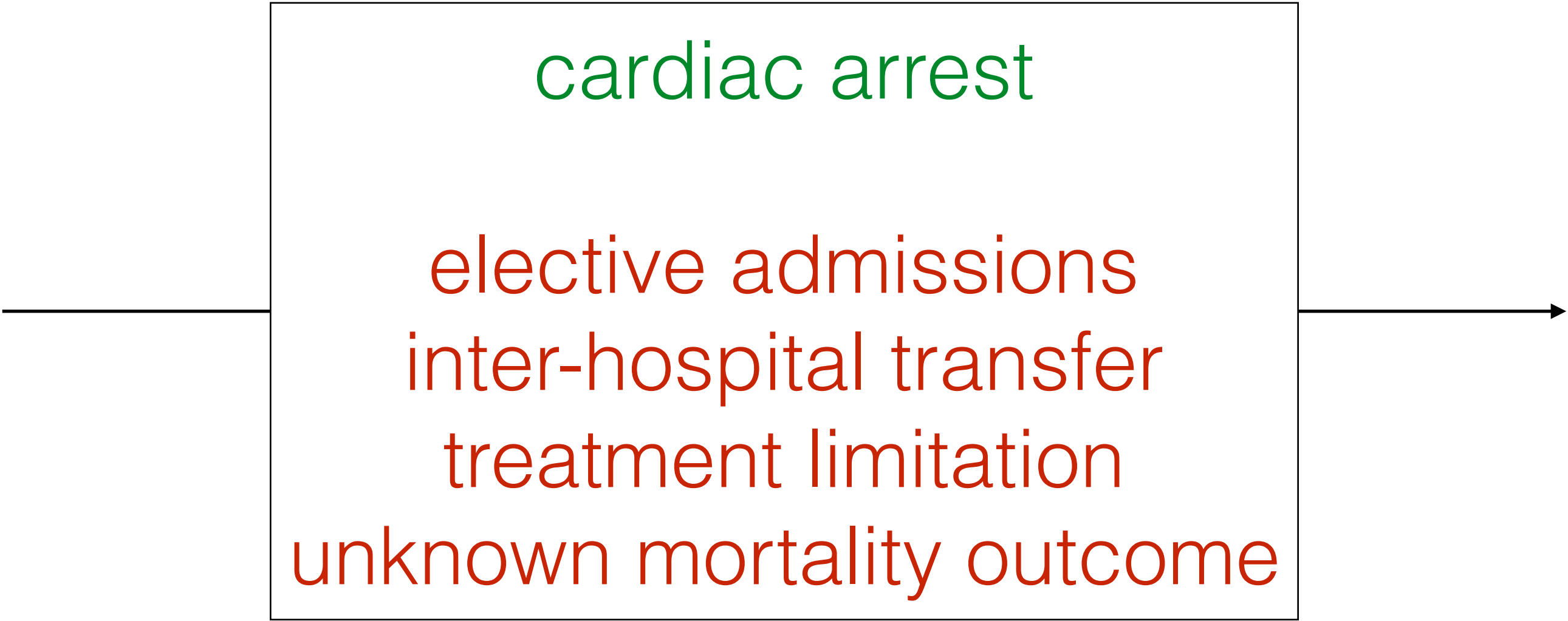
As we automate more and more decisions, being able to understand how an AI thinks is increasingly important.

by Jackie Snow November 7, 2017

ANZICS

Adult Patient Database.

1,484,536
patients



39,566
patients

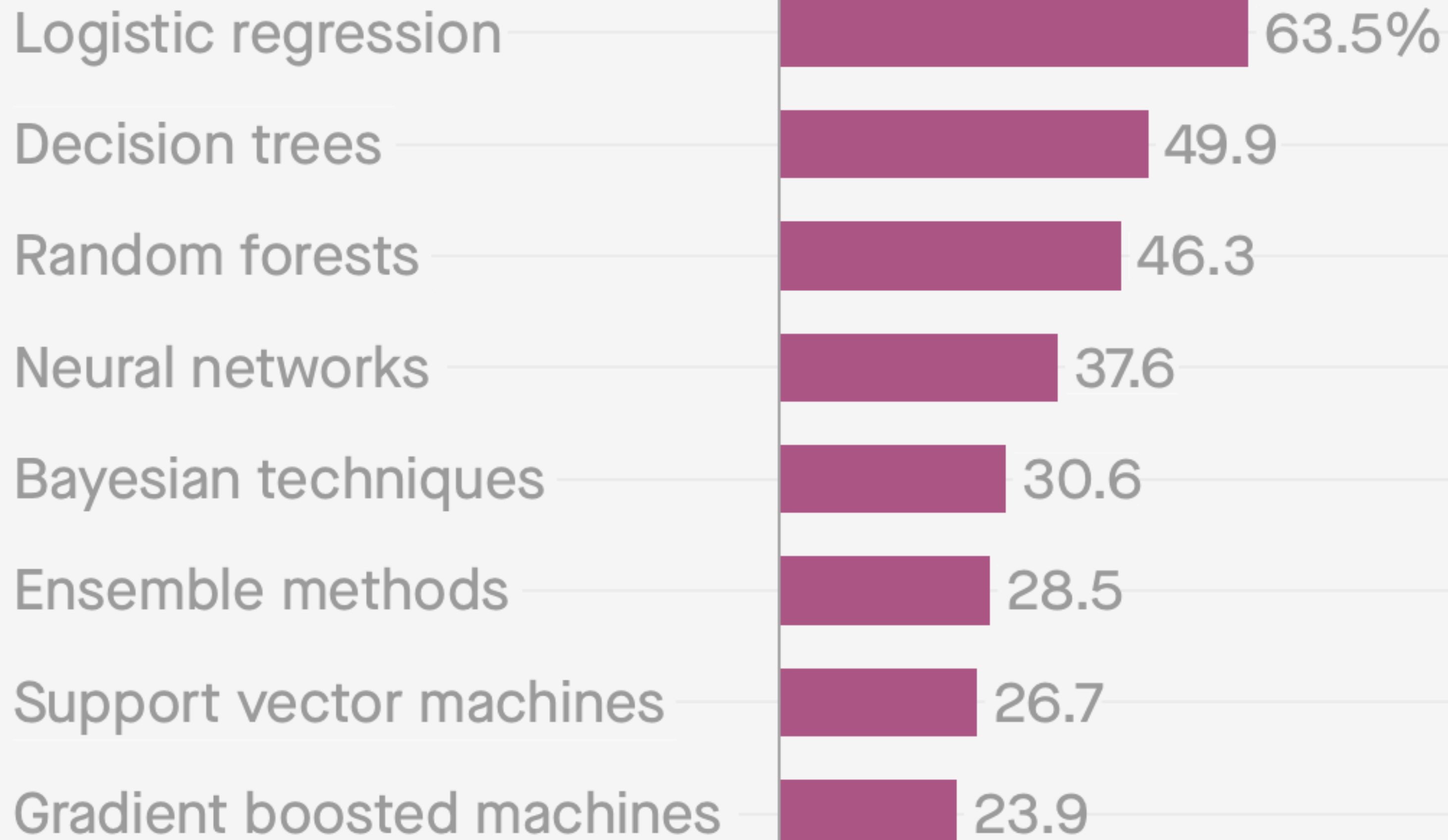
Baseline characteristics

186 ICUs, 39,566 patients.

n	21,547	18,019	p value
	survivors	non-survivors	
age	63 [50-73]	66 [52-77]	<0.001
sex (% male)	14,255 (66.2%)	11,629 (64.5%)	0.001
peak temperature	37.1±1.0	36.8±1.5	<0.001
peak creatinine	101 [76-151]	146 [102-198]	<0.001

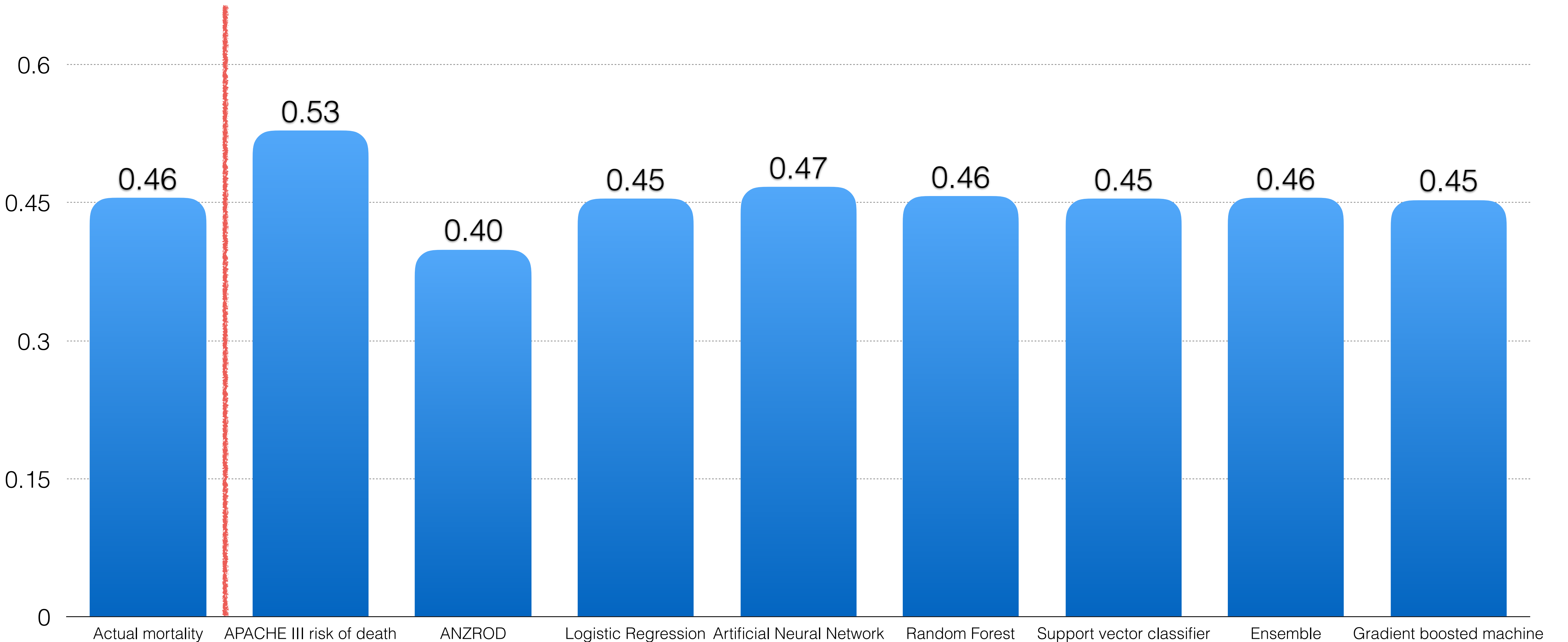
Most popular prediction methods used by data scientists

(Based on a survey of Kaggle users)



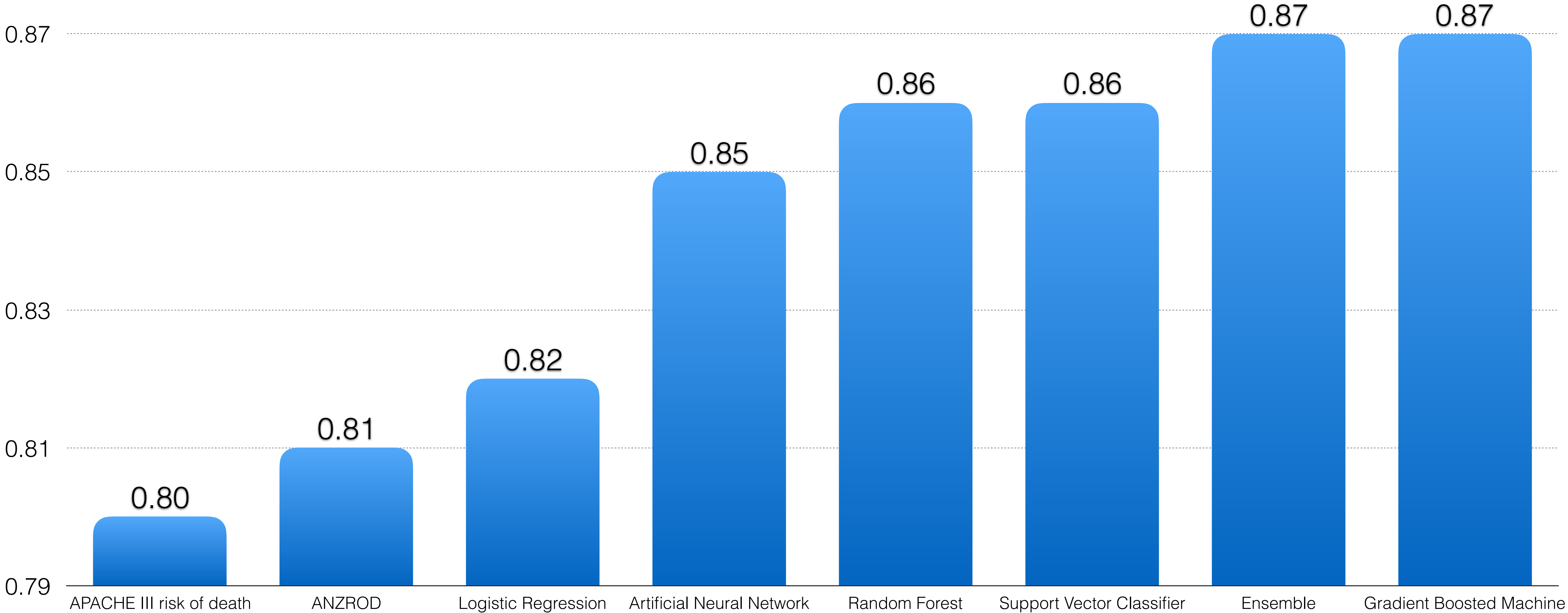
Comparison of model accuracy

Predicted mortality | existing scores, logistic regression, and machine learning models.



Comparison of model accuracy (AUROC)

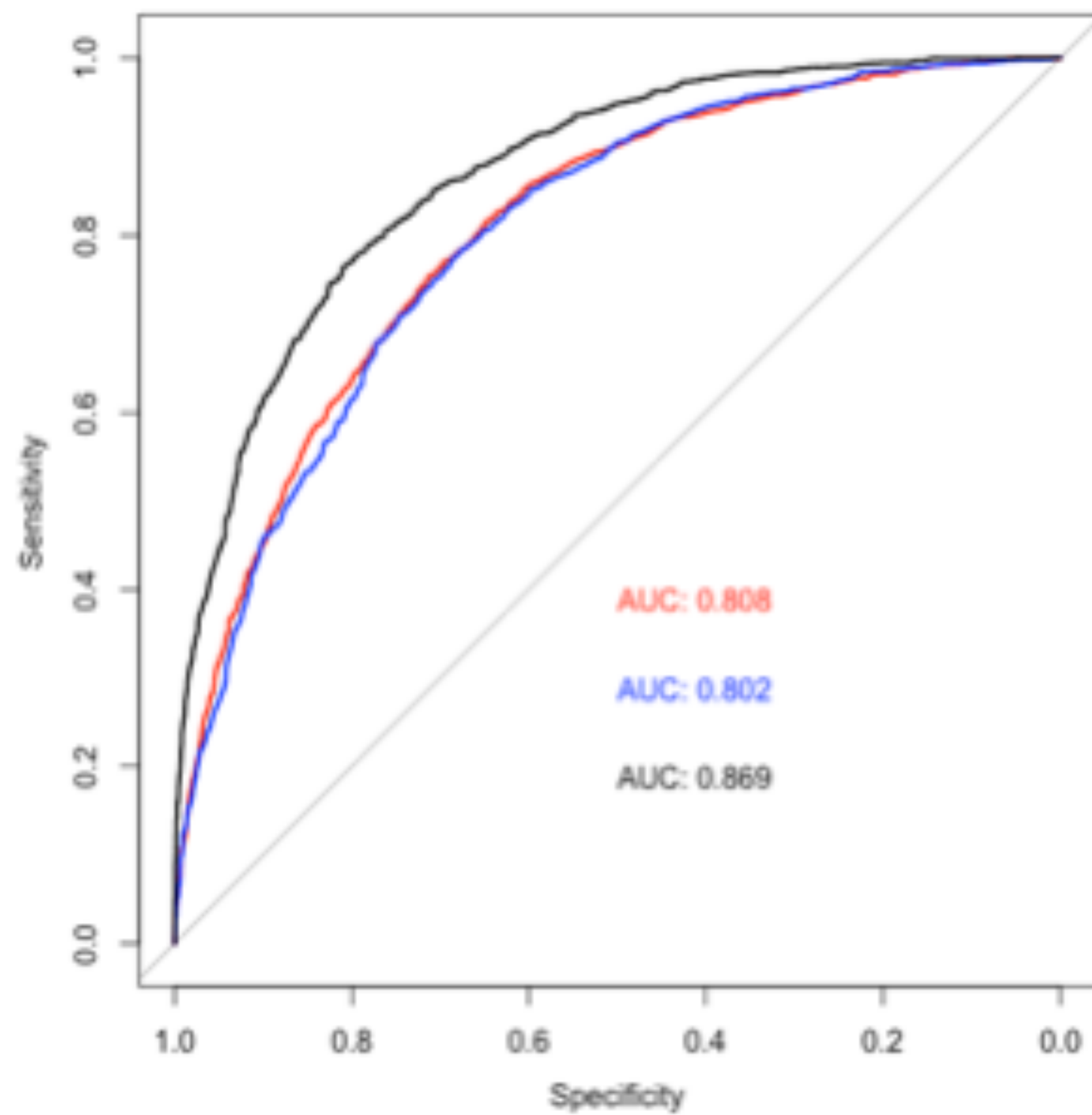
Area under the receiver operating curve | Existing scores, logistic regression, and machine learning models.



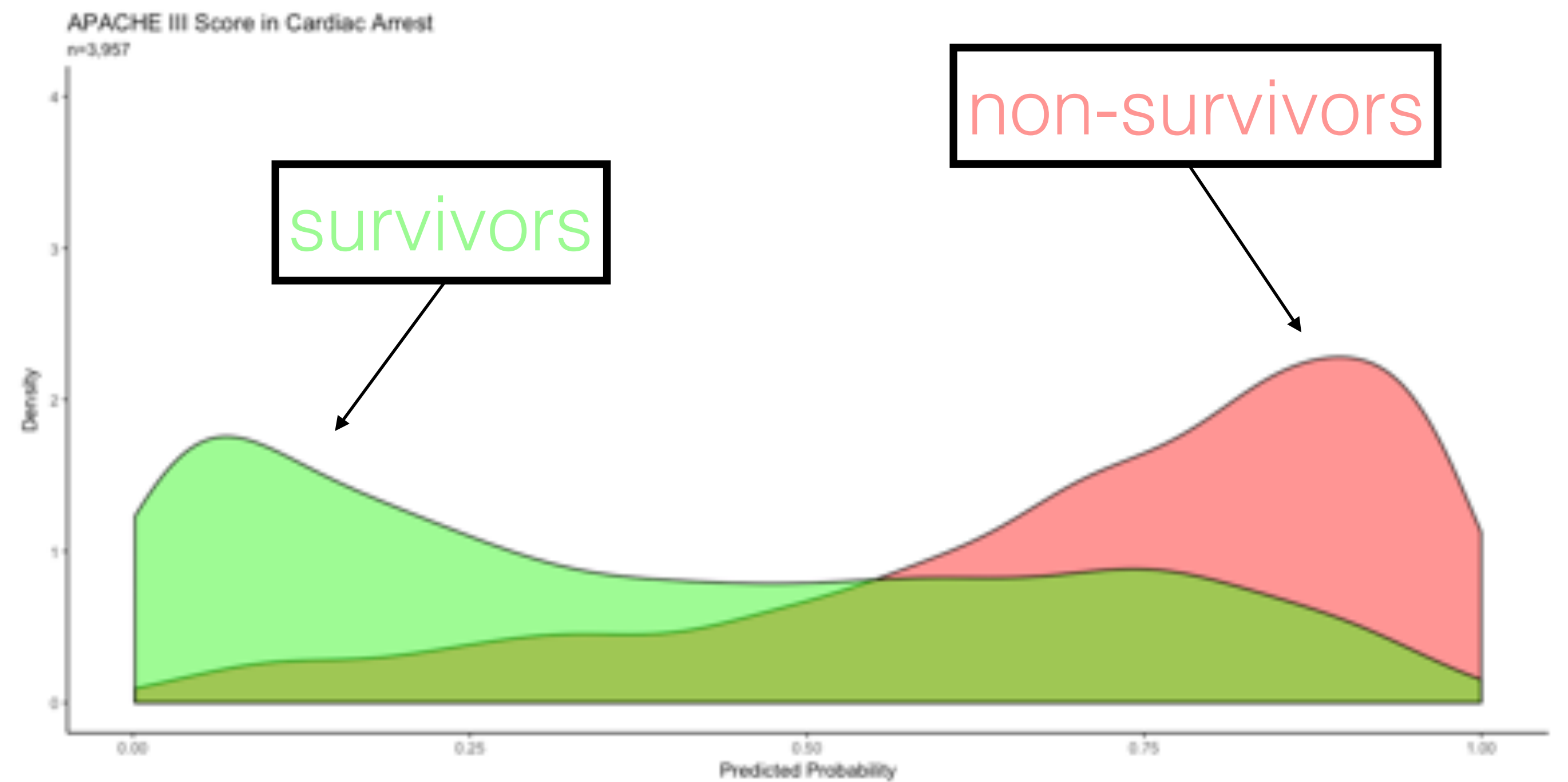
Comparison of model accuracy

Existing scores, logistic regression, and machine learning models.

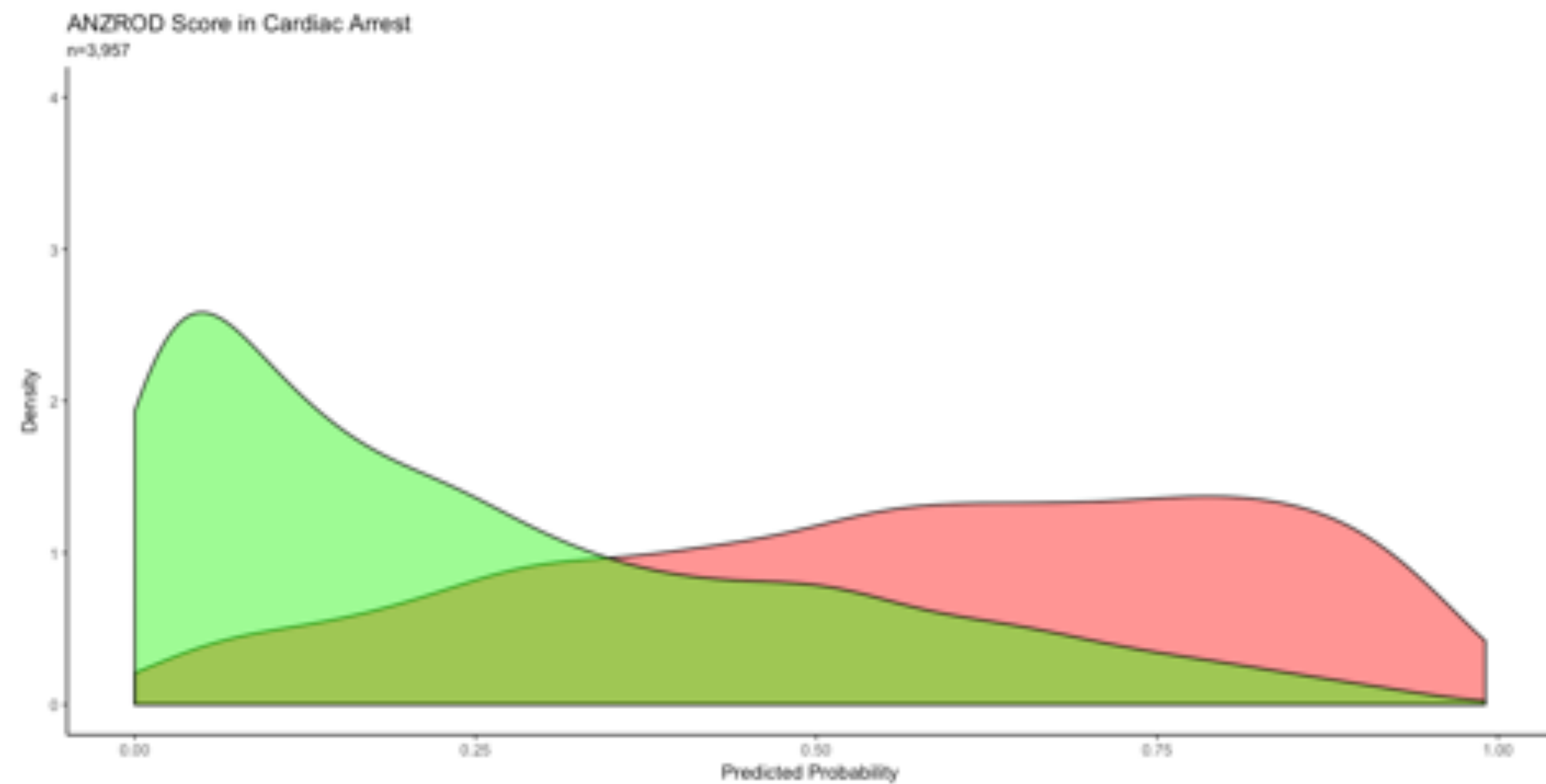
Model	Prediction	AUC (95% CI)	Brier Score	Log Loss
Actual mortality	45.5%			
APACHE III risk of death	52.8%	0.80 (0.79-0.82)	0.190	0.57
ANZROD	39.9%	0.81 (0.80-0.82)	0.182	0.55
Logistic Regression	45.4%	0.82 (0.81-0.83)	0.170	0.51
Artificial Neural Network	46.7%	0.85 (0.84-0.86)	0.158	0.48
Random Forest	45.7%	0.86 (0.84-0.87)	0.156	0.47
Support vector classifier	45.4%	0.86 (0.85-0.87)	0.153	0.47
Ensemble	45.5%	0.87 (0.86-0.88)	0.148	0.45
Gradient boosted machine	45.3%	0.87 (0.86-0.88)	0.147	0.45



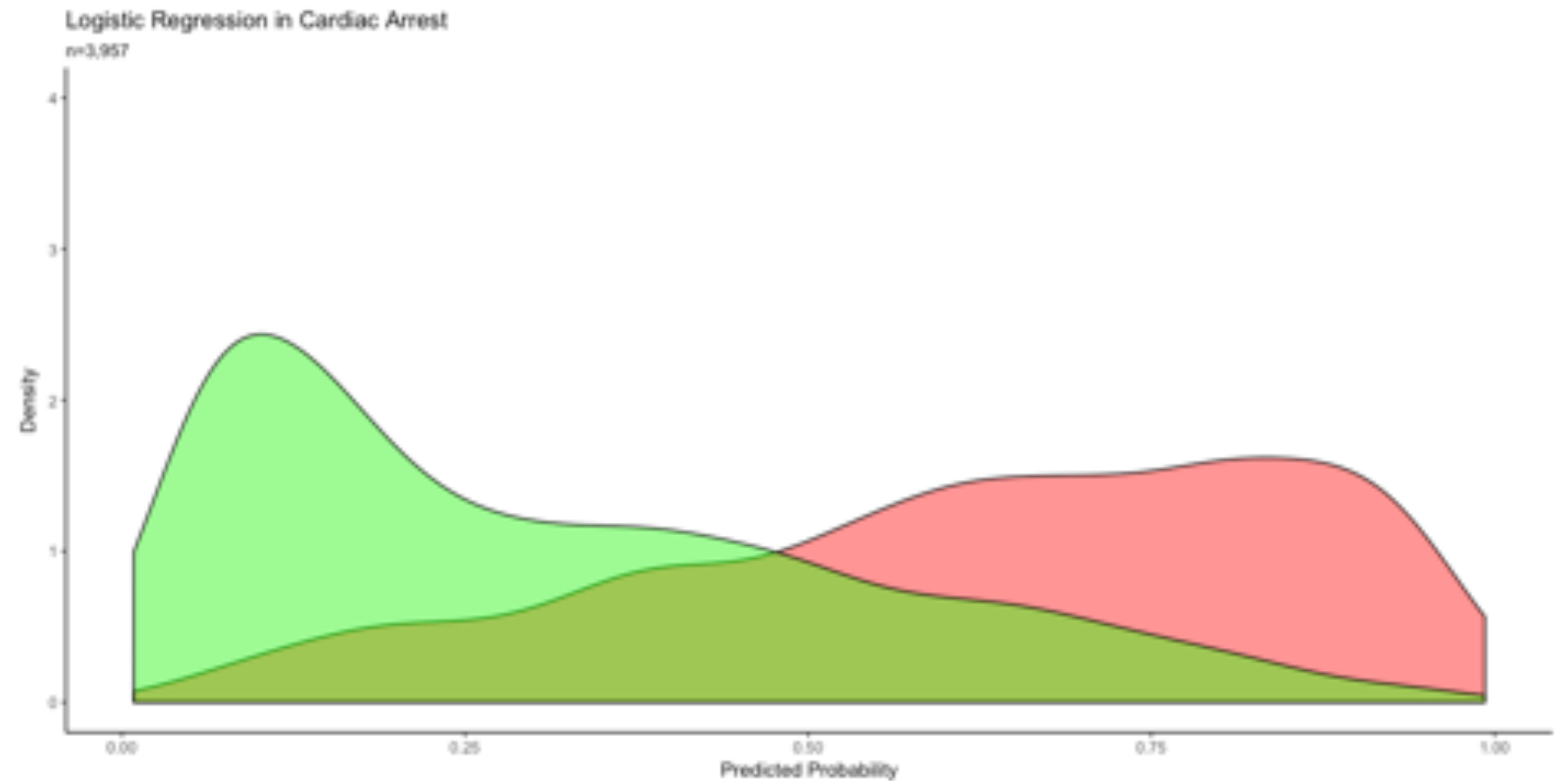
APACHE III



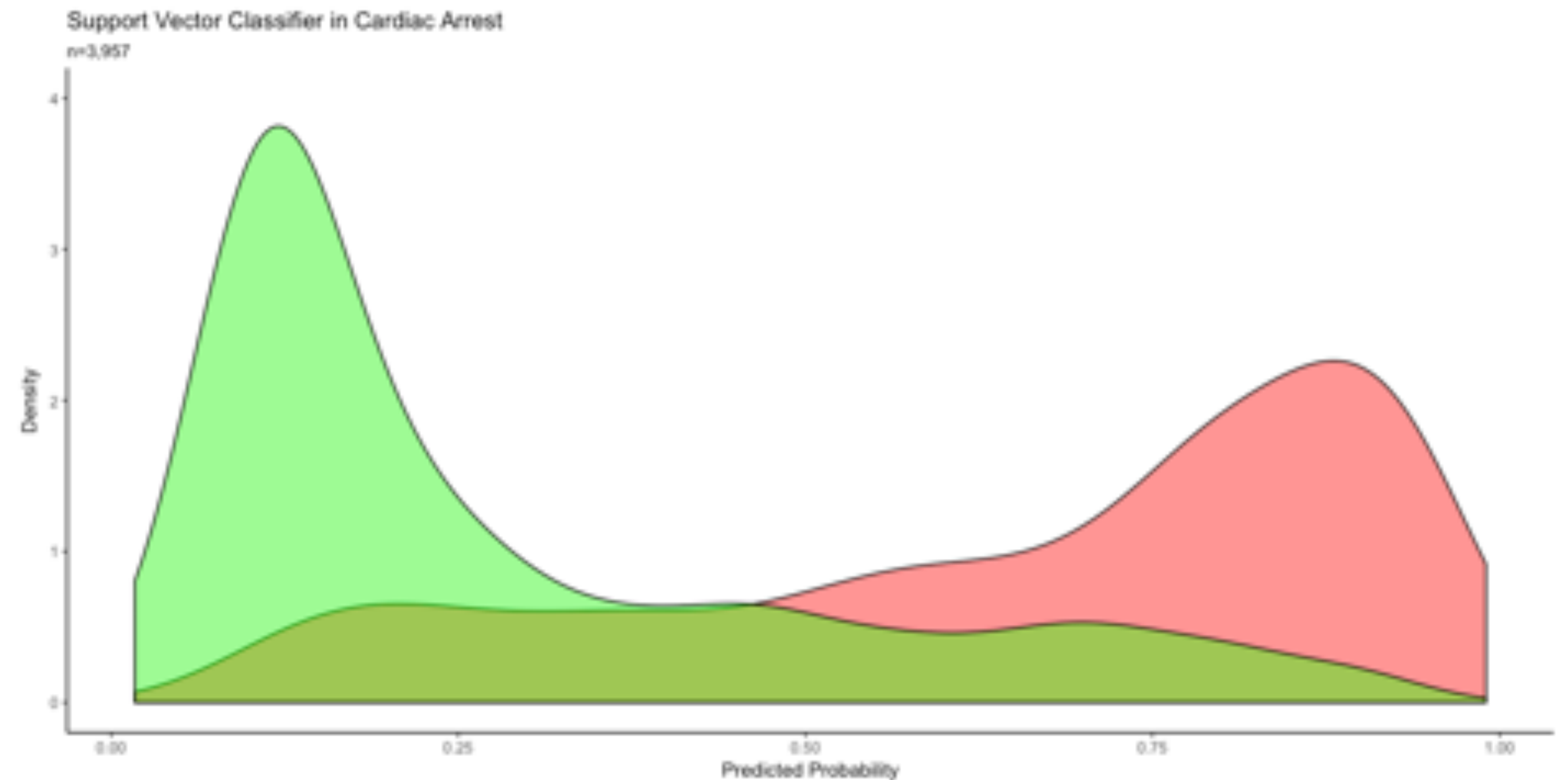
ANZROD



Logistic Regression

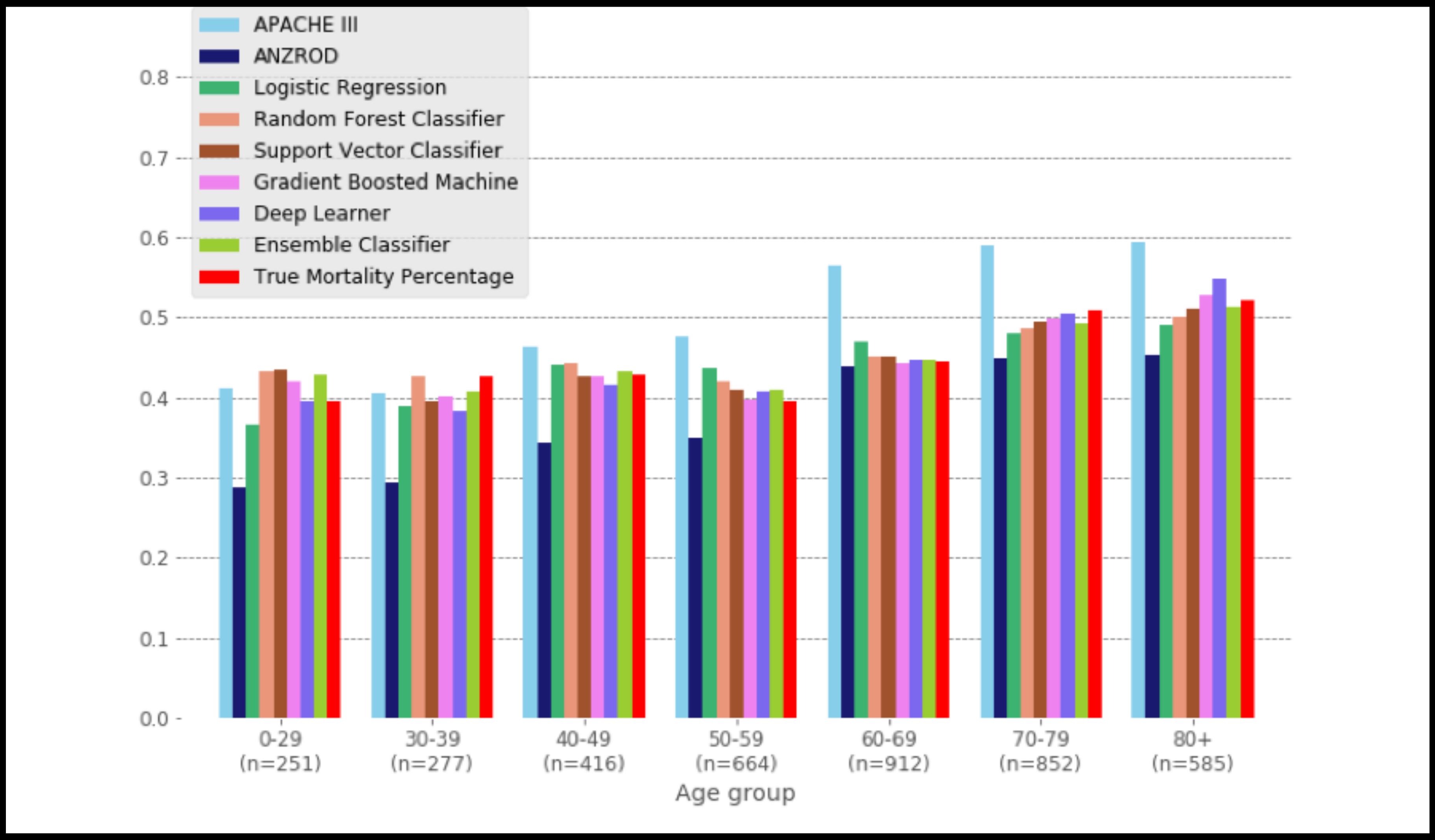


Support Vector Machine



Predicted mortality

Comparing age groups.

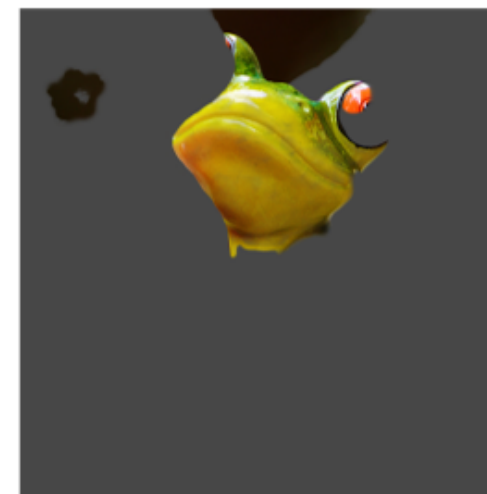


LIME¹ is a sensitivity analysis that reveals which parts of an input matter most to the eventual output.



Turning off all but a few interpretable components of this image reveals the probability that the model will identify ...

... a tree frog
54%



... billiard balls
7%



... a balloon
5%



Attention shines a spotlight on where the model is looking when it makes a particular decision.

Words relevant to **food quality** ...

The fajita we tried was **tasteless** and **burned** and the **mole sauce** was **way too sweet**.

... or to **service**

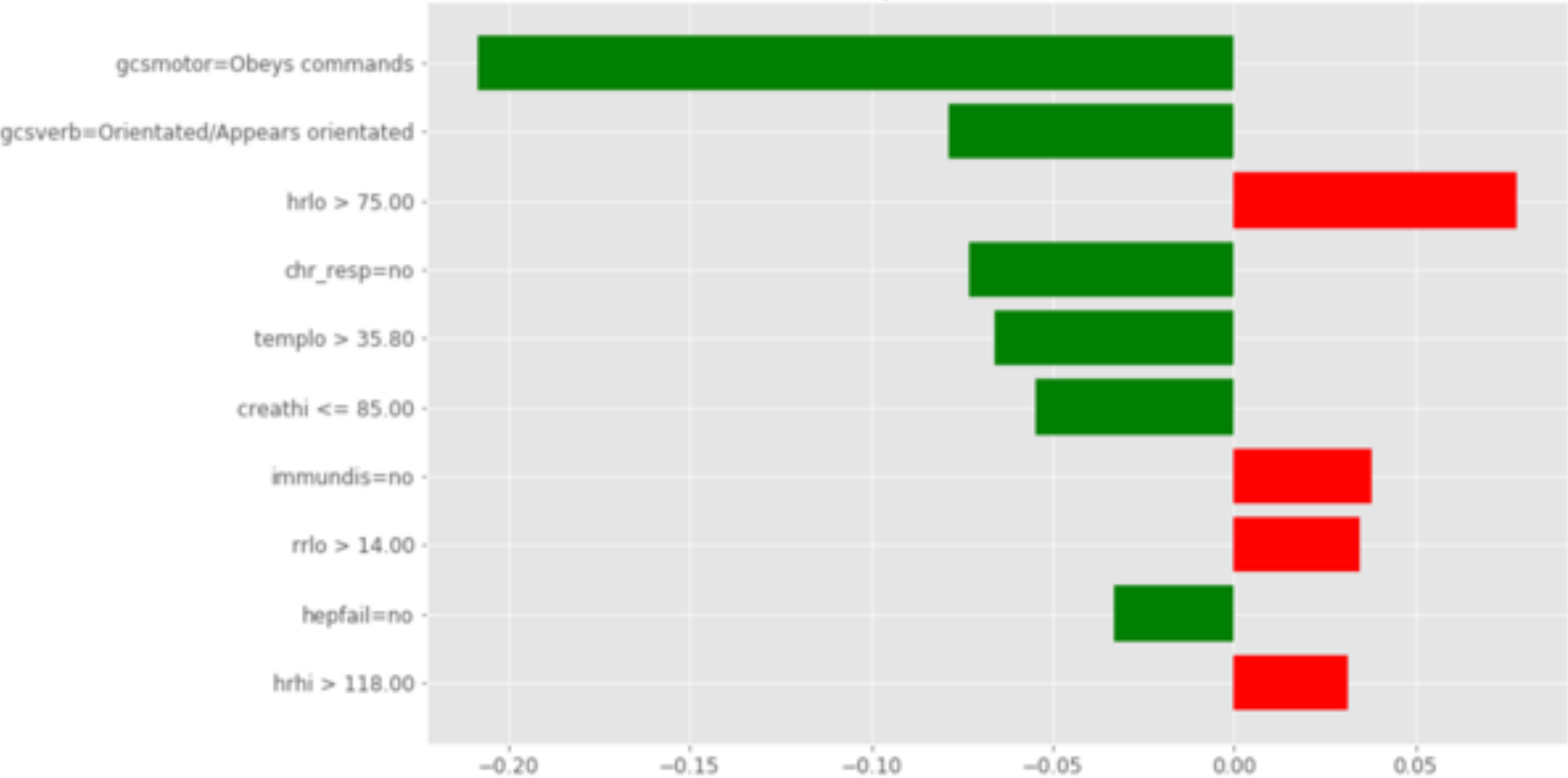
They have one of the **fastest delivery times** in the **city**.

¹LIME = local-interpretable-model-agnostic explanations.

Source: Carlos Guestrin, Marco Tulio Ribeiro, and Sameer Singh, "Introduction to local interpretable model-agnostic explanations (LIME)," August 12, 2016, O'Reilly, oreilly.com; Minlie Huang, Yequan Wang, Li Zhao, and Xiaoyan Zhu, *Attention-based LSTM for aspect-level sentiment classification*, Tsinghua University; Pixabay

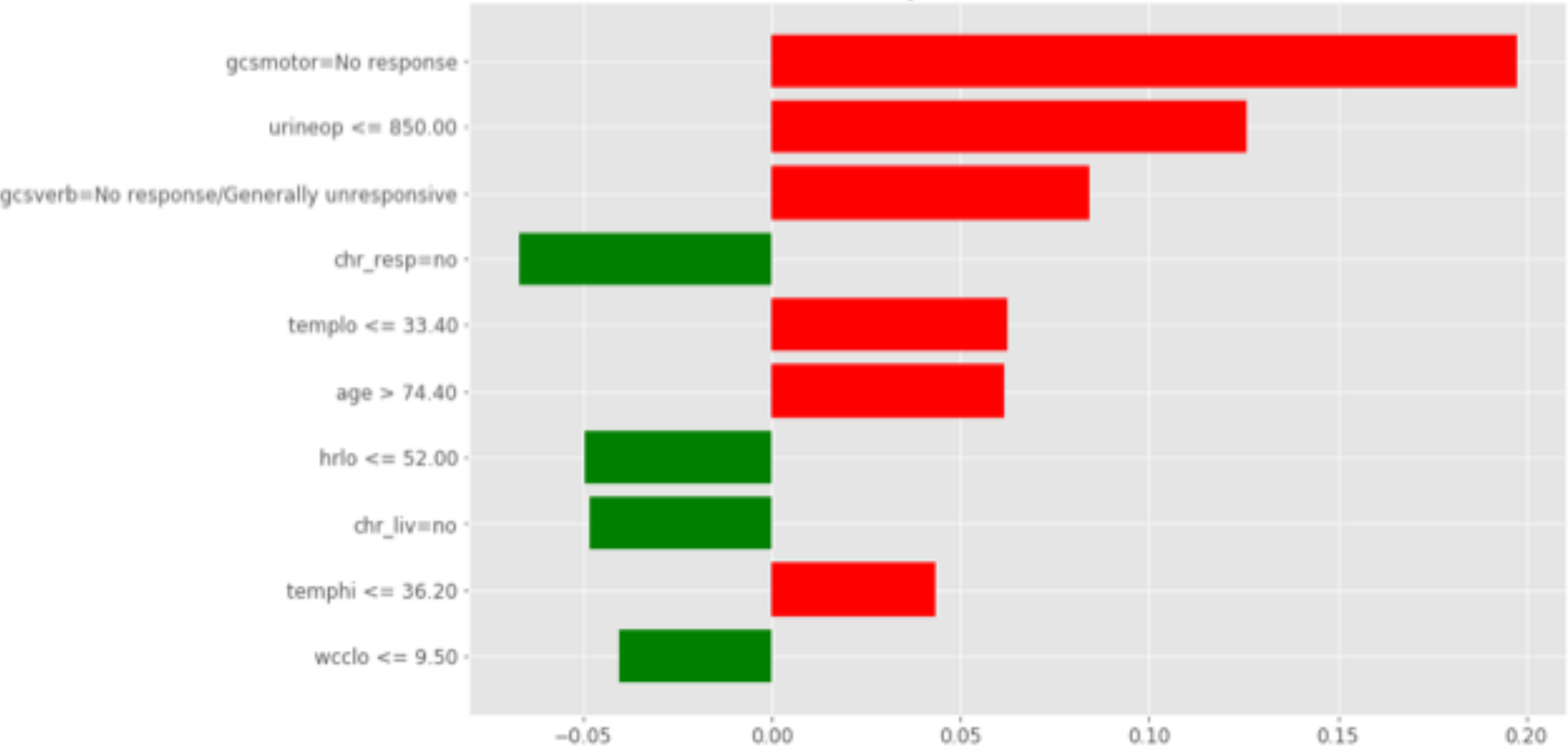
Explaining the model

High probability of survival (83%).



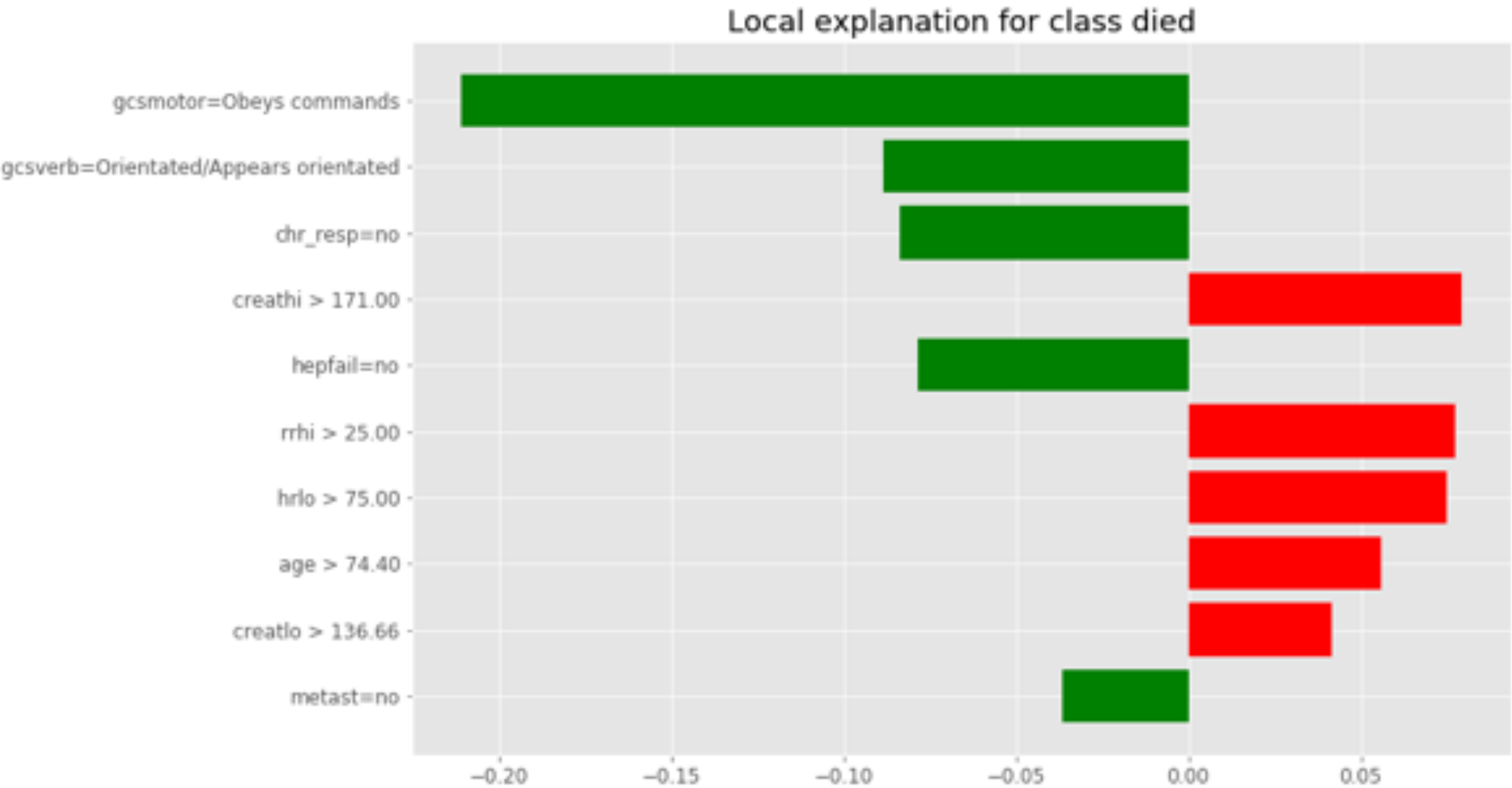
Explaining the model

Low probability of survival (27%).



Explaining the model

Incorrect prediction; 78% probability of survival, however the patient did not survive.



Conclusions.

Predicting, explaining, and future directions.

able to advance on existing critical care models and statistical methods

able to add explainability