



Machine Learning for Healthcare

6.7930, HST.956

Lecture 5:

Feb 23, 2023

Peter Szolovits

(with many slides from David Sontag)



Massachusetts
Institute of
Technology

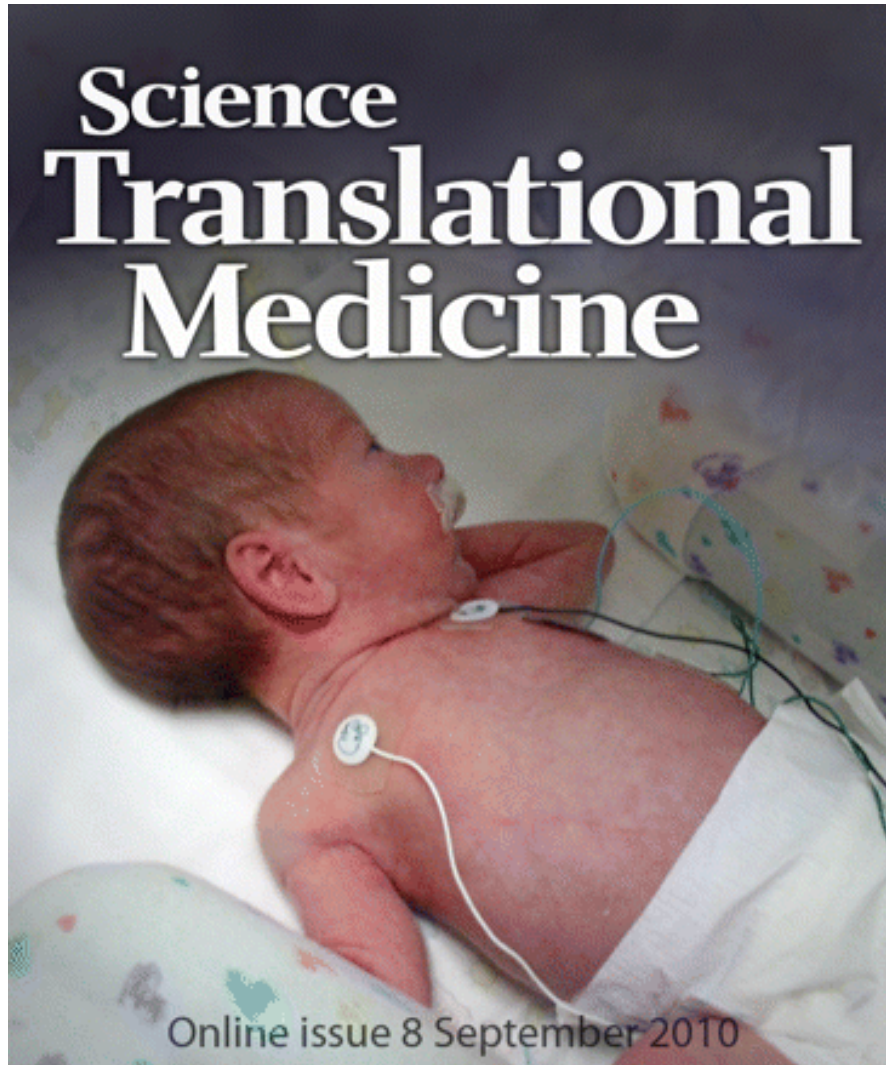
Outline for today's class

- 1. Introduction to risk stratification**
2. Case study: Early detection of Type 2 diabetes
 - Encoding longitudinal structured health data
3. Framing as supervised learning problem
 - Deriving labels from EHR

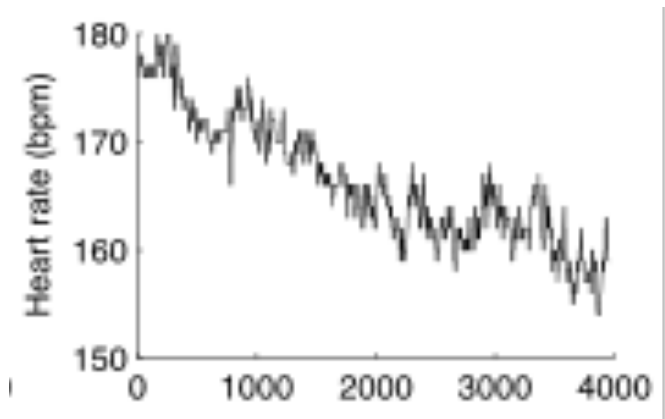
What *is* risk stratification?

- Separate a patient population into **high-risk** and **low-risk** of having an outcome
 - Predicting something in the future
- Coupled with **interventions** that target high-risk patients
- Goal is typically to reduce cost and improve patient outcomes

Examples of risk stratification



Preterm infant's risk of severe morbidity?



(Saria et al., Science Translational Medicine 2010)

Examples of risk stratification



Does this patient need to be admitted to the coronary-care unit?

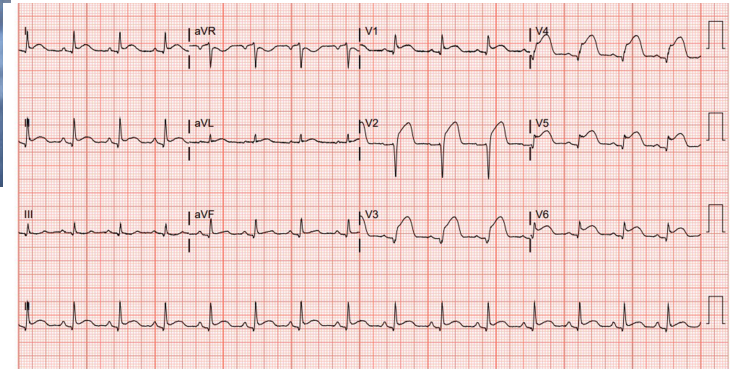


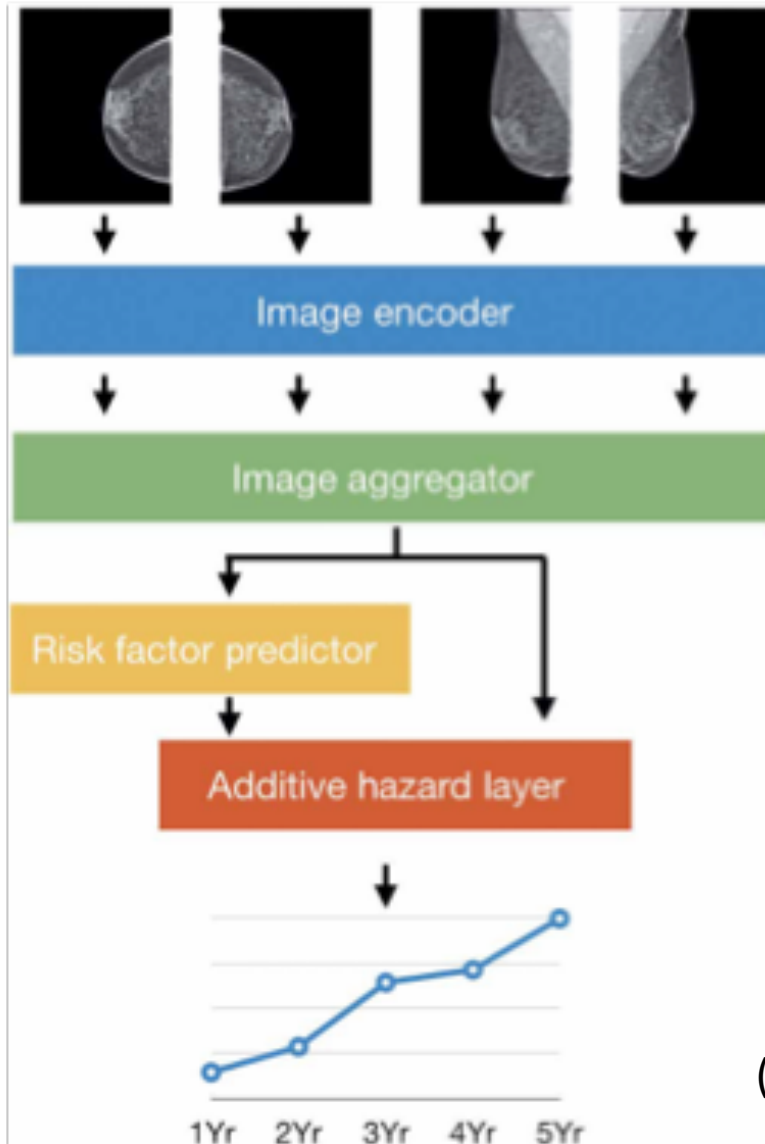
Figure sources:

<https://www.drmani.com/heart-attack/> (top)

<https://www.emra.org/emresident/article/acute-mi-case-report/> (right)

(Pozen et al., NEJM 1984)

Examples of risk stratification



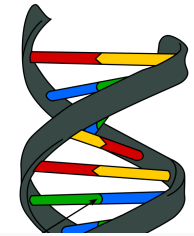
Will this woman develop breast cancer in the next 5 years?

(Yala et al., Science Translational Medicine 2021)

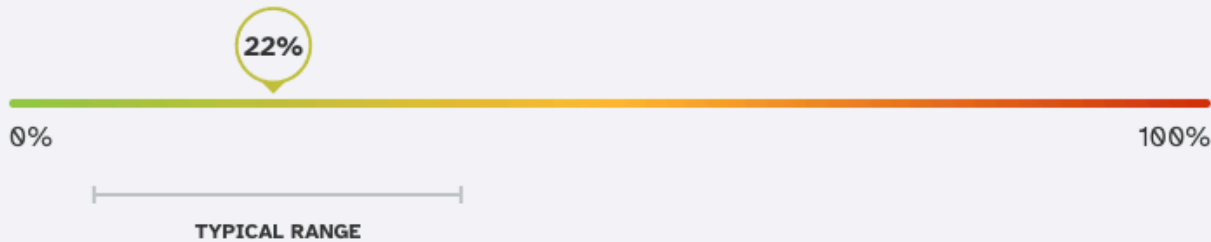
Examples of risk stratification



David, your genetics are associated with a **typical likelihood** of developing type 2 diabetes.



Based on data from 23andMe research participants, people of European descent with genetics like yours have an estimated **22% chance** of developing type 2 diabetes at some point **between the ages of 37 (your current age) and 80.**



ETHNICITY	AUC VALUE
European	0.652
South Asian	0.603
Hispanic/Latino	0.638
East Asian	0.609
African	0.588

DNA
Deoxyribonucleic acid

Summary

This report is based on a statistical model that estimates the likelihood of developing type 2 diabetes by looking at genetic variants at 1,244 places in your DNA. We identified these variants and created this model using data from more than 1,110,000 23andMe research participants of European descent.

How does risk stratification differ from differential diagnosis?

Differential diagnosis	Risk stratification
Usually iterative/active	Usually passive
Often considers a large set of conditions	Often just one condition
Has to consider rare conditions (needs hybrid knowledge/ML approaches)	Often focuses on settings where there is enough training data

Old vs. New

- Traditionally, risk stratification was based on simple scores using human-entered data

APGAR SCORING SYSTEM

	0 Points	1 Point	2 Points	Points totaled
Activity (muscle tone)	Absent	Arms and legs flexed	Active movement	↓
Pulse	Absent	Below 100 bpm	Over 100 bpm	
Grimace (reflex irritability)	Flaccid	Some flexion of Extremities	Active motion (sneeze, cough, pull away)	
Appearance (skin color)	Blue, pale	Body pink, Extremities blue	Completely pink	
Respiration	Absent	Slow, irregular	Vigorous cry	

Severely depressed	0-3
Moderately depressed	4-6
Excellent condition	7-10

Old vs. New

- Traditionally, risk stratification was based on simple scores using human-entered data
- Now, based on machine learning on high-dimensional data
 - Fits more easily into workflow
 - Higher accuracy
 - Quicker to derive (can special case)
- **But, ML approach comes with new challenges**
 - **to be discussed**

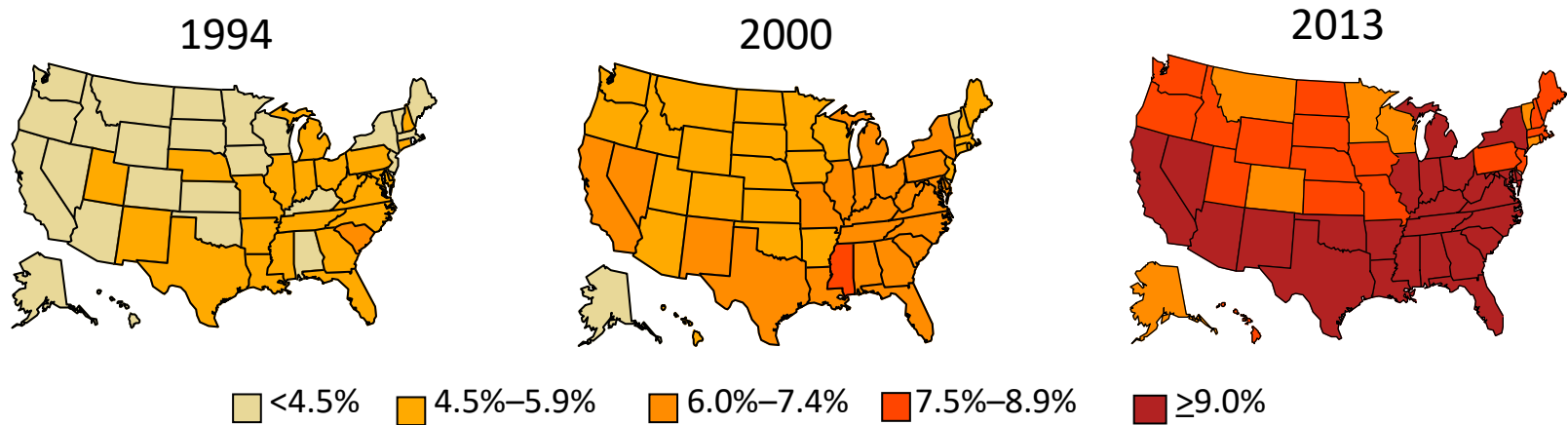
So, what do we need?

- Specification of prediction time / index date
- A way of encoding the data we have on the patient
 - CNN for images
 - Bag of words for text document
 - Longitudinal structured data ...
- A target, typically derived from the EHR
- Choice of appropriate supervised ML algorithm
 - Regression? Classification?

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Type 2 Diabetes: A Major public health challenge



\$245 billion: Total costs of diagnosed diabetes in the United States in 2012

\$831 billion: Total fiscal year federal budget for healthcare in the United States in 2014

- CDC 2022 estimate:
 - 11.3% of adults: 28.7M diagnosed, 8.5M undiagnosed
- Racial disparities among adults (20+)
 - non-Hisp White: 7.5%
 - non-Hisp Asian: 9.2%
 - non-Hisp Black: 11.7%
 - Hispanic: 12.5%
 - Native American: 14.7%


Type 2 Diabetes Can Be Prevented *

Requirement for successful large scale prevention program:

1. Detect/reach truly at risk population
2. Improve the interventions
3. Lower the cost of intervention

Traditional Risk Prediction Models

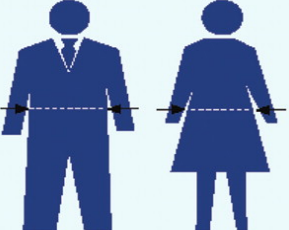
- Successful Examples
 - ARIC
 - KORA
 - FRAMINGHAM
 - AUSDRISC
 - FINDRISC
 - San Antonio Model
- Easy to ask/measure in the office, or for patients to do online
- Simple model: can calculate scores by hand

 Finnish Diabetes Association

TYPE 2 DIABETES RISK ASSESSMENT FORM

Circle the right alternative and add up your points.

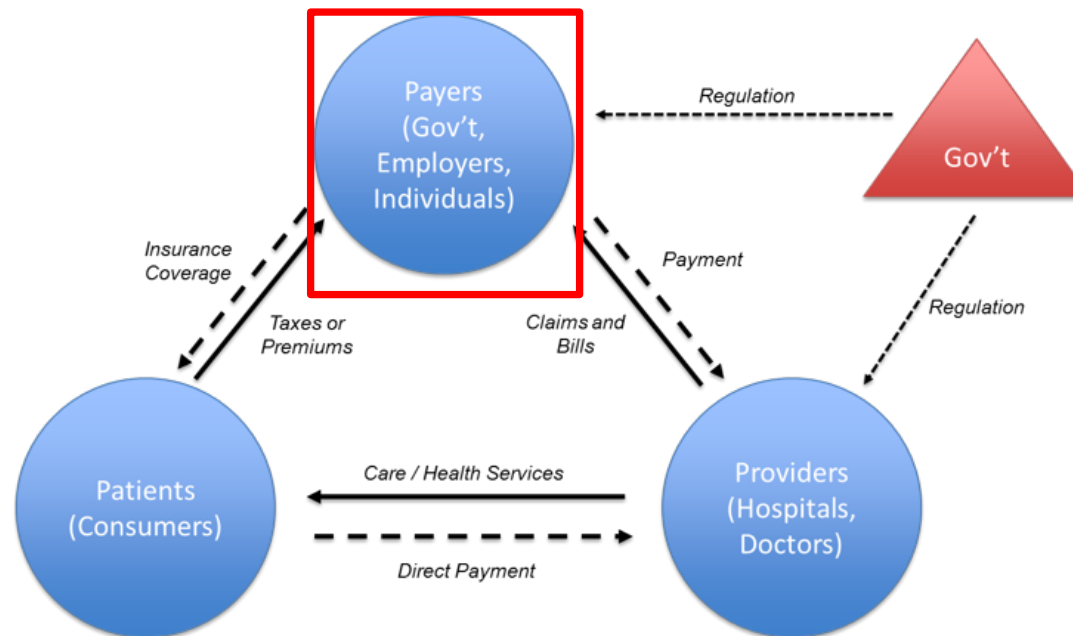
<p>1. Age</p> <p>0 p. Under 45 years 2 p. 45–54 years 3 p. 55–64 years 4 p. Over 64 years</p> <p>2. Body-mass index (See reverse of form)</p> <p>0 p. Lower than 25kg/m² 1 p. 25–30 kg/m² 3 p. Higher than 30 kg/m²</p> <p>3. Waist circumference measured below the ribs (usually at the level of the navel)</p> <table border="0" style="width: 100%;"> <tr> <td style="text-align: center;">MEN</td> <td style="text-align: center;">WOMEN</td> </tr> <tr> <td>0 p. Less than 94cm</td> <td>Less than 80cm</td> </tr> <tr> <td>3 p. 94–102cm</td> <td>80–88cm</td> </tr> <tr> <td>4 p. More than 102cm</td> <td>More than 88cm</td> </tr> </table>	MEN	WOMEN	0 p. Less than 94cm	Less than 80cm	3 p. 94–102cm	80–88cm	4 p. More than 102cm	More than 88cm	<p>6. Have you ever taken anti-hypertensive medication regularly?</p> <p>0 p. No 2 p. Yes</p> <p>7. Have you ever been found to have high blood glucose (e.g. in a health examination, during an illness, during pregnancy)?</p> <p>0 p. No 5 p. Yes</p> <p>8. Have any of the members of your immediate family or other relatives been diagnosed with diabetes (type 1 or type 2)?</p> <p>0 p. No 3 p. Yes: grandparent, aunt, uncle or first cousin (but no own parent, brother, sister or child) 5 p. Yes: parent, brother, sister or own child</p>
MEN	WOMEN								
0 p. Less than 94cm	Less than 80cm								
3 p. 94–102cm	80–88cm								
4 p. More than 102cm	More than 88cm								



<p>4. Do you usually have daily at least 30 minutes of physical activity at work and/or during leisure time (including normal daily activity)?</p> <p>0 p. Yes 2 p. No</p> <p>5. How often do you eat vegetables, fruit' or berries?</p> <p>0 p. Every day 1 p. Not every day</p>	<p>Total risk score</p> <p><input type="checkbox"/> The risk of developing type 2 diabetes within 10 years is</p> <p>Lower than 7 Low: estimated 1 in 100 will develop disease</p> <p>7–11 Slightly elevated: estimated 1 in 25 will develop disease</p> <p>12–14 Moderate: estimated 1 in 6 will develop disease</p> <p>15–20 High: estimated 1 in 3 will develop disease</p> <p>Higher than 20 Very high: estimated 1 in 2 will develop disease</p> <p style="text-align: right; font-size: small;">Please turn over</p>
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Population-Level Risk Stratification

- Key idea: Use readily available administrative, utilization, and clinical data

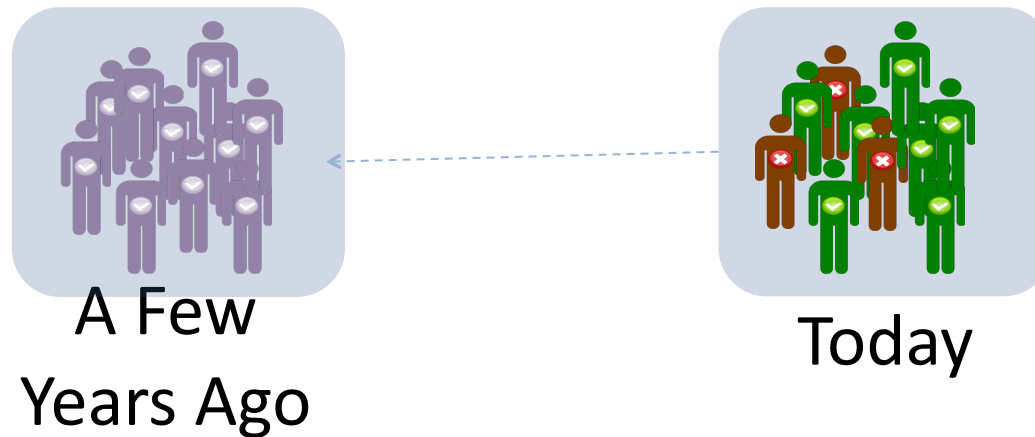


Population-Level Risk Stratification

- Key idea: Use readily available administrative, utilization, and clinical data
- Machine learning will find surrogates for risk factors that would otherwise be missing
- Perform risk stratification at the population level – millions of patients

A Data-Driven approach on Longitudinal Data-Based Prediction

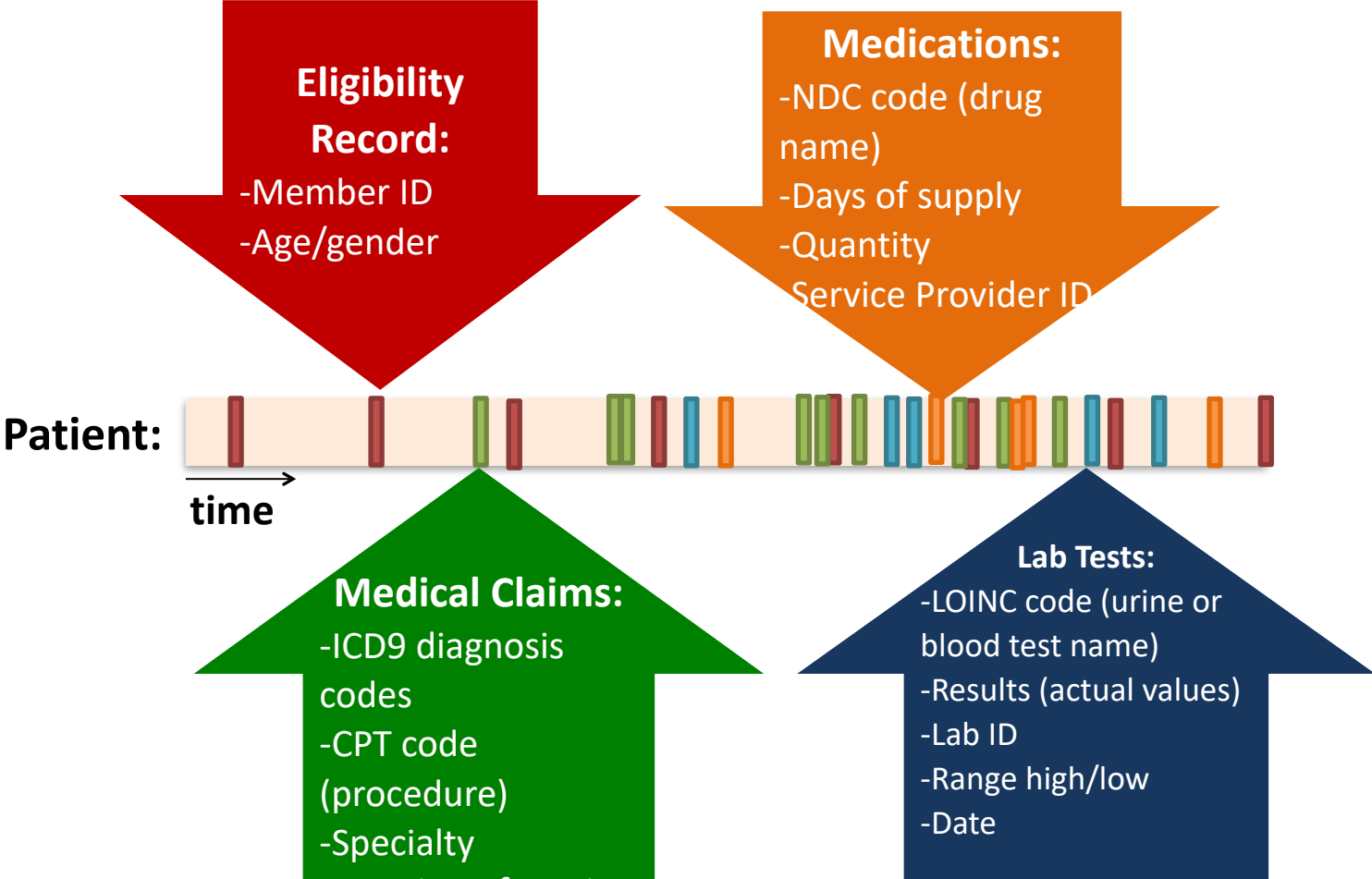
- Looking at individuals who got diabetes *today*, (compared to those who didn't)
 - Can we infer which variables in their record could have predicted their health outcome?

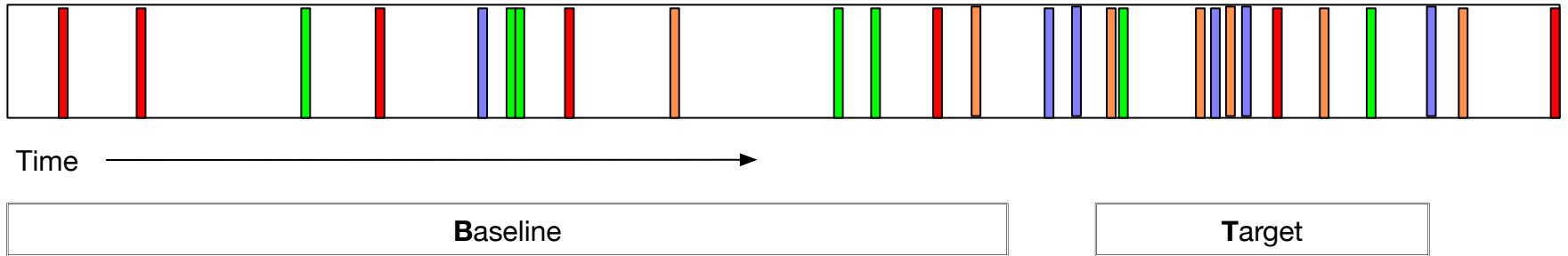


Risk Stratification from Structured Health Data

Reading: Razavian N, Blecker S, Schmidt AM, Smith- McLallen A, Nigam S, Sontag D (2015) [Population-level prediction of type 2 diabetes from claims data and analysis of risk factors](#). Big Data 3:4, 277–287, DOI: 10.1089/big.2015.0020.

Administrative & Clinical Data





$$\text{Target} = f(\text{Baseline})$$

- How to represent Baseline and Target?
- What class of models do we consider for f ?

Claims Data Characteristics

- Sparse data
 - Vast coding spaces for diagnoses, symptoms, procedures, medications, labs
 - Most patients don't have most of these
- Visit-level temporality
 - Data collected only at interactions with the health care system; highly variable intervals
- Long-term dependencies
 - How to encode these? LSTM, bi-RNN, CNN, attention, ...
 - simpler trends

Top diagnosis codes

Disease	count
401.1 Benign hypertension	447017
272.4 Hyperlipidemia NEC/NOS	382030
401.9 Hypertension NOS	372477
250.00 DMII wo cmp nt st uncntr	339522
272.0 Pure hypercholesterolem	232671
272.2 Mixed hyperlipidemia	180015
V72.31 Routine gyn examination	178709
244.9 Hypothyroidism NOS	169829
780.79 Malaise and fatigue NEC	149797
V04.81 Vaccin for influenza	147858
724.2 Lumbago	137345
V76.12 Screen mammogram NEC	129445
V70.0 Routine medical exam	127848

Disease	count
530.81 Esophageal reflux	121064
427.31 Atrial fibrillation	113798
729.5 Pain in limb	112449
414.01 Crnry athrsl natve vssl	104478
285.9 Anemia NOS	103351
786.50 Chest pain NOS	91999
599.0 Urin tract infection NOS	87982
V58.69 Long-term use meds NEC	85544
496 Chr airway obstruct NEC	78585
477.9 Allergic rhinitis NOS	77963
414.00 Cor ath unsp vsl ntv/gft	75519

Disease	count
719.47 Joint pain-ankle	28648
300.4 Dysthymic disorder	28530
268.9 Vitamin D deficiency NOS	28455
V72.81 Preop cardiovscrl exam	27897
724.3 Sciatica	27604
787.91 Diarrhea	27424
V2.21 Supervis oth normal preg	27320
365.01 Opn angl brderln lo risk	26033
379.21 Vitreous degeneration	25592
424.1 Aortic valve disorder	25425
616.10 Vaginitis NOS	24736
702.19 Other sborheic keratosis	24453
380.4 Impacted cerumen	24046

Out of 135K patients who had laboratory data

Top lab test results

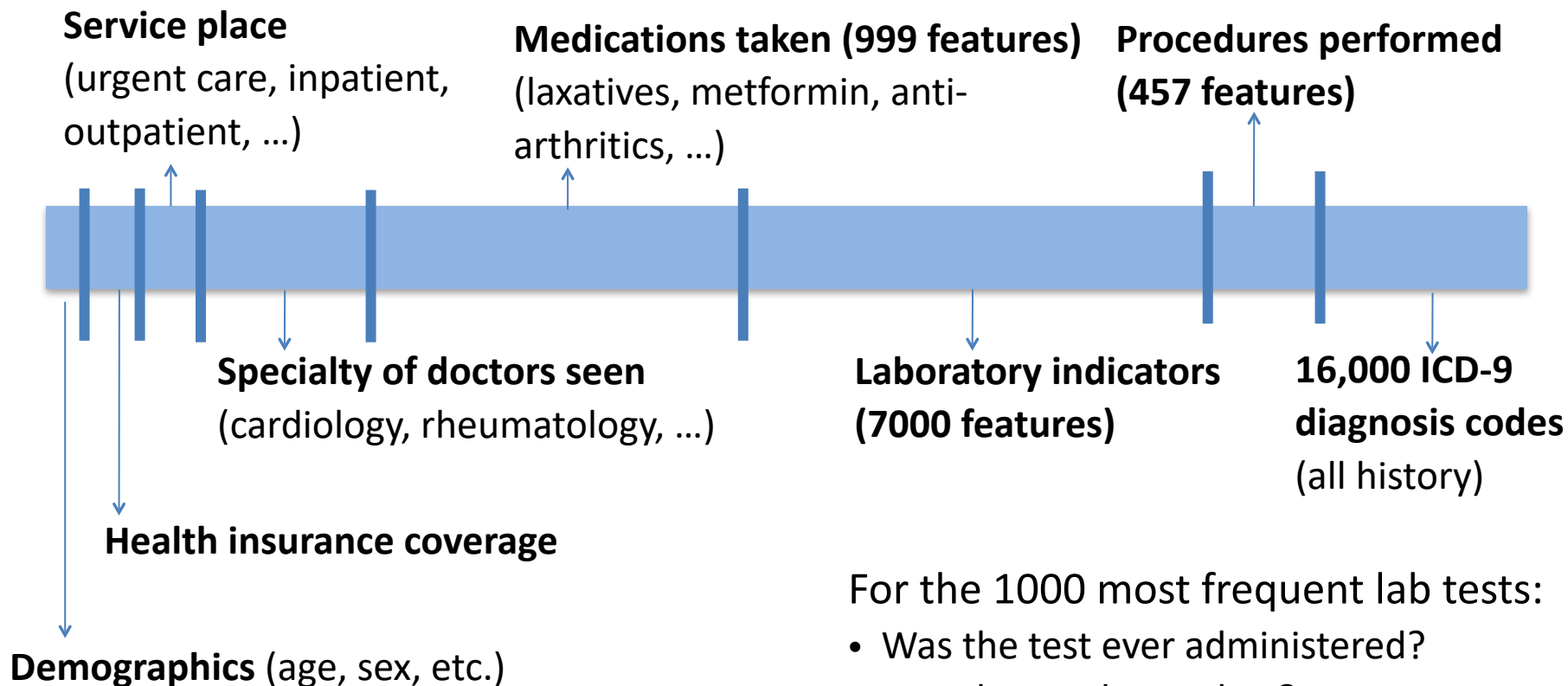
Lab test	
2160-0 Creatinine	1284737
3094-0 Urea nitrogen	1282344
2823-3 Potassium	1280812
2345-7 Glucose	1299897
1742-6 Alanine aminotransferase	1187809
1920-8 Aspartate aminotransferase	1187965
2885-2 Protein	1277338
1751-7 Albumin	1274166
2093-3 Cholesterol	1268269
2571-8 Triglyceride	1257751
13457-7 Cholesterol.in LDL	1241208
17861-6 Calcium	1165370
2951-2 Sodium	1167675

Lab test	
2085-9 Cholesterol.in HDL	1155666
718-7 Hemoglobin	1152726
4544-3 Hematocrit	1147893
9830-1 Cholesterol.total/Cholesterol.in HDL	1037730
33914-3 Glomerular filtration rate/1.73 sq M.predicted	561309
785-6 Erythrocyte mean corpuscular hemoglobin	1070832
6690-2 Leukocytes	1062980
789-8 Erythrocytes	1062445
787-2 Erythrocyte mean corpuscular volume	1063665

Lab test	
770-8 Neutrophils/100 leukocytes	952089
731-0 Lymphocytes	943918
704-7 Basophils	863448
711-2 Eosinophils	935710
5905-5 Monocytes/100 leukocytes	943764
706-2 Basophils/100 leukocytes	863435
751-8 Neutrophils	943232
742-7 Monocytes	942978
713-8 Eosinophils/100 leukocytes	933929
3016-3 Thyrotropin	891807
4548-4 Hemoglobin A1c/Hemoglobin.total	527062

Count of the test result (ever)

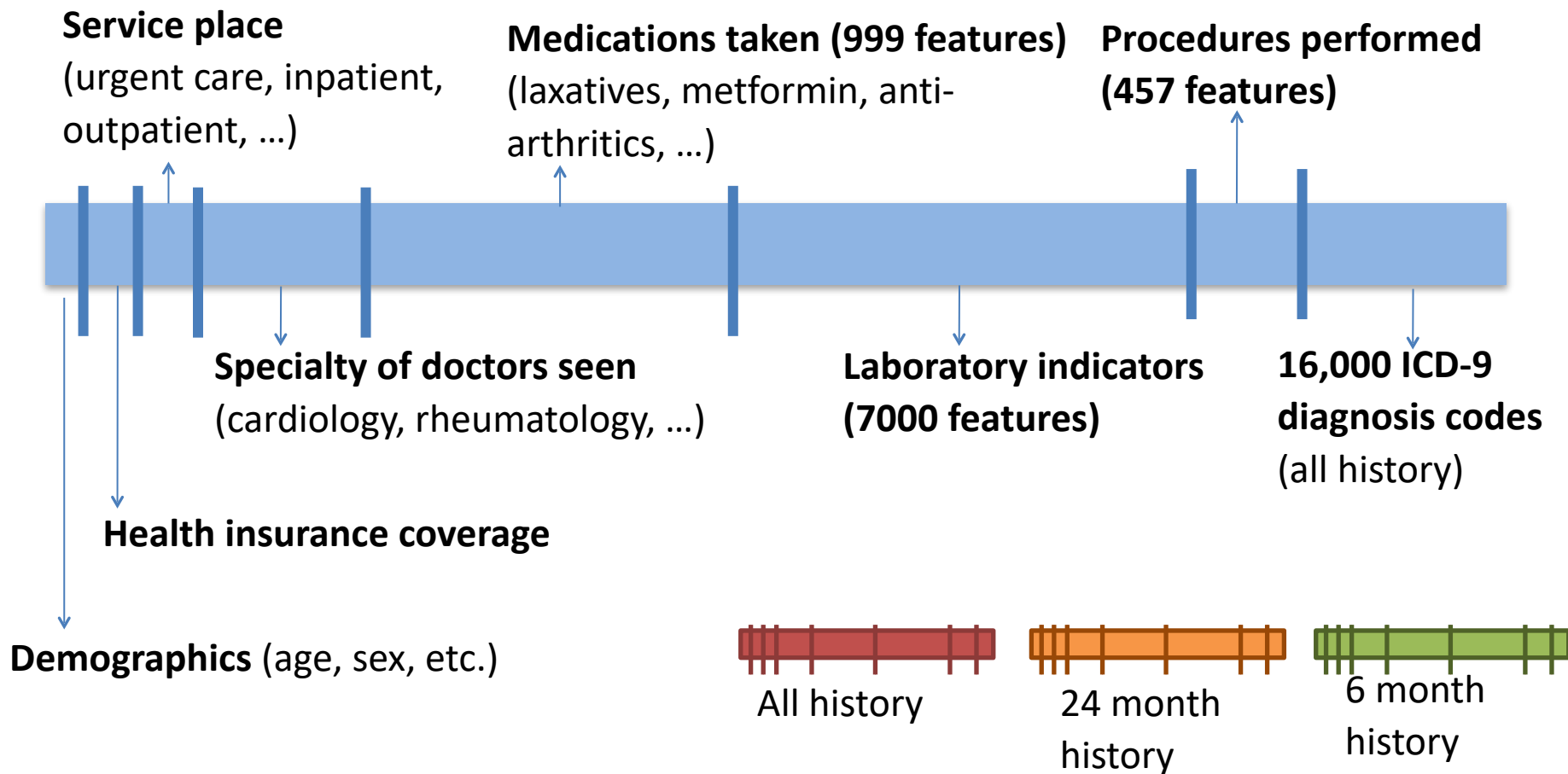
Encoding the longitudinal health data



For the 1000 most frequent lab tests:

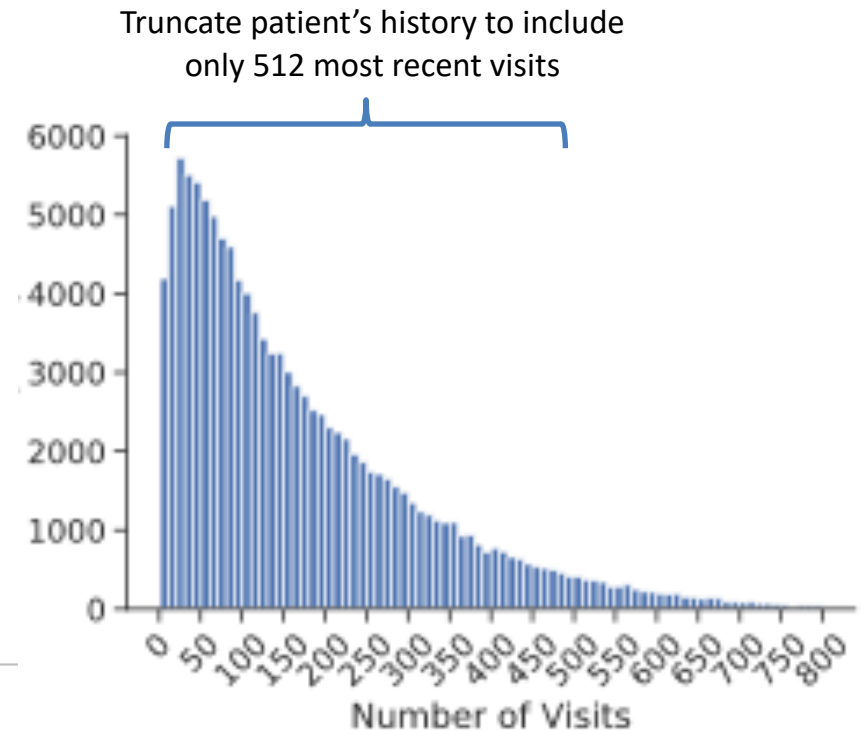
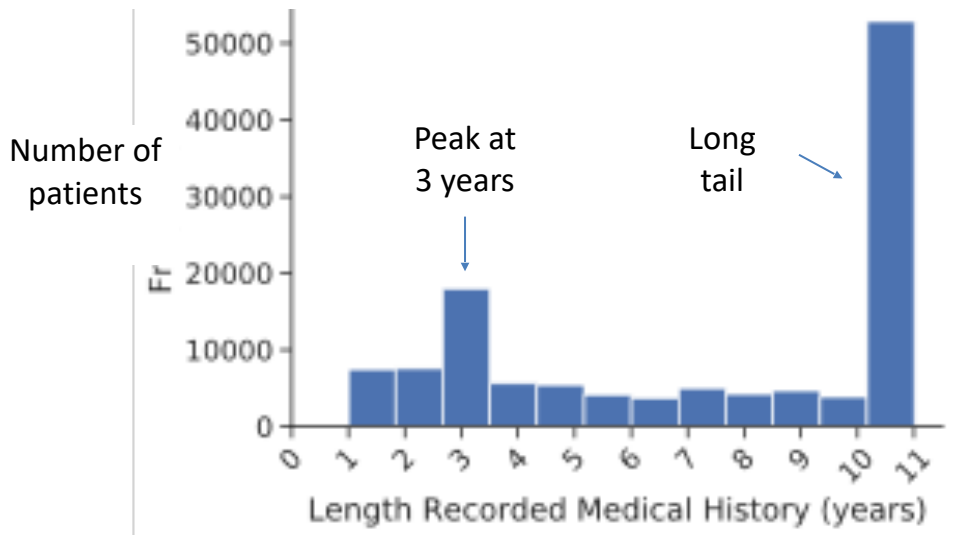
- Was the test ever administered?
- Was the result ever low?
- Was the result ever high?
- Was the result ever normal?
- Is the value increasing?
- Is the value decreasing?
- Is the value fluctuating?

Encoding the longitudinal health data

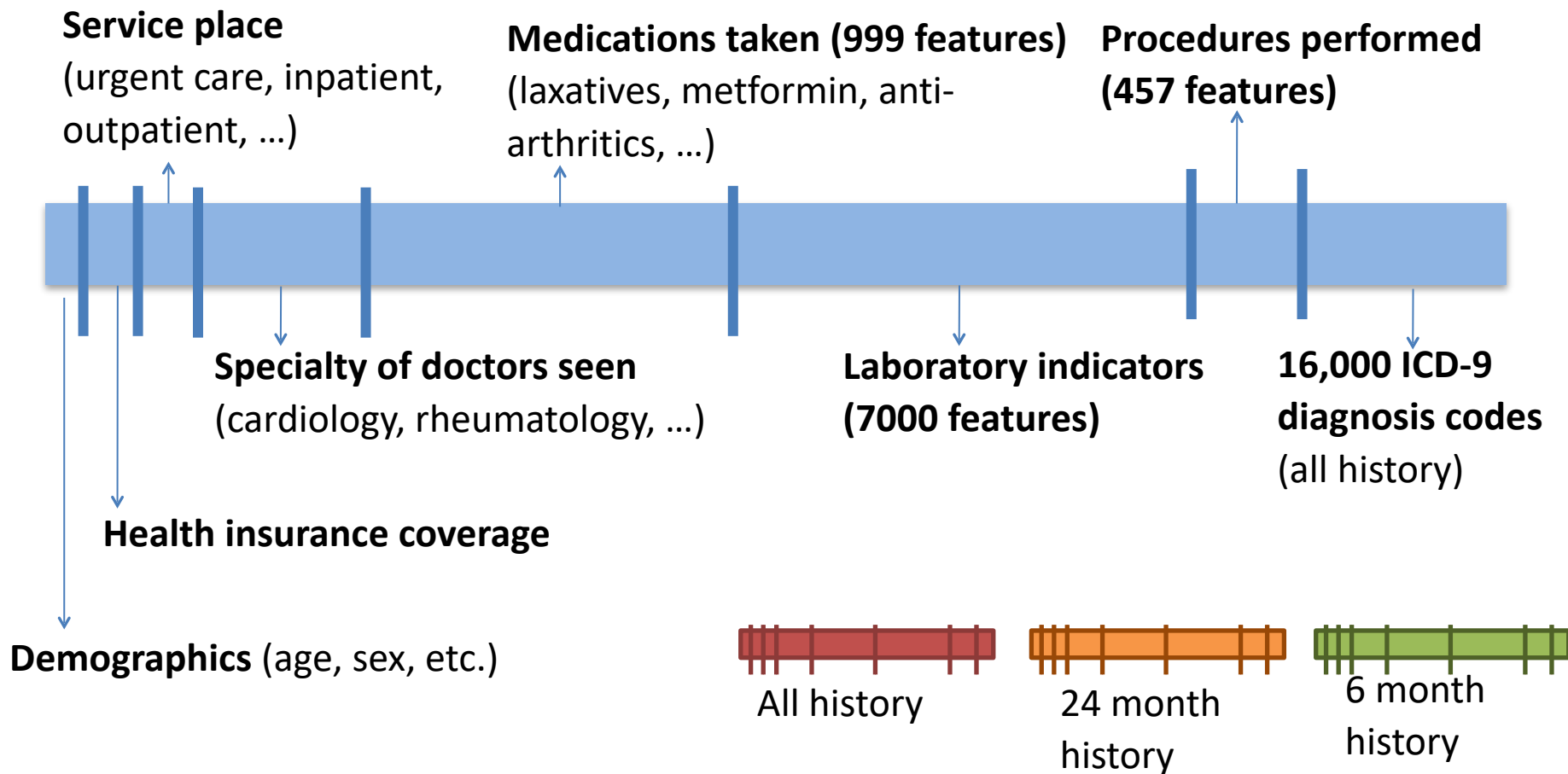


10s-100s of thousands of features

There may be a varying amount of history per patient



Encoding the longitudinal health data



How does this deal with missing data? What are its limitations?

Combining Multi-Modal Data

ML approach

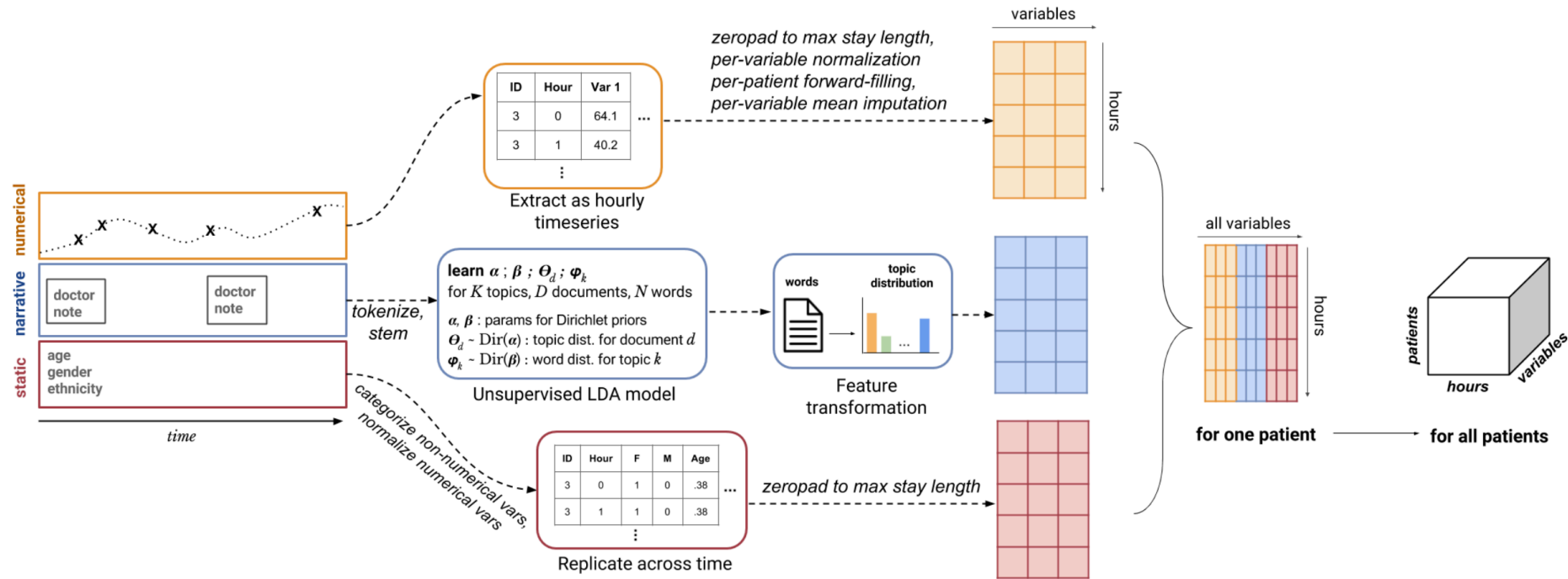
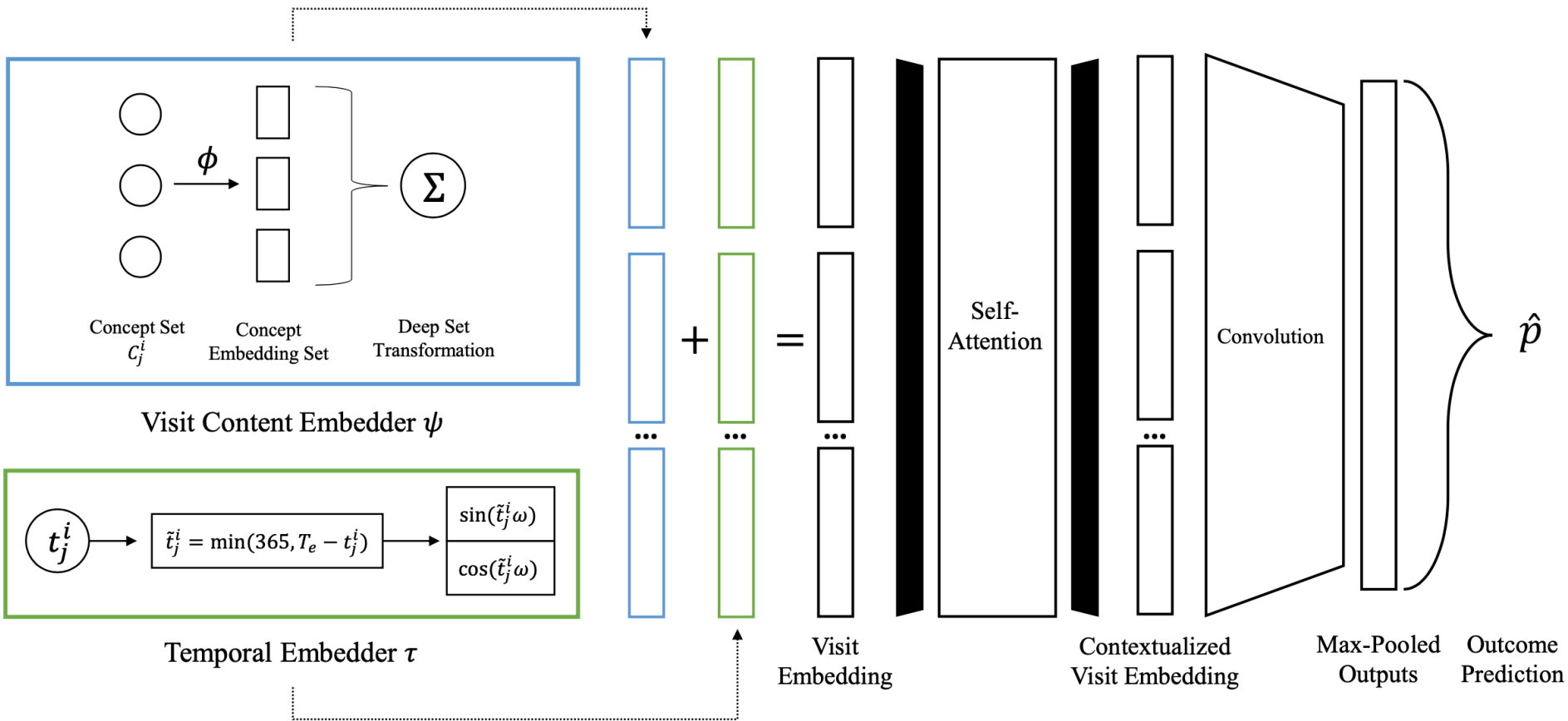
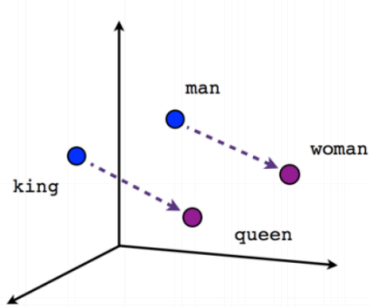


Figure 1: A visual representation of the data used. 1) Numerical data, including vitals and lab tests. The timestamp for each data point is rounded to the nearest hour, and hours with multiple measurements for a variable are assigned the average of those measurements. Each measurement is normalized according to the min and max for that var and each patient's data are zero-padded to the maximum stay length (240 hours). To fill in missing values, we forward-fill values for each patient, and mean-impute for any remaining missing values. 2) **Narrative data**, which consists of unstructured text notes. After preprocessing, LDA is used to obtain underlying topics and we then represent each note as a distribution over these topics. We forward-fill and aggregate these topic vectors across time, mean-imputing any values that are still missing. 3) **Static Data**, including variables recorded at admission such as sex, age, and ethnicity. Categorical variables such as ethnicity and ICU type are transformed into one-hot vectors containing each possible type. We replicate this data across time so that we are able to feed in this information at every timestep. We normalize numerical values and use forward-filling and imputation as before.

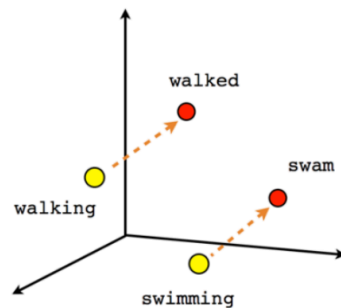
Alternative encoding using self-attention / transformers



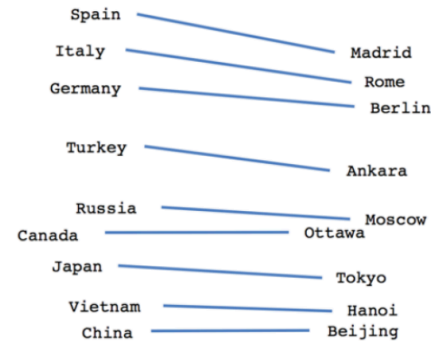
The latter can make use of unsupervised learning of concept embeddings



Male-Female



Verb tense



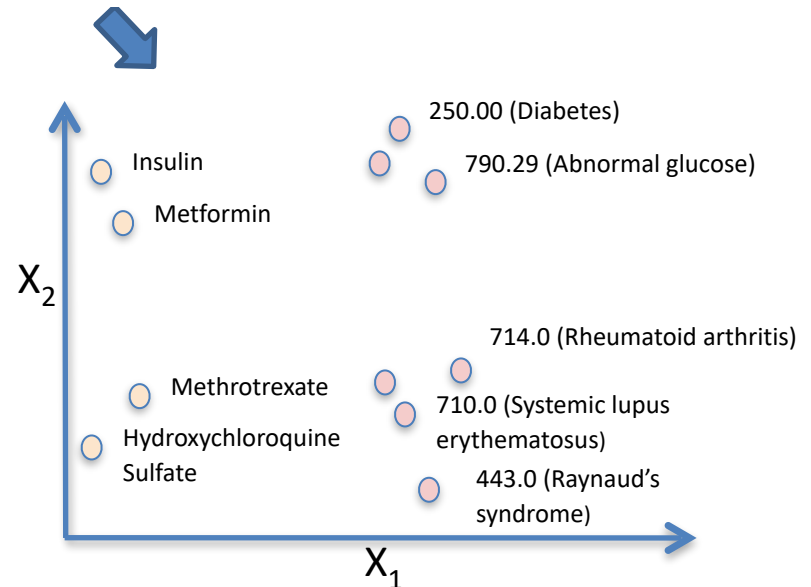
Country-Capital

Mikolov et al., Efficient Estimation of Word Representations in Vector Space, ICLR '13

Figure: <https://cbail.github.io/texasdata/word2vec/rmarkdown/word2vec.html>

Choi, Chiu, Sontag. Learning low-dimensional representations of medical concepts. AMIA Summits on Translational Science Proceedings, '16
<https://github.com/clinicalml/embeddings>

Beam et al., Clinical Concept Embeddings Learned from Massive Sources of Multimodal Medical Data. PSB '20



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Where do the labels come from?

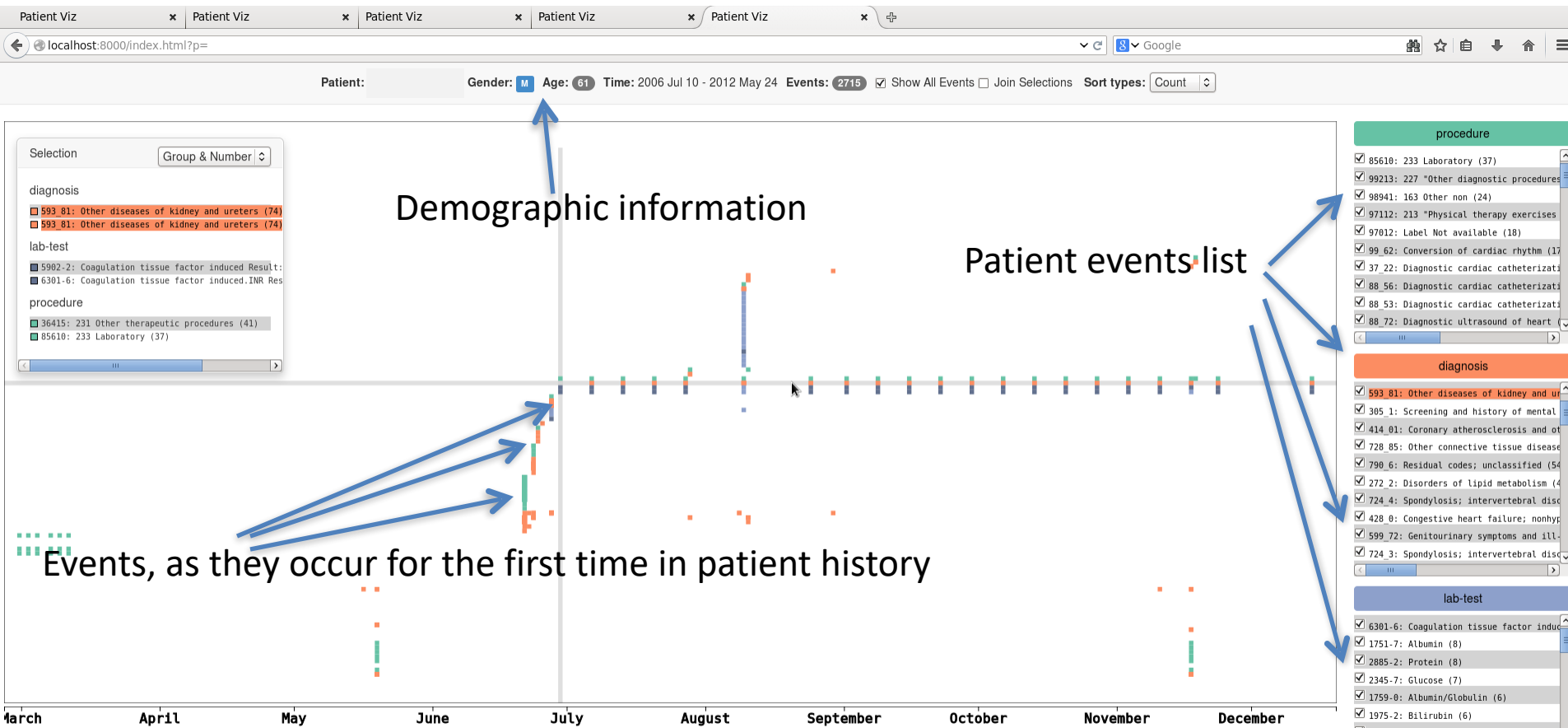
Typical pipeline:

1. Manually label several patients' data by "chart review"
2. A) Come up with a simple rule to automatically derive label for all patients, **or**

B) Use machine learning to get the labels themselves

Step 1:

Visualization of individual patient data is an important part of chart review

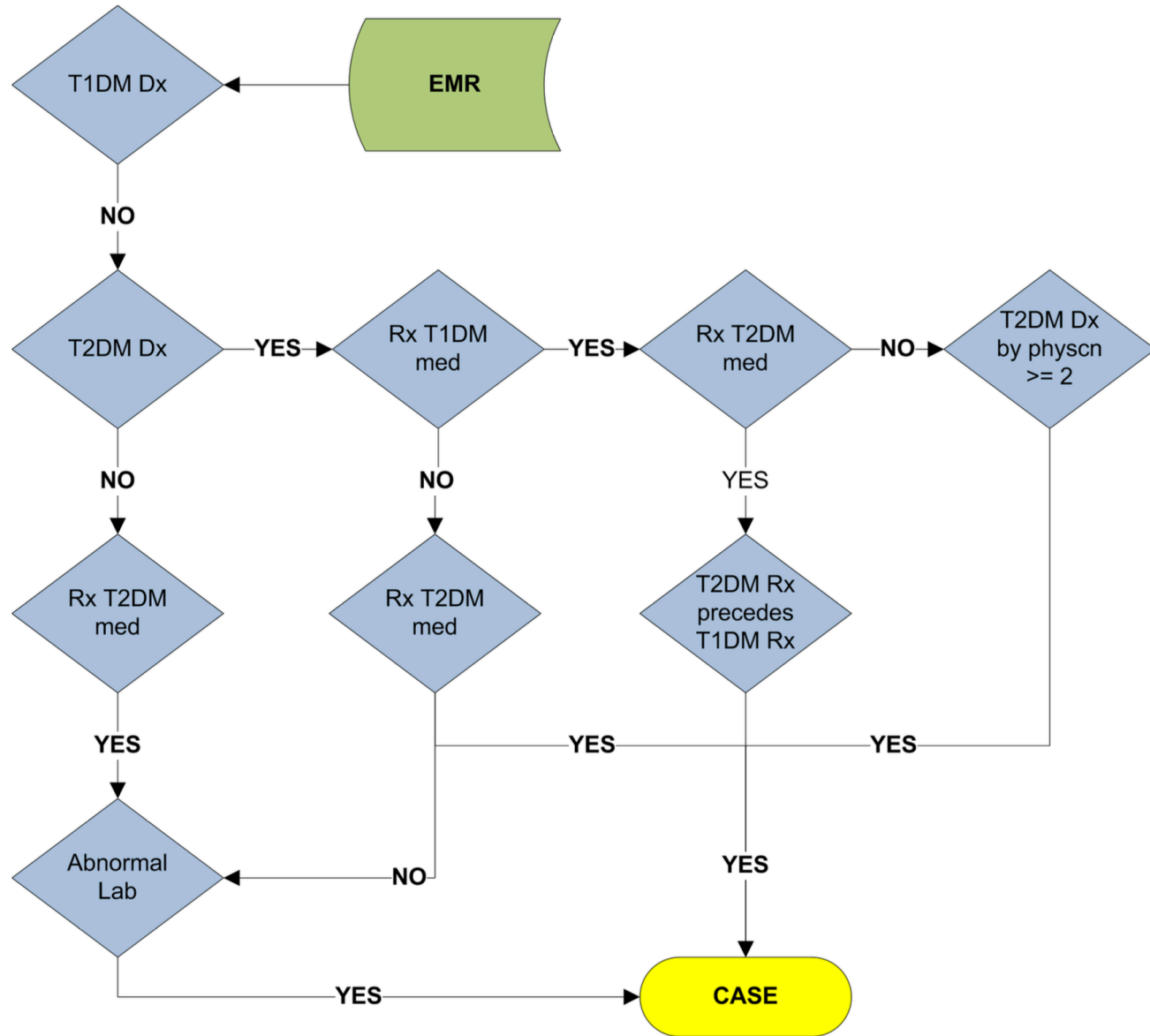


<https://github.com/nyuvis/patient-viz>

<https://github.com/BenGlicksberg/PatientExploreR>

Figure 1: Algorithm for identifying T2DM cases in the EMR.

Step 2:
Example of
a rule-based
phenotype
(Northwestern U.)



Step 2: Example of a rule-based phenotype

Coverage of Different Diabetes Outcome Definitions on Claims Data

Condition	Percentage
Have 250.x diagnosis, or have been on diabetic medication, or have any HbA1c \geq 6.5	100 %
Have been diagnosed 250.xx	89.9 %
Have been on diabetic medications	15.0 %
Have HbA1c values \geq 6.5	20.9 %
Have 250.xx diagnosis on more than one distinct date	40.0 %
(Have 250.xx diagnosis, or have been on diabetic medication, or have any HbA1c \geq 6.5) on more than one distinct date	44.0 %
(Have 250.xx diagnosis, or have been on diabetic medication, or have any HbA1c \geq 6.5) on two dates separated by at least a week	41.1 %

The earliest date the rule triggers is defined as the date of diabetes diagnosis

Definition selected



Public Phenotypes

Public Collaboration

Public phenotypes are believed to be complete and final by their authors. When you are logged in you can view and edit phenotypes in your groups that are not public and in various stages of development.

Login To View Private Group Phenotypes

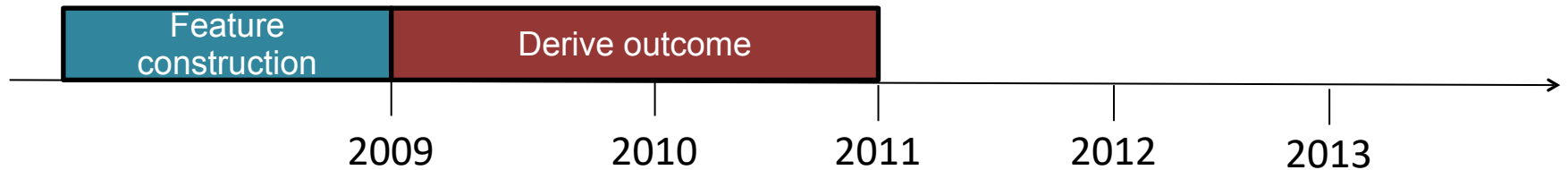
Institution Type of Phenotype Owner Phenotyping Groups View Phenotyping Groups

Data Model

Title	Institution	Data Modalities and Methods Used	Owner Phenotyping Groups	View Groups	Has new content	Status	Type
Abdominal Aortic Aneurysm (AAA)	Geisinger	CPT Codes, ICD 9 Codes, Vital Signs	eMERGE Geisinger Group	eMERGE Geisinger Group, eMERGE Phenotype WG		Final	Disease or Syndrome
ADHD phenotype algorithm	CHOP	ICD 9 Codes, Medications, Natural Language Processing	eMERGE CHOP Group	eMERGE Phenotype WG		Final	Disease or Syndrome
Appendicitis	Cincinnati Children's Hospital Medical Center	CPT Codes, ICD 9 Codes, Medications, Natural Language Processing	eMERGE CCHMC/BCH Group	eMERGE Phenotype WG		Final	Disease or Syndrome
Atrial Fibrillation - Demonstration Project	Vanderbilt University	CPT Codes, ICD 9 Codes, Natural Language Processing	Vanderbilt - SD/RD Group	Vanderbilt - SD/RD Group		Final	Disease or Syndrome
Autism	Cincinnati Children's Hospital Medical Center	ICD 9 Codes, Medications, Natural Language Processing	eMERGE CCHMC/BCH Group	eMERGE Phenotype WG		Final	Disease or Syndrome
Cataracts	Marshfield Clinic Research Foundation	CPT Codes, ICD 9 Codes, Medications, Natural Language Processing	eMERGE Marshfield Group	eMERGE Phenotype WG		Final	Disease or Syndrome
Crohn's Disease -	Vanderbilt University	ICD 9 Codes, Medications,	Vanderbilt -	Vanderbilt -		Final	Disease or Syndrome

Step 2:
Example of a
rule-based
phenotype

Framing for supervised machine learning



Exclusion criteria:

- Diabetes diagnosis (according to our rule) observed prior to January 1, 2009
- Less than 6 months of enrollment in feature construction window
- Member left health insurance prior to Jan. 1, 2011

What if someone is diagnosed with diabetes in 2012?

Why not model as “patient develops diabetes anytime after 2009”?

Framing for supervised machine learning



Exclusion criteria:

- Diabetes diagnosis (according to our rule) observed prior to January 1, ~~2009~~ 2011
- Less than 6 months of enrollment in feature construction window
- Member left health insurance prior to Jan. 1, ~~2011~~ 2013

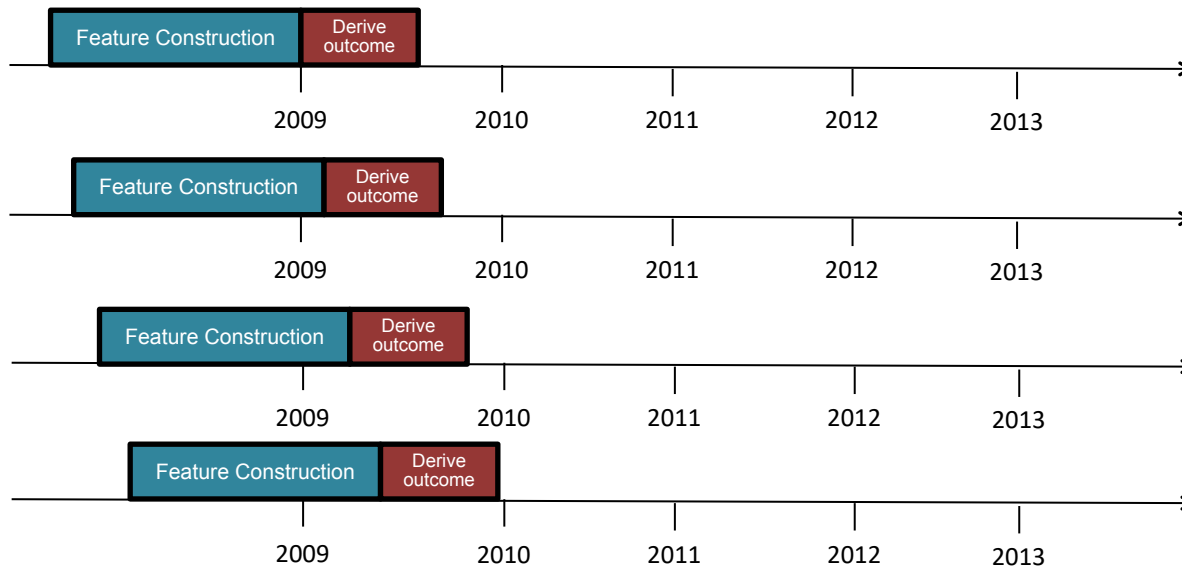
Framing for supervised machine learning



- Suppose we want to run the above model in August 2009. It may not have good performance due to *non-stationarity* in the data
- We now have data through 2021. Using a fixed prediction time / index date of Jan. 1, 2009 is ignoring most of the diabetes onsets!

Framing for supervised machine learning

- We can instead create *many* data points from each patient, using e.g. every month as an index date:



- **Important:** If multiple data points per patient, make sure each patient's data is in *only* train, validate, or test