

Machine Learning for Healthcare

HST.956, 6.S897

Lecture 4: Risk stratification

David Sontag



Course announcements

- Recitation Friday at 2pm (4-153) – optional
- No class this Tuesday
- Problem set 1 due next Thursday, Feb 21
- Sign up for lecture scribing or MLHC community consulting
- Readings will be posted several days ahead
- All course communication through Piazza

Outline for today's class

1. Risk stratification
2. Case study: Early detection of Type 2 diabetes
 - Framing as supervised learning problem
 - Deriving labels
 - Evaluating risk stratification algorithms
3. Subtleties with ML-based risk stratification
4. Discussion with Leonard D'Avolio (Assistant Professor at HMS, CEO @ Cyft)

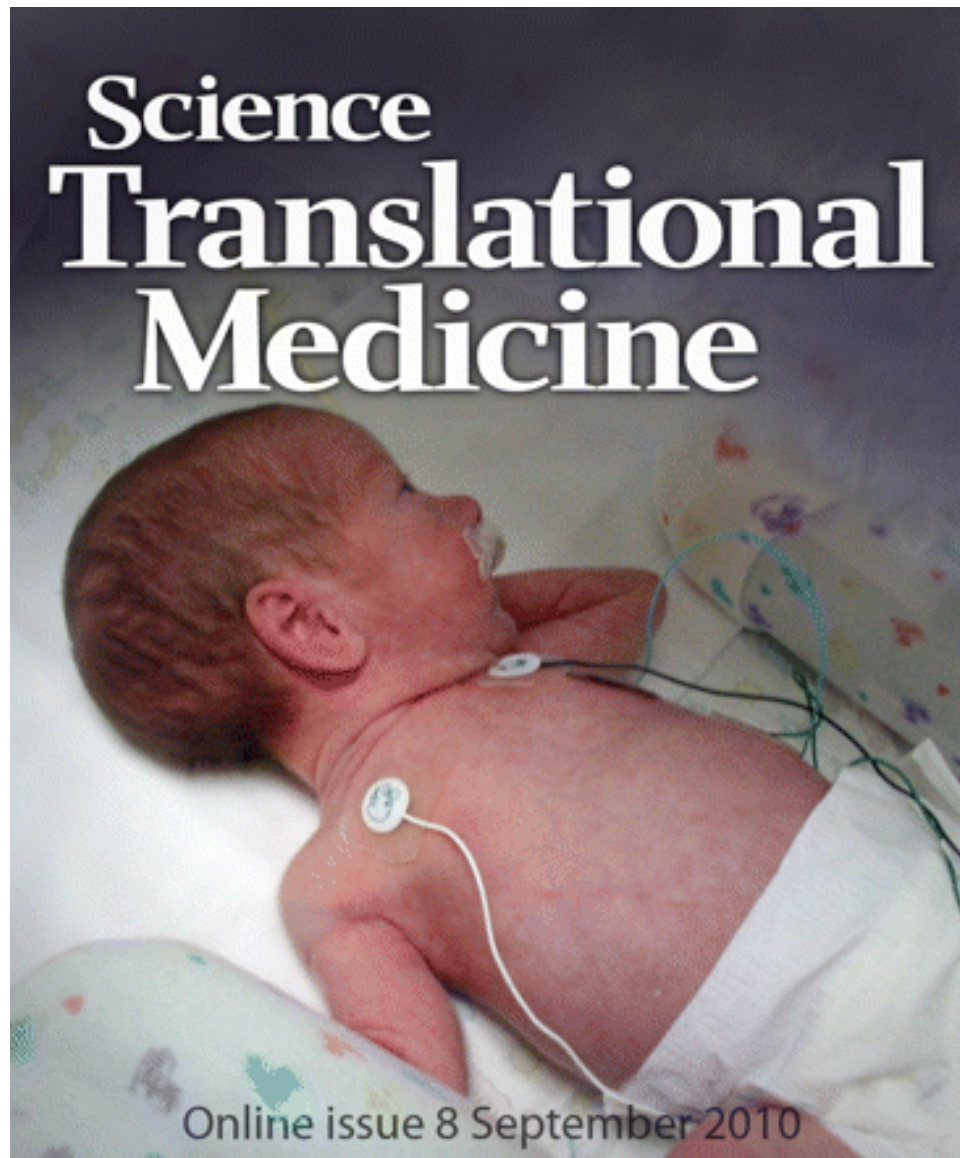
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What *is* risk stratification?

- Separate a patient population into **high-risk** and **low-risk** of having an outcome
 - Predicting something in the future
 - Goal is different from diagnosis, with distinct performance metrics
- 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



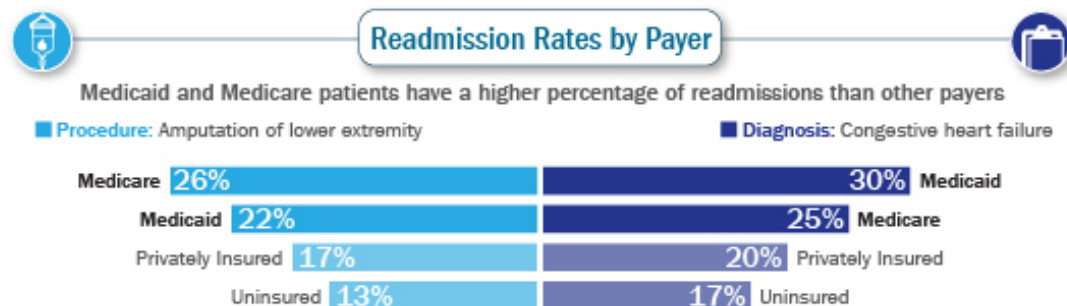
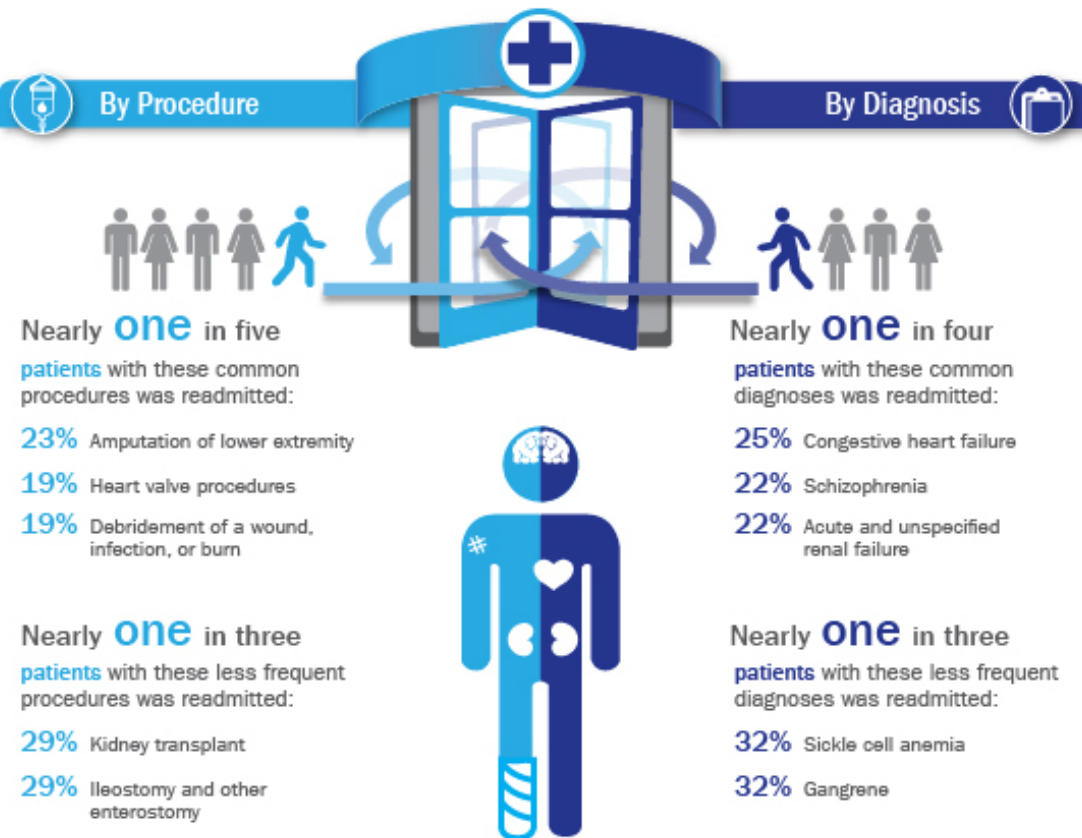
Does this patient need to be admitted to the coronary-care unit?

Figure source: <https://www.drmani.com/heart-attack/>

(Pozen et al., NEJM 1984)

30-DAY READMISSION RATES TO U.S. HOSPITALS

Healthcare Cost and Utilization Project (HCUP) data from 2010 provide the most comprehensive national estimates of 30-day readmission rates for specific procedures and diagnoses.* Examples include:



*Readmissions were for all causes and did not necessarily include the same procedure or diagnosis as the original admission (index stay).

Source: HCUP Statistical Briefs #153 and #154:
<http://www.hcup-us.ahrq.gov/reports/statbriefs/statbriefs.jsp>



AHRQ
 Agency for Healthcare Research and Quality
 Advancing Excellence in Health Care • www.ahrq.gov


Likelihood of hospital readmission?

Figure source:
<https://www.air.org/project/revolving-door-u-s-hospital-readmissions-diagnosis-and-procedure>

Old vs. New

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

APGAR SCORING SYSTEM

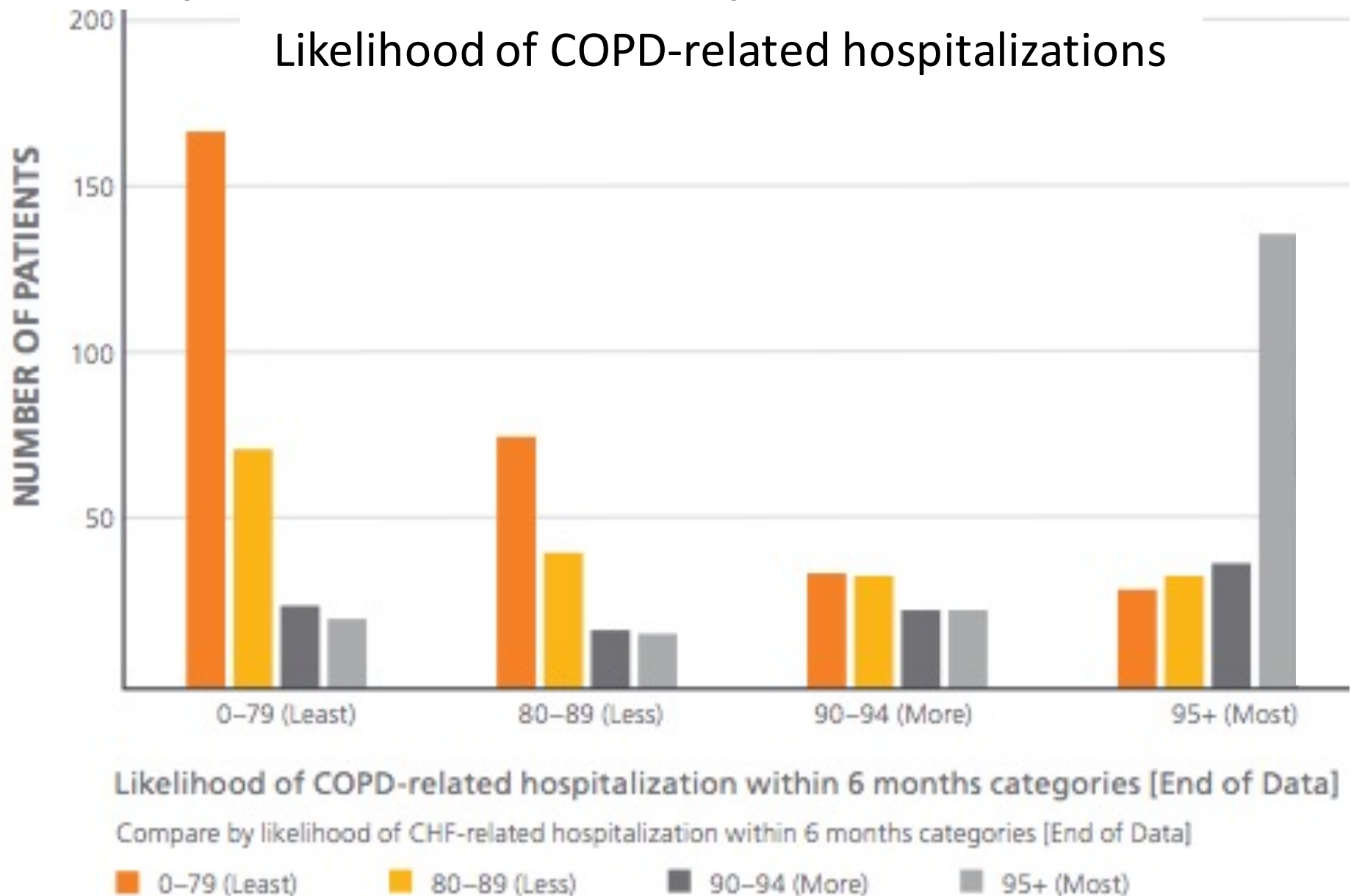
	0 Points	1 Point	2 Points	Points totaled
Activity (muscle tone)	Absent	Arms and legs flexed	Active movement	
Pulse	Absent	Below 100 bpm	Over 100 bpm	
Grimace (reflex irritability)	Flaccid	Some flexion of Extremities	Active motion (sneeze, cough, pull away)	
Appearance (skin color)	Blue, pale	Body pink,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

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

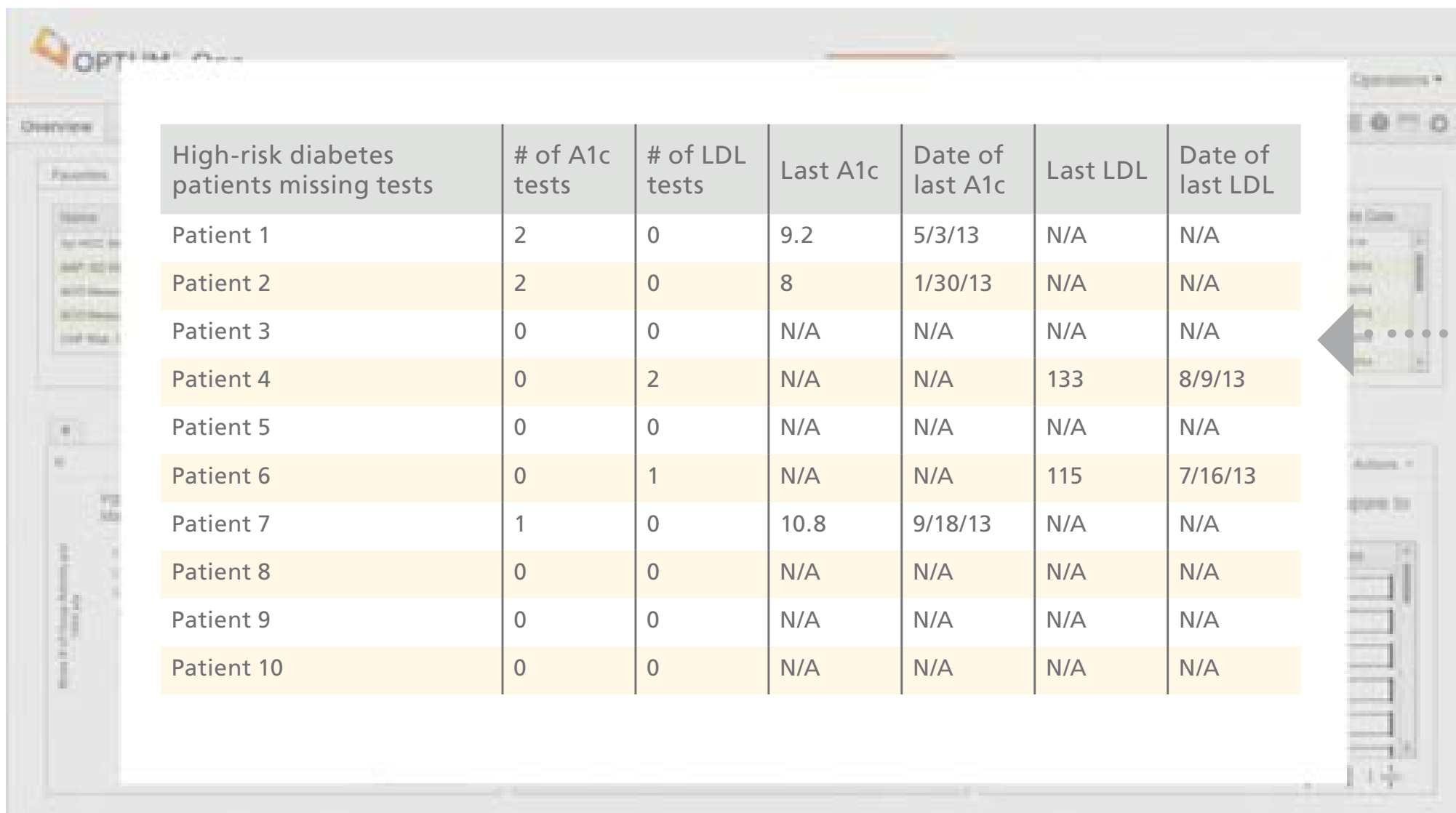
Old vs. New

- 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, new dangers introduced with ML approach – to be discussed**

Example commercial product



Example commercial product



The screenshot shows the OPTIM software interface. On the left is a sidebar with 'Overview' and 'Patients' tabs. The main area displays a table with 7 columns: 'High-risk diabetes patients missing tests', '# of A1c tests', '# of LDL tests', 'Last A1c', 'Date of last A1c', 'Last LDL', and 'Date of last LDL'. The table lists data for 10 patients. Rows for Patient 2, Patient 4, Patient 6, Patient 8, and Patient 10 are highlighted in yellow. On the right, there is a vertical toolbar with various icons, and a grey arrow points from the table area towards it.

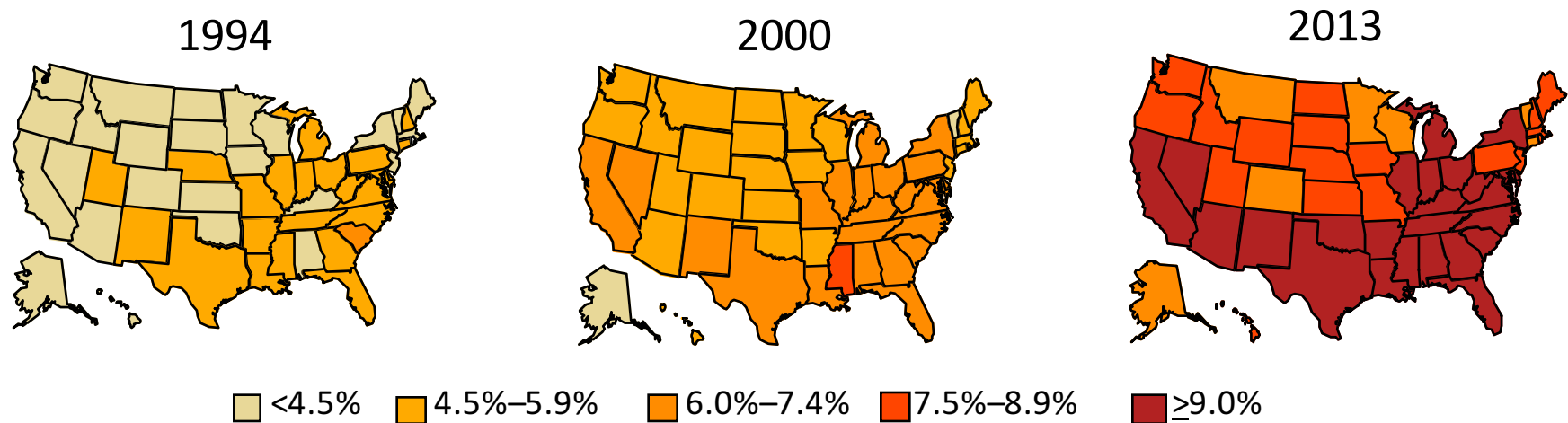
High-risk diabetes patients missing tests	# of A1c tests	# of LDL tests	Last A1c	Date of last A1c	Last LDL	Date of last LDL
Patient 1	2	0	9.2	5/3/13	N/A	N/A
Patient 2	2	0	8	1/30/13	N/A	N/A
Patient 3	0	0	N/A	N/A	N/A	N/A
Patient 4	0	2	N/A	N/A	133	8/9/13
Patient 5	0	0	N/A	N/A	N/A	N/A
Patient 6	0	1	N/A	N/A	115	7/16/13
Patient 7	1	0	10.8	9/18/13	N/A	N/A
Patient 8	0	0	N/A	N/A	N/A	N/A
Patient 9	0	0	N/A	N/A	N/A	N/A
Patient 10	0	0	N/A	N/A	N/A	N/A

Optum Whitepaper, "Predictive analytics: Poised to drive population health"

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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

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

* Diabetes Prevention Program Research Group. "Reduction in the incidence of type 2 diabetes with lifestyle intervention or metformin." The New England journal of medicine 346.6 (2002): 393.

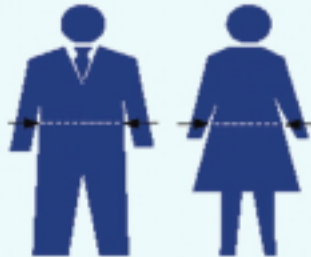
Traditional Risk Prediction Models

- 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 25 kg/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%;"> <thead> <tr> <th style="text-align: left;">MEN</th> <th style="text-align: left;">WOMEN</th> </tr> </thead> <tbody> <tr> <td>0 p. Less than 94 cm</td> <td>Less than 80 cm</td> </tr> <tr> <td>3 p. 94–102 cm</td> <td>80–88 cm</td> </tr> <tr> <td>4 p. More than 102 cm</td> <td>More than 88 cm</td> </tr> </tbody> </table> <p></p> <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>	MEN	WOMEN	0 p. Less than 94 cm	Less than 80 cm	3 p. 94–102 cm	80–88 cm	4 p. More than 102 cm	More than 88 cm	<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 94 cm	Less than 80 cm								
3 p. 94–102 cm	80–88 cm								
4 p. More than 102 cm	More than 88 cm								

Total risk score

☐ The risk of developing type 2 diabetes within 10 years is

Lower than 7	Low: estimated 1 in 100 will develop disease
7–11	Slightly elevated: estimated 1 in 25 will develop disease
12–14	Moderate: estimated 1 in 6 will develop disease
15–20	High: estimated 1 in 3 will develop disease
Higher than 20	Very high: estimated 1 in 2 will develop disease

Please turn over

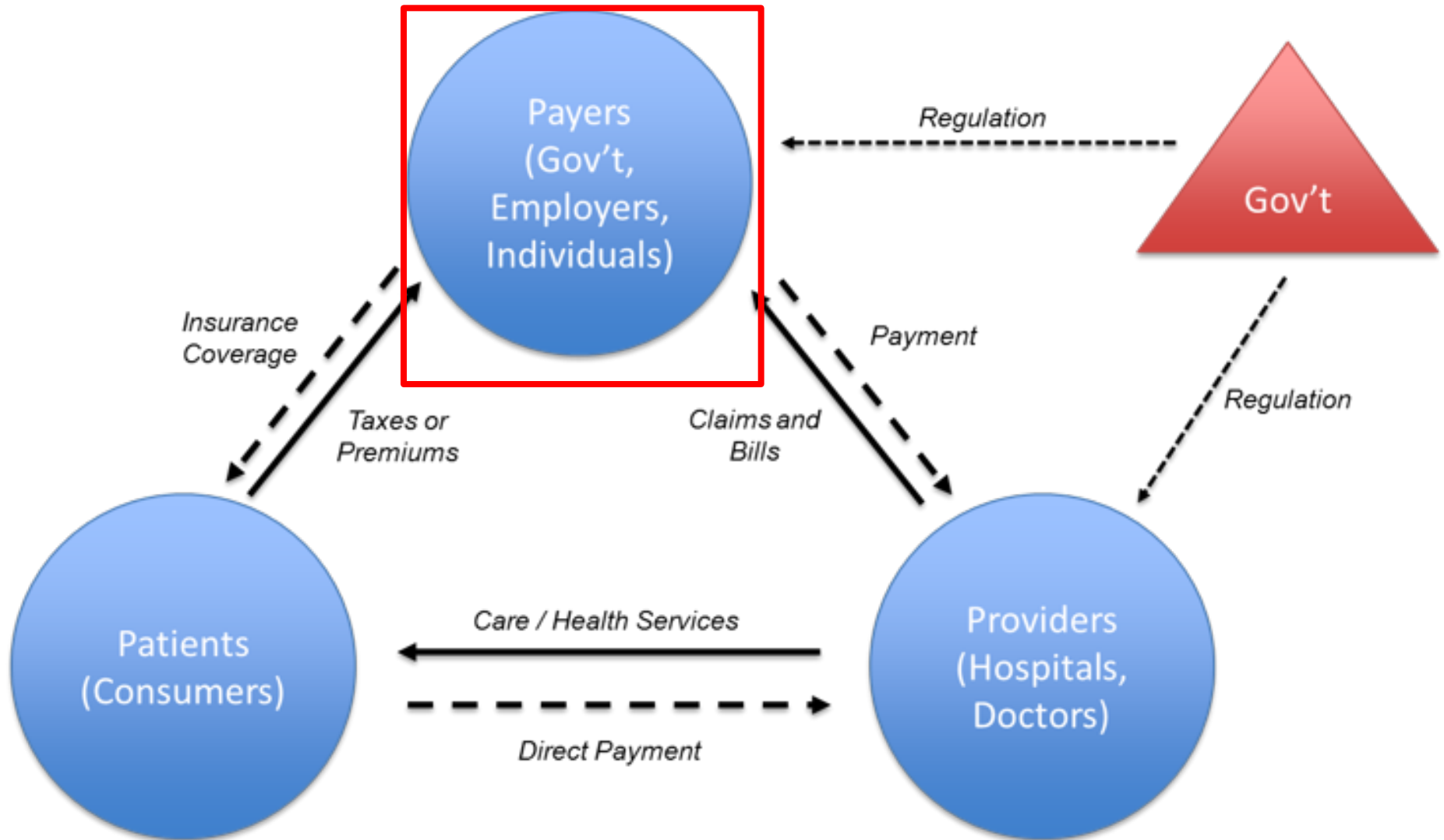
Challenges of Traditional Risk Prediction Models

- A screening step needs to be done for every member in the population
 - Either in the physician's office or as surveys
 - Costly and time-consuming
 - Infeasible for regular screening for millions of individuals
- Models not easy to adapt to multiple surrogates, when a variable is missing
 - Discovery of surrogates not straightforward

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

Health stakeholders

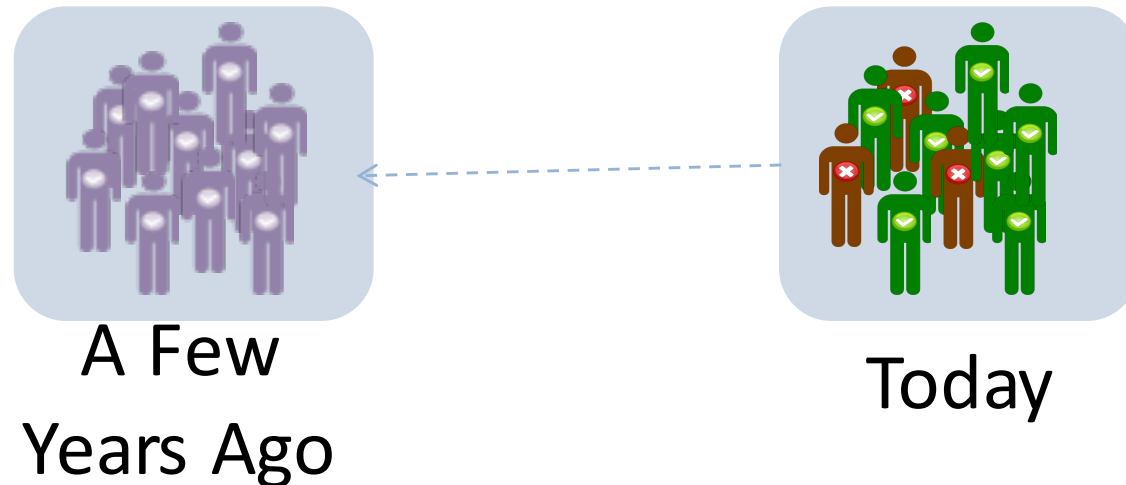


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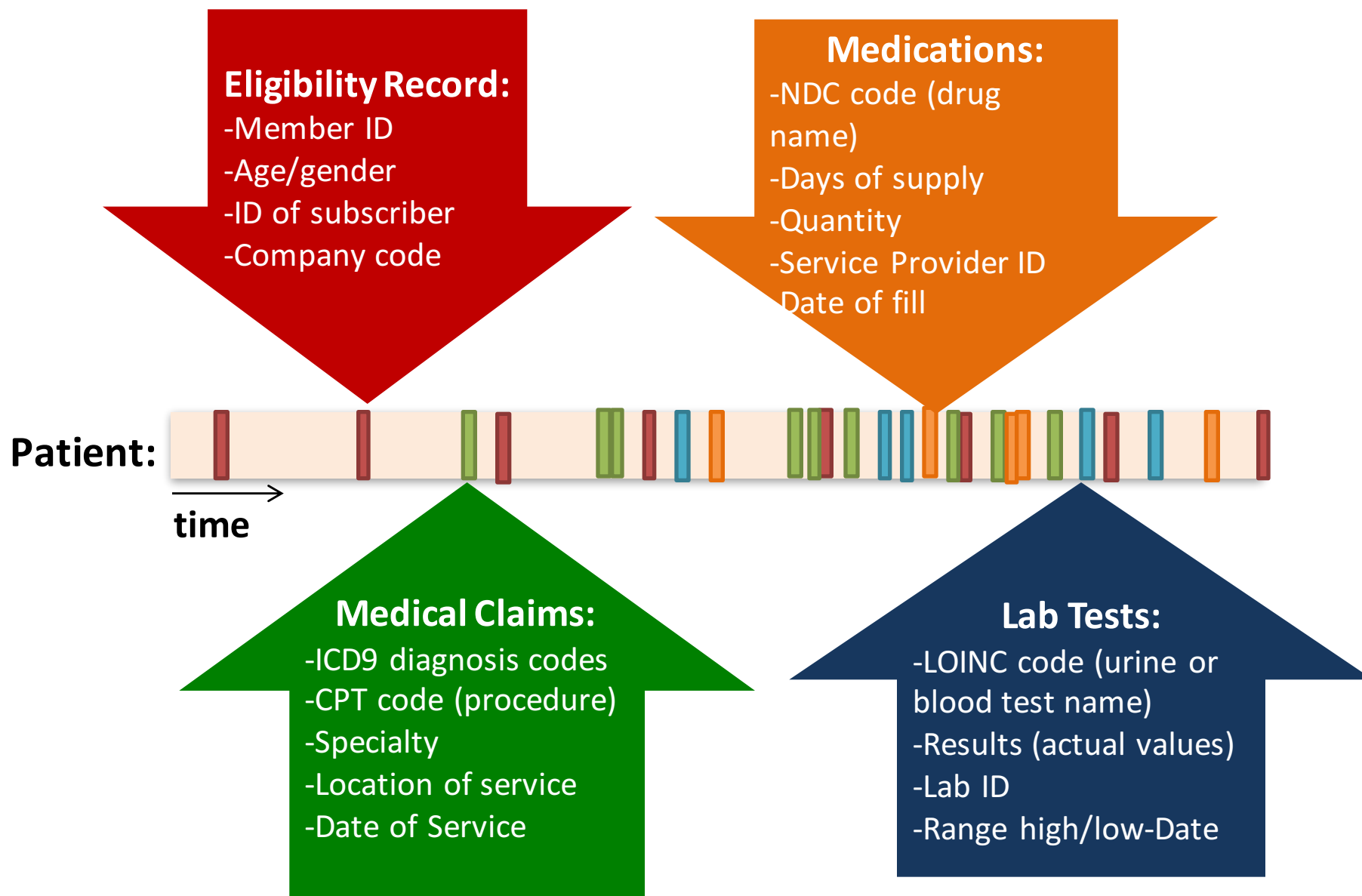
<http://www.mahesh-vc.com/blog/understanding-whos-paying-for-what-in-the-healthcare-industry>

A Data-Driven approach on Longitudinal Data

- 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?



Administrative & Clinical Data



Top diagnosis codes

Disease	count
4011 Benign hypertension	447017
2724 Hyperlipidemia NEC/NOS	382030
4019 Hypertension NOS	372477
25000 DMII wo cmp nt st uncuntr	339522
2720 Pure hypercholesterolem	232671
2722 Mixed hyperlipidemia	180015
V7231 Routine gyn examination	178709
2449 Hypothyroidism NOS	169829
78079 Malaise and fatigue NEC	149797
V0481 Vaccin for influenza	147858
7242 Lumbago	137345
V7612 Screen mammogram NEC	129445
V700 Routine medical exam	127848

Disease	count
53081 Esophageal reflux	121064
42731 Atrial fibrillation	113798
7295 Pain in limb	112449
41401 Crnry athrscd natve vssl	104478
2859 Anemia NOS	103351
78650 Chest pain NOS	91999
5990 Urin tract infection NOS	87982
V5869 Long-term use meds NEC	85544
496 Chr airway obstruct NEC	78585
4779 Allergic rhinitis NOS	77963
41400 Cor ath unsp vsl ntv/gft	75519

Disease	count
71947 Joint pain-ankle	28648
3004 Dysthymic disorder	28530
2689 Vitamin D deficiency NOS	28455
V7281 Preop cardiovsclr exam	27897
7243 Sciatica	27604
78791 Diarrhea	27424
V221 Supervis oth normal preg	27320
36501 Opn angl brderln lo risk	26033
37921 Vitreous degeneration	25592
4241 Aortic valve disorder	25425
61610 Vaginitis NOS	24736
70219 Other sborheic keratosis	24453
3804 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

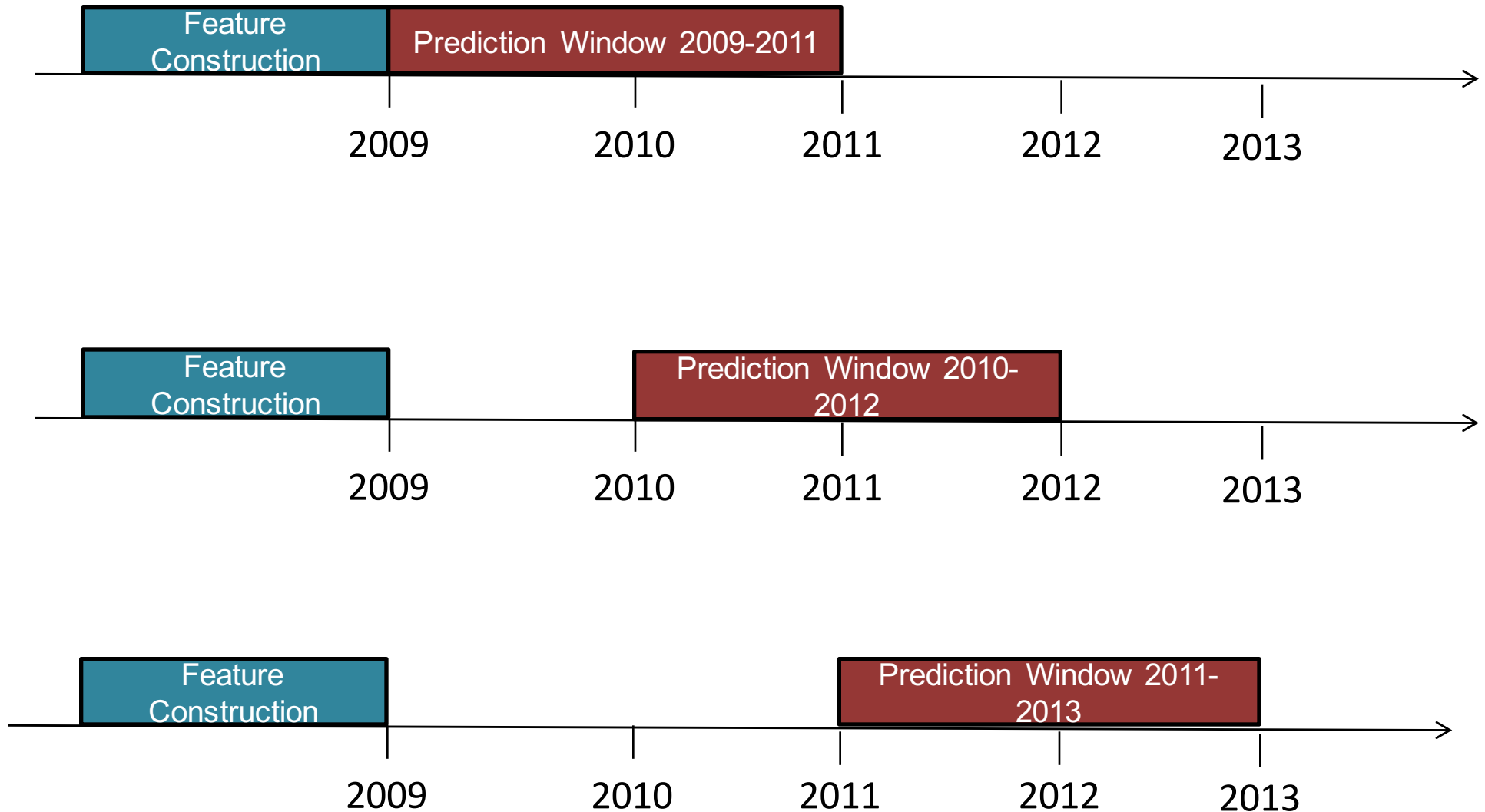
Lab test	
770-8 Neutrophils/100 leukocytes	952089
731-0 Lymphocytes	943918
704-7 Basophils	863448
711-2 Eosinophils	935710
5905-5 Monocytes/100 leukocytes	943764
706-2 Basophils/100 leukocytes	863435
751-8 Neutrophils	943232
742-7 Monocytes	942978
713-8 Eosinophils/100 leukocytes	933929
3016-3 Thyrotropin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

Count of people who have the test result (ever)

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Framing for supervised machine learning



Gap is important to prevent label leakage

Framing for supervised machine learning

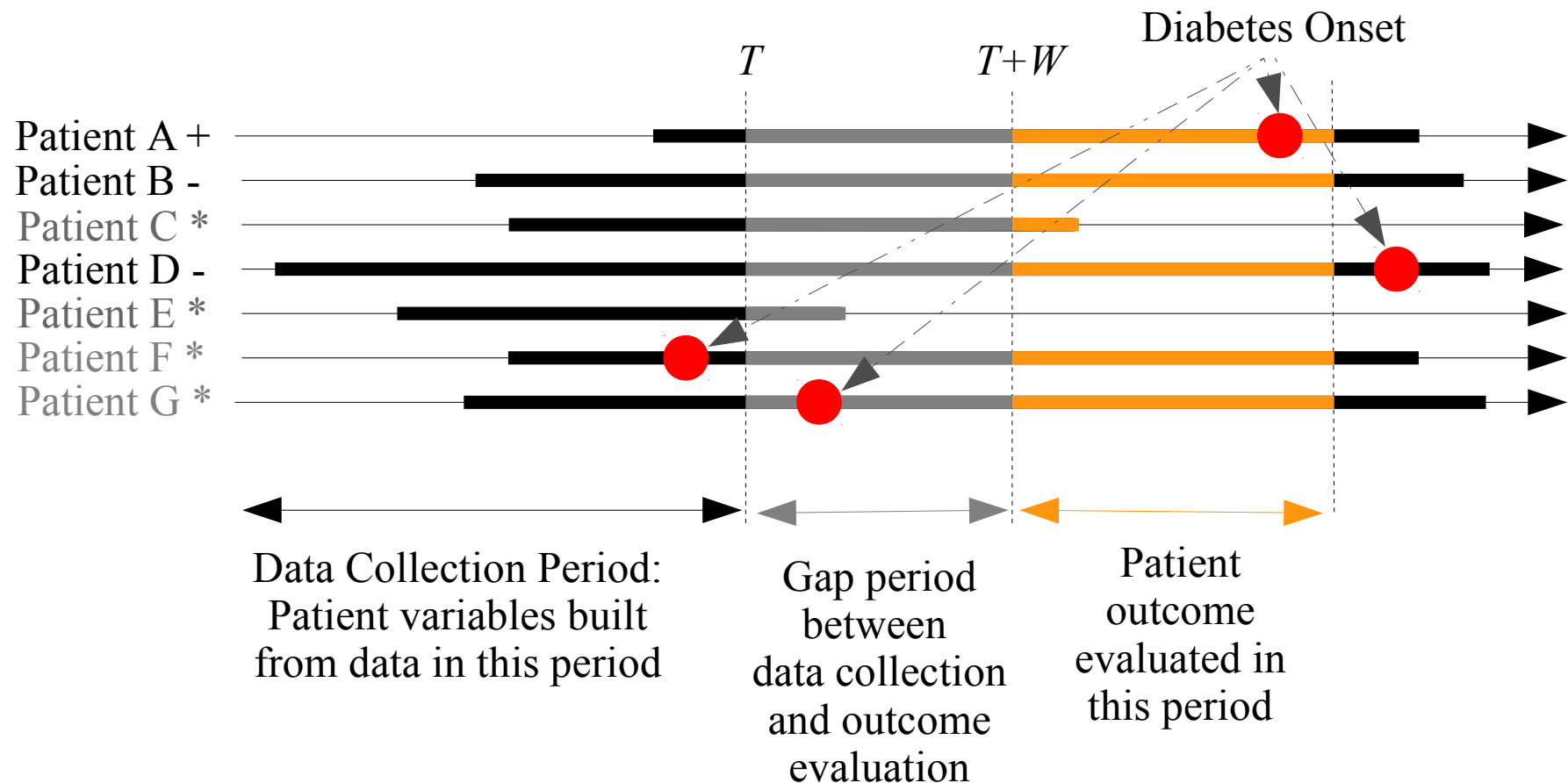


Problem: Data is censored!

- Patients change health insurers frequently, but data doesn't follow them
- *Left censored*: may not have enough data to derive features
- *Right censored*: may not know label

Reduction to binary classification

Exclude patients that are left- and right-censored.



This is an example of alignment by *absolute time*

Alternative framings

- Align by relative time, e.g.
 - 2 hours into patient stay in ER
 - Every time patient sees PCP
 - When individual turns 40 yrs old
- Align by data availability

NOTE:

- If multiple data points per patient, make sure each patient in *only* train, validate, or test

Methods

- L1 Regularized Logistic Regression
 - Simultaneously optimizes predictive performance *and*
 - Performs feature selection, choosing the subset of the features that are most predictive
- This prevents overfitting to the training data

L1 regularization

- Penalizing the L1 norm of the weight vector leads to *sparse* (read: many 0's) solutions for w .

$$\min_w \sum_i \ell(x_i, y_i; w) + \lambda ||w||_1 \qquad ||\vec{w}||_1 = \sum_d |w_d|$$

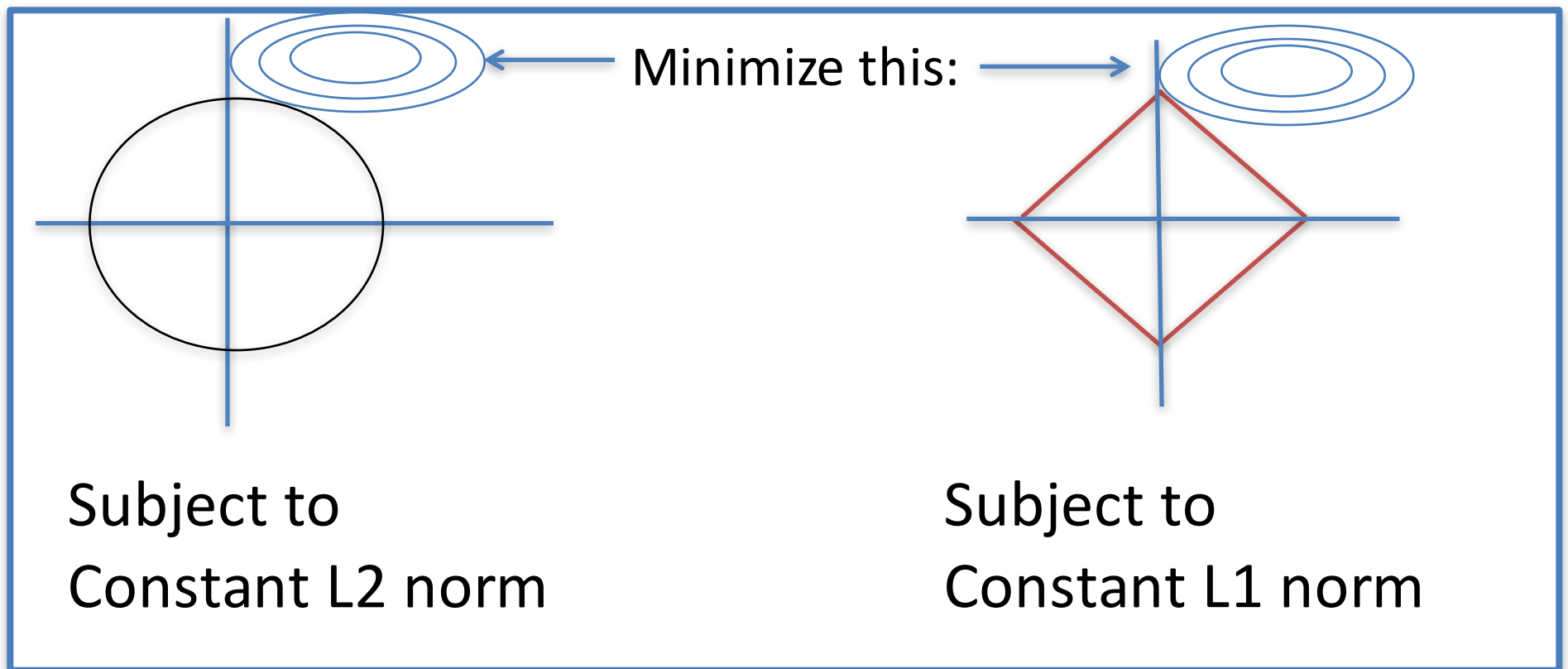
instead of

$$\min_w \sum_i \ell(x_i, y_i; w) + \lambda ||w||_2^2 \qquad ||\vec{w}||_2^2 = \sum_d w_d^2$$

- Why?

L1 regularization

- Penalizing the L1 norm of the weight vector leads to *sparse* (read: many 0's) solutions for w .



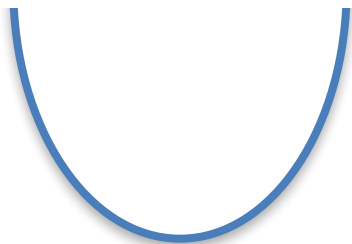
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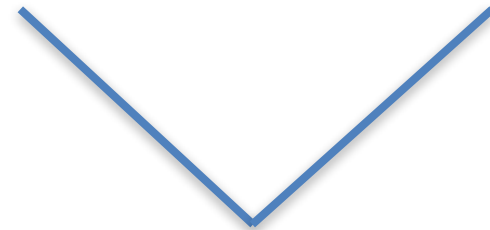
Intuition #2 – w.w.g.d.d

(What would gradient descent do?)

$$\frac{d}{dw_i} \lambda ||w||_2^2 = \pm \lambda w_i$$



$$\frac{d}{dw_i} \lambda |w| = \pm \lambda$$



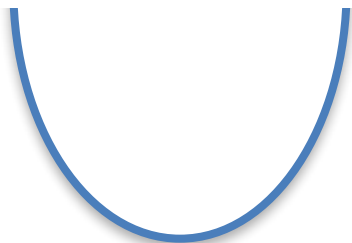
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Intuition #2 – w.w.g.d.d

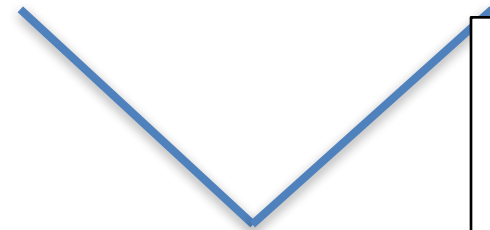
(What would gradient descent do?)

$$\frac{d}{dw_i} \lambda ||w||_2^2 = \pm \lambda w_i 2$$



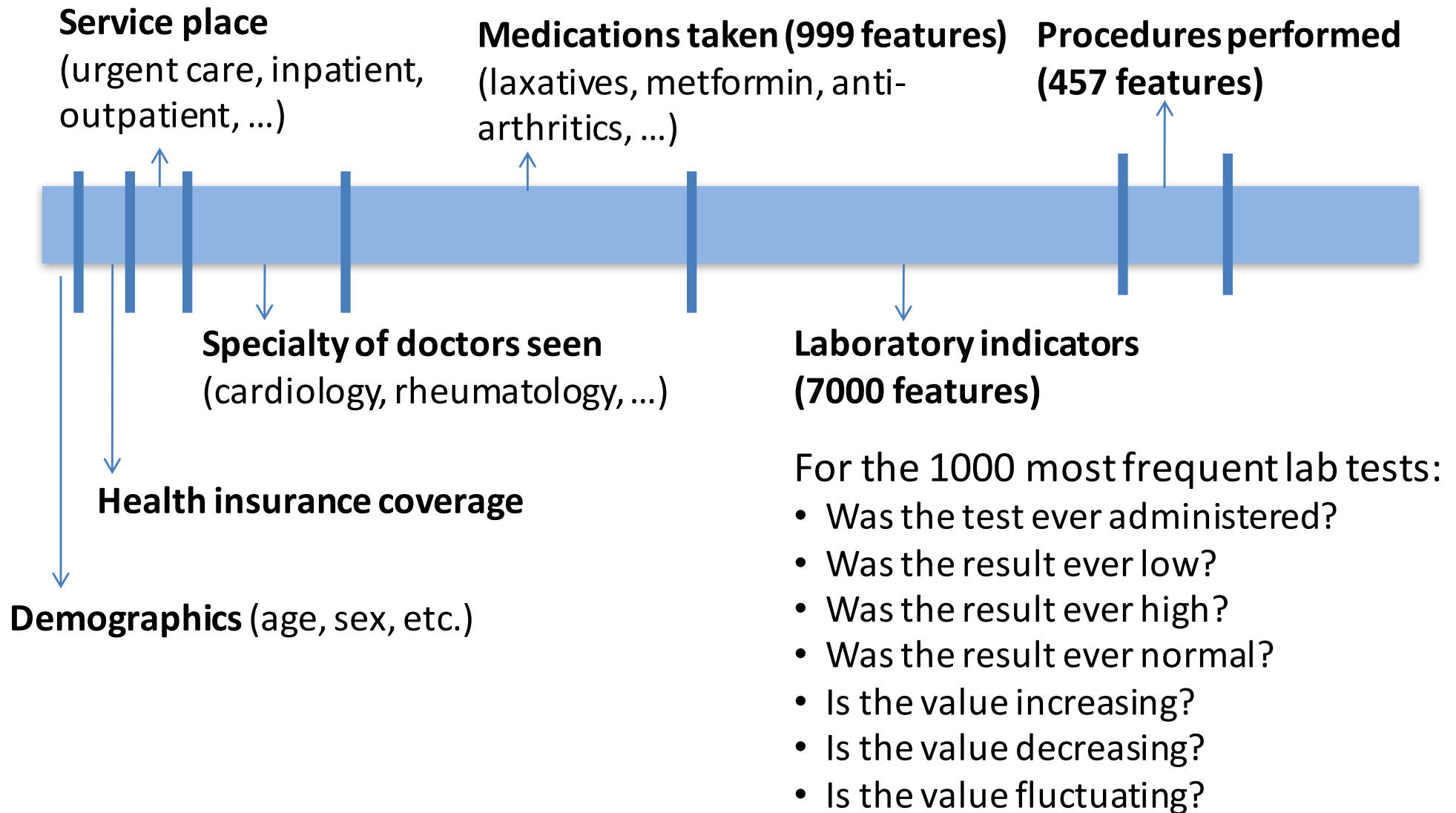
The push
towards 0 gets
weaker as w_i
gets smaller

$$\frac{d}{dw_i} \lambda |w| = \pm \lambda$$

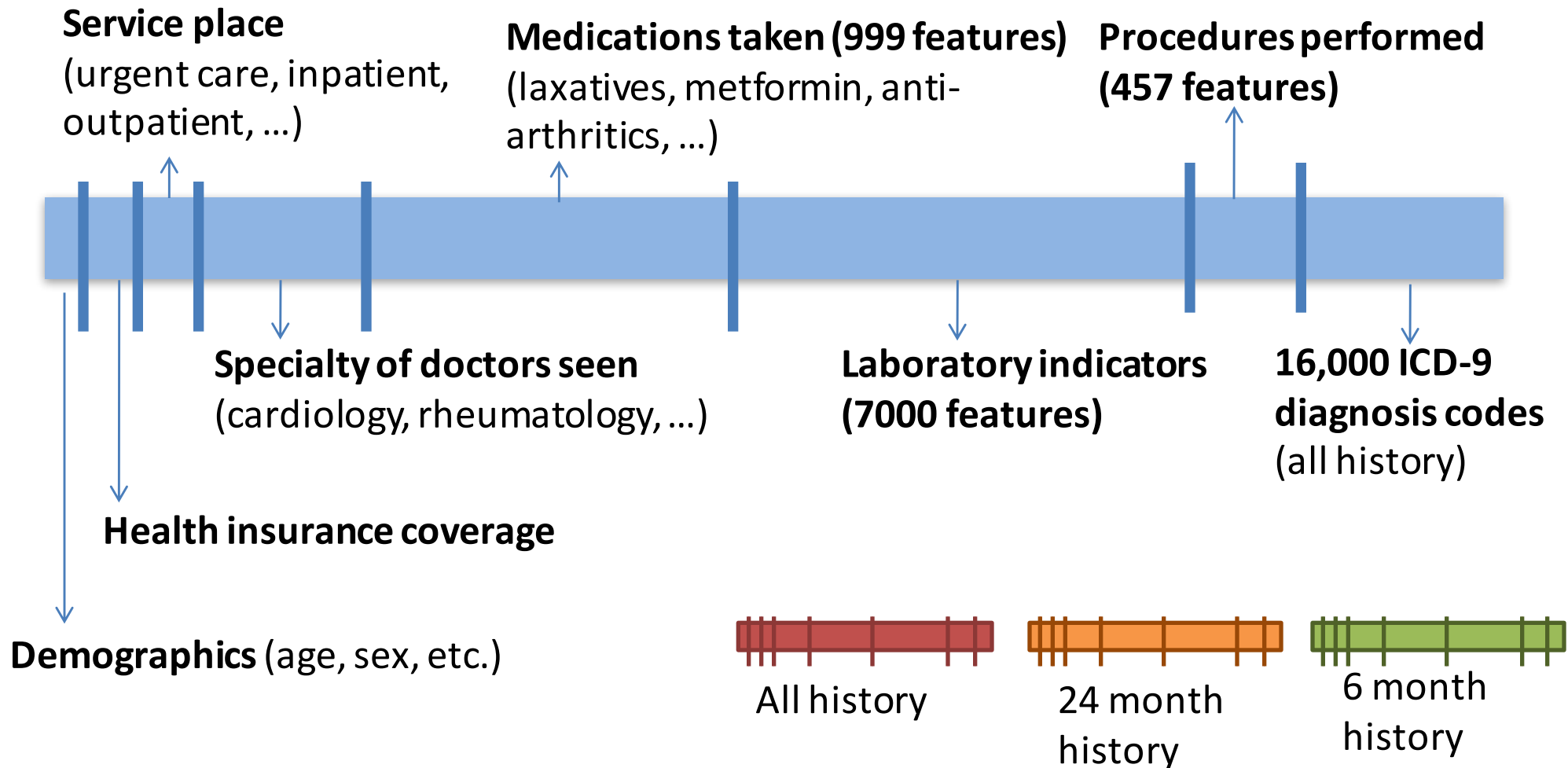


Always
pushes
elements of
 w_i towards 0

Features used in models



Features used in models

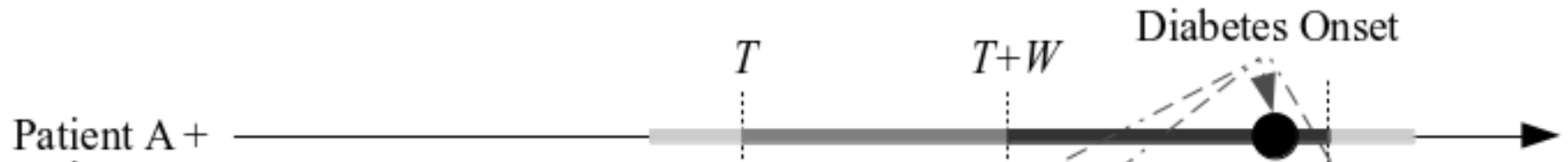


Total features per patient: 42,000

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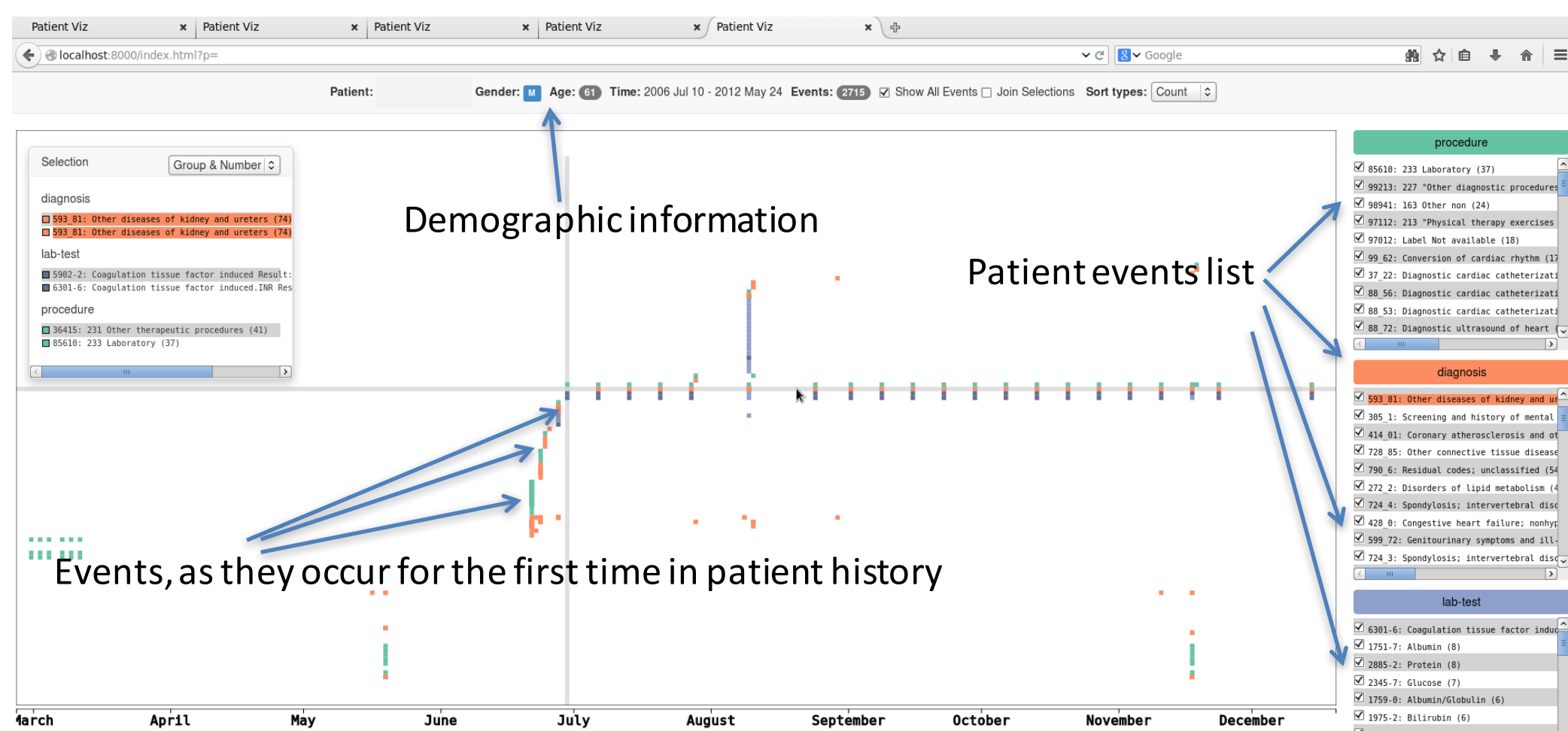
Where do the labels come from?



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

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





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 non public and in various stages of development.

[Login To View Private Group Phenotypes](#)

Institution

▶

Type of Phenotype

Disease or Syndrome ▼

Owner Phenotyping Groups

▶

View Phenotyping Groups

▶

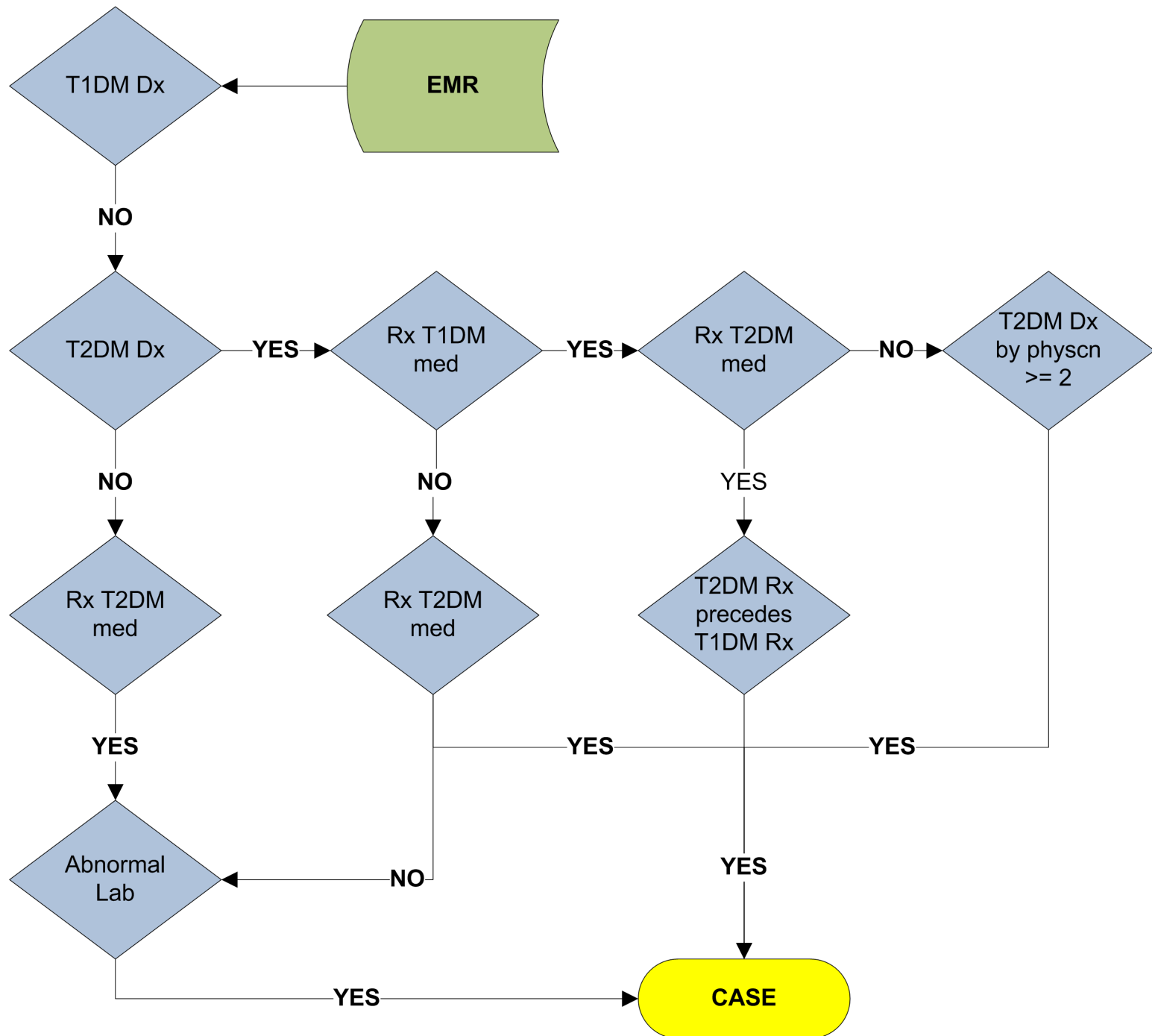
Data Model

- Any - ▼

Apply

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

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



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What are the Discovered Risk Factors?

- 769 variables have non-zero weight

Top History of Disease	Odds Ratio
Impaired Fasting Glucose (Code 790.21)	4.17 (3.87 4.49)
Abnormal Glucose NEC (790.29)	4.07 (3.76 4.41)
Hypertension (401)	3.28 (3.17 3.39)
Obstructive Sleep Apnea (327.23)	2.98 (2.78 3.20)
Obesity (278)	2.88 (2.75 3.02)
Abnormal Blood Chemistry (790.6)	2.49 (2.36 2.62)
Hyperlipidemia (272.4)	2.45 (2.37 2.53)
Shortness Of Breath (786.05)	2.09 (1.99 2.19)
Esophageal Reflux (530.81)	1.85 (1.78 1.93)

Diabetes
1-year gap

What are the Discovered Risk Factors?

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Top History of Disease

Impaired Fasting Glucose (Code

Abnormal Glucose NEC (790.29)

Hypertension (401)

Obstructive Sleep Apnea (327.23)

Obesity (278)

Abnormal Blood Chemistry (790.6

Hyperlipidemia (272.4)

Shortness Of Breath (786.05)

Esophageal Reflux (530.81)

Additional Disease Risk Factors Include:

Pituitary dwarfism (253.3),

Hepatomegaly(789.1), Chronic Hepatitis C

(070.54), Hepatitis (573.3), Calcaneal

Spur(726.73), Thyrotoxicosis without

mention of goiter(242.90), Sinoatrial Node

dysfunction(427.81), Acute frontal sinusitis

(461.1), Hypertrophic and atrophic

conditions of skin(701.9), Irregular

menstruation(626.4), ...

(1.99 2.19)

1.85

(1.78 1.93)

Diabetes
1-year gap

What are the Discovered Risk Factors?

- 769 variables have non-zero weight

Top Lab Factors	Odds Ratio
Hemoglobin A1c /Hemoglobin.Total (High - past 2 years)	5.75 (5.42 6.10)
Glucose (High- Past 6 months)	4.05 (3.89 4.21)
Cholesterol.In VLDL (Increasing - Past 2 years)	3.88 (3.53 4.27)
Potassium (Low - Entire History)	2.58 (2.24 2.98)
Cholesterol.Total/Cholesterol.In HDL (High - Entire History)	2.29 (2.19 2.40)
Erythrocyte mean corpuscular hemoglobin concentration -(Low - Entire History)	2.25 (1.92 2.64)
Eosinophils (High - Entire History)	2.11 (1.82 2.44)
Glomerular filtration rate/1.73 sq M.Predicted (Low -Entire History)	2.07 (1.92 2.24)
Alanine aminotransferase (High Entire History)	2.04 (1.89 2.19)

Diabetes
1-year gap

What are the Discovered Risk Factors?

- 769 variables have non-zero weight

Top Lab Factors

Hemoglobin A1c /Hemoglobin.Total (High

Glucose (High- Past 6 months)

Cholesterol.In VLDL (Increasing - Past 2 y

Potassium (Low - Entire History)

Cholesterol.Total/Cholesterol.In HDL (High

Additional Lab Test Risk Factors Include:

Albumin/Globulin (Increasing -Entire history), Urea nitrogen/Creatinine -(high - Entire History), Specific gravity (Increasing, Past 2 years), Bilirubin (high -Past 2 years),...

Erythrocyte mean corpuscular hemoglobin concentration -(Low - Entire History)

Eosinophils (High - Entire History)

Glomerular filtration rate/1.73 sq M.Predicted (Low -Entire History)

Alanine aminotransferase (High Entire History)

(2.10 2.10)

2.25
(1.92 2.64)

2.11
(1.82 2.44)

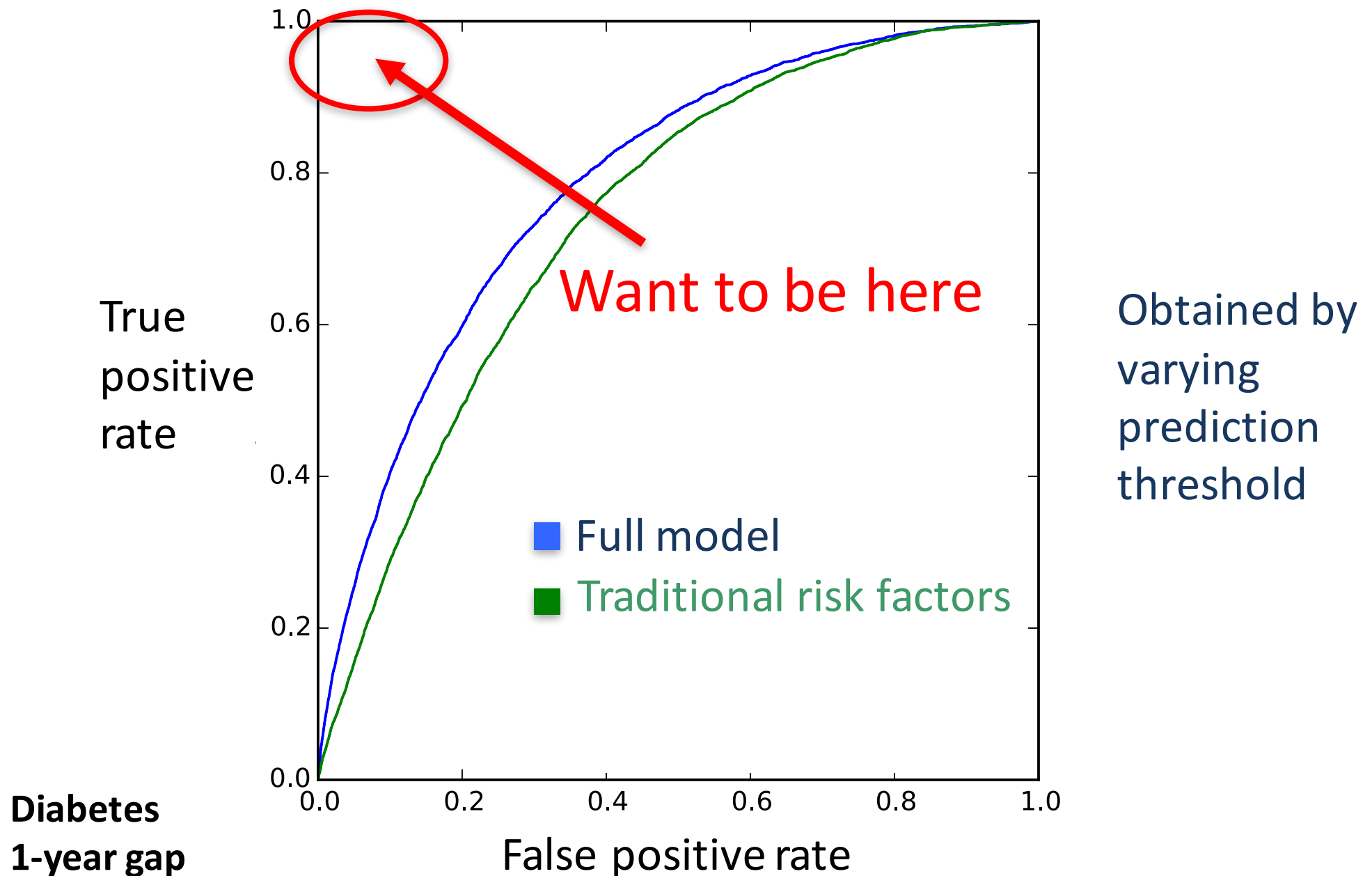
2.07
(1.92 2.24)

2.04
(1.89 2.19)

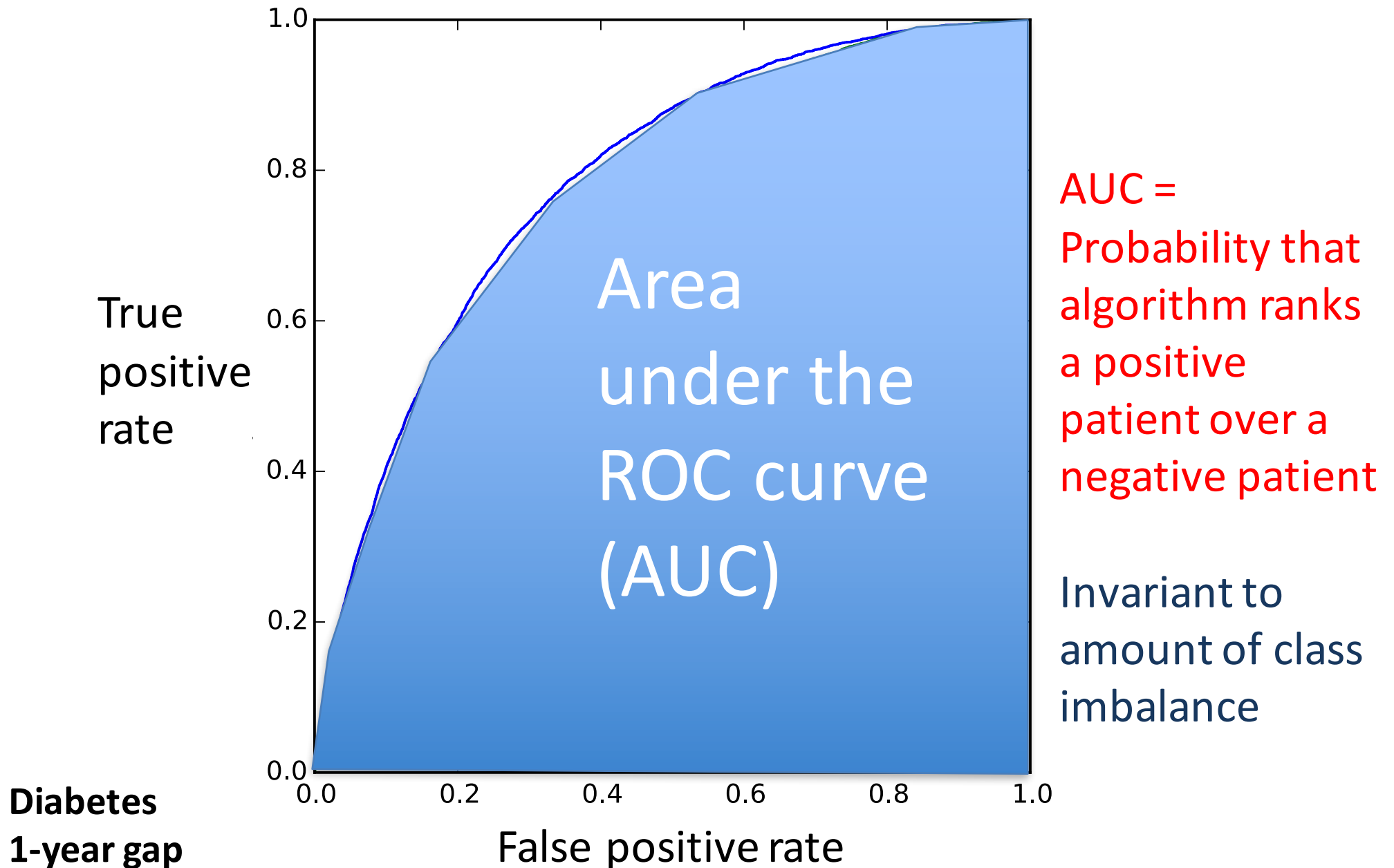
Diabetes

1-year gap

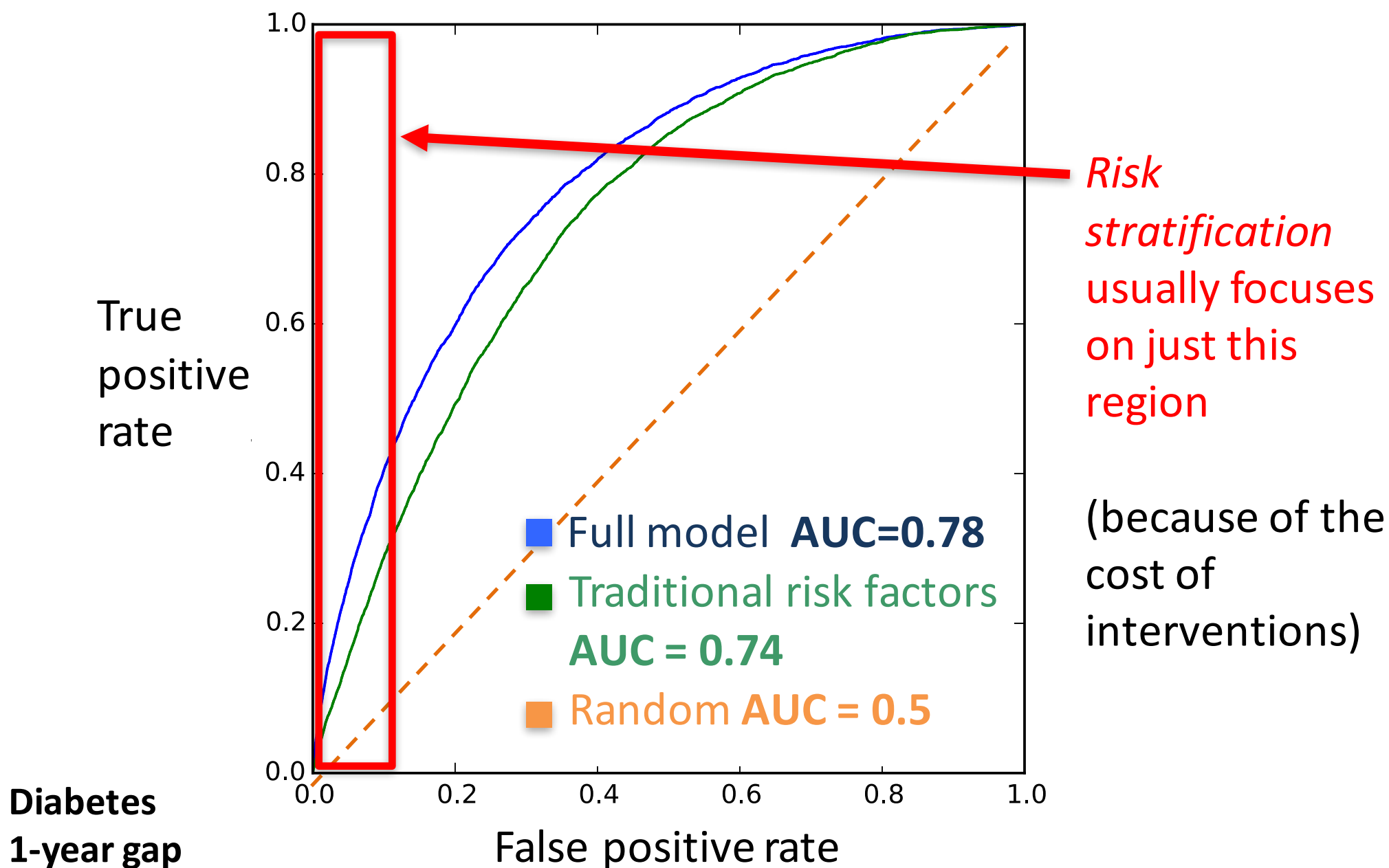
Receiver-operator characteristic curve



Receiver-operator characteristic curve

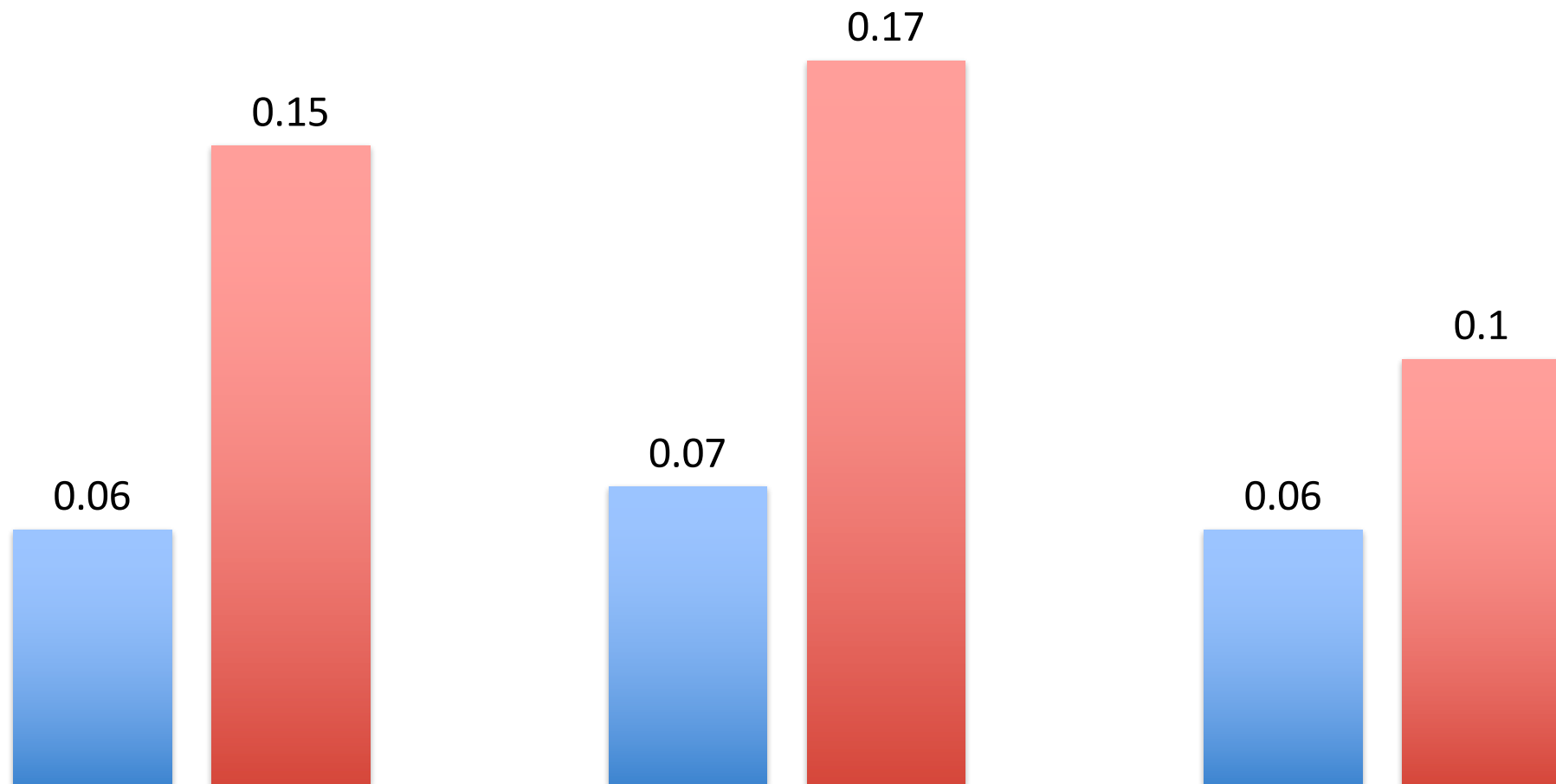


Receiver-operator characteristic curve



Positive predictive value (PPV)

■ Traditional risk factors ■ Full model



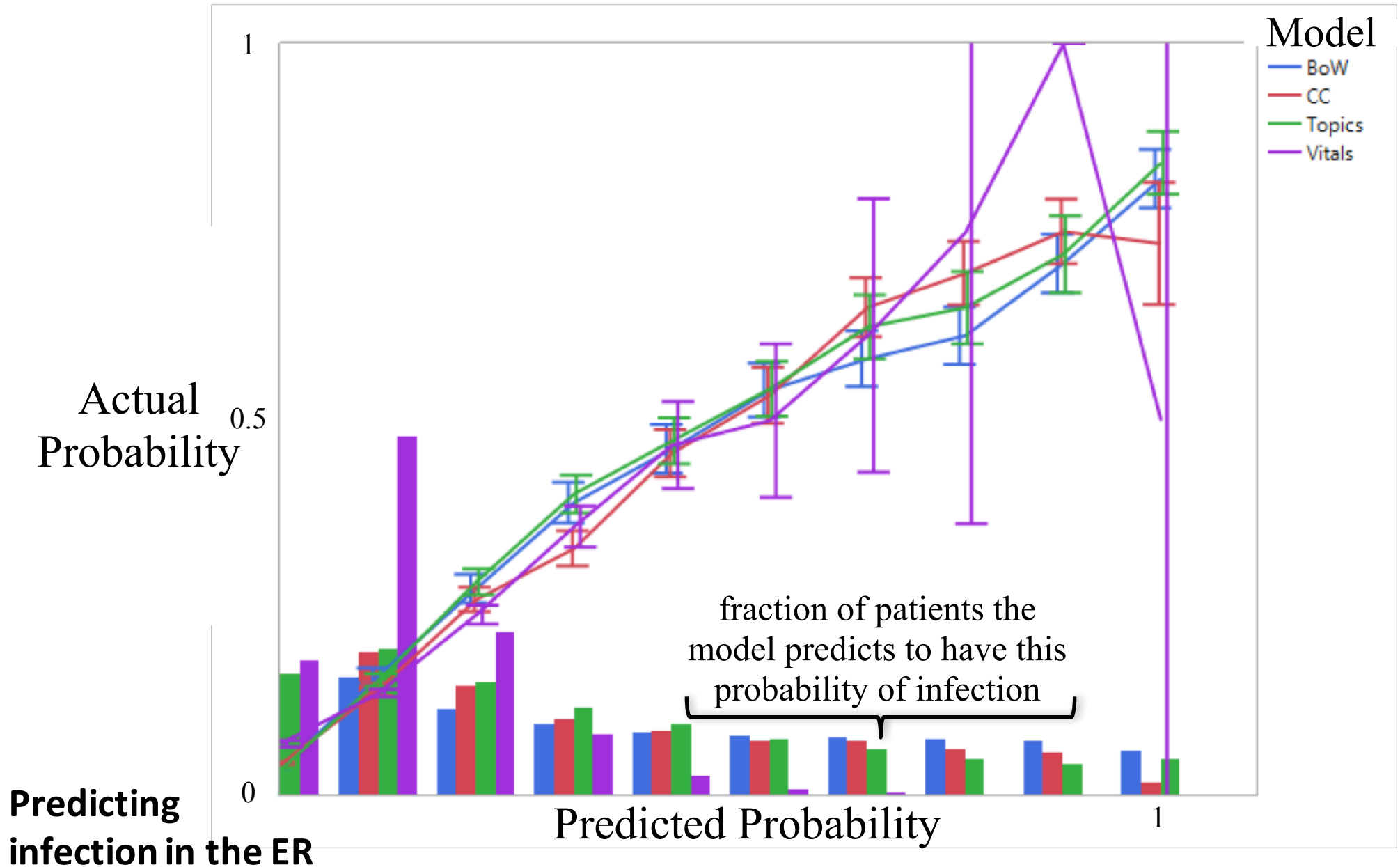
Top 100 Predictions

Top 1000 Predictions

Top 10000 Predictions

Diabetes 1-year gap

Calibration (*note: different dataset*)



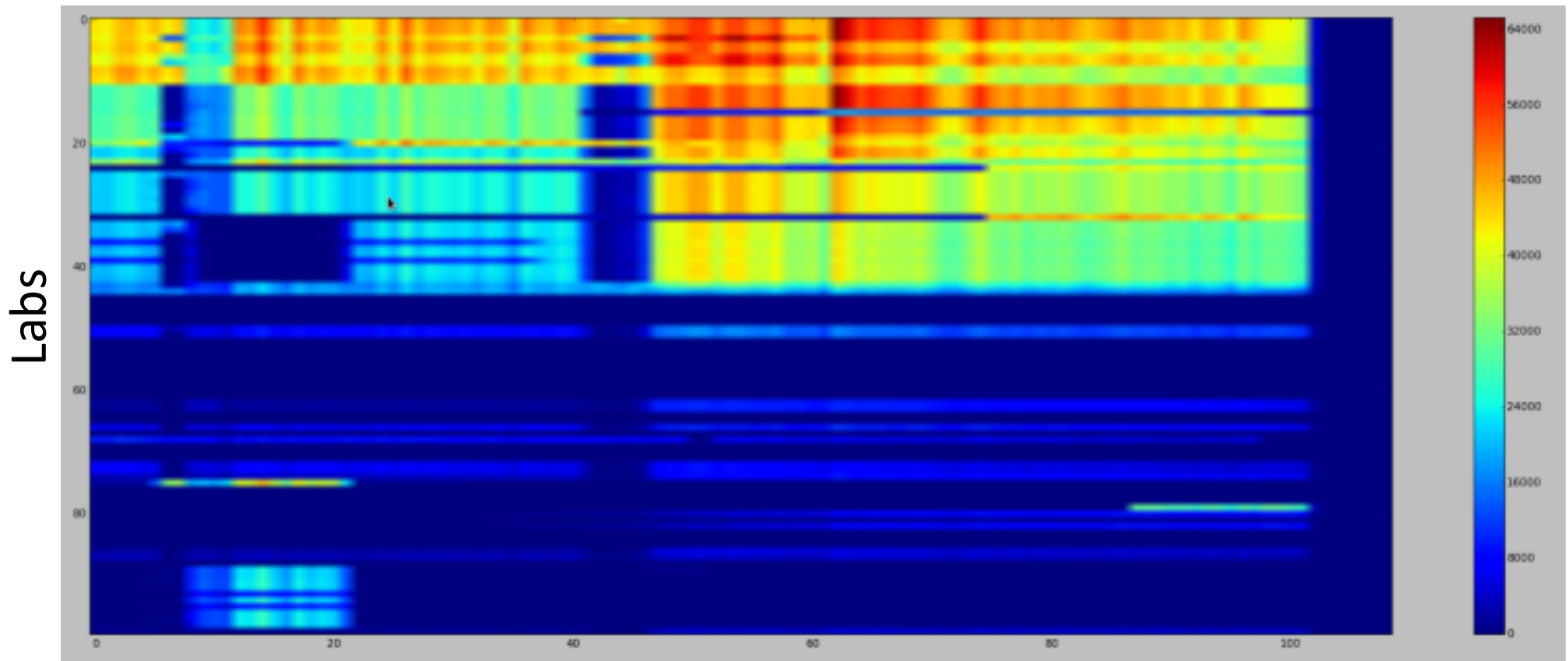
Outline for today's class

1. Risk stratification
2. Case study: Early detection of Type 2 diabetes
 - Framing as supervised learning problem
 - Deriving labels
 - Evaluating risk stratification algorithms
3. **Subtleties with ML-based risk stratification**
4. Discussion with Leonard D'Avolio (Assistant Professor at HMS, CEO @ Cyft)

Major challenge: non-stationarity

- ICD10 rolled out in 2015: predictive models learned using ICD9 features are no longer useful!
- Logistical issues => some features may not be available!
- Prevalence and significance of features may change over time
- Automatically derived labels may change meaning

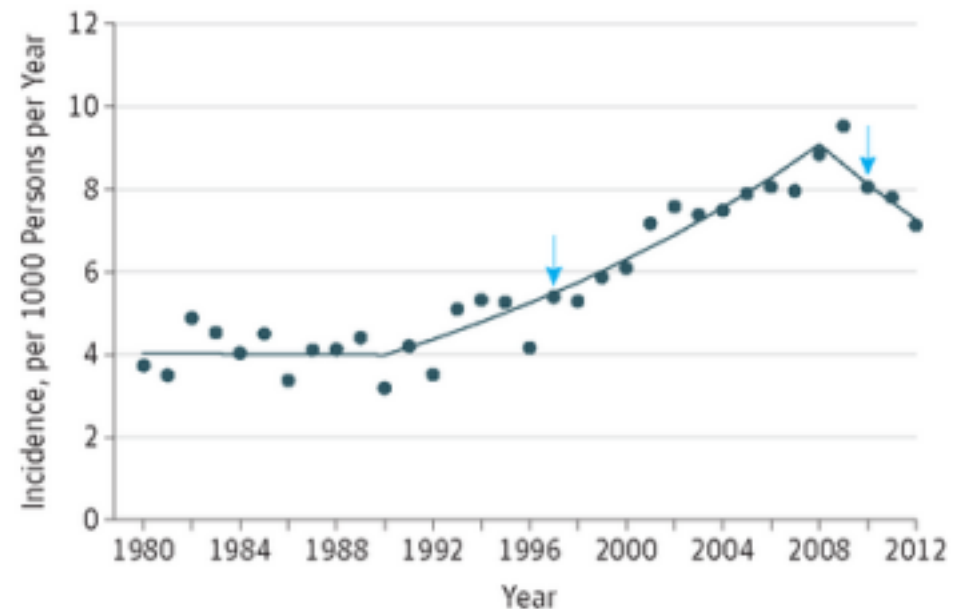
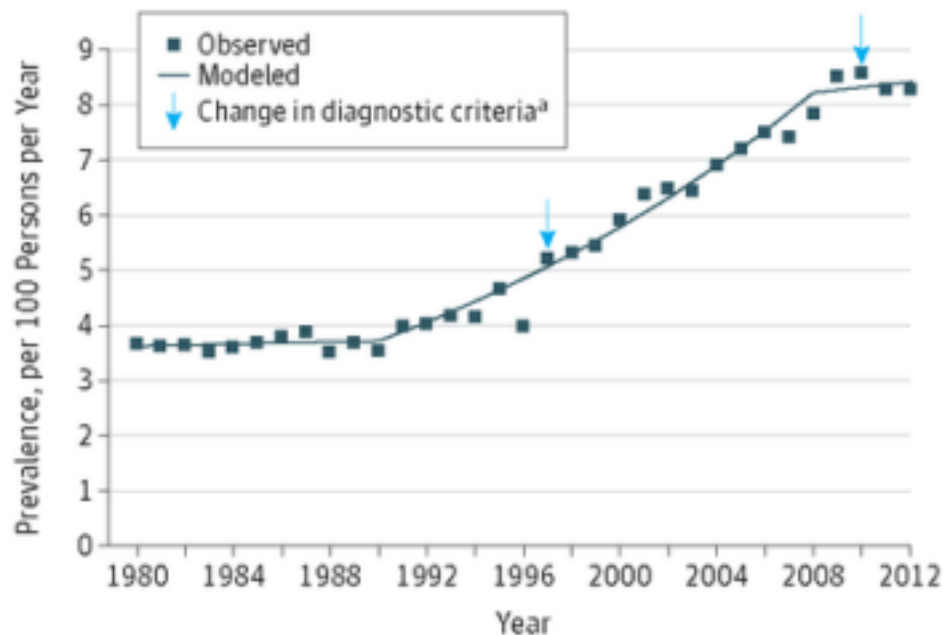
Top 100 lab measurements over time



Time (in months, from 1/2005 up to 1/2014)

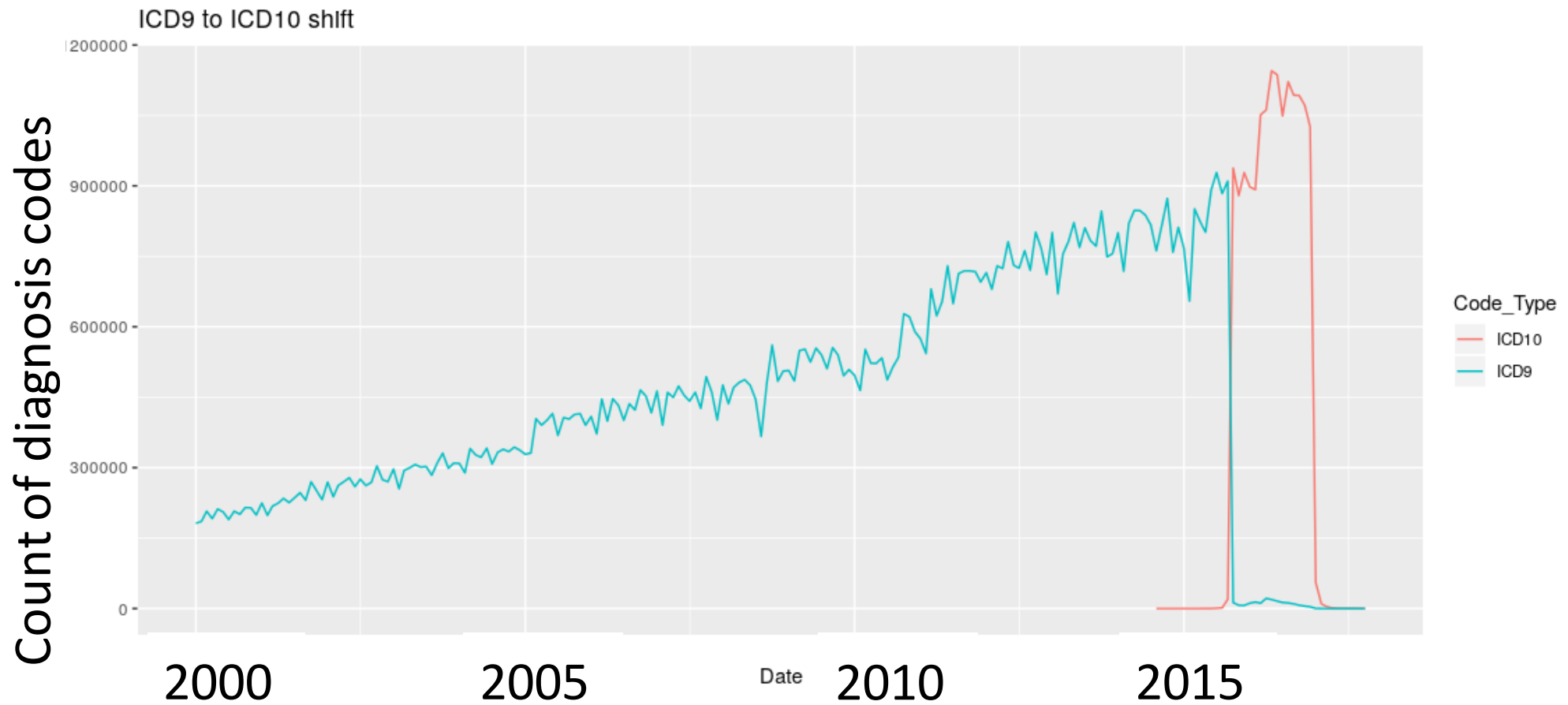
[Figure credit: Narges Razavian]

Diabetes Onset after 2009



Geiss LS, Wang J, Cheng YJ, et al. Prevalence and Incidence Trends for Diagnosed Diabetes Among Adults Aged 20 to 79 Years, United States, 1980-2012. *JAMA*. 2014;312(12):1218-1226.

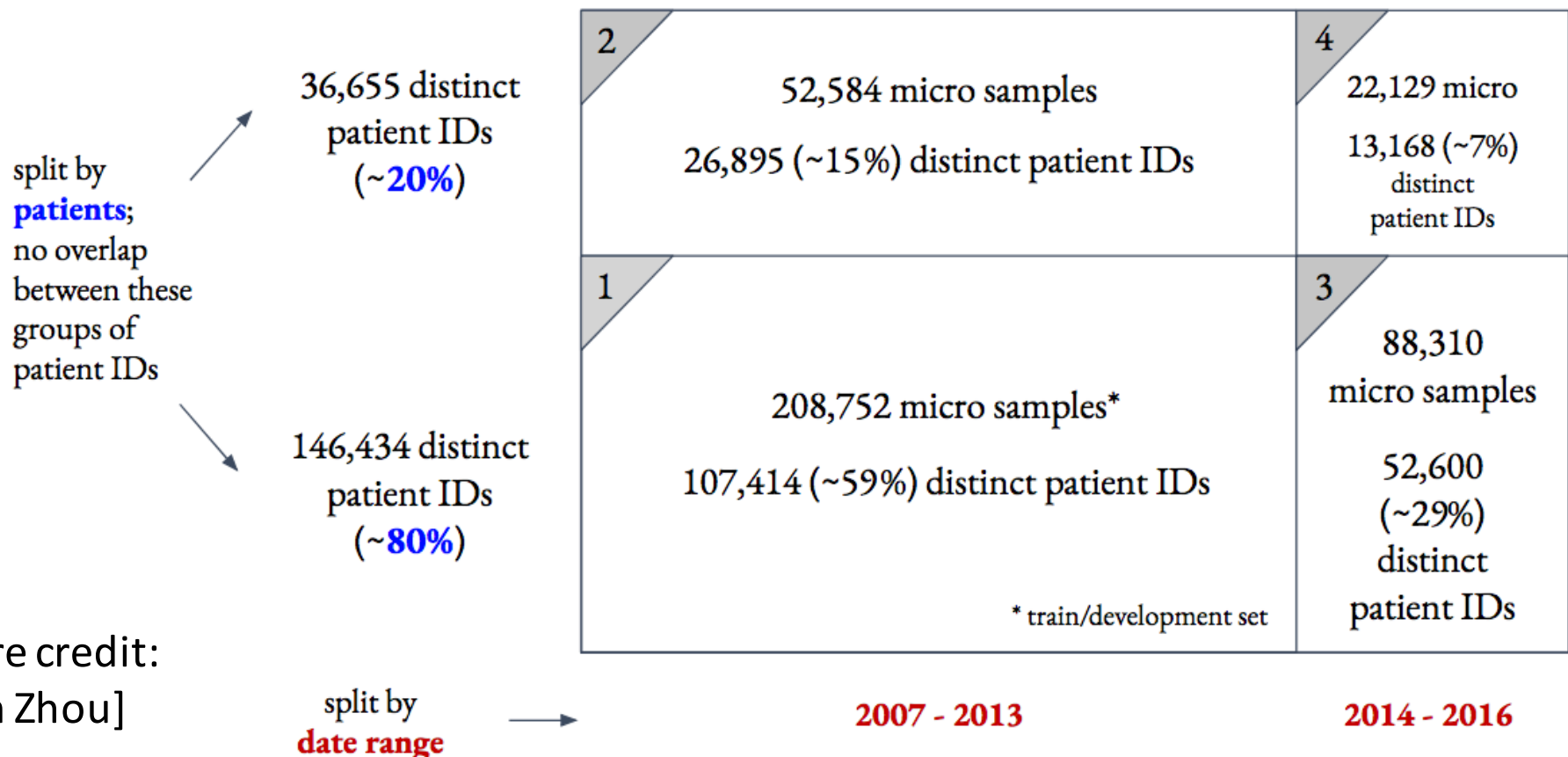
ICD-9 to ICD-10 shift



[Figure credit: Mike Oberst]

Re-thinking evaluation in the face of non-stationarity

- Given all this, do you notice anything wrong with how we evaluated the diabetes model?
- Good practice is to let the test data be from a future year:



[Figure credit:
Helen Zhou]

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