



# Prognostic and Predictive Classification Approaches for Disease Prediction Modeling

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**Chetan Paul**

Vice President – Technology & Innovation  
Leidos Civilian Health Services

## Highlights

- ▶ Advance Mission-solutions, research and outcomes for federal healthcare agencies (FDA, NIH, CDC, and CMS) through application of Emerging Technologies and HPC.
- ▶ Practitioner in Deep Learning, image processing, parallel computing GPU architectures for scientific and regulatory applications.
- ▶ Education: Strategic Leadership (Harvard), AI/ML Nanodegree (Stanford University), Masters Degree in Computers Science and Engineering
- ▶ Awards: “SASE” Technical Achievement, ACT-IAC Blockchain Innovation, G2Xchange Disruptive Tech, ACT-IAC Innovation, FDA Scientific Computing



**Dr. Ravichandran Sarangan**

Bioinformatics & Data Science Lead  
Leidos Civilian Health Services

## Highlights

- ▶ Bioinformatician and data scientist with extensive computing experience in analyzing and modeling public health and biomedical sciences data.
- ▶ Expert in developing statistical, Machine-Learning, and deep-learning models for high-dimensional Omics and health-focused (Real-world) data. Extensive experience collaborating and managing biomedical, and genetic diseases projects;
- ▶ Authored 54 peer-reviewed publications and a patent.
- ▶ Education: Ph.D. Computational Chemistry
- ▶ Awards: FNLCR Annual achievement team award; Annual performance award, ABCC; FDA Scientific Computing

# The Quest to Predict the “Future”

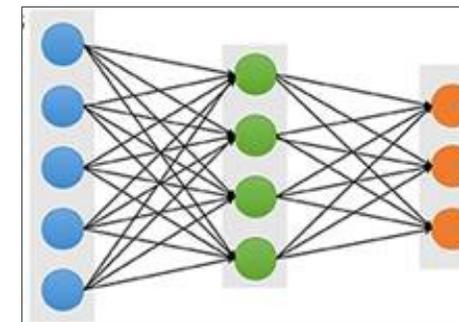
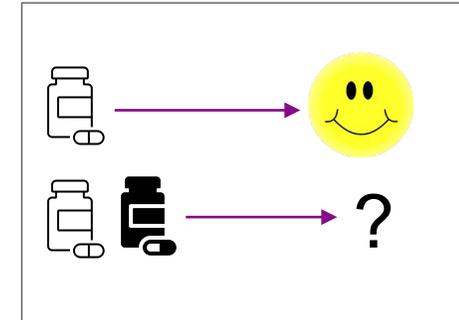


The *Palantiri*, also known as the **Seven Seeing-stones**, were used for intelligence gathering and could show visions of the future

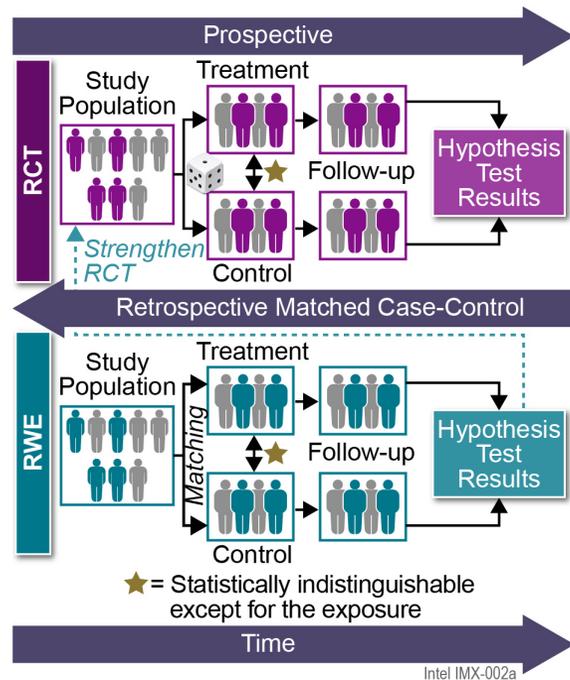


Prognostic and Predictive Modeling methods and approaches allow an estimation of future events which one can make by incorporating and casting forward data related to the past in a pre-determined and systematic manner

- ▶ **Causal Inference (Drug Repurposing Study)**
  - Population-based; sub-population-based
- ▶ **Prognostic Modeling (Long COVID Study)**
  - Population (group) based: Survival Analysis
  - Patient focused: Hazard Modeling; Random Survival Forest
- ▶ **Data Harmonization/Quality**
- ▶ **Predictive modeling pipeline**
- ▶ **Synthetic Data**
- ▶ **Deep-Learning Approaches and Transfer Learning**
- ▶ **AI Trust and Explainability**

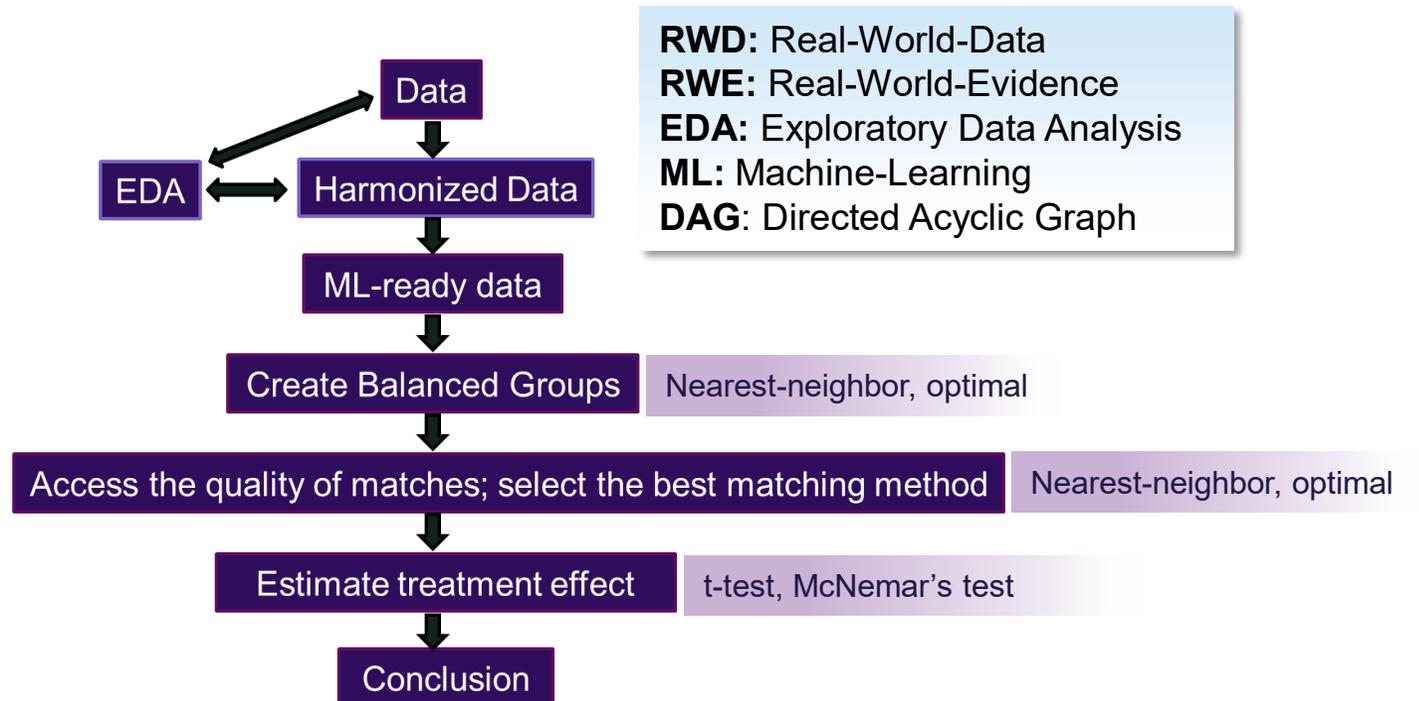


Using Real-World-Data (RWD), develop in-silico models for rapidly identifying repurposed drugs that can lower the risk of death due to disease/infection.



Areas of focus	Solution
Identify study population, randomization, and removing confounding	Inclusion-exclusion, matching, and DAG

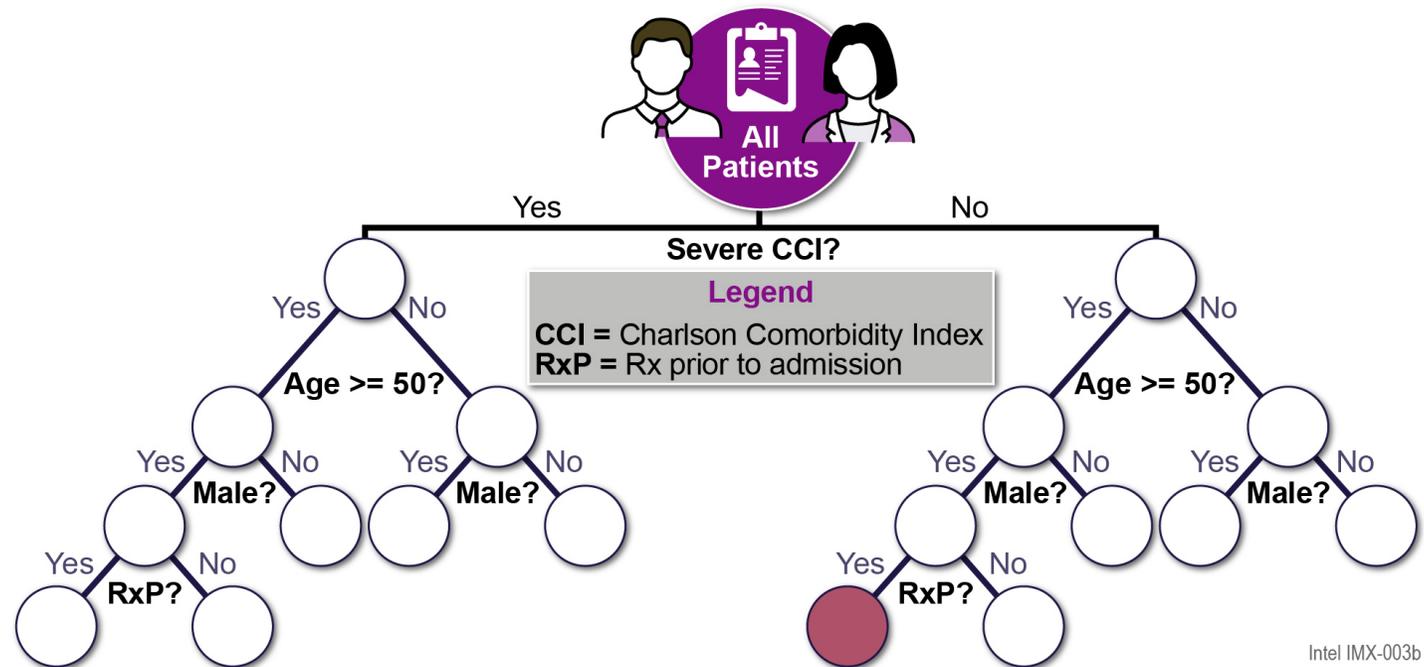
Leidos process of Simulating in-silico Randomized Clinical Trials Using Real-World-Data (RWD) to Model Causal Effect of Repurposed Drugs



# Subpopulation Based Association

Using Real-World-Data (RWD), develop in-silico models for rapidly the efficacy of drugs in subpopulations

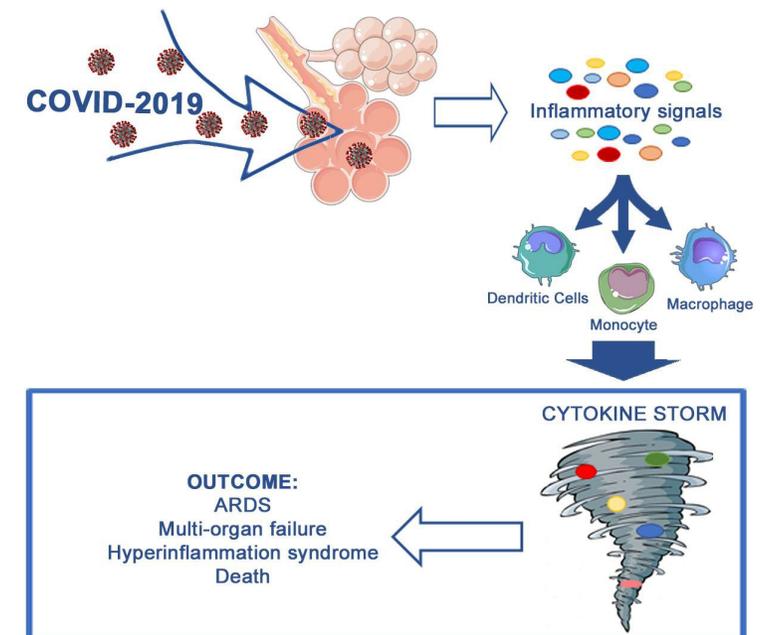
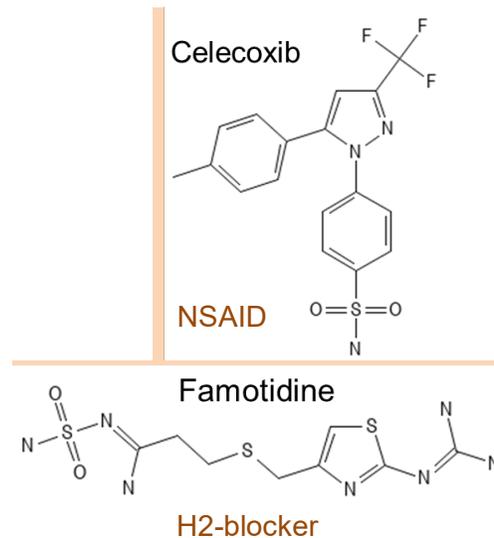
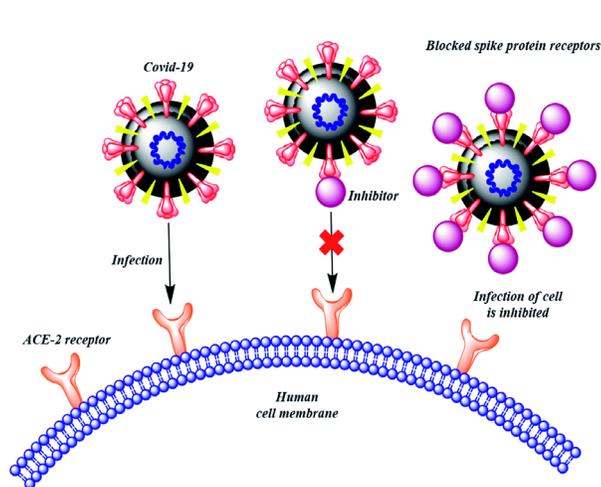
At each node of the tree, the population was split into two groups, and a two-sample proportion test was performed to determine whether any significant differences exist between patients who took medication A versus those who took medication B.



Intel IMX-003b

# In-silico Simulated Randomized Clinical Trials Using Real-World-Data (RWD) to Model Causal Effect of Repurposed Drugs Study

- ▶ Using Real-World-Data (RWD), develop in-silico models for rapidly identifying repurposed drugs that can lower the risk of death due to Sars-CoV-2 infection
- ▶ Risk of death due to COVID-19 is predominantly due to hyperactive host inflammatory responses resulting from infection



# COVID-19 Real World Data (RWD)



## Charge Data Master (CDM)

CDM In-patient and out-patient having COVID19 diagnoses from 6/1/2020 - 1/31/2021

### CDMVisitLocationFact

DISCHG\_MONTH\_ID  
VST\_ID  
DAY\_IN\_VST\_NBR  
LOCATION\_ID  
LOCATION\_DESC

### CDMFacility

FCLT\_ID  
REGION\_NM  
RURAL\_URBAN\_CD  
TCHG\_IND  
BED\_SIZE\_DESC

### Procedure

PRC\_CD  
PRC\_VERS\_TYP\_ID  
PRC\_TYP\_CD  
PRC\_SHORT\_DESC  
PRC\_DESC

### Patient

PATIENT\_ID  
PAT\_BRTH\_YR\_NBR  
PAT\_GENDER\_CD

### Diagnosis

DIAG\_CD  
DIAG\_VERS\_TYP\_ID  
DIAG\_SHORT\_DESC  
DIAG\_DESC

### CDMVisitDiagnosisFact

DISCHG\_MONTH\_ID  
VST\_ID  
DIAG\_CD  
DIAG\_VERS\_TYP\_ID  
PRI\_DIAG\_IND

### CDMVisitFact

DISCHG\_MONTH\_ID  
VST\_ID  
PATIENT\_ID  
ADMT\_CATG\_DESC  
ADMT\_DIAG\_CD  
ADMIT\_DIAG\_VERS\_TYP\_ID  
DISCHG\_STATUS\_DESC  
TOTAL\_CRG\_AMT  
ADMT\_DT  
FCLT\_ID  
IP\_OP\_IND  
PAY\_TYP\_DESC  
DISCHG\_DT

### CDMVisitResourceFact

DISCHG\_MONTH  
VST\_ID  
SVC\_DT  
REV\_CD  
BILLG\_DESC  
PRC\_CD  
PRC\_VERS\_TYP\_ID  
RSRC\_QTY  
RSRC\_TOTAL\_CRG\_AMT

## Longitudinal Rx (LRx)

Rx data for 120-day lookback from each patient's earliest CDM visit with a COVID-19 diagnosis

### RxFact

MONTH\_ID  
SVC\_DT  
PATIENT\_ID  
CHNL\_CD  
CLAIM\_ID  
RX\_TYP\_CD  
PROVIDER\_ID  
PAY\_TYP\_DESC  
PRODUCT\_ID  
AUTH\_RFLL\_NBR  
DSPNSD\_QTY  
DAYS\_SUPPLY\_CNT

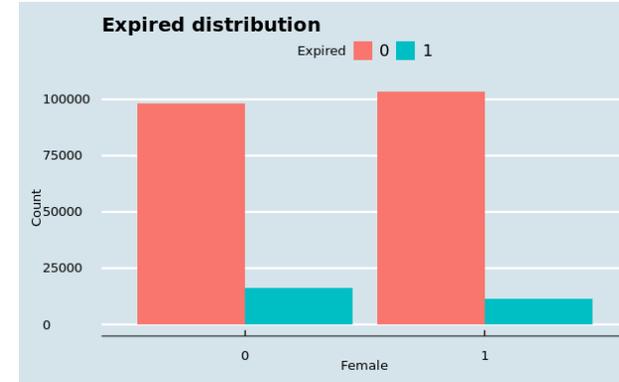
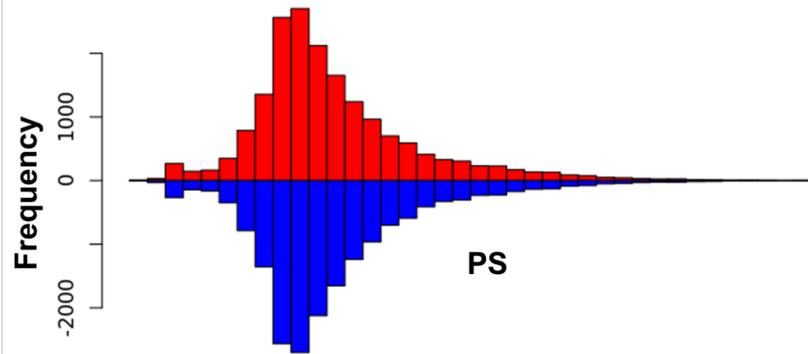
### RxProduct

PRODUCT\_ID  
NDC\_CD  
MKTED\_PROD\_NM  
STRNT\_DESC  
DOSAGE\_FORM\_NM  
LBLER\_NM  
LBLER\_TYP\_CD

CDM and LRx Data  
Source: IQVIA Inc

# Results and Findings

Propensity Score (PS), Red: Treatment vs Blue: No-Treatment



Results shown for Famotidine; Celecoxib results are similar

$$SMD = \frac{\bar{x}_{treatment} - \bar{x}_{control}}{\sqrt{\frac{s_{treatment}^2 + s_{control}^2}{2}}}$$

## McNemar's Exact Test Results

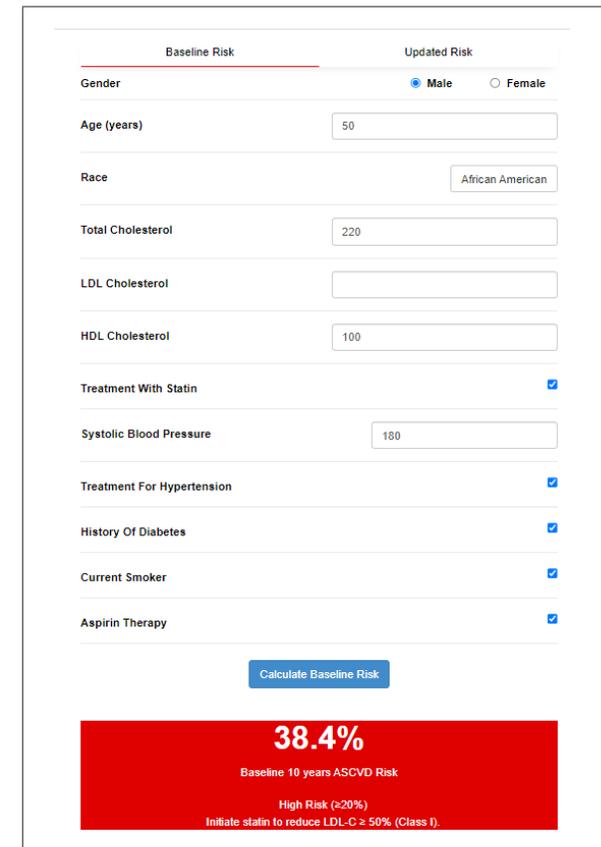
Run	Celecoxib				Famotidine			
	N	OR	CI (95%)	P-value	N	OR	CI (95%)	P-value
1	1013	2.3870	1.5498, 3.7573	3.276e-05	17916	2.400	2.2254, 2.5898	< 2.2e-16
2	999	4.5882	2.6903, 8.2730	1.642e-10	17892	2.5143	2.3304, 2.7145	< 2.2e-16
3	1019	2.0000	1.3148, 3.0927	8.200e-04	17622	2.5978	2.4045, 2.8085	< 2.2e-16
4	1026	2.3636	1.5545, 3.6669	2.326e-05	17897	2.4851	2.3029, 2.6833	< 2.2e-16
5	1046	2.4838	1.6175, 3.9002	1.115e-05	17916	2.5967	2.4056, 2.8050	< 2.2e-16

Our matched case-control study results for both treatment options Celecoxib and Famotidine show Odds Ratio OR > 1 indicating that the Famotidine and Celecoxib did not provide protective effects for COVID-19 patients. Please note that conclusions need more strengthening with follow-up studies

# Prognostic Modeling?

- ▶ **What is Prognostic Modeling?**
  - Predicting risk of a future event
    - Heart attack, death
- ▶ **Applications:**
  - Useful for finding the survival with a disease (ex., brain tumor)
  - What is the risk of a disease?
  - Effective for creating treatment guidance
    - Who is eligible for end-of-life care

## 2018 Prevention Guidelines Tool CV risk calculator (American Heart Association)



The screenshot shows a web-based risk calculator interface. It has two tabs: 'Baseline Risk' (selected) and 'Updated Risk'. The form includes the following fields and options:

- Gender:  Male,  Female
- Age (years): 50
- Race: African American
- Total Cholesterol: 220
- LDL Cholesterol: (empty)
- HDL Cholesterol: 100
- Treatment With Statin:
- Systolic Blood Pressure: 180
- Treatment For Hypertension:
- History Of Diabetes:
- Current Smoker:
- Aspirin Therapy:

A 'Calculate Baseline Risk' button is located below the form. The result is displayed in a red box: **38.4%** Baseline 10 years ASCVD Risk. Below this, it states 'High Risk (≥20%)' and 'Initiate statin to reduce LDL-C ≥ 50% (Class I)'.

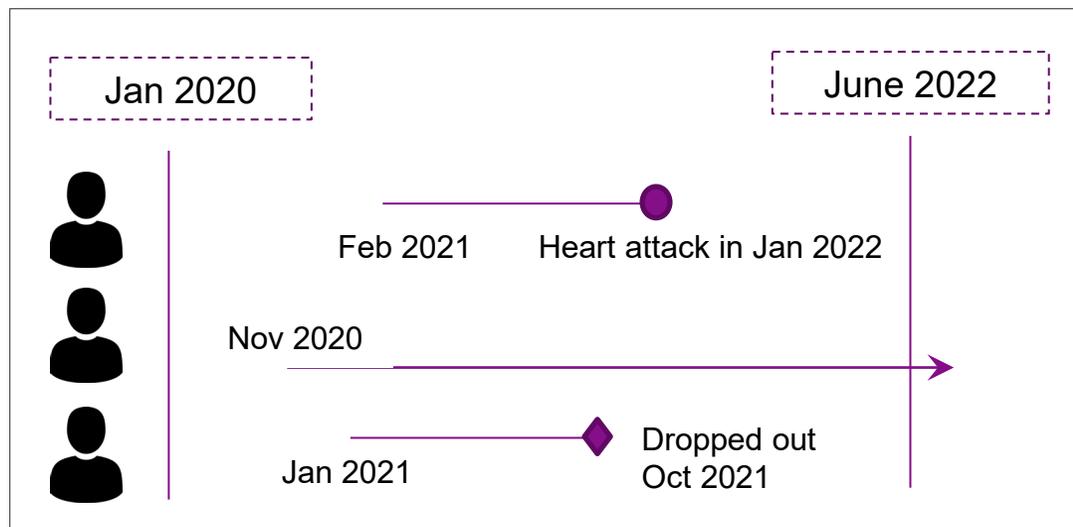
<http://static.heart.org/riskcalc/app/index.html#!/baseline-risk>

## ► Survival Models

- Computes the probability of survival past any time 't'
- Group based analysis
  - Ex., stage-1 cancer vs stage-4 cancer patients' survival
  - Ex.,  $P(\text{time to death} > 2 \text{ years}) = 0.8$

## ► Evaluating models

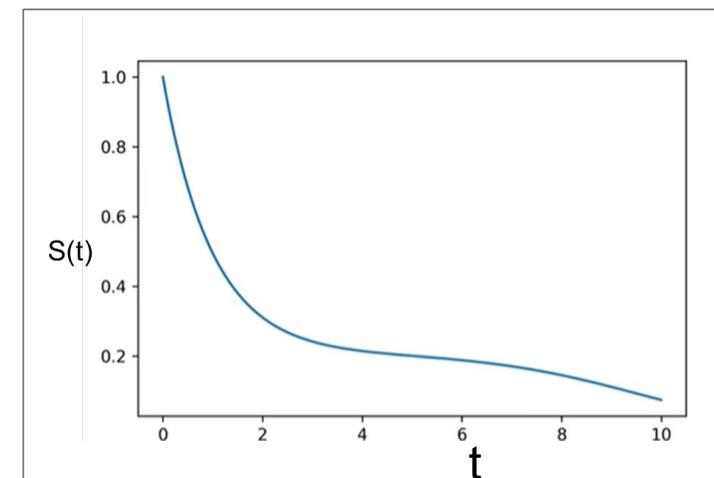
- C-index (Concordance index)



+ indicates censored patients

i	$T_i$	Event
1	10	1
2	55+	0
3	20	1
4	15+	0
5	30+	0
...	...	...

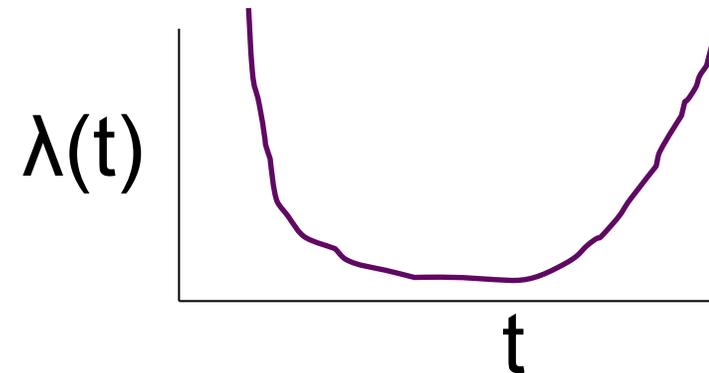
$S(t) = Pr(T > t)$ ; T is the time to an event



► **Cox Proportional Hazard Model:**

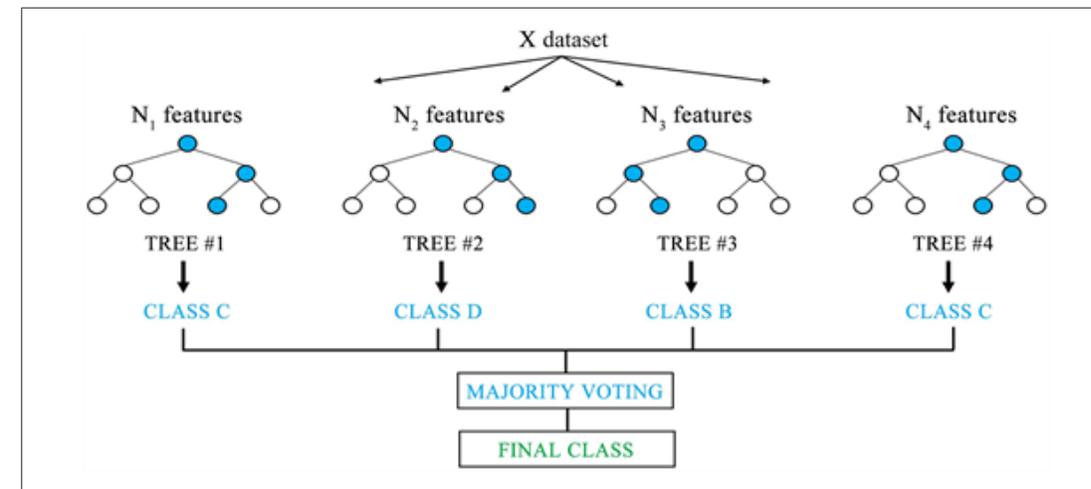
- Popular choice for right-censored time-to-event data
- Individual patient prediction

*Hazard Function:  $H(t)$  or  $\lambda(t) = \Pr(T=t \mid T \geq t)$ ;  $T$  is the time to an event*



► **Random Survival Forest:**

- Alternative approach to Cox Proportional Hazard Models
- Tree-based method for analysis of right censored time-to-event data
- Extension of Random Forest
- Approach to model subpopulations

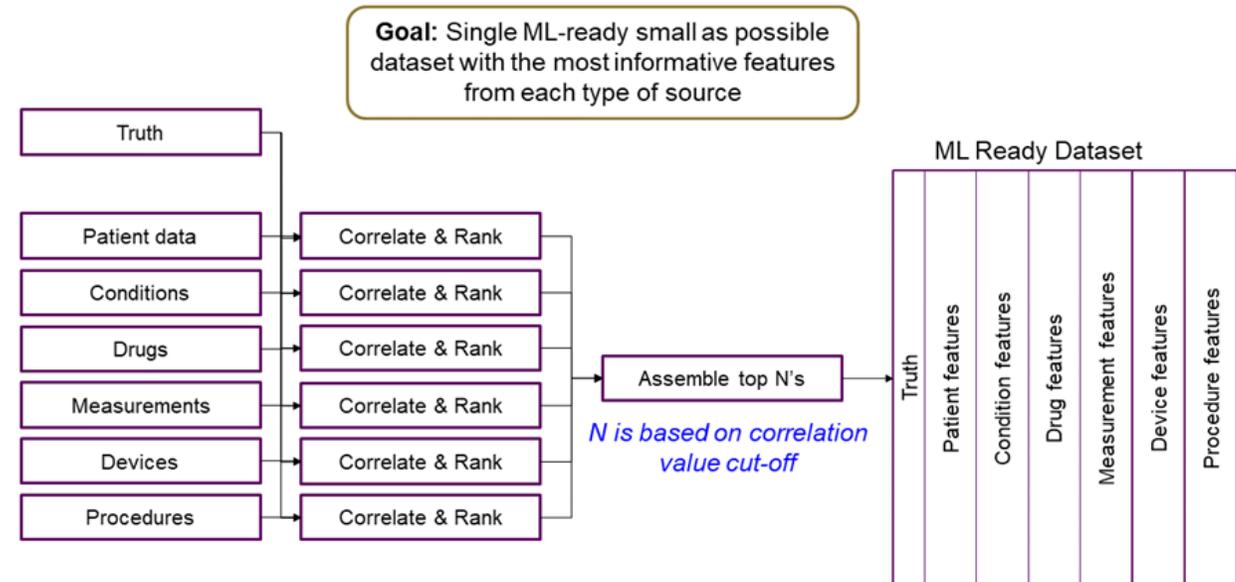


## **Predicting the likelihood of developing Long COVID is essential to identify at-risk patients and to provide timely treatment options**

- ▶ **Data:** The EHR data contained COVID-19 infected patients with information such as demographics, procedures, medical conditions, physical measurements, lab results, and many more factors. The dataset contains more than 15 million patients, including more than 5 million COVID-19 infected patients, and over 17.5 billion rows of raw data (Source: N3C)
- ▶ **Assumptions:** Data missing 'time-to-event'  $T_i$  for censored patients. Couldn't determine if these patients left the study before the end or survived until the end of the study. Because majority of the population was censored ( $> 80\%$ ), chose to retain by assigning a maximum value for time-to-event  $T_i$ .

# Feature Engineering & Selection for High-Dimensional Real-World Data

- ▶ Correlations of each feature with the presence of outcome were tabulated and ranked from most to least informative.
- ▶ Approach provided substantial improvement to prognostic and predictive classification models.
- ▶ Applications
  - Reduce dimensionality, especially for high-dimensional Real-World-Data

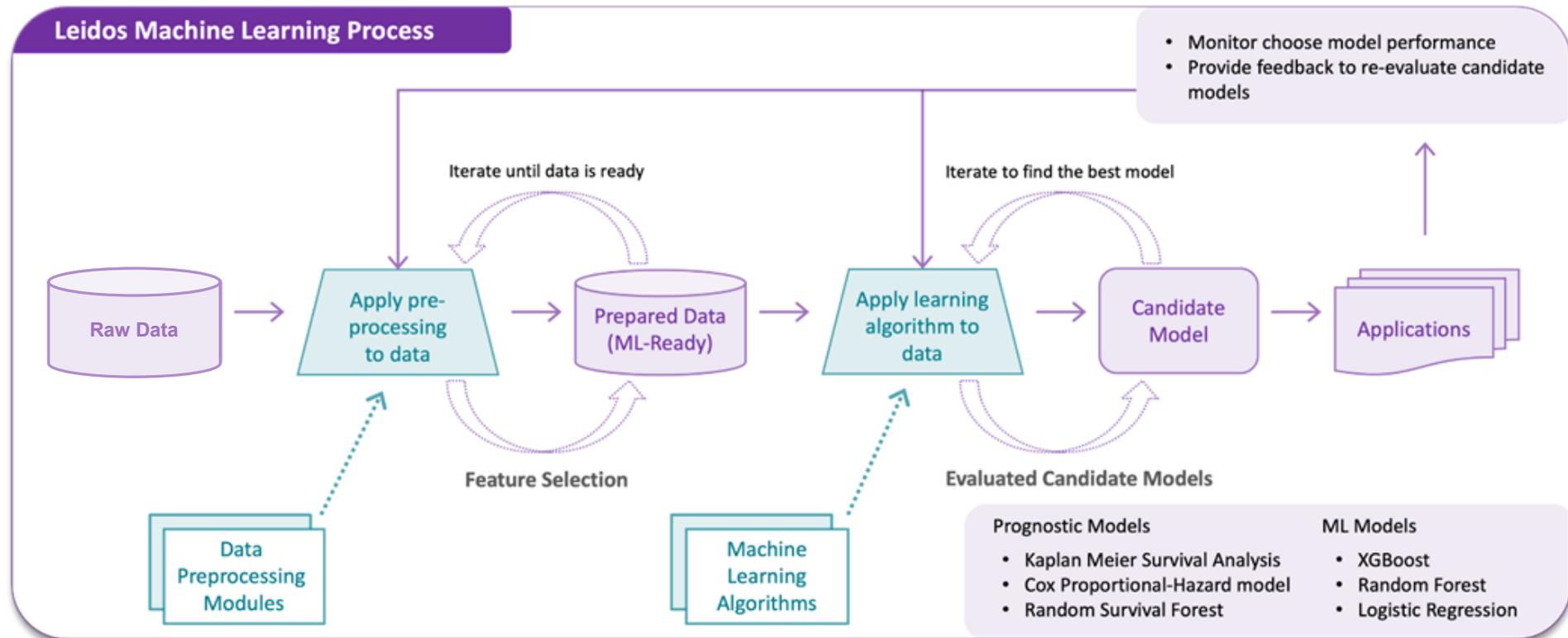


## Correlation method (binary-binary)

- Tetrachoric correlation
- Based on Chi-squared
- Theil's U

# Modeling Methodology Pipeline

- Includes Feature Selection and Modeling (Prognostic and Machine-Learning models)



# Prognostic Modeling: Long COVID prediction

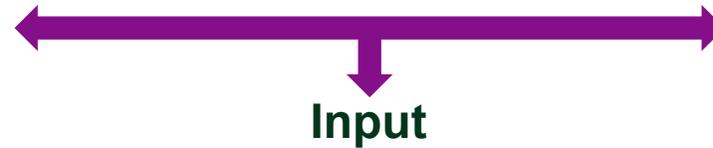
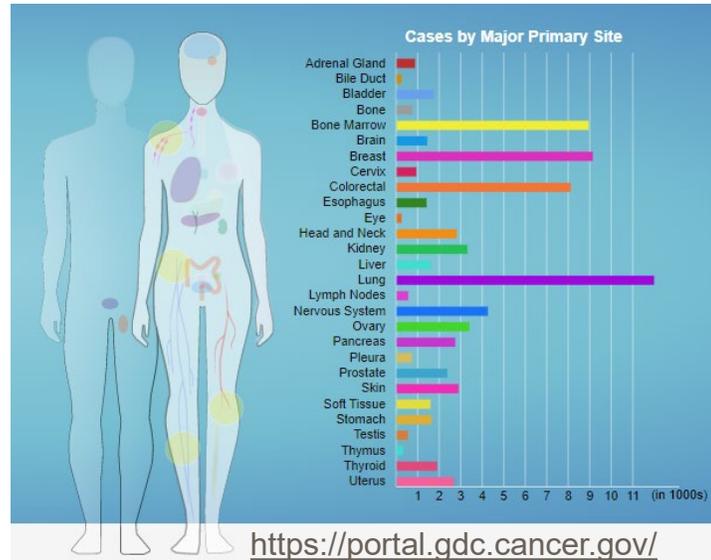


## ► Results and Findings

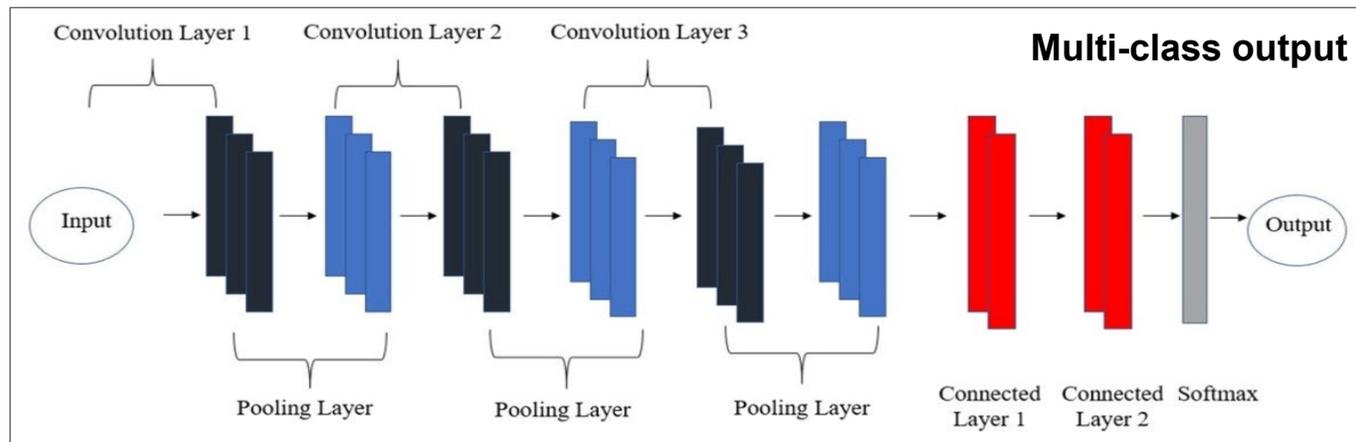
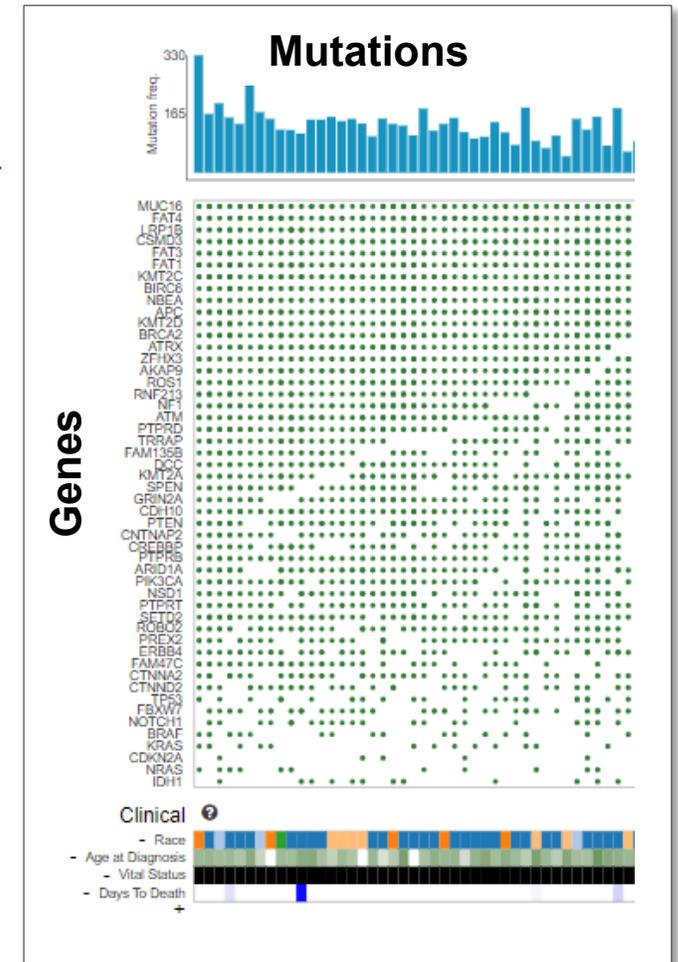
- Performance analysis of predictive classification models showed XGBoost as best with AUC = 0.93 and a balanced accuracy of 0.75.
- Feature selection based on measuring feature power provided substantial improvement to prognostic and predictive classification models.
- The most powerful features had correlations with Long COVID diagnoses as great as 0.45 and were substantially better than the correlations of 0.15 obtained on the original feature set

Models	Performance	
	AUC	Other
XGBoost	0.9260	-
XGBoost_upsampling	0.9236	-
XGBoost_downsampling	0.9104	-
RandomForest	0.9114	-
Logistic Regression	0.8500	-
Odds Ratio (exposed: gender)	-	OR(CI): 1.27 (1.21, 1.33)
Kaplan-Meier (log-rank test; gender group test)	-	p-value: 3.835e-23
Cox Proportional Hazard	-	Harrell C-index: 0.7939
SurvivalRandomForest	-	Harrell C-index: 0.8624

# Genomic Expression Data Modeling Study

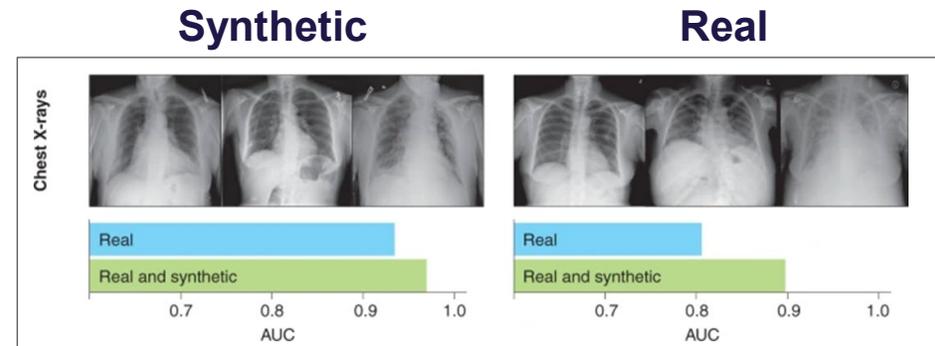


<https://portal.gdc.cancer.gov/>

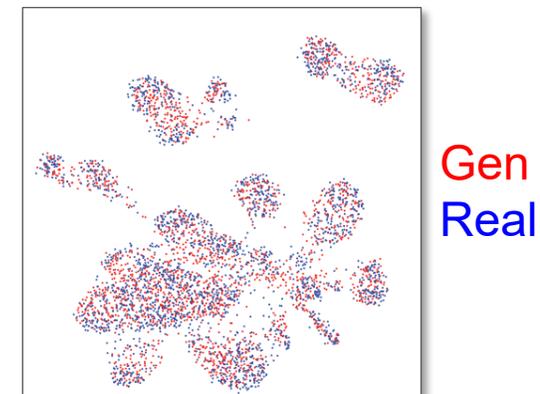


# Synthetic Data & Synthea

- ▶ Having access to good-quality Healthcare data is a bottle-neck for research, training, and technology development
- ▶ The scientific community is struggling to strike a balance between the competing interests of protecting patient privacy/confidentiality and making data public
- ▶ Can we generate/share realistic synthetic healthcare data that statistically reflects the concerned population, and protecting patient privacy and confidentiality?



<https://www.nature.com/articles/s41551-021-00751-8>



doi: 10.1093/bioinformatics/btab035

- ▶ Synthetic data can be used as a test for Bias detection
- ▶ It is the data and not the algorithm that is biased
- ▶ If we want to eliminate bias from our AI systems, then we need to remove the bias from the data before we use it to build models
- ▶ Bias mitigation strategies:
  - Before training: rebalance the data by collecting representative datasets (not easy);
  - During training: Data augmentation, adversarial training
  - After training: Model outcomes can be post-processed based on sub-groups

<https://www.nature.com/articles/s43856-021-00028-w>;

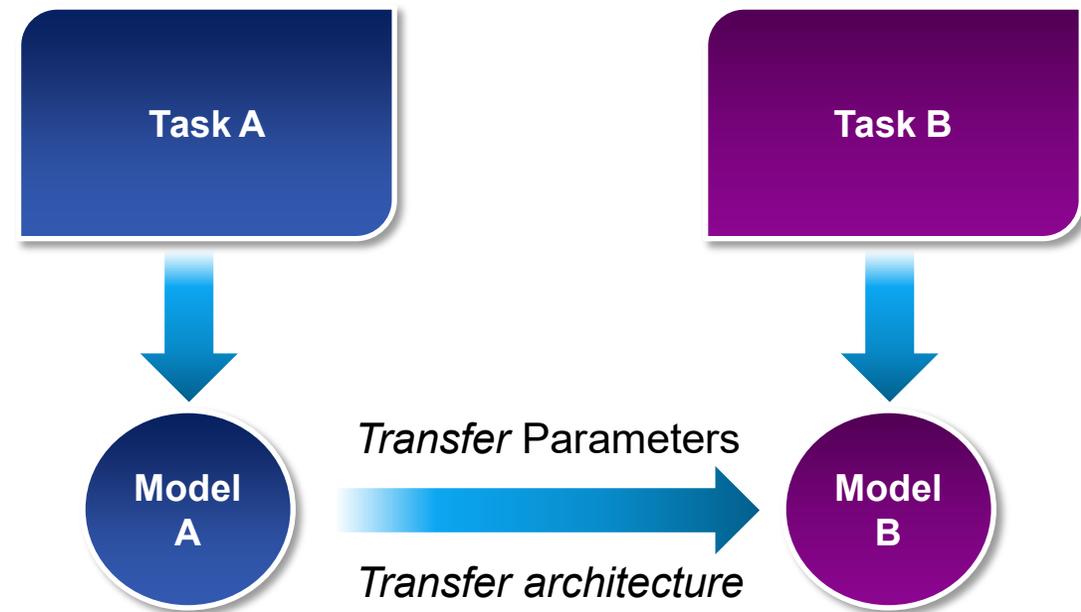
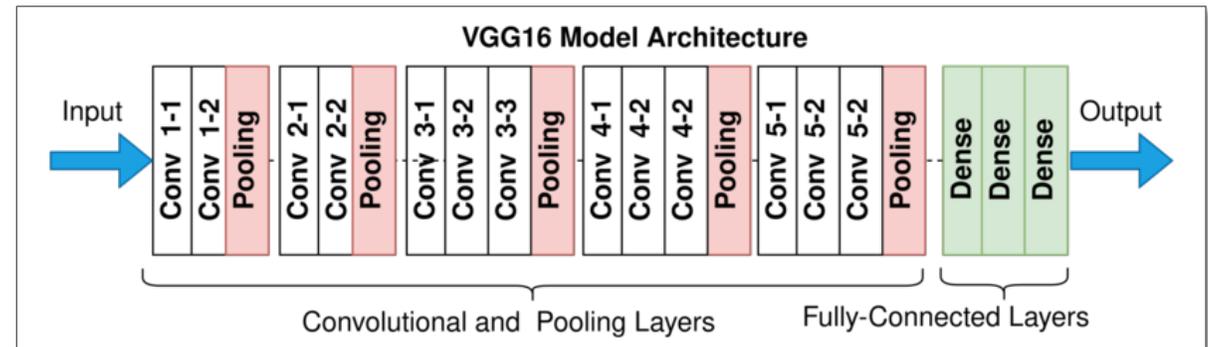
<https://www.nature.com/articles/s41467-022-32186-3>

<https://github.com/synthetichealth/synthea>

# Transfer Learning

- ▶ We could use image (Chest X-ray) dataset and train a CNN to classify a presence or absence of disease (ex., pneumonia).
- ▶ This model predictions can help human radiologist to speed up the predictions.
- ▶ One could take the technology and submit it to the FDA for 510(k) clearance as Software as a Medical Device.

## CNN: Convolutional Neural Network



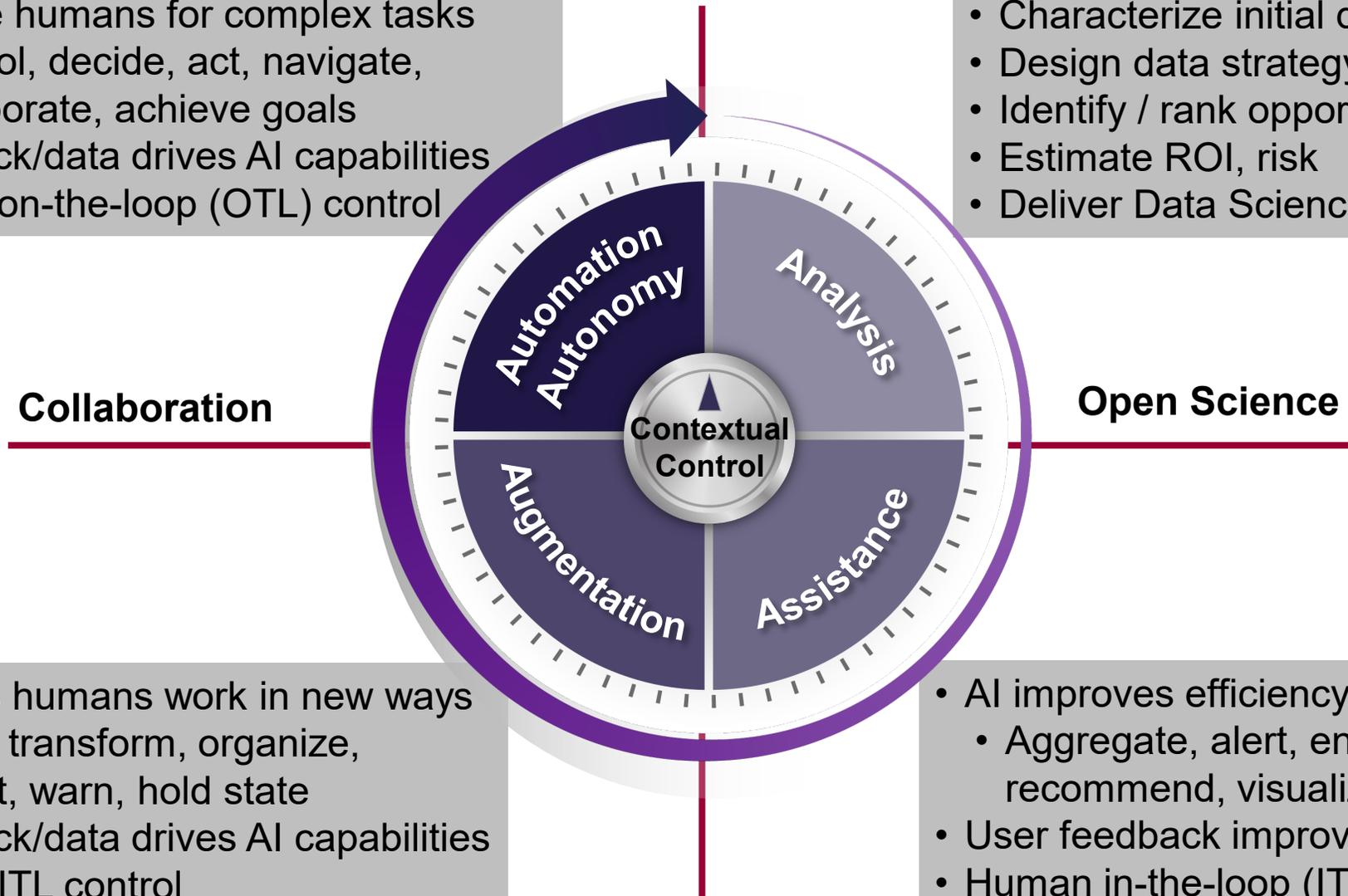
- ▶ How to handle confounding due to unobserved features? (causal modeling)
- ▶ Best method to combine RCTs and observational data? (causal modeling)
- ▶ Patient self-selection is a problem in observational studies. How to control this issue?
- ▶ Feature selection and dimensionality reduction in high-dimensional RWD
- ▶ Bias detection modeling/evaluation for healthcare data is complex. If the prevalence of the target disease is different between different groups, then what `selection fairness condition` would be appropriate?

- ▶ What is the best Deep-Learning architecture?
- ▶ Create Interpretable Neural Network models? (Explainability)
- ▶ Data normalizations for biological expression data?
- ▶ Model repositories, storage/retrieval? (Sharing Models, Outcomes, Collaboration)

# 4A Methodology for AI Trust and Explainability

- Replace humans for complex tasks
  - Control, decide, act, navigate, collaborate, achieve goals
- Feedback/data drives AI capabilities
- Human on-the-loop (OTL) control

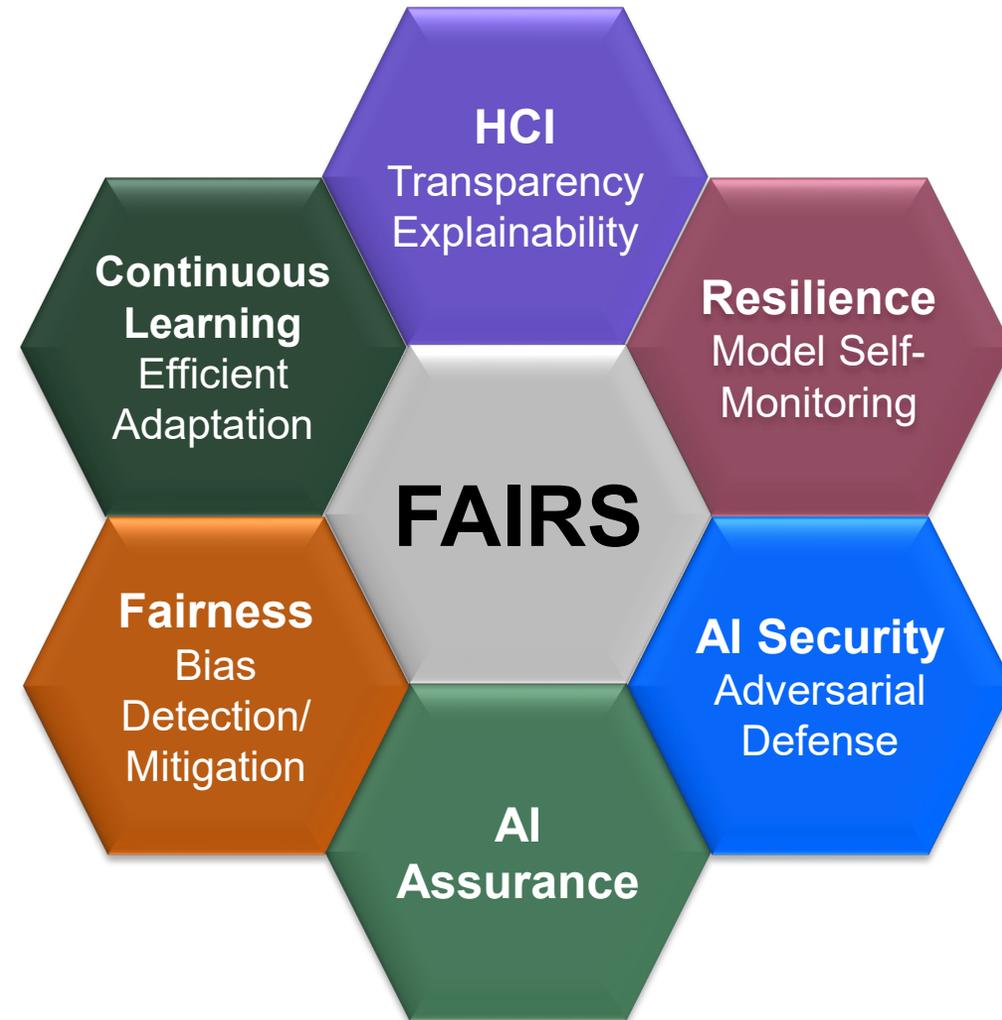
- Characterize initial data
- Design data strategy
- Identify / rank opportunities for AI
- Estimate ROI, risk
- Deliver Data Science capabilities



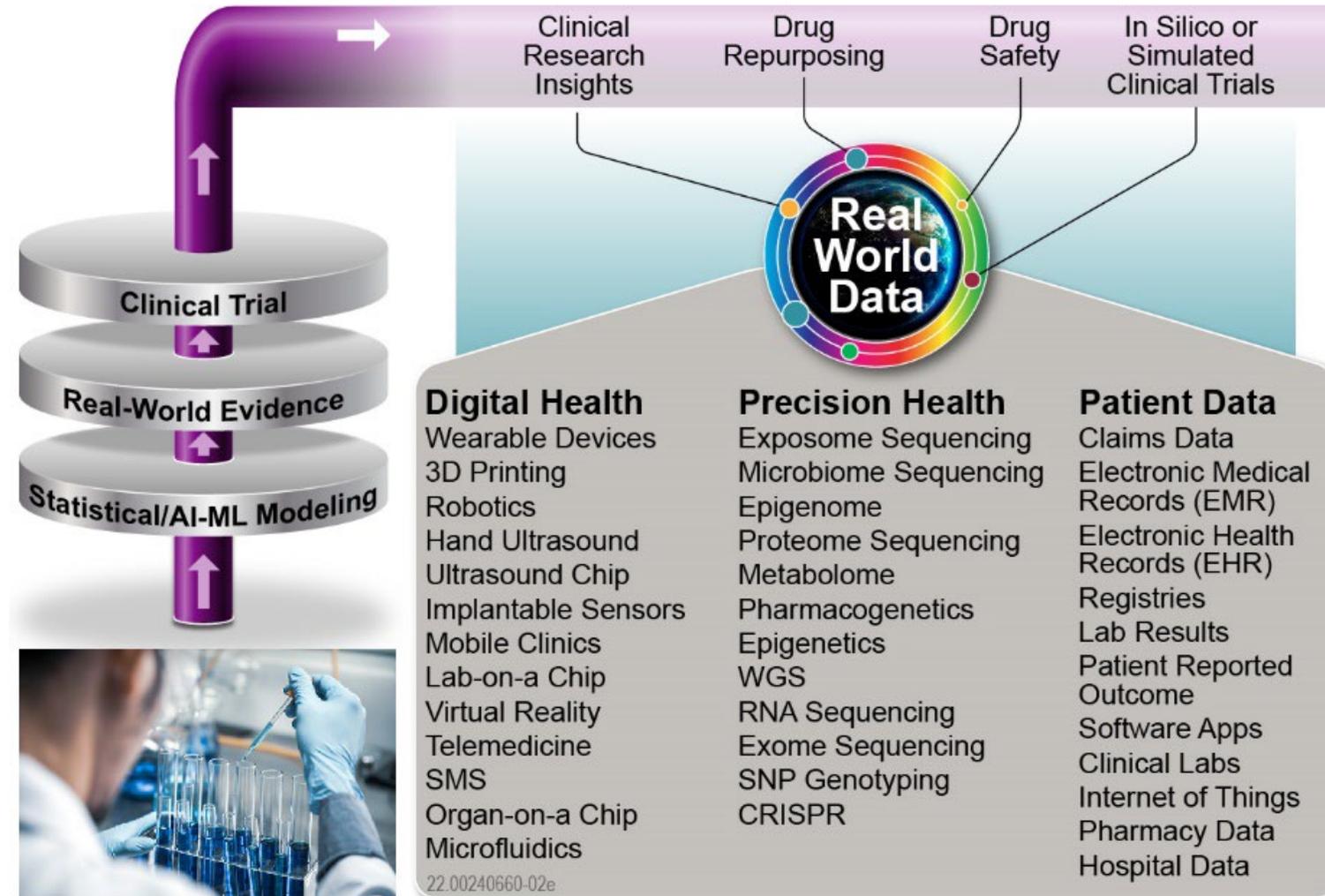
- AI helps humans work in new ways
  - Track, transform, organize, predict, warn, hold state
- Feedback/data drives AI capabilities
- Human ITL control

- AI improves efficiency of workflows
  - Aggregate, alert, enrich, recommend, visualize
- User feedback improves accuracy
- Human in-the-loop (ITL) control

- **FAIRS** provides a broad set of complementary and mutually reinforcing capabilities for AI Trust in a unified framework
- **FAIRS** components packaged as microservices, allow flexible deployment, configuration with speed, scale and security



# Future Directions



# Contact Info

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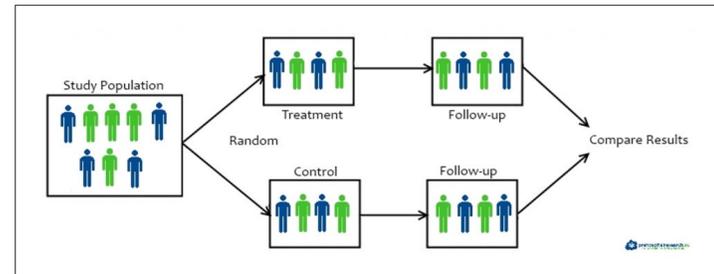
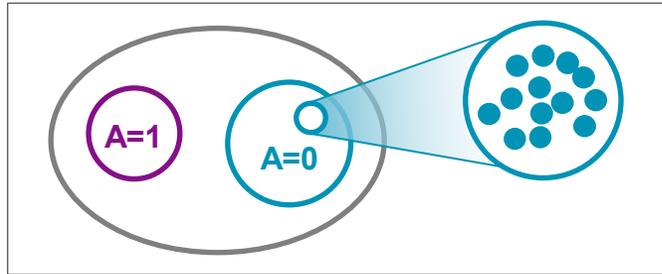
Dr. Ravichandran Sarangan

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# Backup Slides

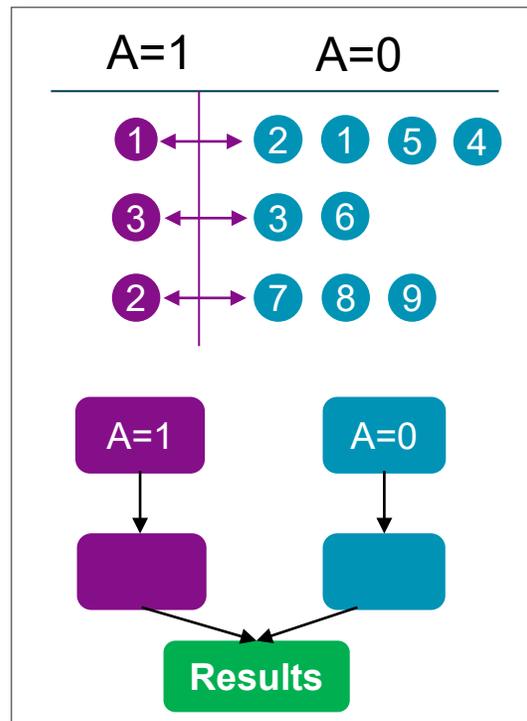
# RCT vs RWE



This Photo by Unknown  
Author is licensed under [CC BY](#)

← Time

→ Time



Id	Age	Female	BC	CC
1	36	1	1	1
2	37	1	1	0
3	36	0	0	0

	1	2	3
1	0	$\sqrt{2}$	$\sqrt{3}$
2		0	$\sqrt{3}$
3			0

Distance →  
Matching using  
Nearest Neighbor

**Treated**

Weight:

$$\frac{1}{P(A=1|X=1)} = \frac{1}{0.1} = 10$$

**Control**

Weight:

$$\frac{1}{P(A=0|X=1)} = \frac{1}{0.9} = \frac{10}{9}$$

Inverse Probability  
of Treatment  
Weighting (IPTW)

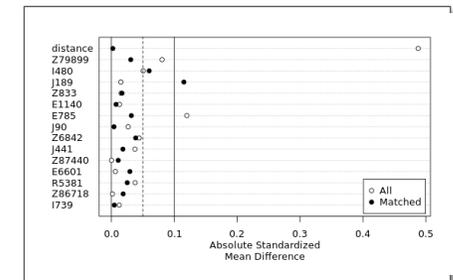
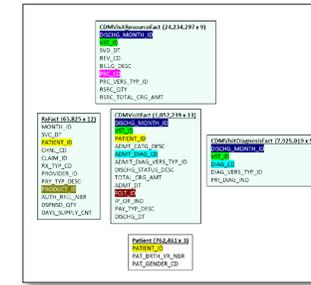
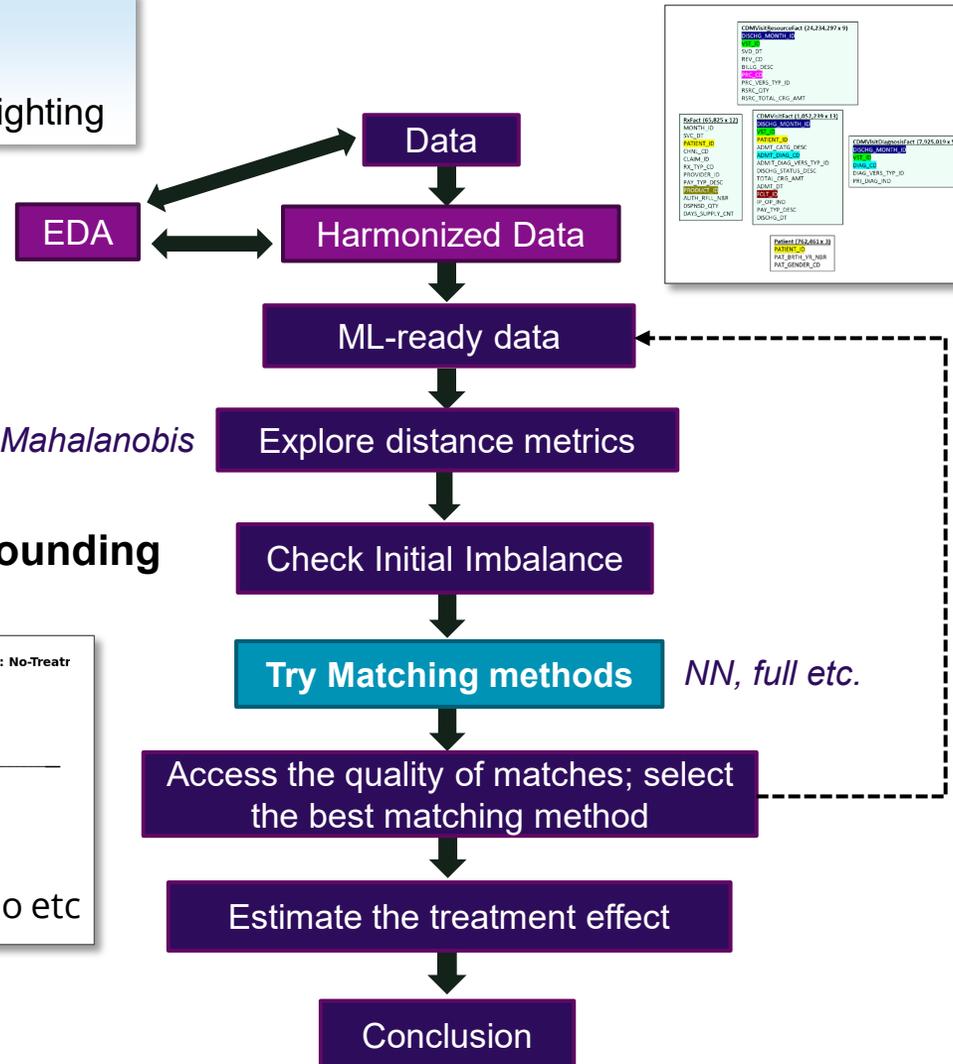
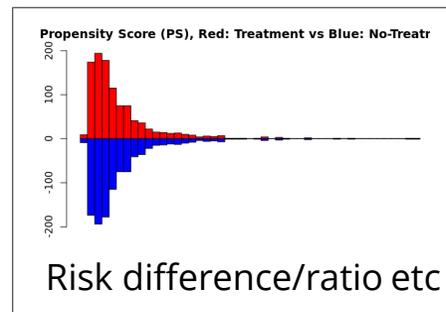
# Overview of RWE Modeling

**EDA:** Exploratory Data Analysis  
**RCT:** Randomized Control Trial  
**IPTW:** Inverse Probability Treatment Weighting

$$smd = \left( \frac{\bar{x}_{treatment} - \bar{x}_{control}}{\sqrt{\frac{s_{treatment}^2 + s_{control}^2}{2}}} \right)$$

*Mahalanobis*

**Use IPTW reduce confounding**



Hypothetical Data to Illustrate Matching Methods

Treated individuals		Comparison individuals	
Individual	Income (in \$10,000)	Individual	Income (in \$10,000)
A	42	a	44
B	35	b	42
C	24	c	37
D	22	d	34
		e	23

Dev Psychol. 2008 March ; 44(2): 395–406. doi:10.1037/0012-1649.44.2.395.

Method	Matched Pair	Global Distance
Greedy (or Nearest-Neighbor)	{Ab}, {Bd}, {Ce}, {Dc}	17 = (0 + 1 + 1 + 15)
Optimal (1:1) matching	{Ab}, {Bc}, {Cd}, {De}	13 = (0 + 2 + 10 + 1)
Full Matching	{Aab}, {Bcd}, {Cde}	7 = (2 + 0 + 2 + 1 + 1 + 1)