

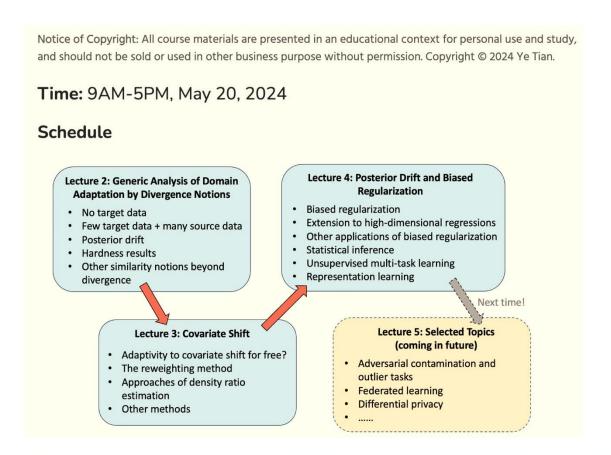
Minghe Wang Biostatistics MS student



## Acknowledgement

This tutorial is modified based on the Theoretical Covariate Shift Lecture from Ye Tian

A (Selective) Introduction to the Statistics Foundations of Transfer Learning

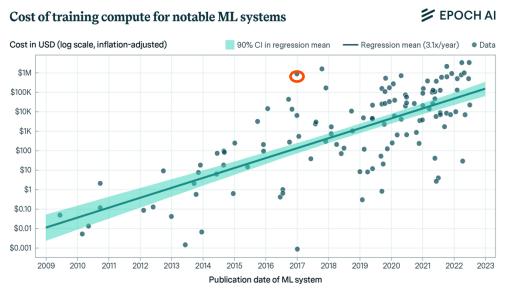


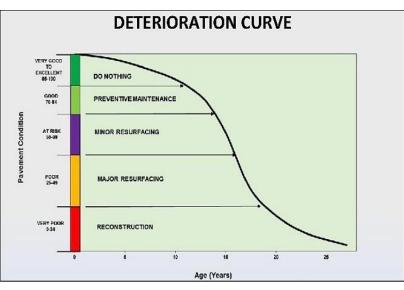
https://www.columbia.edu/~yt2661/STL.html

## Machine Learning Dilemma

Machine learning is powerful but **expensive**, because it requires:

- large labeled datasets
- significant computing power and training time





Ruiz, Victor M. et al.

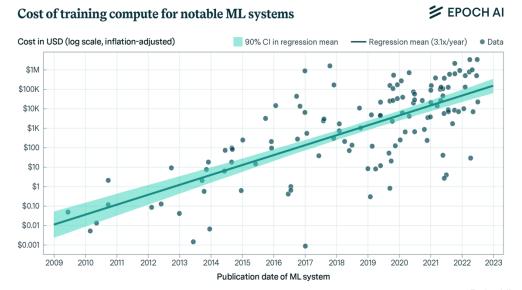
The Journal of Thoracic and Cardiovascular Surgery, Volume 164, Issue 1, 211 - 222.e3

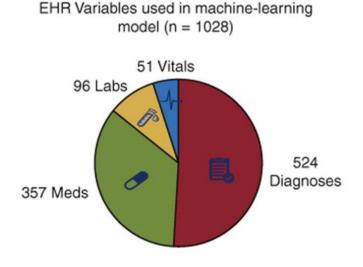
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Annotation (eg. chart review) by doctors is costly





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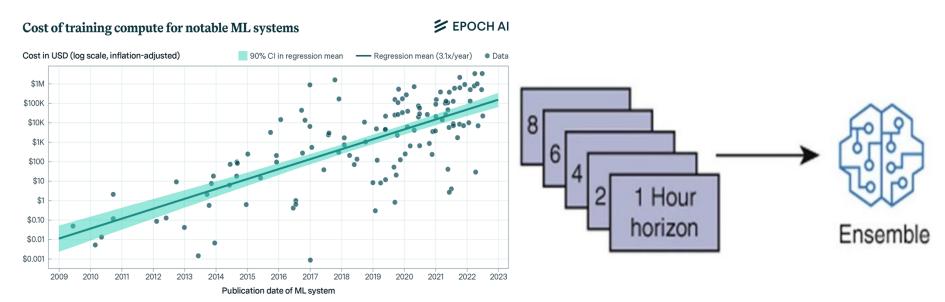
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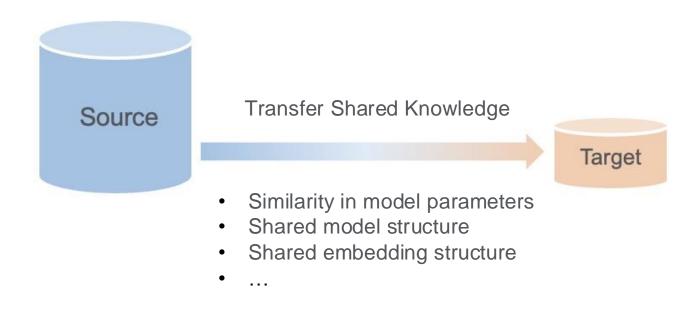
  Annotation (eg. chart review) by doctors is costly
  - significant computing power and training time

    Complex models trained on free text diagnosis



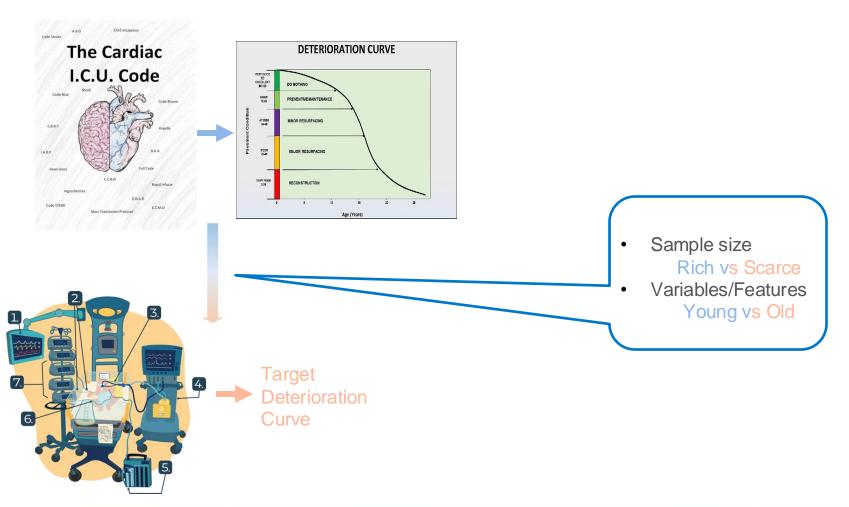
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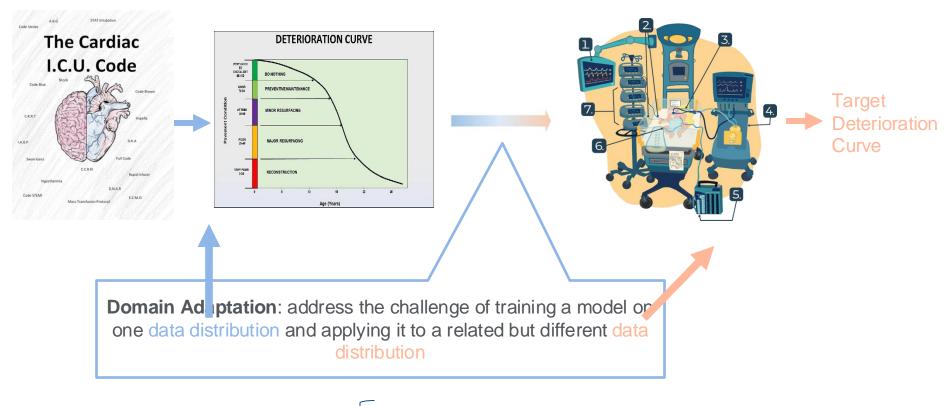
Leverage knowledge from a model trained for one task and reused to help with a similar, related task



Leverage knowledge from a model trained for one task and reused to help with a similar, related task Annotation(eg. chart review) by doctor is costly Complex models trained on free text diagnosis Transfer Shared Knowledge Source **Target** The Cardiac **DETERIORATION CURVE** I.C.U. Code VERY GOOD TO EXCOLLENT 85-100 **Target** AT RISK 10-09 Deterioration ff FOOR Curve

Difference between source and target dataset:





Domain Adaptation (a solution) -

Model finetuning
Covariate shift adjustment
Learning invariant representations

#### **Distributional Shift**

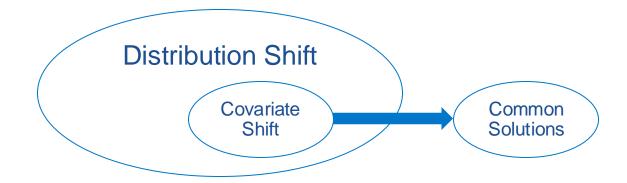
- Target T,  $(X,Y)_T$ Source S,  $(X,Y)_S$
- Probability Chain Rule: P(X,Y) = P(Y|X) P(X)
- Distribution Shift types:
  - Joint Shift:  $P_S(X,Y) \neq P_T(X,Y)$

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  - Covariate Shift:  $P_S(X) \neq P_T(X)$



#### **Tutorial Plan**

- Separate Density Estimation
  - Kernel Density Estimation
  - Histogram-based Method
- Estimating Weight as a whole
  - Kernel Mean Matching
  - Least Square Method
  - Kullback-Leibler Method
  - Discriminative Learning
  - Profile Likelihood Method
- Beyond Reweighting Technique and Covariate Shift
  - Marginal Transfer Learning
  - Domain Invariant Method
  - Optimal Transformation

April 18th

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April 25<sup>th</sup>

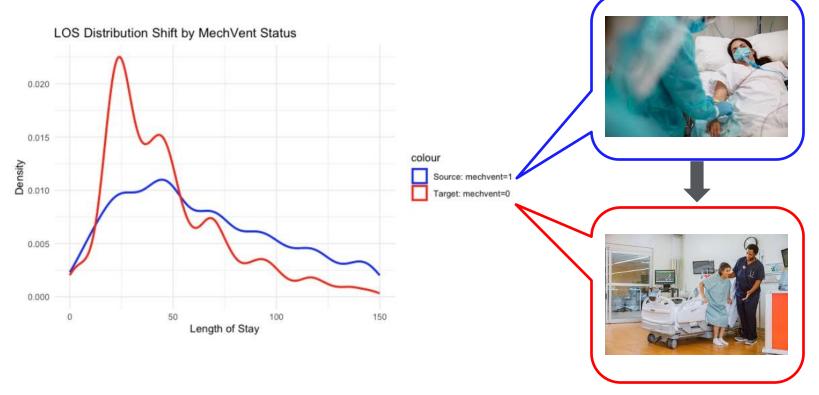
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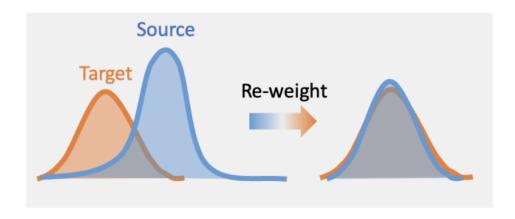
Fall 2025

- Profile Likelihood Method
- Beyond Reweighting Technique and Covariate Shift
  - Marginal Transfer Learning
  - **Domain Invariant Method**
  - **Optimal Transformation**

#### **Covariate Shift**



## Importance Weighting Framework



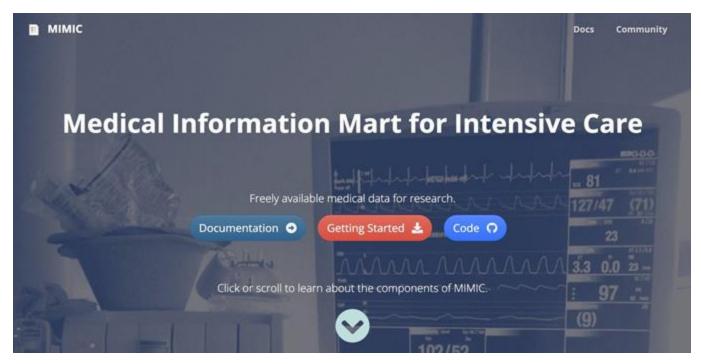
Density ratio model: 
$$\omega(X) = \frac{P^T(X)}{P^S(X)}$$

$$E^{\mathcal{T}}[f(\mathbf{X})] = E^{\mathcal{S}}[\boldsymbol{\omega}(\mathbf{X})f(\mathbf{X})]$$

holds for any f(X)

$$\int f(\mathbf{X}) P^{T}(\mathbf{X}) dX = \int \frac{P^{T}(\mathbf{X})}{P^{S}(\mathbf{X})} f(\mathbf{X}) P^{S}(\mathbf{X}) dX$$

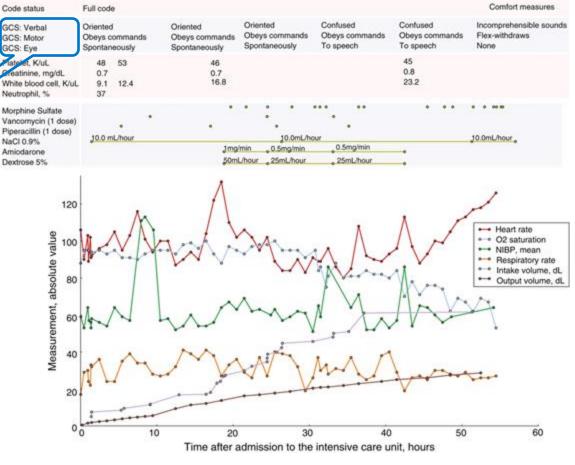




- PhysioNet: <a href="https://physionet.org/">https://physionet.org/</a>
  - The Research Resource for Complex Physiologic Signals
  - Need to get credentialed and trained

Sample data for a single patient stay in a medical intensive care Unit including:

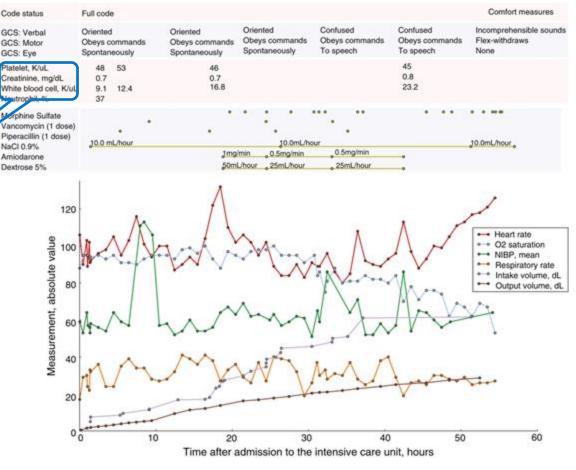
- Consciousness
- Blood test result
- Drug
- Vital sign measurement



https://www.nature.com/articles/sdata201635#Fig2

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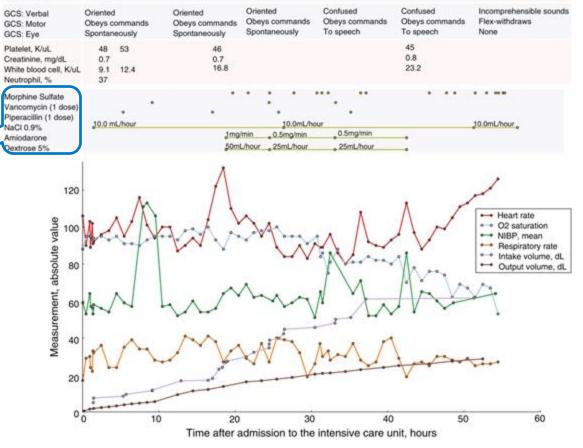
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Code status

Full code

Sample data for a single patient stay in a medical intensive care Unit including:

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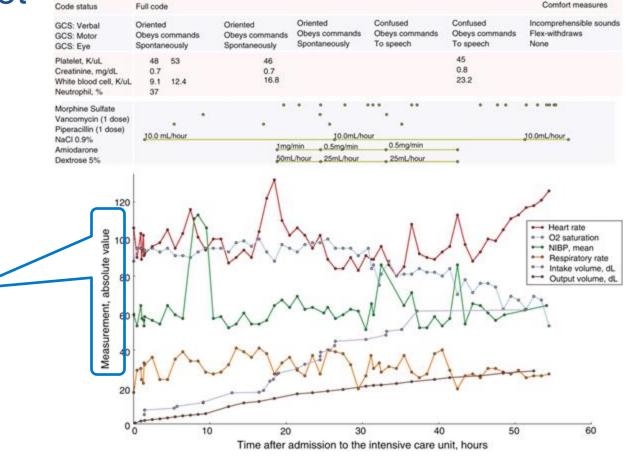
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Comfort measures

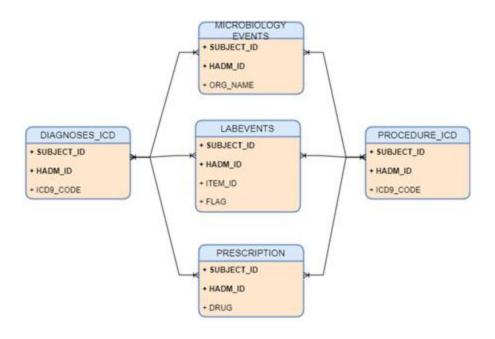
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https://www.nature.com/articles/sdata201635#Fig2



- **'Transfer'**: Physical locations for patients throughout their hospital stay
- 'service' (med\_service\_only): Lists services that a patient was admitted/transferred under.
- **'Callout'**: Provides information when a patient was READY for discharge from the ICU, and when the patient was actually discharged from the ICU.
- 'Patients': Defines each SUBJECT\_ID in the database, i.e. defines a single patient.(demographic)
- 'Elixhauser': Elixhauser Comorbidity Index (ECI) is a tool used to assess the severity and number of chronic health conditions in a patient
- 'Diagnoses\_icd': Definition table for ICD diagnoses.
- '**Drgcodes**': Contains diagnosis related groups (DRG) codes for patients.

https://physionet.org/content/mimiciii-demo/1.4/

Kernel Density Estimation

April 18th

Histogram-based Method

Estimating Weight as a whole

Kernel Mean Matching

Least Square Method

Kullback-Leibler Method

$$\omega(X) = \frac{P^{T}(X)}{P^{S}(X)} = \frac{f_{T}(X)}{f_{S}(X)} -$$

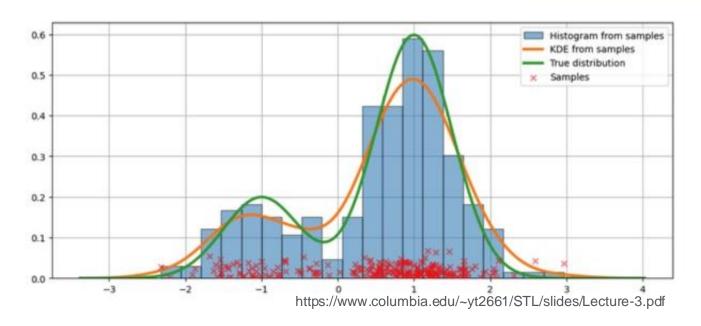
**Separate Density Estimation** 

Parametric density estimation

Non-parametric density estimation

ue and Covariate Shift

- Marginal Transfer Learning
- **Domain Invariant Method**
- **Optimal Transformation**

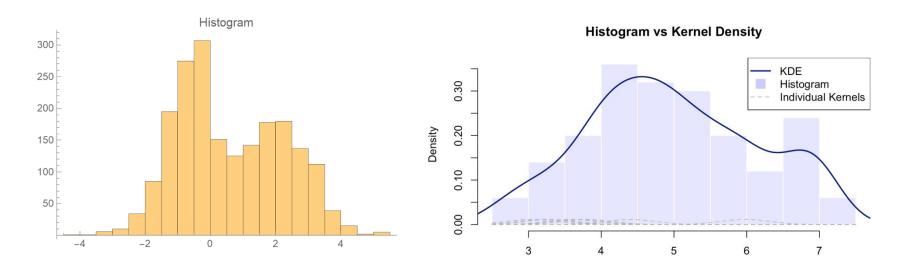


https://minghe4419.github.io/covariate\_shift.github.io/separate\_density\_estimation.html

**COLUMBIA** 

## Kernel Density Estimation (KDE)

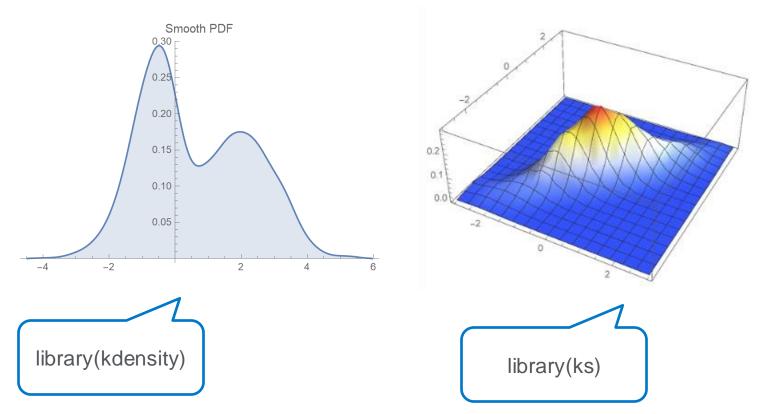
A non-parametric probability density estimation.



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# Kernel Density Estimation (KDE)

A non-parametric probability density estimation.



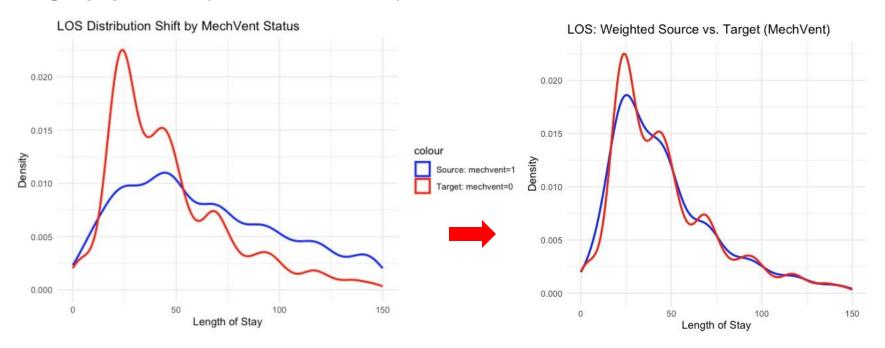
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**Univariate Density Estimation** 

'library(kdensity)'

**Source population**: patients experienced Mechanical Ventilation

**Target population**: patients did NOT experience Mechanical Ventilation



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$$\omega(\mathbf{X}) = \frac{P^{T}(\mathbf{X})}{P^{S}(\mathbf{X})} = \frac{f_{T}(\mathbf{X})}{f_{S}(\mathbf{X})}$$

#### **Univariate Density Estimation**

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```
kde_source_hd <- kde(x = as.matrix(source_hd))
kde_target_hd <- kde(x = as.matrix(target_hd))

# 3. Evaluate densities at the source data points
p_source_hd <- predict(kde_source_hd, x = as.matrix(source_hd))
p_target_hd <- predict(kde_target_hd, x = as.matrix(source_hd))

# 4. Compute the density ratio weights: w(x) = P_T(x) / (P_S(x) + epsilon)
epsilon_val <- 1e-10
weights_hd <- p_target_hd / (p_source_hd + epsilon_val)</pre>
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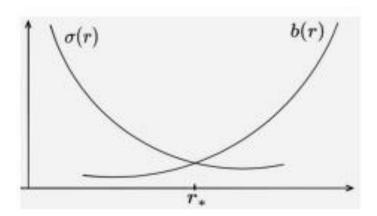
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```

Histogram-based Method

$$\widetilde{\omega}(x) = \frac{P_T(B(x,r))}{P_S(B(x,r))} = \frac{n_T^{-1} \sum_{i=1}^{n_T} I(\left| \left| x - x_{T,i} \right| \right| \le r)}{n_S^{-1} \sum_{i=1}^{n_S} I(\left| \left| x - x_{S,i} \right| \right| \ge r)}$$



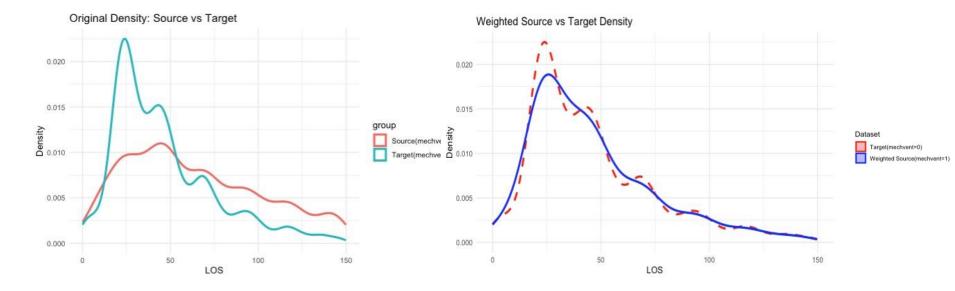
number of target point in circle  $\omega(x) =$ number of source point in circle  $\widehat{\omega}(x) = \widetilde{\omega}(x)I(P_T(B(x/r)) \ge \alpha$ Neighborhood Ball B(x, r Red = Target, Blue = Source

Feature 1

Feature 2

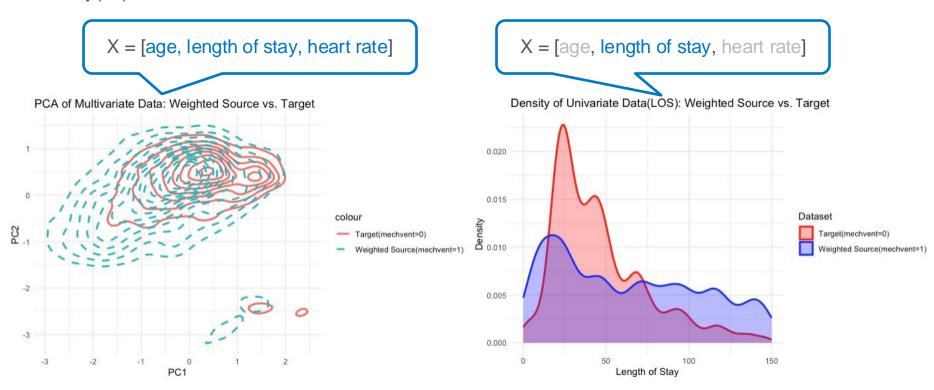
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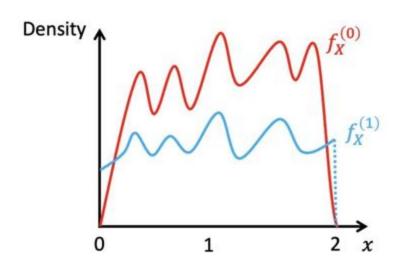
#### **Multivariate Density Estimation**

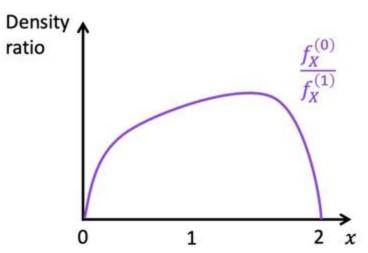
'library(ks)'



Issue: Worse smoothness in density ratio

- Densities are rough while the density ratio is smooth
- Resulting large variance in weights





## **Summary**

- Transfer learning and domain adaptation
- Importance weighting framework
- Separate density estimation
  - KDE
  - Histogram-based
- Helpful sources
  - Our website about Covariate Shift:
     <a href="https://minghe4419.github.io/covariate-shift.github.io/index.html">https://minghe4419.github.io/covariate-shift.github.io/index.html</a>
    - Summary of methods to interact with more helpful resources
  - PhysioNet: <a href="https://physionet.org/">https://physionet.org/</a>
    - The Research Resource for Complex Physiologic Signals
    - Need to get credentialed and trained

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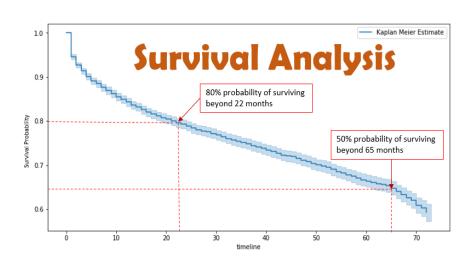
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## Example: Survival Analysis & Cox Model

Survival Analysis: model the time until an event of interest occurs

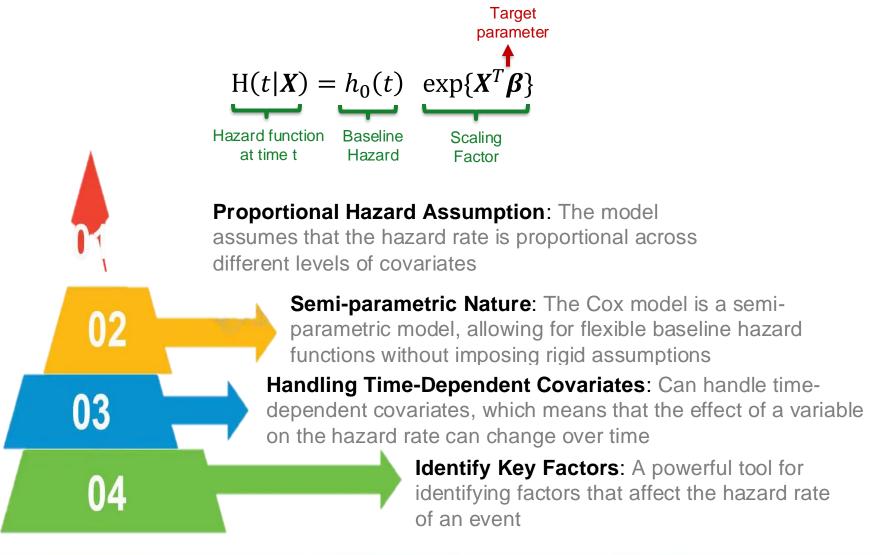
- Medical Research
- Finance
- Social Sciences



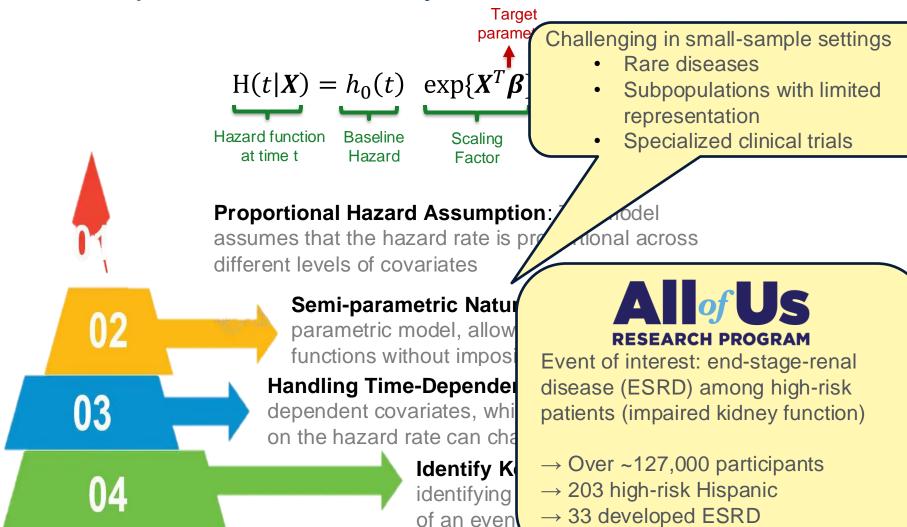


https://medium.com/data-science/survival-analysis-intuition-implementation-in-python-504fde4fcf8e

# Example: Survival Analysis & Cox Model



## Example: Survival Analysis & Cox Model



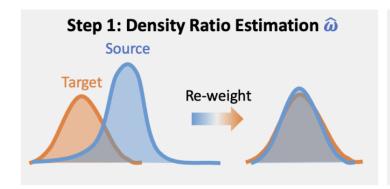
https://fastercapital.com/content/Cox-proportional-hazards

# Transfer Learning in Survival Analysis What are the gaps

- Covariate (X) Shift: Systematic differences between source & target datasets, which will introduce biases, reducing model robustness, especially in the Cox model when model misspecification exists
  - Cox model is used as a "working model" to analyze the relationship between covariates and the hazard rate, even when its assumptions—such as proportional hazards—do not strictly hold
- Limited Adaptation: Existing methods do not fully address cross-population heterogeneity, including baseline hazard variation and coefficient difference

**Q** Our contribution: A robust transfer learning framework that accounts for multi-level data shifts in Cox model

## Proposed 2-step CoxTL



**Step 2: Calibrate Log-likelihood for Joint Estimation** 

$$\boldsymbol{l}^{TL}(\boldsymbol{\beta}) = \boldsymbol{l}^{T}(\boldsymbol{\beta}; X_{T}, Y_{T}) + \boldsymbol{\nu} \ \widehat{\boldsymbol{\omega}} \ \boldsymbol{l}^{S}(\boldsymbol{\beta}; X_{S}, Y_{S})$$
Target Prevent Calibrated Source Negative Transfer

Obtain  $\boldsymbol{\beta}^{TL}$  by maximizing  $\boldsymbol{l}^{TL}(\boldsymbol{\beta})$ 

Adjust the difference in covariate X

- Adjust the difference in parameter β
- Prevent negative effects from heterogeneous sources