

# CS 260: Foundations of Data Science

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Spring 2025



**Haverford**  
COLLEGE

# Outline

- Midterm 2 Review
  - Logistic regression and cross entropy
  - Naive Bayes
  - Disparate impact
- Less focus
  - tSNE
  - t-tests

# How can we tell if an algorithm is biased?

D: dataset with attributes  $X$ ,  $Y$

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Direct discrimination:  $C = f(X)$

- \* Female instrumentalist not hired for orchestra
- \* Some ethnic groups not allowed to eat at a restaurant

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Indirect discrimination:  $C = f(Y)$

- \* but strong correlation between  $X$  and  $Y$
- \* Ex: housing loans
- \* Ex: programming experience

# Notation

features  $\left\{ \begin{array}{l} X: \text{protected feature} \\ Y: \text{other features} \end{array} \right\}$

$X=0$  minority group  
 $X=1$  majority group

9, 10, 6

label  $\left\{ \begin{array}{l} C: \text{binary outcome} \\ \text{if not: } C=0 \end{array} \right\}$  (hired, admitted)  
 $C=1$

## Disparate Impact

$$P(C=1 | X=0) \leq 0.8 P(C=1 | X=1)$$

Idea: if we can predict  $X$  from  $Y$   
there could be disparate impact

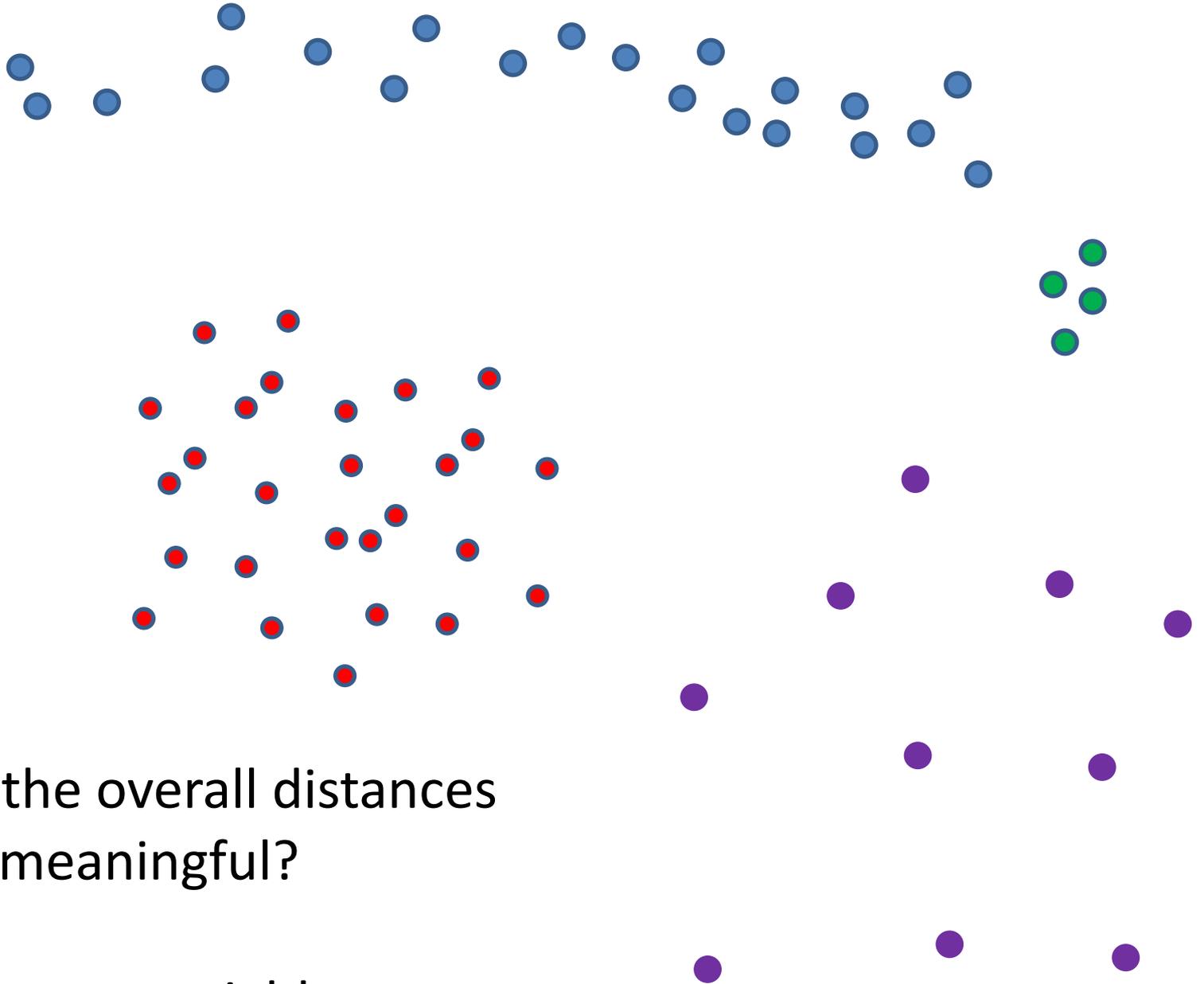
① train  $f: Y \rightarrow X$  ( $f$  is a classifier)  
input (features)      output (label)

② use to predict  $X \rightarrow$  get BER  
balanced error rate

③ if BER is too low, could be disparate impact.

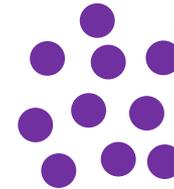
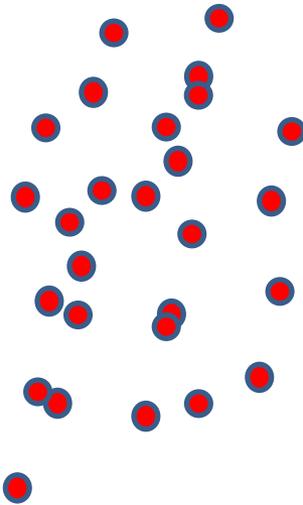
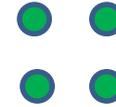
*See video tutorial on Piazza!*

# Bootstrap demo



What if the overall distances  
are not meaningful?

Focus on your neighbors

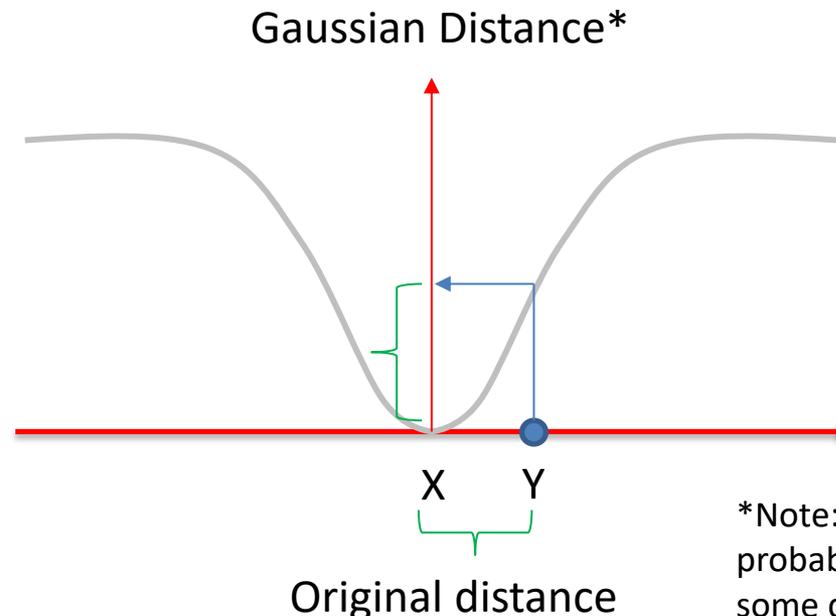


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# tSNE (t-distributed Stochastic Neighborhood Embedding)

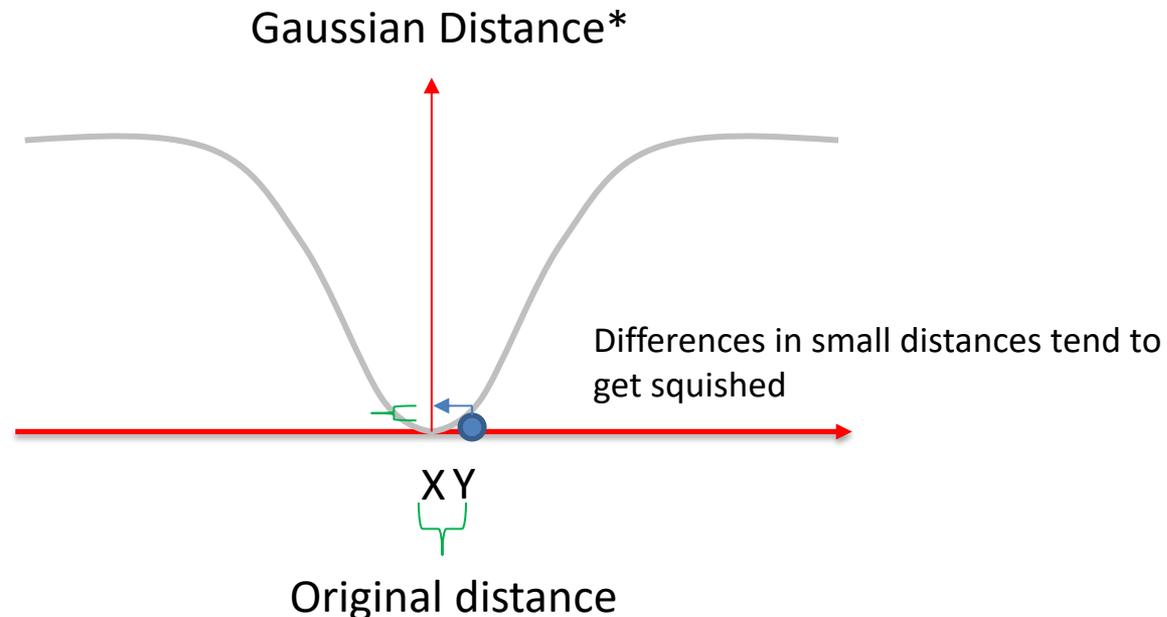
- Define distances between a point X to a point Y by a Gaussian function centered at X



\*Note: the actual algorithm uses notions of probability (i.e., probability of finding Y at some distance from X). I use notion of distance as a proxy

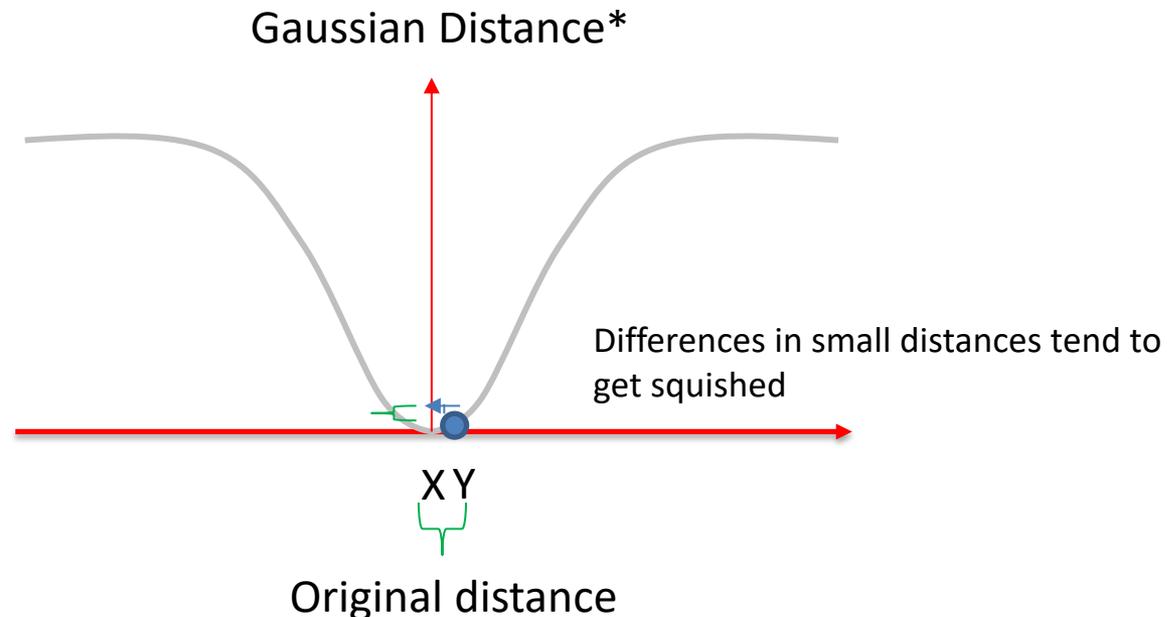
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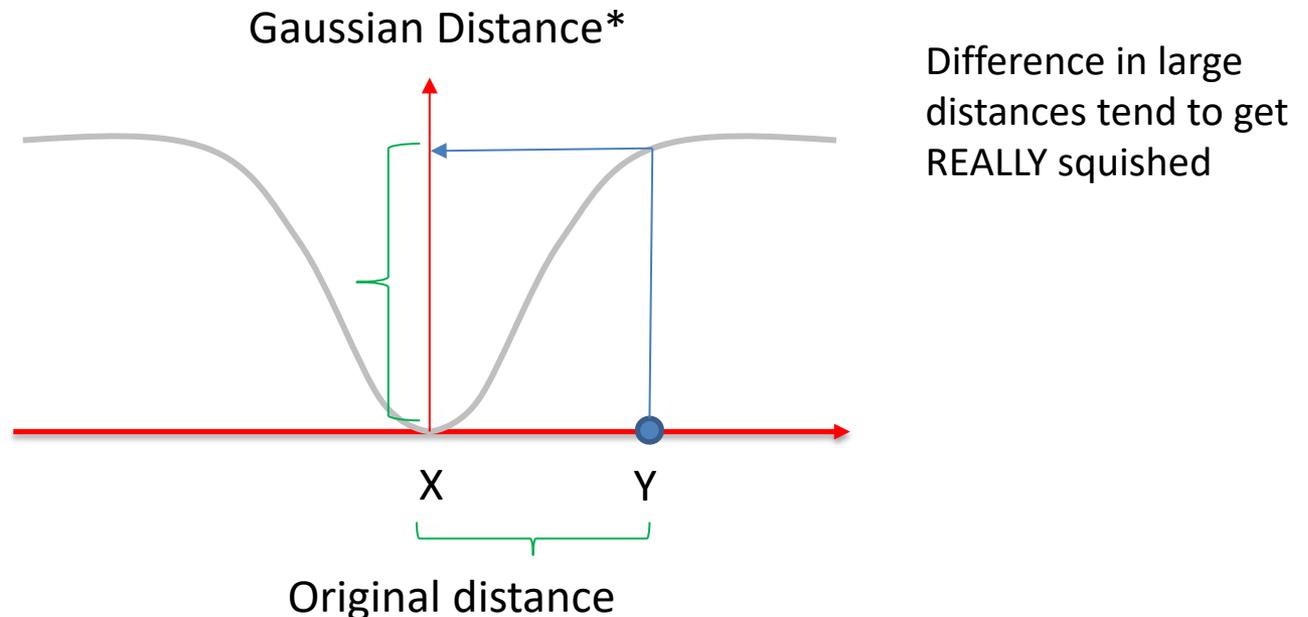
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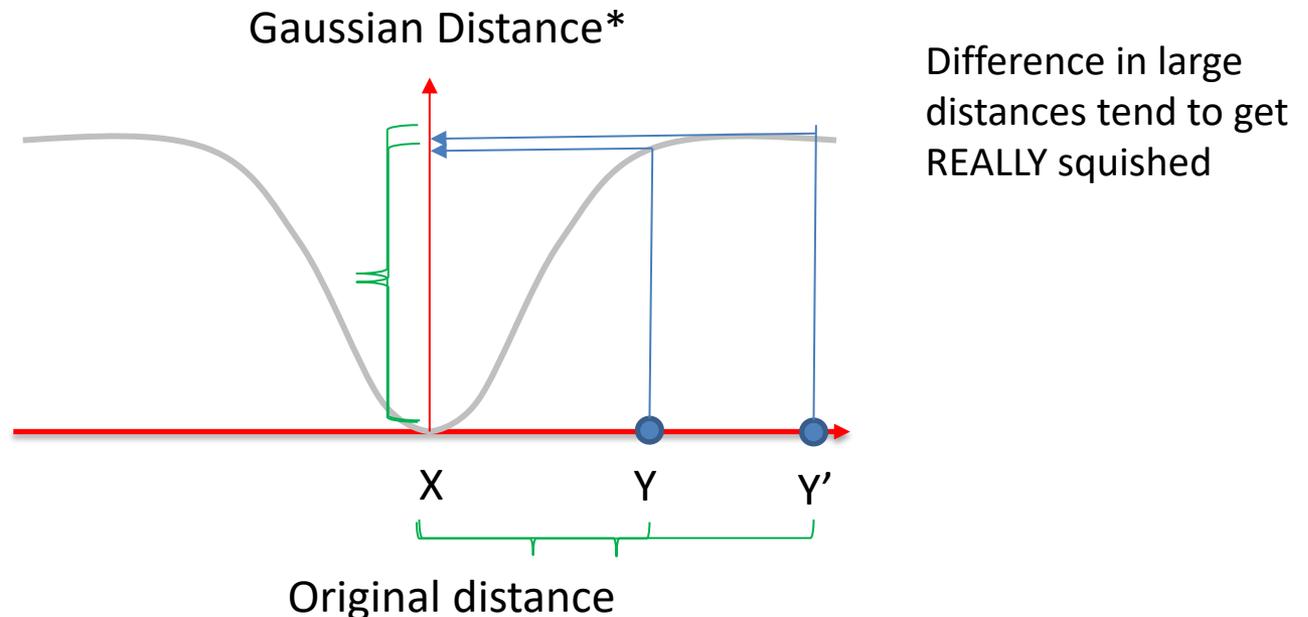
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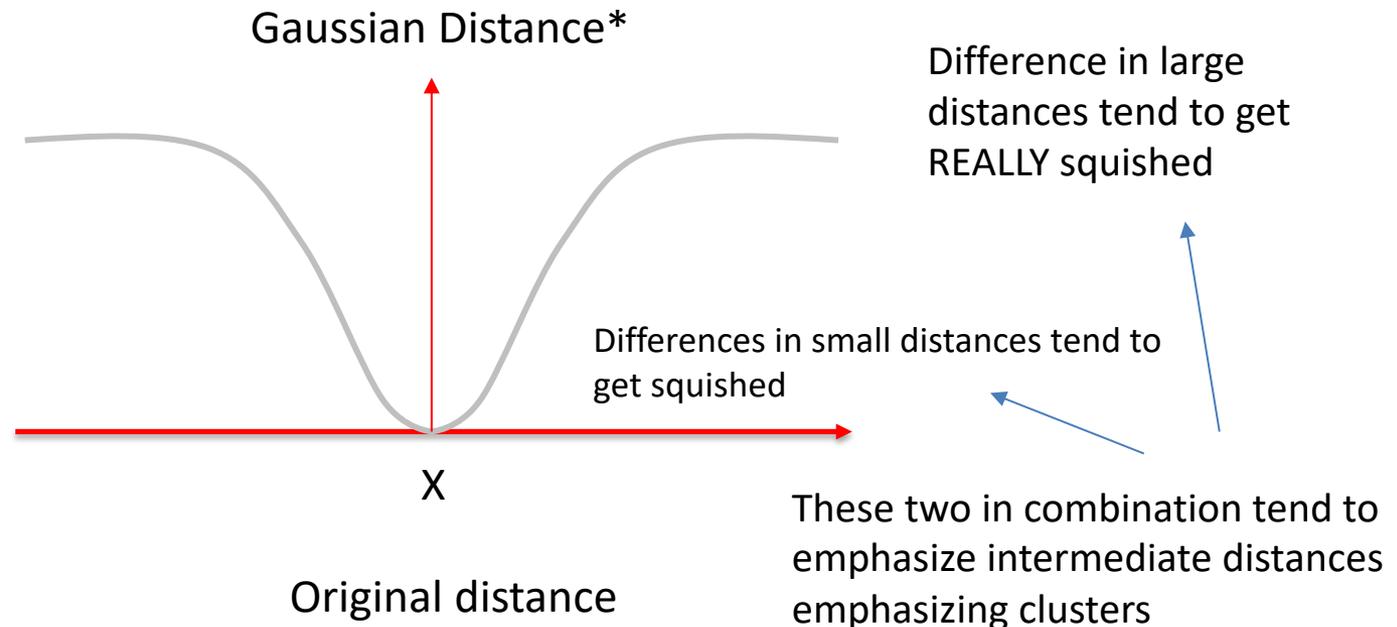
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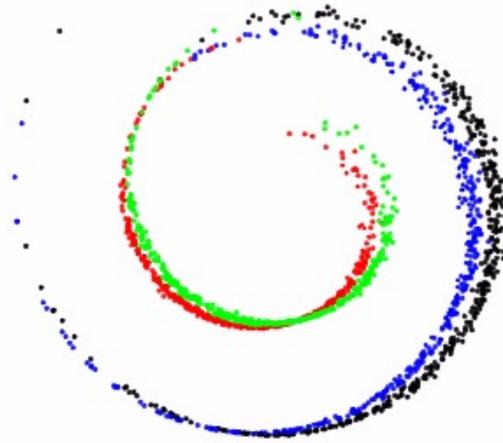


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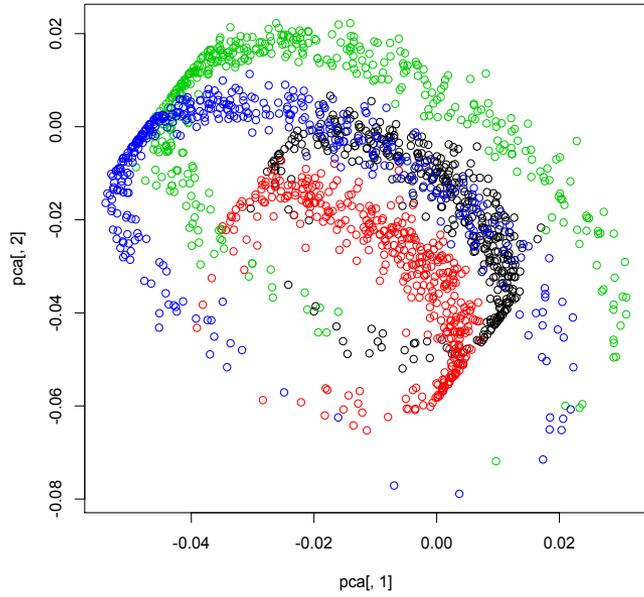
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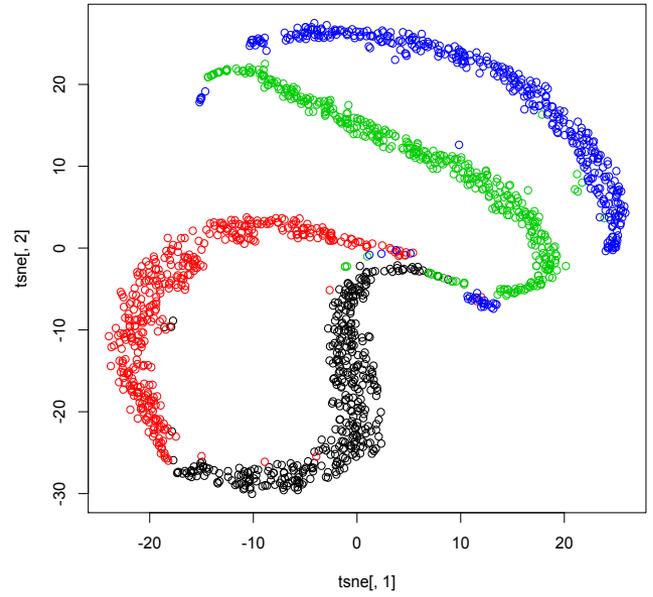
Original data



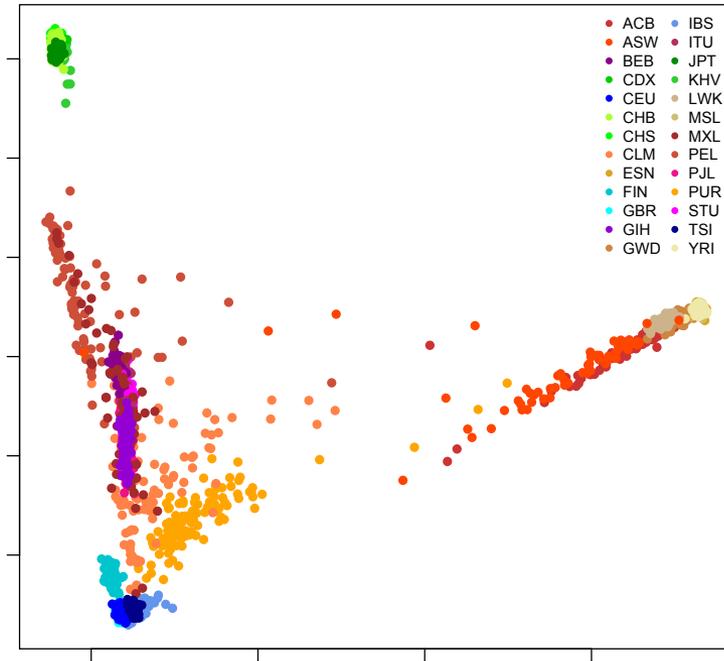
PCA



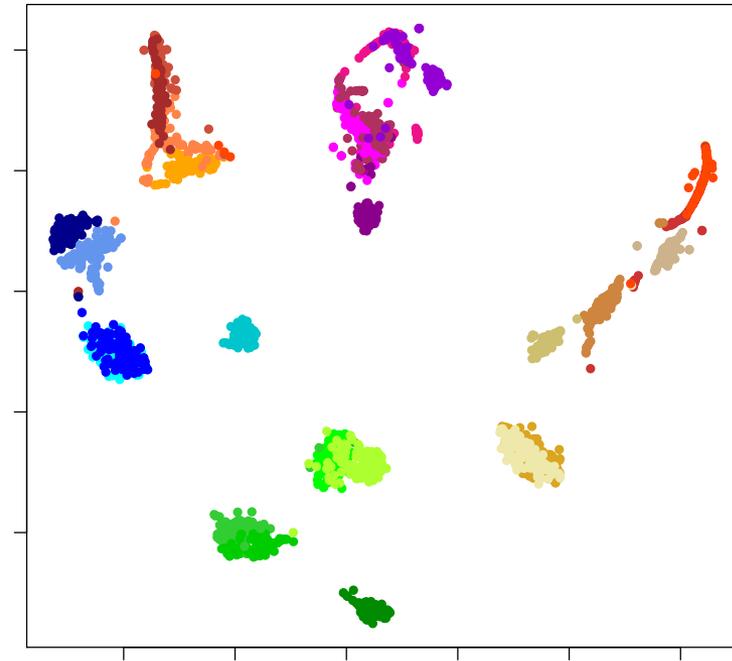
t-SNE



PCA

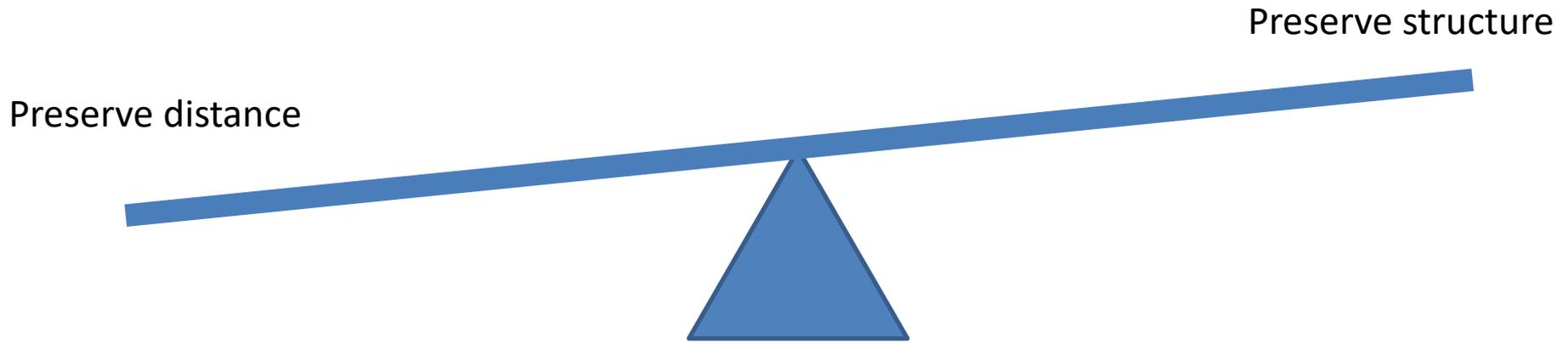


t-SNE



<b>CHB</b>	Han Chinese in Beijing, China
<b>JPT</b>	Japanese in Tokyo, Japan
<b>CHS</b>	Southern Han Chinese
<b>CDX</b>	Chinese Dai in Xishuangbanna, China
<b>KHV</b>	Kinh in Ho Chi Minh City, Vietnam
<b>CEU</b>	Utah Residents (CEPH) with Northern and Western European Ancestry
<b>TSI</b>	Toscani in Italia
<b>FIN</b>	Finnish in Finland
<b>GBR</b>	British in England and Scotland
<b>IBS</b>	Iberian Population in Spain
<b>YRI</b>	Yoruba in Ibadan, Nigeria
<b>LWK</b>	Luhya in Webuye, Kenya
<b>GWD</b>	Gambian in Western Divisions in the Gambia

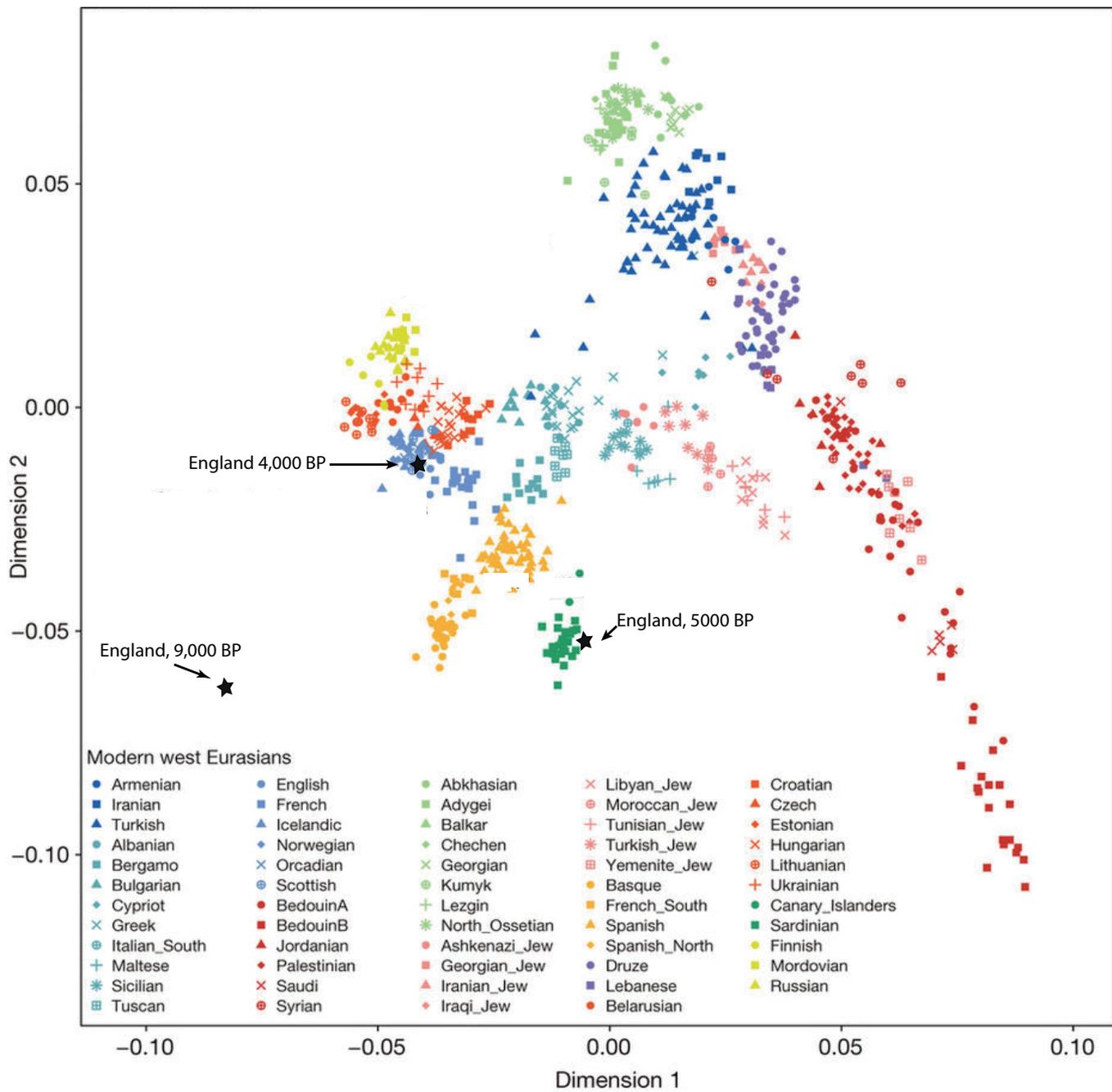
<b>MSL</b>	Mende in Sierra Leone
<b>ESN</b>	Esan in Nigeria
<b>ASW</b>	Americans of African Ancestry in SW USA
<b>ACB</b>	African Caribbeans in Barbados
<b>MXL</b>	Mexican Ancestry from Los Angeles USA
<b>PUR</b>	Puerto Ricans from Puerto Rico
<b>CLM</b>	Colombians from Medellin, Colombia
<b>PEL</b>	Peruvians from Lima, Peru
<b>GIH</b>	Gujarati Indian from Houston, Texas
<b>PJL</b>	Punjabi from Lahore, Pakistan
<b>BEB</b>	Bengali from Bangladesh
<b>STU</b>	Sri Lankan Tamil from the UK
<b>ITU</b>	Indian Telugu from the UK



How to visualize data always depends on the data, and the question

There is rarely if ever a single correct approach





P = pos  
N = neg

H = healthy  
D = disease

$$P(D|P) = 0.8$$

$$P(N|H) = P(P|D) = \frac{P(P)P(D|P)}{P(D)} = \frac{P(P) \cdot 0.8}{\frac{1}{500}} = x$$

$$x = (x + 499(1-x)) \cdot 0.8$$

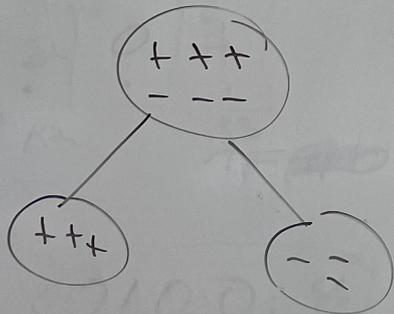
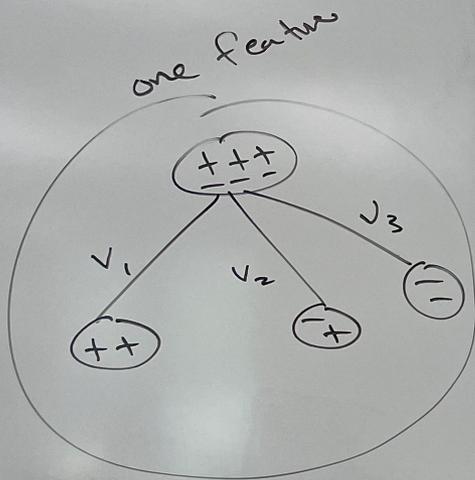
$$\frac{5}{4}x = x + 499 - 499x$$

$$\frac{5}{4}x = 499 - 498x$$

$$x \approx 0.9995$$

$$\begin{aligned} P(P) &= P(P, D) + P(P, H) \\ &= P(D)P(P|D) + P(H)P(P|H) \\ &= \frac{1}{500}x + \frac{499}{500} \left( 1 - \underbrace{P(N|H)}_x \right) \end{aligned}$$

$$x = \left[ \frac{1}{500}x + \frac{499}{500}(1-x) \right] \cdot 0.8$$



$$H(Y) = - \sum_{c \in \text{vals}(Y)} p(Y=c) \log_2 p(Y=c)$$

$$= - p(Y=+) \log_2 p(Y=+) - p(Y=-) \log_2 p(Y=-)$$

$$= - \frac{1}{2} \log_2 \frac{1}{2} - \frac{1}{2} \log_2 \frac{1}{2}$$

$$= + \frac{1}{2} + \frac{1}{2} = 1$$

$$H(Y|X) = \sum_u p(X=u) H(Y|X=u) \left. \vphantom{\sum_u} \right\} \begin{array}{l} \text{weighted avg} \\ \text{of leaf entropies} \end{array}$$

one feature

$$H(Y|X=u) = - \sum_c p(Y=c|X=u) \log_2 p(Y=c|X=u)$$

(9)