### **Dimensionality reduction**

# Example: MNIST http://yann.lecun.com/exdb/mnist/



https://github.com/cazala/mnist

- The "hello world" of deep networks.
- MNIST ZIP Code handwritten images.
- Input 28x28 pixel images [0,255].
- Output: label as integer classes [0-9]
- Training set: 60,000 samples
- Test set: 10,000 samples

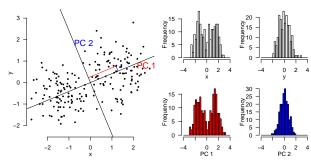
### What does 784-D space look like?

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## Dimensionality reduction / PCA

What: Maximise variance of encoding.

How: Eigenvectors of covariance matrix of inputs. Fractions of variance given by eigenvalues.



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## Multidimensional scaling (MDS)

• For each point  $X_i$  in some high-dim space, we have an equivalent point  $Y_i = (y_{i1}, \dots y_{id})$  in some low (d= 2 or 3) dim space.

$$o_{ij} = dist(X_i, X_j)$$
 (fixed)
$$d_{ij} = dist(Y_i, Y_j) = (\sum_{k=1}^{d} (y_{ik} - y_{jk})^2)^{0.5}$$

- Project down into lower dimension so that  $o_{ij} \approx d_{ij}$ .
- $o_{ij}$  are fixed, so we vary points  $y_{ij}$  to minimise stress term:

$$S^2 = \frac{\sum_i \sum_j (d_{ij} - o_{ij})^2}{\sum_i \sum_j o_{ij}^2}$$

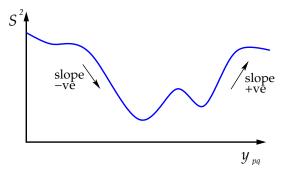
## Relationship between MDS and PCA

- MDS and 'PCA are equivalent when using Euclidean distance measure with stress measure:  $S^2 = \sum_{i,j} (d_{ij} o_{ij})^2$ .
- Other stress measures can be used (e.g. Sammon, next slide) to emphasise certain aspects of data.
- Non-metric versions (NMDS) do not consider absolute distances, but preserve only ranking of distances.

### MDS: Minimisation of Stress term

• Starting from some guess for initial points  $o_{ij}$ , calculate  $S^2$  and then evaluate gradient for each parameter  $y_{pq}$ .

$$\Delta y_{pq} = -\alpha \frac{\partial S^2}{\partial y_{pq}}$$



• Iteratively update  $y_{pq}$  until local minimum (gradient is zero).

Sammon mapping

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• The Sammon mapping uses the error measure:

$$E = \sum_{i < j} \frac{(o_{ij} - d_{ij})^2}{o_{ij}}$$

See separate video on how we apply gradient descent so that we take the derivative of E with respect to each element of each y. As we move each y, the  $d_{ij}$  are recalculated such that E decreases.

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### eurodist: the distances between European cities

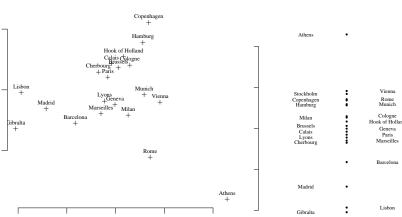
The eurodist dataset (a lower triangular matrix) tells you the road distance in km between 21 cities:

#### > eurodist

	Athens	${\tt Barcelona}$	${\tt Brussels}$	${\tt Calais}$	Cherbourg	Colo
Barcelona	3313					
Brussels	2963	1318				
Calais	3175	1326	204			
Cherbourg	3339	1294	583	460		
Cologne	2762	1498	206	409	785	
Copenhagen	3276	2218	966	1136	1545	į

### MDS example: European distances





(R notes: eurodist data from MASS package, with cmdscale() and sammon())

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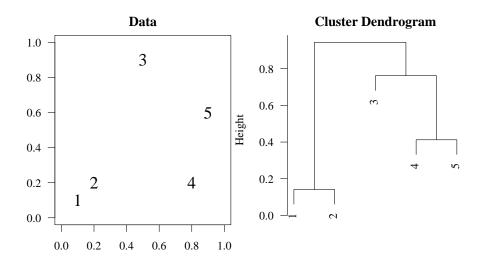
## t-SNE: t-distributed stochastic neighbour embedding

- 1. Another gradient descent approach (van der Maarten and Hinton, 2008).
- 2. Probabilistic approach, with distances scaled by local density of data (locality controlled by parameter "perplexity"). Measure p(i,j) as probability of being neighbours in high-D space, and fix. q(i,j) is adjusted in low-D space by gradient descent on KL divergence.
- 3. Computationally expensive (UMAP has mathematical and computational advantages). Used in genomics a lot, but beware of limitations about intepretation. essential reading: Wattenberg et al (2016) *How to use t-SNE effectively*

https://distill.pub/2016/misread-tsne/).

https://pair-code.github.io/understanding-umap/

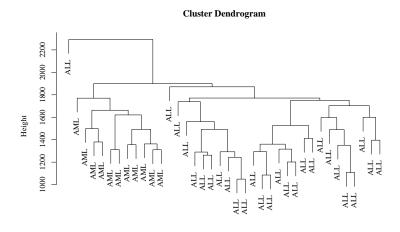
## Hierarchical clustering



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### AML / ALL example (Golub et al. 1999)

Here we have information (type of cancer) associated with each input vector (gene-expression data).

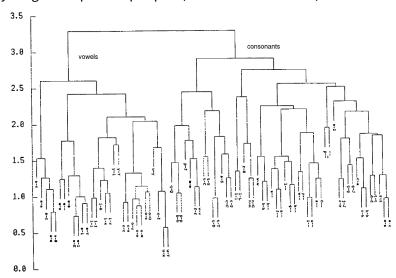


dgTr average linkage, manhattan distance, scaled arrays, 3,051 genes

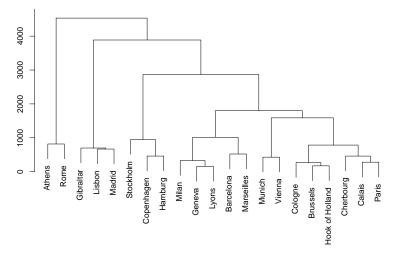
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## NetTalk: Hidden unit analysis

What features are the network extracting? Compute 80-d vector of average activity for given input–output pair (of which there are 79).



## **European cities**



plot(hclust(eurodist), main=',')

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## So, what does 784D space look like?

Excellent online resource for visualisation:

http://colah.github.io/posts/2014-10-Visualizing-MNIST/

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