# Features Dimensionality Reduction



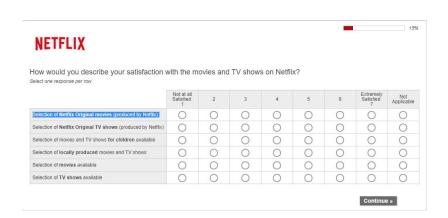
- It's generally a good idea to extract as many diverse features from data as possible. After all, any ML model utilizes the patterns in the features for training. More number of diverse features may provide more number of unique patterns.
- It is also good to extract more features because a fewer number of features may just not be enough for training efficiently any ML model.
- But, having too many features may also pose some problems because the ML model may not work very well on a large dimensional space.







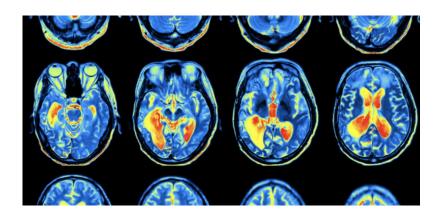
Thousands of documents
Millions of words!
Millions of contexts between words!



Thousands of movies and webseries.
Each with tens of questions in a survey!
Each with hundreds of features based on cast, content, genre, etc.







Hundreds of fMRI brain scans. Each scan with thousands of locations (voxels)! Many scans over time!



Millions of seconds of financial data.

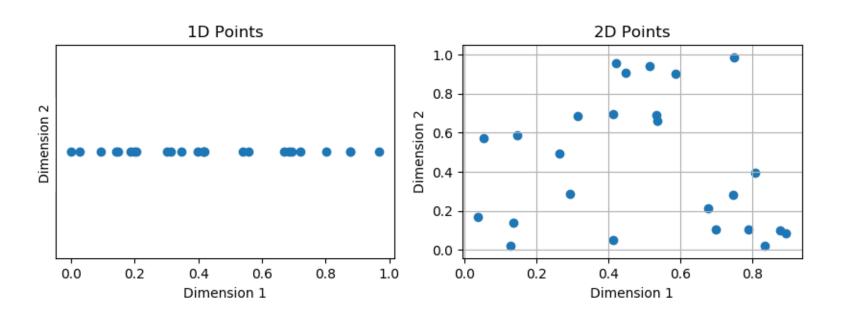
Each datapoint with many features from multiple stocks!

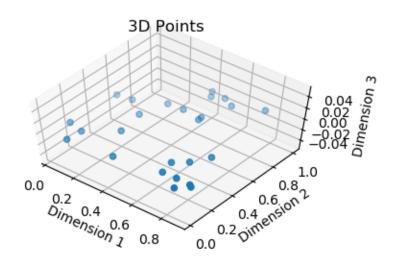
Each stock with many features from the company's declarations!





As we increase the number of features, the data starts to become sparse.

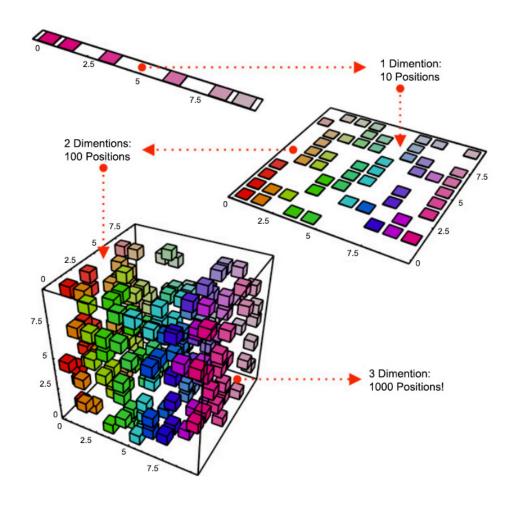




So, we need to go and collect even more data as the number of features increase so that the ML model does not overfit.







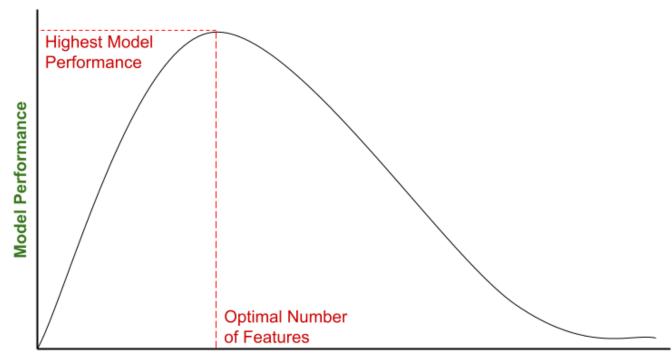
To have the same average distance between the datapoints, we need to add more and more datapoints (i.e., collect more data) as we increase the number of features i.e., dimensionality.





## The Curse of Dimensionality

High number of features may not always be optimal to train an ML model.



No. of Dimensions (Features)

In general, the highest model performance may not be achieved by using a large number of features.

This is the curse of dimensionality.

Training a model with a large number of features is also more time-consuming and may lead to overfitting.

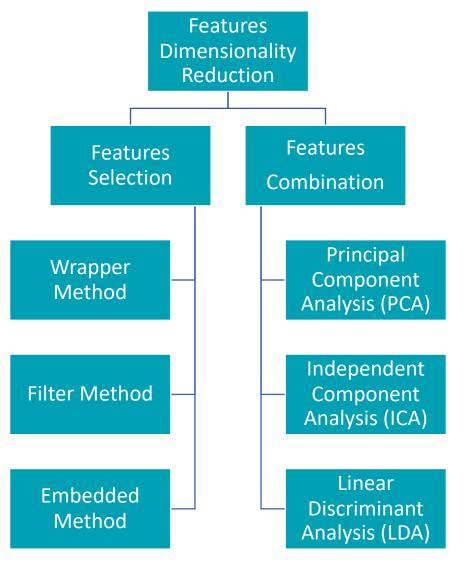
Thus, there is a reduce the number of features for "optimal" model performance.





#### The Solution

The idea here is: Choose a few features from many features.



The idea here is: Combine many features to create a few "superfeatures".

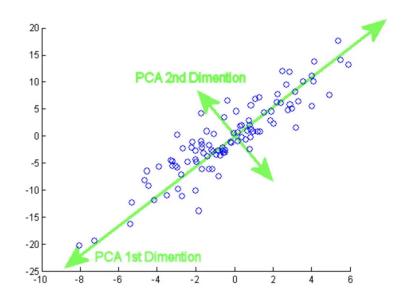




# Principal Component Analysis (PCA)

A technique to extract variance structure from high-dimensional data.

PCA is an orthogonal projection of data into a lower-dimensional subspace such that the variance of the projected data is maximized.

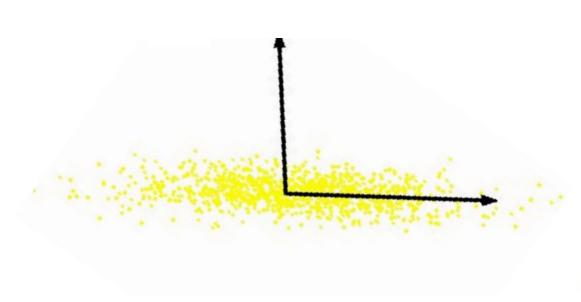


PCA is an unsupervised algorithm!

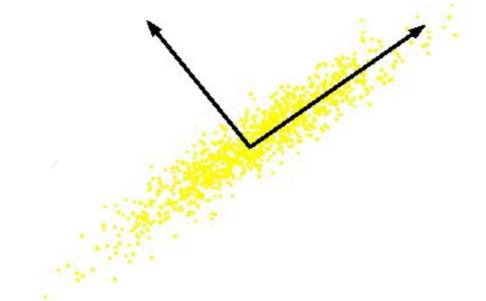




# Principal Component Analysis (PCA)



Two features here Only one is relevant since only one has high variance.



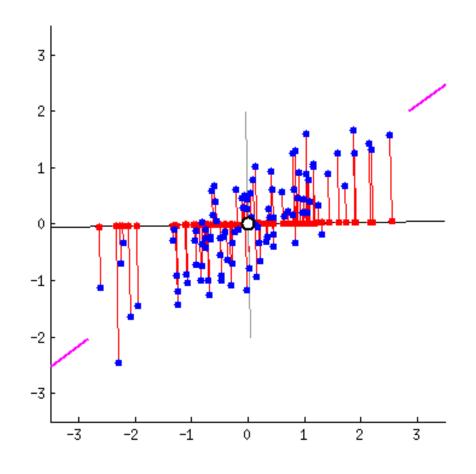
Both features are relevant here. However, only one coordinate is more relevant since high variance only along one coordinate.

How can we transform the features so that the relevant "feature" along the coordinate is preserved?





#### How does PCA work?



Principal components are constructed to account for the variance in features.

The first principal component account for the largest possible variance in the dataset.

In this graphic, the first principal component is the line that matches the purple marks.

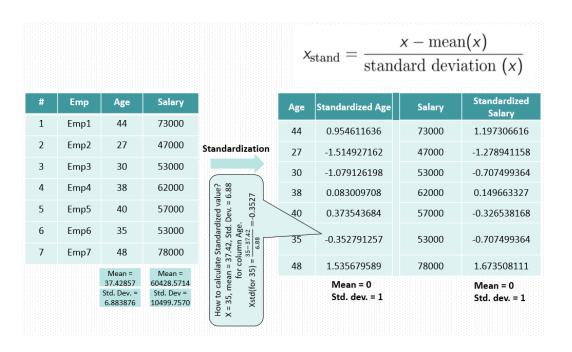
This is because it is the line that maximizes the variance. The variance is calculated as the average of the squared distances from the projected points (red dots) to the origin.

The second principal component is calculated similarly with the condition that it is orthogonal to the first, and so on!





#### Step 1: Standardization



Features are scaled such that mean = 0, std = 1.

Good for Gaussian data.

More robust to outliers.

PCA works under the assumption that the data is Gaussian, thus standardization is important so that large differences between range of variables could be minimized.





Step 2: Compute the Covariance matrix

$$Cov(x,y) = rac{\sum_{i}^{n}(x_{i}-\overline{x})\cdot(y_{i}-\overline{y})}{N-1}$$

Sample Covariance for two independent variables (features) x & y

Covariance is positive if both variables (features) increase.

Covariance is negative if when one variable increases, the other decreases.

Covariance is zero, when there is no direct relation.





Step 3: Compute Eigenvalues and Eigenvectors of the Covariance Matrix For a nxn square matrix A and a non-zero vector X

$$AX = \lambda X$$

for some scalar values  $\lambda$ , known as the Eigenvalues of matrix A.

$$AX - \lambda X = 0$$

$$(A - \lambda I)X = 0$$

where I is an nxn identity matrix. We can solve for

$$|A - \lambda I| = 0$$

to get the Eigenvalues  $\lambda$ , and get corresponding eigenvectors then by solving

$$AX = \lambda X$$



Step 4: Sort the Eigenvalues and corresponding Eigenvectors in descending order We sort the eigenvalues  $\lambda_1$ ,  $\lambda_2$ ,  $\lambda_3$ , ...,  $\lambda_N$  from largest to smallest and sort the eigenvectors accordingly.

Eigenvectors represent the direction in which the data varies the most, and eigenvalues represent the amount of variation along the corresponding direction.

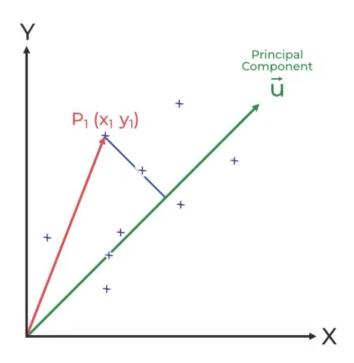
Since these eigenvalues represent the variance in the data, we can pick P eigenvalues from N that explain most variance.

Generally, we taken P to be as many eigenvalues as explain 95% variance in the data. This P then becomes the P number of principal components.





Step 5: Project the data onto the P selected Principal Components

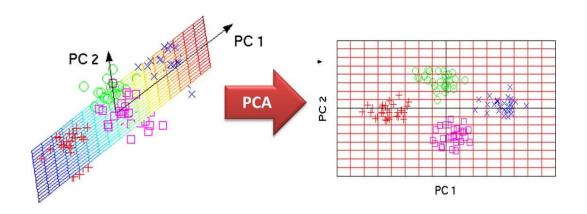


$$Proj_{P_1}\vec{u} = \frac{P_1 \cdot \vec{u}}{|u|}$$
$$= P_1 \cdot \vec{u} \quad .....\vec{u} \rightarrow Unit \ Vector$$

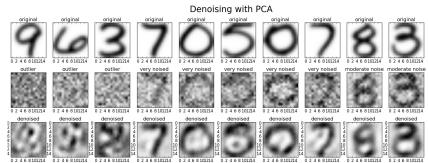




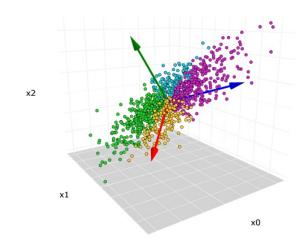
## **PCA** Applications



**Dimensionality Reduction** 



Denoising (if noise is distributed along PCs with low variance)



**Data Visualization** 



**Data Compression** 





#### PCA Disadvantages

- Interpretation of PCs: It's often difficult to interpret PCs as they are a combination of the original features. Hard to know which features contributed how much for PCs.
- Information Loss: Some information will always be lost when we will choose a few PCs from all. Thus, important to know how many to choose.
- Linear assumption: PCA works by assuming that the features have a linear relationship between them. If it is not linear, PCA may now work well.
- Computational Complexity: This will come into play if the number of variables is very large.





Let's say we have a small dataset of five datapoints with four features.

f1	f2	f3	f4
1	2	3	4
5	5	6	7
1	4	2	3
5	3	2	1
8	1	2	2

We want to go from this four-dimensional dataset to a lower dimensional one.





#### Step 1: Standardization

		f1	f2	f3	f4
μ	=	4	3	3	3.4
σ	=	3	1.58114	1.73205	2.30217

f1	f2	f3	f4
-1	-0.63246	0	0.26062
0.33333	1.26491	1.73205	1.56374
-1	0.63246	-0.57735	-0.17375
0.33333	0	-0.57735	-1.04249
1.33333	-1.26491	-0.57735	-0.60812



























#### Step 2: Compute the covariance matrix

	f1	f2	f3	f4
f1	var(f1)	cov(f1,f2)	cov(f1,f3)	cov(f1,f4)
f2	cov(f2,f1)	var(f2)	cov(f2,f3)	cov(f2,f4)
f3	cov(f3,f1)	cov(f3,f2)	var(f3)	cov(f3,f4)
f4	cov(f4,f1)	cov(f4,f2)	cov(f4,f3)	var(f4)

$$Cov(x,y) = rac{\sum_{i}^{n}(x_{i}-\overline{x})\cdot(y_{i}-\overline{y})}{N-1}$$

	f1	f2	f3	f4
f1	0.8	-0.25298	0.03849	-0.14479
f2	-0.25298	0.8	0.51121	0.4945
f3	0.03849	0.51121	0.8	0.75236
f4	-0.14479	0.4945	0.75236	0.8





























Step 3: Compute Eigenvalues and Eigenvectors of the Covariance Matrix

$$|A - \lambda I| = 0$$

	f1	f2	f3	f4
f1	0.8 - λ	-0.25298	0.03849	-0.14479
f2	-0.25298	0.8- λ	0.51121	0.4945
f3	0.03849	0.51121	0.8 - λ	0.75236
f4	-0.14479	0.4945	0.75236	0.8 - λ

 $\lambda = 2.51579324$ , 1.0652885, 0.39388704, 0.02503121





Step 3: Compute Eigenvalues and Eigenvectors of the Covariance Matrix

$$(A - \lambda I)X = 0$$

$$\begin{pmatrix} 0.8000000 - \lambda & -(0.252982) & 0.038490 & -(0.144791) \\ -(0.252982) & 0.8000000 - \lambda & 0.511208 & 0.494498 \\ 0.038490 & 0.511208 & 0.8000000 - \lambda & 0.752355 \\ -(0.144791) & 0.494498 & 0.752355 & 0.8000000 - \lambda \end{pmatrix} \times \begin{pmatrix} v_1 \\ v_2 \\ v_3 \\ v_4 \end{pmatrix} = 0$$

These are our eigenvectors.



Step 4: Sort the Eigenvalues and corresponding Eigenvectors in descending order

```
\lambda = 2.51579324, 1.0652885
```

```
e1 e2
0.161960 -0.917059
-0.524048 0.206922
-0.585896 -0.320539
-0.596547 -0.115935
```

We choose two eigenvalues and the corresponding two eigenvectors.





Step 5: Project the data onto the P selected Principal Components

```
f1
                 f2
                                                       e1
                                                                  e2
                                                                              nf1
                                                                                        nf2
                                                                           0.014003
                                                                                      0.755975
-1.000000 -0.632456
                     0.000000
                                                 0.161960 -0.917059
                                0.260623
                                                                           -2.556534 -0.780432
0.333333
           1,264911
                     1.732051
                                                -0.524048 0.206922
                                1.563740
                                                                           -0.051480
                                                                                      1.253135
                                                -0.585896 -0.320539
-1.000000
           0.632456 -0.577350 -0.173749
                                                                           1.014150
                                                                                      0.000239
                                                -0.596547 -0.115935
           0.000000 -0.577350 -1.042493
0.333333
                                                                           1.579861 -1.228917
1.333333 -1.264911 -0.577350 -0.608121
                                                   (4,2)
                                                                             (5,2)
                                  (5,4)
```

PCA allowed us to reduce the dimensionality of the data from four to two!





# Independent Component Analysis (ICA)

ICA can find new representation to transform data just like PCA.

It can be used for dimensionality reduction like PCA but instead of finding the components that explain maximum variance, it finds the independent components making up the data. Generally, it is used for separating the multivariate signals into components that are maximally independent.

ICA assumes that the subcomponents of the features are non-Gaussian and are statistically independent.

ICA is also an unsupervised method!





#### PCA vs. ICA

#### • PCA

- Focuses on uncorrelated and Gaussian components
- Second-order statistics (variance)
- Orthogonal transformation

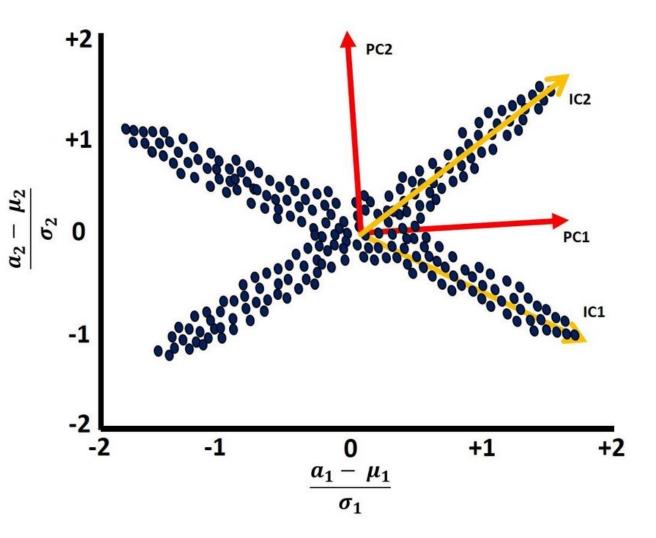
#### • ICA

- Focuses on independent and non-Gaussian components
- Higher-order statistics (kurtosis)
- Non-orthogonal transformation





#### PCA vs. ICA



In PCA, we find the principal components that explain the maximum variance in the data. The PCs are orthogonal to each other.

In ICA, we find the independent components that form the data. They are non-orthogonal.

What are independent components? If one component cannot be estimated from the other component, then it is independent.

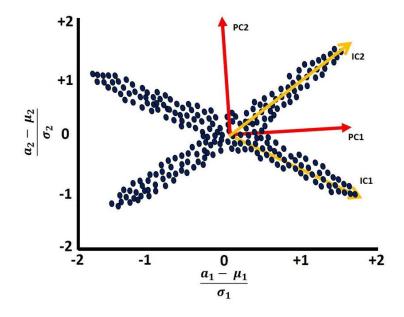




#### ICA Assumption

By Central Limit Theorem, a sum of two independent random variables is more Gaussian than original variables.

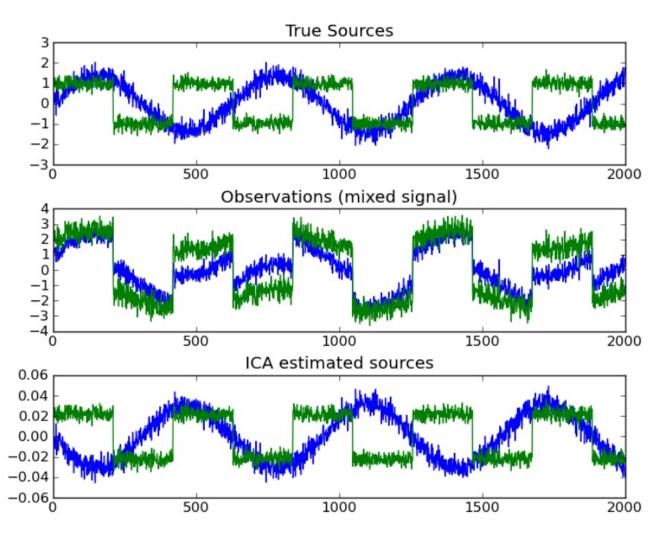
Thus, the distribution of independent components are non-Gaussian.







#### Blind Source Separation







# Blind Source Separation

Consider some signal (data) *s* that is generated by *n* independent sources

$$x = As$$

where A is an unknown matrix (called mixing matrix) and x is the received signal

We have repeated observations of x in our dataset  $\{x^{(i)}, i = 1, 2, 3, ..., m\}$ 

Our goal is to recover the original signal  $s^{(i)}$ 





#### ICA Problem Statement

- We have no prior knowledge of the sources or the mixing matrix *A*
- Permutation of the sources is ambiguous
- We assume that the original signals are non-Gaussian, how can we recover the *n* independent sources forming our data *s*?





Suppose the distribution of each sources  $s_i$  is given by a density  $p_s$ .

The joint distribution of the sources  $s_i$  is given by

$$p(s) = \prod_{i=1}^{n} p_s(s_i).$$

Now,  $x = As = W^{-1}s$ 

$$p(x) = \prod_{i=1}^n p_s(w_i^T x) \cdot |W|.$$

The assumption here is that the sources are independent because only then the joint distribution is the product of the individual sources.

We need to find  $W^{-1}$  which is an approximation of A i.e., the original mixing matrix.



- We need to specify a cdf which increases from 0 to 1.
- Sigmoid function is a good candidate

$$g(s) = \frac{1}{(1 + e^{-s})}.$$

- This yields,  $p_s(s) = g'(s)$
- Given a training set  $\{x^{(i)}, i = 1, 2, 3, ..., m\}$ , the log likelihood for our parameter matrix W is

$$\ell(W) = \sum_{i=1}^m \left(\sum_{j=1}^n \log g'(w_j^T x^{(i)}) + \log |W|
ight)$$



• Maximizing this in terms of W, we derive a stochastic gradient ascent learning rule for training example  $\mathbf{x}^{(i)}$ 

$$W := W + lpha \left( egin{bmatrix} 1 - 2g(w_1^T x^{(i)}) \ 1 - 2g(w_2^T x^{(i)}) \ dots \ 1 - 2g(w_n^T x^{(i)}) \end{bmatrix} x^{(i)^T} + (W^T)^{-1} 
ight)$$

where  $\alpha$  is the learning rate.

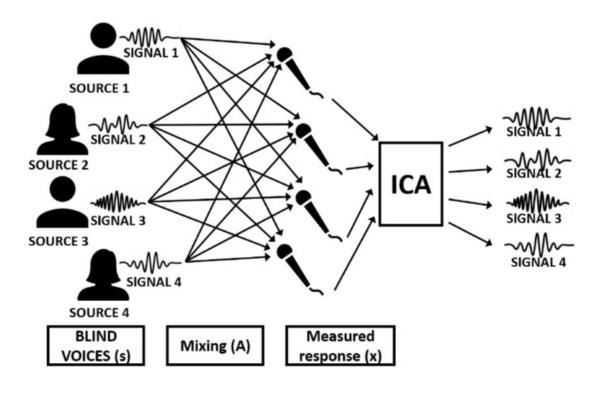
• After the algorithm converges, we get the matrix W to estimate the mixing matrix A.





# ICA and the Cocktail Party Problem



























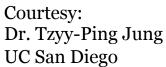






#### ICA in action





















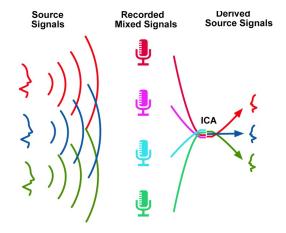








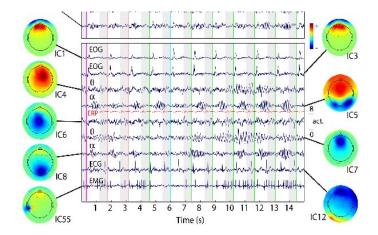
## ICA Applications



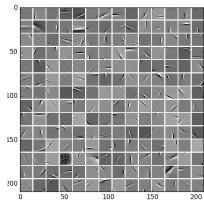
Speech source separation



Finding hidden factors in Financial Data



EEG noise removal (remove contamination due to sources such as eye blinks, heart activity, line noise, etc.



**Image Filtering** 





#### Summary

#### This lecture

- PCA algorithm and its applications
- ICA and its applications

#### Next lecture

- LDA and its applications
- Differences between PCA and LDA, and how LDA could be used for classification.





# Questions?



