



Independent Component Analysis (ICA): A latent variable model

- Find something “interesting” in signals:
‘cocktail party problem’
EEG, ECG signals, FMRI data
- Feature extraction.

Generative Model



$$\mathbf{x}(t) = \mathbf{A}\mathbf{S}(t) + \text{noise}$$

- $\mathbf{x} = (x_1, \dots, x_d)$ vector of observed data (signals, images), $t = \text{index}$
- $\mathbf{S} = (s_1, \dots, s_m)$ vector of statistically independent latent source variables (unknown!)
- \mathbf{A} : $(d \times m)$ Mixing Matrix (unknown parameter !)



Goal:

Demix the signals and recover sources

$$\hat{\mathbf{S}}(t) = \mathbf{W}\mathbf{x}(t)$$

with $\mathbf{W} = \mathbf{A}^{-1}$ for square matrices and no noise.

Ambiguities: Permutation of Sources, Scaling $s_i \rightarrow \lambda s_i$.

Some Interpretations of ICA

- $x_i(t) = \sum_j A_{ij}s_j(t)$

$x_i(t)$ is signal at sensor i & $s_j(t)$ speaker j at time t .

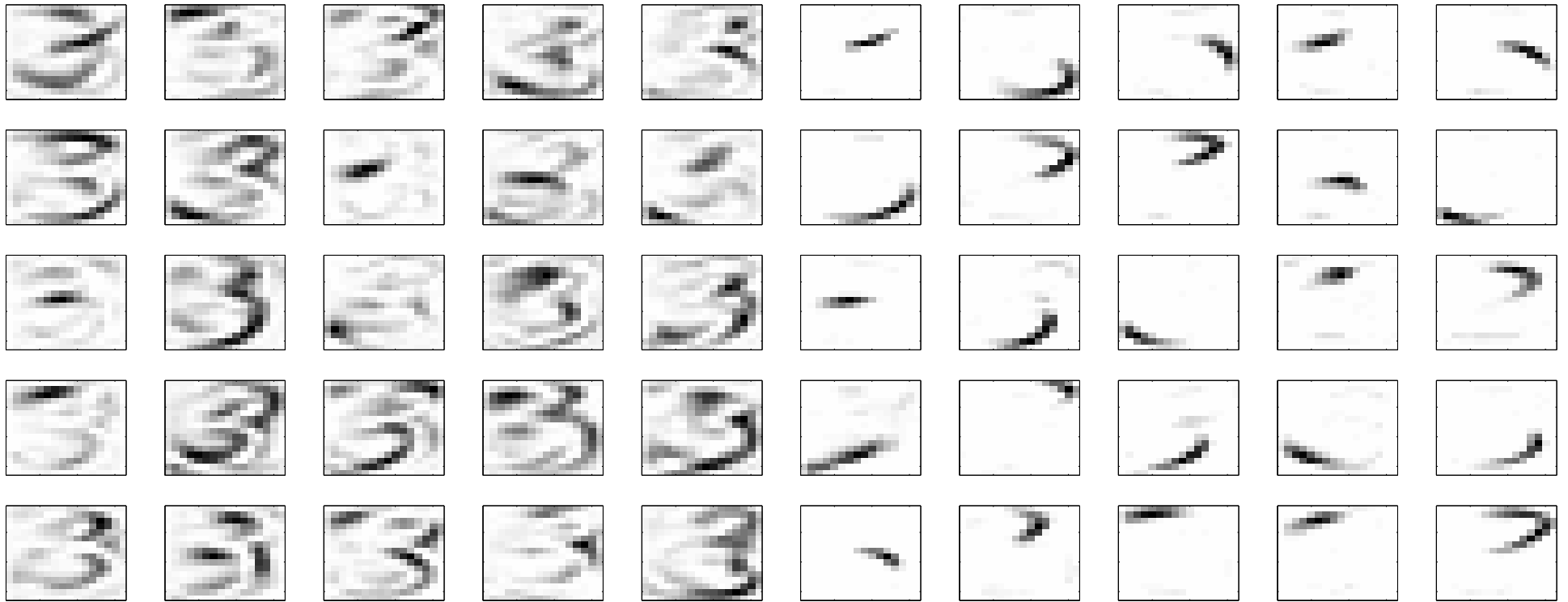
- $x_i(t) = \sum_j A_{ij}s_j(t)$

Vector $x_i(t)$ of pixel intensities of image t is expanded into features $A_{\bullet j}$ and the $s_j(t)$ are the statistically independent coefficients.

- $x_t(i) = \sum_j A_{tj}s_j(i)$

$x_t(i)$ intensity of each pixel i at time t is a time dependent mixture of time independent activity pattern $s_j(i)$.

Feature Extraction

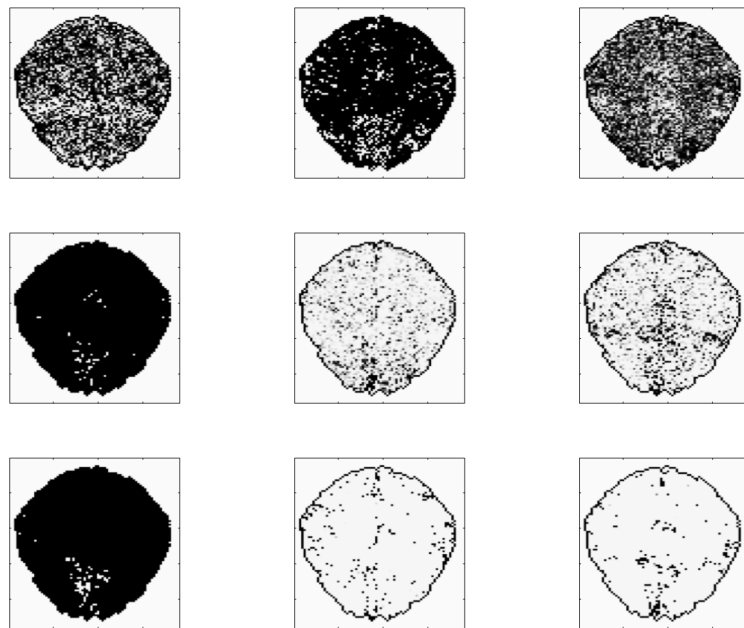


left: unconstrained **right:** constrained (positive) mixing matrix \mathbf{A} .

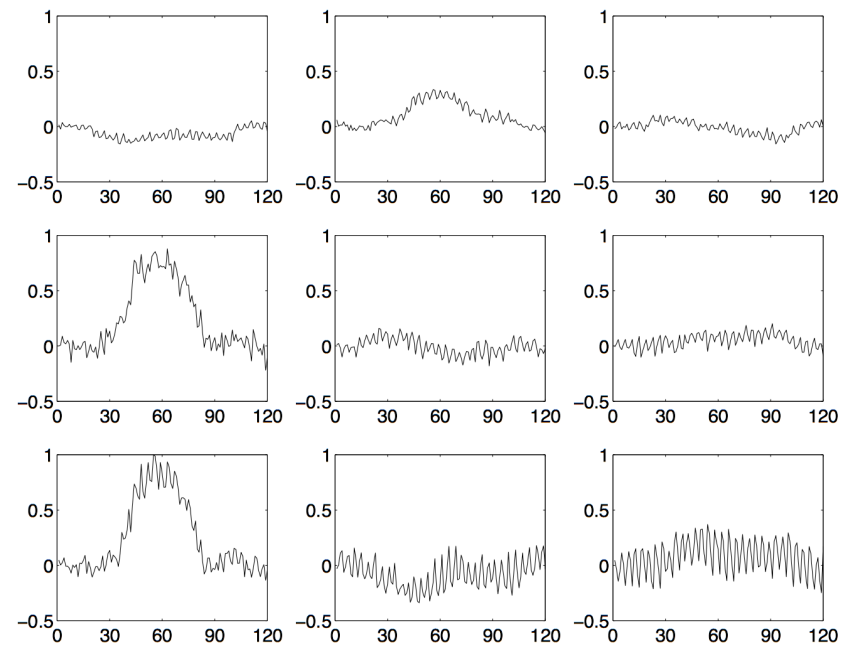
$x_i(t)$ = sequence of 500 images (handwritten '3's). $p(s) = e^{-s}$, $s \geq 0$.
 Shown are the $m = 25$ columns $A_{\bullet j}$ of the matrix \mathbf{A} .

Functional Magnetic Resonance Imaging (fMRI) from: Højen–Sørensen, Hansen & Winther.

left: Posterior mean sources



right: responses $A_{\bullet i}$ for $i = 1, \dots, 9$.



Computing the Likelihood

Assume no noise and $d = m$

- Assume all n data are independent (no temporal structure):

$$p(D|\mathbf{A}) = \prod_{t=1}^n p(\mathbf{x}(t)|\mathbf{A})$$

- Look at a single data point: $p(\mathbf{x}|\mathbf{A}) = \int d\mathbf{S} p(\mathbf{x}|\mathbf{A}, \mathbf{S}) p(\mathbf{S})$

with $p(\mathbf{S}) = \prod_{i=1}^d p_i(s_i)$ (ICA assumption) and

$p(\mathbf{x}|\mathbf{A}, \mathbf{S}) = \prod_{k=1}^d \delta(x_k - (\mathbf{A}\mathbf{S})_k)$ Dirac - δ distributions (i.e. no noise).

The Likelihood cont'd

$$p(\mathbf{x}|\mathbf{A}) = \int d\mathbf{S} \, p(\mathbf{x}|\mathbf{A}, \mathbf{S}) \, p(\mathbf{S}) = \frac{1}{|\det \mathbf{A}|} \prod_{i=1}^d p_i((\mathbf{A}^{-1}\mathbf{x})_i)$$

With $\mathbf{W} = \mathbf{A}^{-1}$, we get for the negative log-likelihood

$$-\ln p(D|\mathbf{W}) = -n \ln |\det \mathbf{W}| - \sum_t \sum_i \ln p_i((\mathbf{W}\mathbf{x}(t))_i)$$

which must be minimized with respect to the matrix \mathbf{W} .

Modeling the sources

- Relation to PCA

Let \mathbf{U} matrix of eigenvectors of covariance matrix, i.e. $\Sigma\mathbf{U} = \mathbf{U}\Lambda$.
If we set $\mathbf{W} = \Lambda^{-\frac{1}{2}}\mathbf{U}^T$, then the vector

$\mathbf{W}\mathbf{x} \doteq \Lambda^{-\frac{1}{2}}\mathbf{U}^T\mathbf{x}$ has decorrelated components with unit variance.

For Gaussian signals: decorrelated = independent!

BUT any $\mathbf{Q}\mathbf{W}$ with orthogonal \mathbf{Q} (i.e. $\mathbf{Q}\mathbf{Q}^T = \mathbf{I}$) will also decorrelate the signal: Estimation of “true” mixing matrix impossible for Gaussian signals/sources. Rotating a spherical Gaussian doesn't change its shape!

- Hence, *assume* non-Gaussian sources like e.g.

the **super-Gaussian** $p_i(s) \propto \frac{1}{e^s + e^{-s}}$.

Disadvantages of Simple Model

- Noise ?
- Constraints on Mixing Matrix (positivity) ?
- Number of sources \neq number of sensors ?
- How many sources are enough ?

Other approaches I: Minimize Mutual Information

Goal: Find \mathbf{W} such that $\mathbf{S} \doteq \mathbf{W}\mathbf{x}$ has independent components.

Minimize Mutual information

$$I = \int d\mathbf{S} p(\mathbf{S}) \ln \frac{p(\mathbf{S})}{\prod_{i=1}^m p_i(s_i)}$$

with respect to \mathbf{W} . Problem: Find good estimate for I from data sample $\mathbf{x}(1), \mathbf{x}(2), \dots, \mathbf{x}(T)$.

Practical Solutions:

- Approximate I using low order cumulants.
- Assume source model, eg $p(s) = \frac{1}{\pi \cosh(s)}$ - equivalent to Maximum Likelihood (Bell & Sejnowski, Cardoso & Laheld, MacKay)

Other approaches II: Non – Gaussianity

Mixing sources $\sim \sum$ of independent random variables \sim Gaussian distribution.

Demixing Make distribution $p(S)$ of $S \doteq \mathbf{W}\mathbf{x} = \mathbf{W}\mathbf{A}\mathbf{x}$ as non Gaussian as possible!

Possible 'contrast functions' for Minimization

- Higher Cumulants such as $\text{kurt}(s) \doteq E[s^4] - 3(E[s^2])^2$
(Hyvärinen's *FastICA*)
- 'Negentropy': $H_{Gauss} - H[\mathbf{S}]$.

More approaches

- Use temporal structure
- Kernel ICA
- ...

The ICA model was an example of

Latent Variable Models

- Simple models (like exponential families) allow for simple analytic parameter estimation by Maximum Likelihood.
- More complex models explain data by hidden (unobserved) variables, the so called latent variables. Such models are very useful in practice.
- However, even Maximum Likelihood (ML) estimation can become a hard computational task.

Overview

- Latent variable models: Definition
- Examples
- ML with the EM Algorithm

Latent variable Models: Definition

y = observed variables.

$\theta = (\theta_y, \theta_x)$ sets of parameters.

x = latent, unobserved variables.

Total likelihood

$$p(y|\theta) = \sum_{\mathbf{x}} p(y|\mathbf{x}, \theta_y) p(\mathbf{x}|\theta_x)$$

If the x 's would be known, ML would often be easy!

Example I: Mixtures of Gaussians

Model for multimodal densities

$$\begin{aligned} p(y|\{\mu_c, \sigma_c, p(c)\}_{c=1}^K) &= \sum_c p(c) \frac{1}{\sqrt{2\pi\sigma_c^2}} \exp\left[-\frac{(y - \mu_c)^2}{2\sigma_c^2}\right] \\ &\equiv \sum_c p(c)p(y|c, \boldsymbol{\theta}) \end{aligned}$$

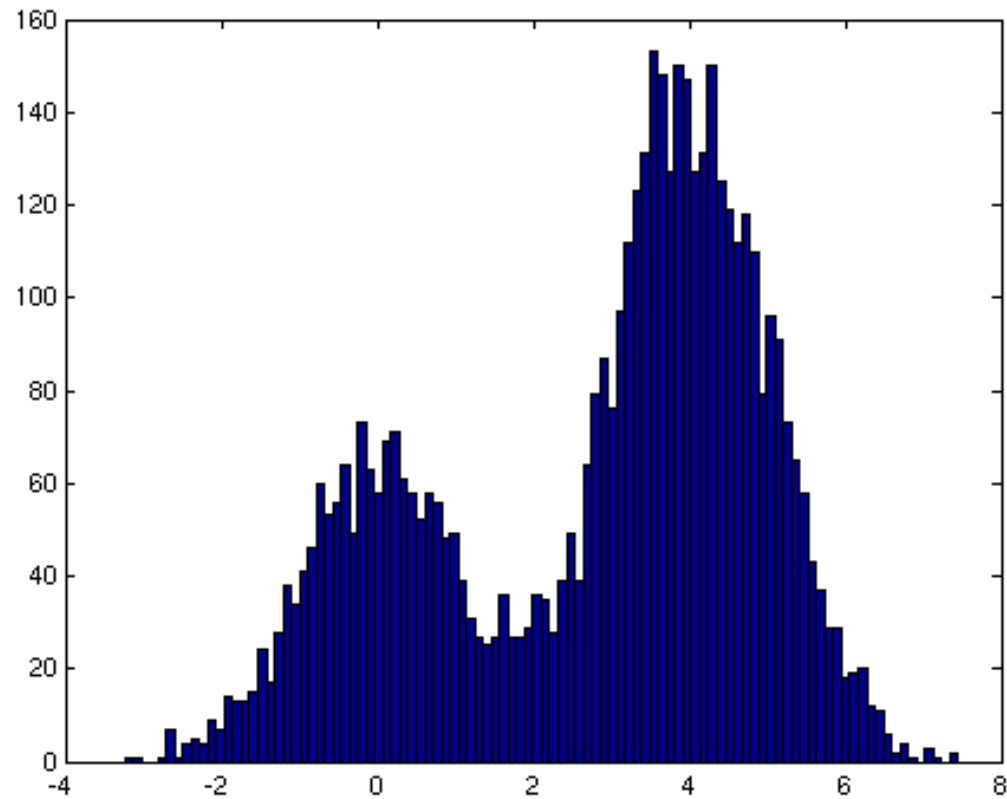
Total likelihood $p(D|\boldsymbol{\theta}) = \prod_i p(y_i|\boldsymbol{\theta})$

y_i observed, component c_i hidden,

$\boldsymbol{\theta} = \{\mu_c, \sigma_c, p(c)\}_{c=1}^K$ parameters to be estimated by ML.

Take $\nabla_{\boldsymbol{\theta}} \ln p(D|\boldsymbol{\theta}) = 0$ results in complicated set of nonlinear equations.

Data from a mixture of 2 Gaussians



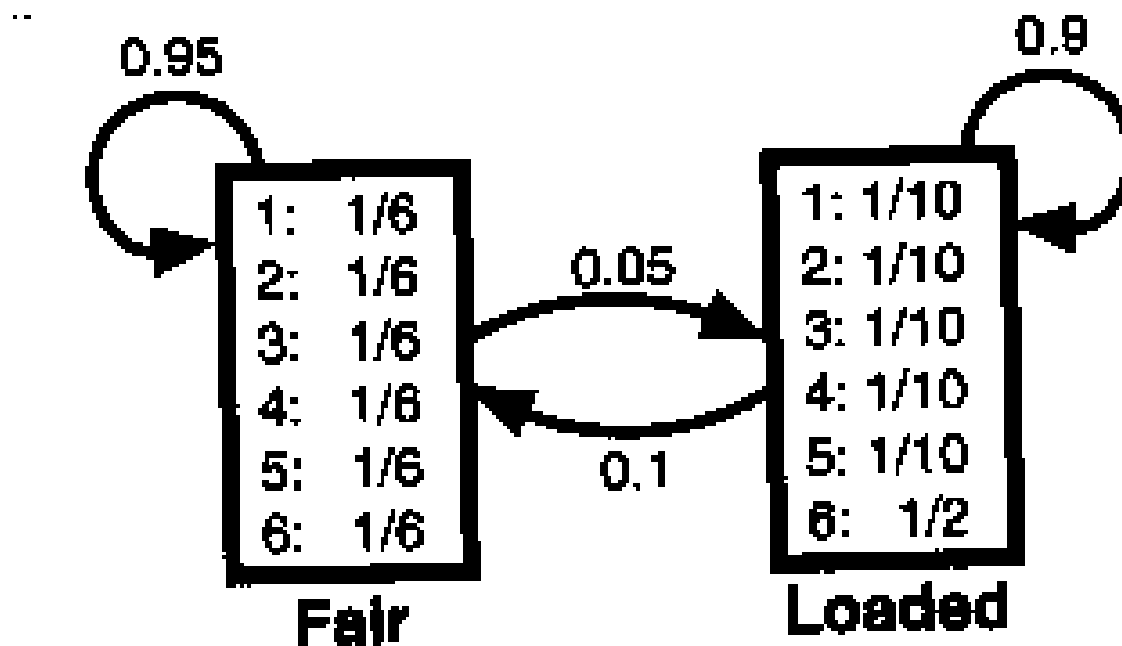
Example II: Hidden Markov Models

Modelling dependencies in one dimensional data structures, eg

- Speech recognition (Word models etc)
- Biosequences (DNA, proteins)

Example: The occasional dishonest casino (Durbin et al)

The HMM



Hidden Markov Models: Definitions

- Observations $\mathbf{y} = (y_1, y_2, \dots, y_T)$ are *independent* given the sequence of states $\mathbf{S} = (s_1, s_2, \dots, s_T)$. ie

$$P(\mathbf{y}|\mathbf{S}) = \prod_{i=1}^T P(y_i|s_i) = \prod_{i=1}^T b_{s_i}(y_i)$$

with the matrix of *emission probabilities* $b_k(l) = P(y = l | s = k)$.

- States are not observed (hidden) and generated from a *Markov chain*

$$P(\mathbf{S}) = \pi_{s_1} P(s_2|s_1) P(s_3|s_2) \dots P(s_T|s_{T-1}) .$$

- The total probability of the observed sequences is obtained by marginalization of the joint probability $P(\mathbf{y}, \mathbf{S}) = P(\mathbf{y}|\mathbf{S})P(\mathbf{S})$ over the states

$$P(\mathbf{y}) = \sum_{\mathbf{S}} P(\mathbf{y}|\mathbf{S})P(\mathbf{S})$$

For N states, there are N^T different paths in the sum!!