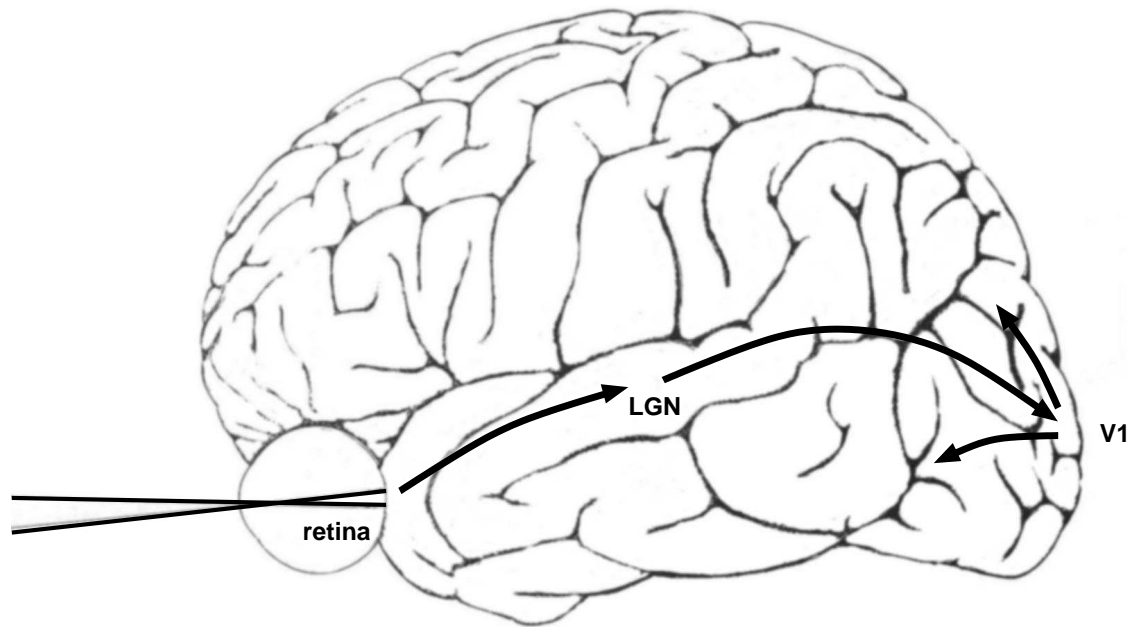


Natural Image Statistics

— A probabilistic approach to
modelling early visual processing in the cortex

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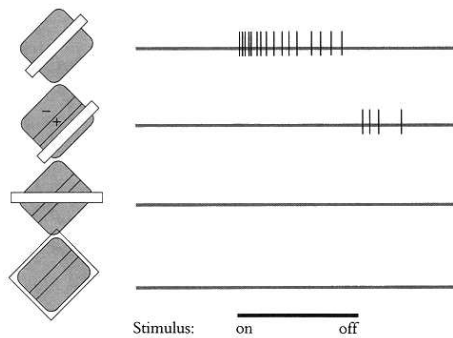
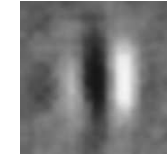
Early visual processing



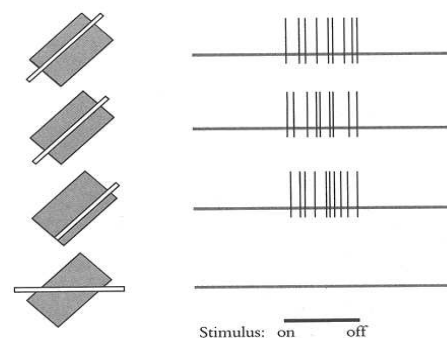
From the eye to the primary visual cortex (V1)

Simple and complex cells

- Basic dichotomy of neurons in primary visual cortex (V1)
- Simple cells modelled as linear functions of input: “Receptive fields” can be simply plotted
- Complex cells considered strongly nonlinear
 - Invariance (tolerant) to location (phase) of input
 - Modelled as sum of squares of simple cell outputs



Simple cells:
Selective to orientation
and location of the bar



Complex cells:
Tolerant to exact location

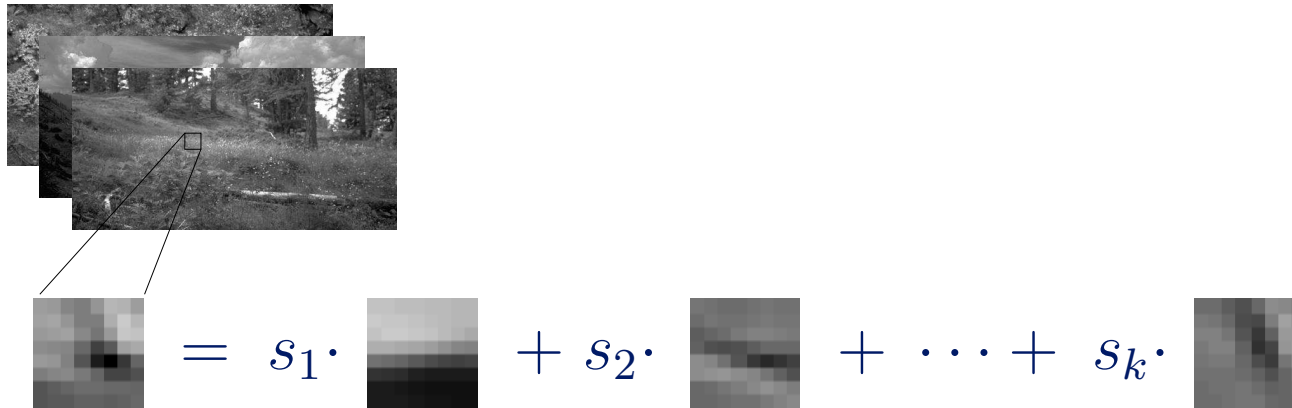
Theories of response properties of visual neurons

- Edge detection
- Joint localization in space and frequency
- Texture classification
- But: the above give only vague predictions.
- Here: Statistical-ecological approach (Barlow, 1972)
 - What is important in a real environment?
 - Natural images have statistical regularities.
 - Can we “explain” receptive fields by basic statistical properties of natural images?
 - **Emergence** : a lot of precise predictions from only a couple statistical assumptions.
- Extremely relevant to image processing /engineering

Outline of this talk:

- Statistical models that account for some properties of the (primary) visual cortex.
 - Independent Component Analysis / Sparse Coding
 - Various extensions
- Properties in visual cortex explained
 - simple cells
 - complex cells
 - spatial organization (topography)
- Multi-layer approach can predict properties beyond V1.

Linear statistical models of images



- Denote by $I(x, y)$ the gray-scale values of pixels.
- Model as a linear sum of basis vectors:

$$I(x, y) = \sum_i A_i(x, y) s_i \quad (1)$$

- What are the “best” basis vectors for natural images?

Independent Component Analysis (Jutten and Héroult, 1991)

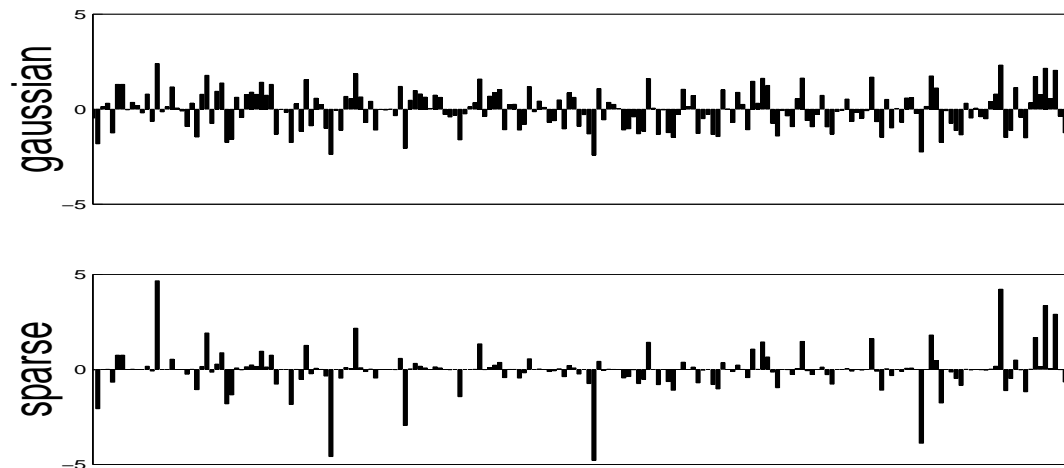
- Linear model:

$$I(x, y) = \sum_i A_i(x, y) s_i \quad (2)$$

- In ICA, we assume that
 - The s_i are mutually statistically independent
 - The s_i are **nongaussian**, e.g. sparse
 - For simplicity: number of A_i equals number of pixels
- Then, the actual basis vectors A_i can be estimated, if the data is actually generated using the linear model (Comon, 1994).
- Thus we get the best basis vectors from one statistical viewpoint.

Sparsity

- A form of nongaussianity often encountered in natural signals
- A random variable is “active” only rarely



- Outputs of linear filters are usually sparse when input is natural images.

Sparse coding and ICA

- Sparse coding (Barlow 1972): Find linear representation

$$I(x, y) = \sum_i A_i(x, y) s_i \quad (3)$$

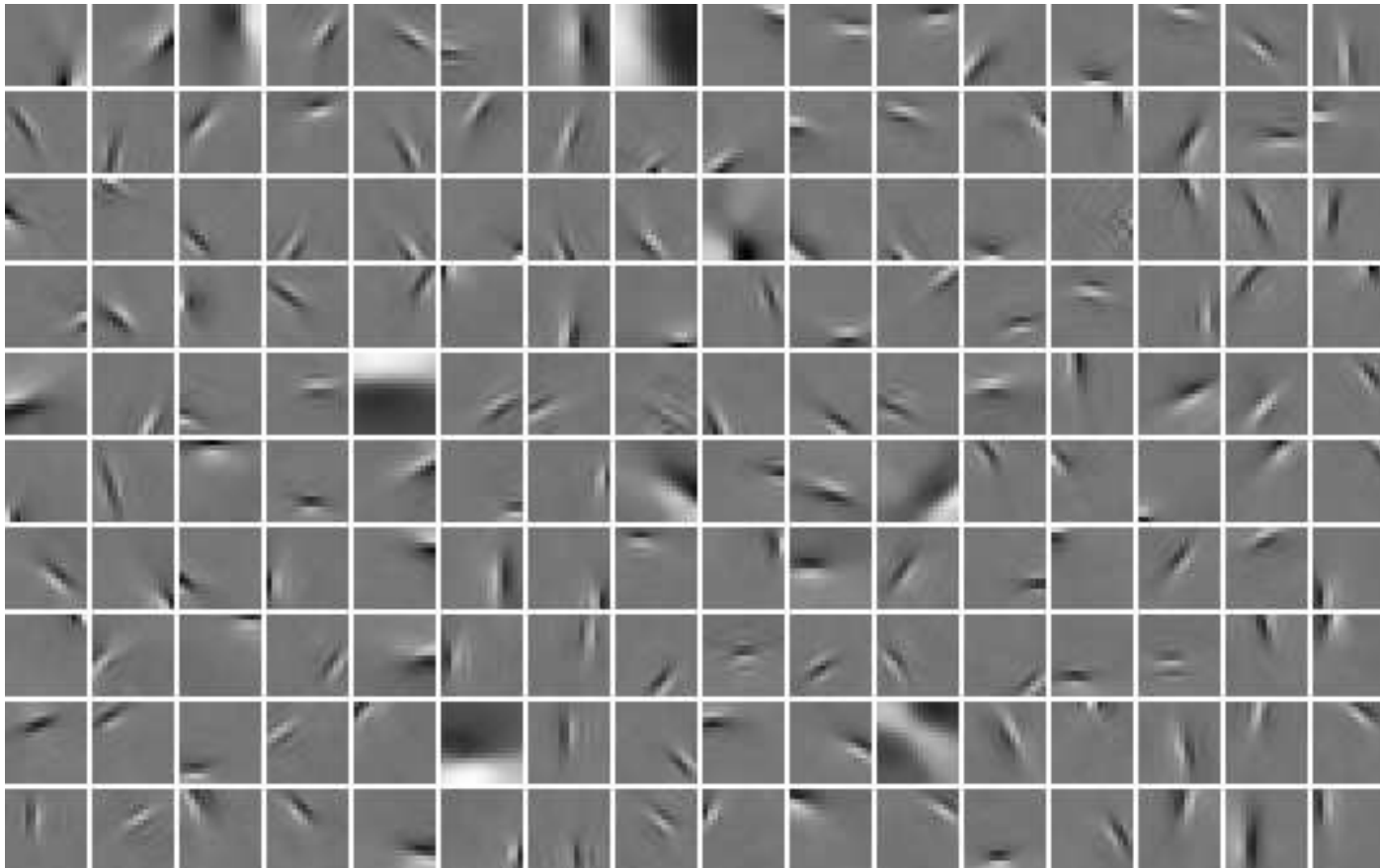
so that the s_i are as sparse as possible.

- Important property: a given data point is represented using only a limited number of “active” (clearly non-zero) components s_i .
- In contrast to PCA, active components change from image patch to patch.
- Deep result: For images, **ICA is sparse coding**.
- Vectorizing whitened image as \mathbf{x} , denoting inverse system by \mathbf{w}_i :

$$\min_{\text{orthog } \mathbf{w}_1, \dots, \mathbf{w}_n} \hat{E}\{|\mathbf{w}_i^T \mathbf{x}|\} \quad (4)$$

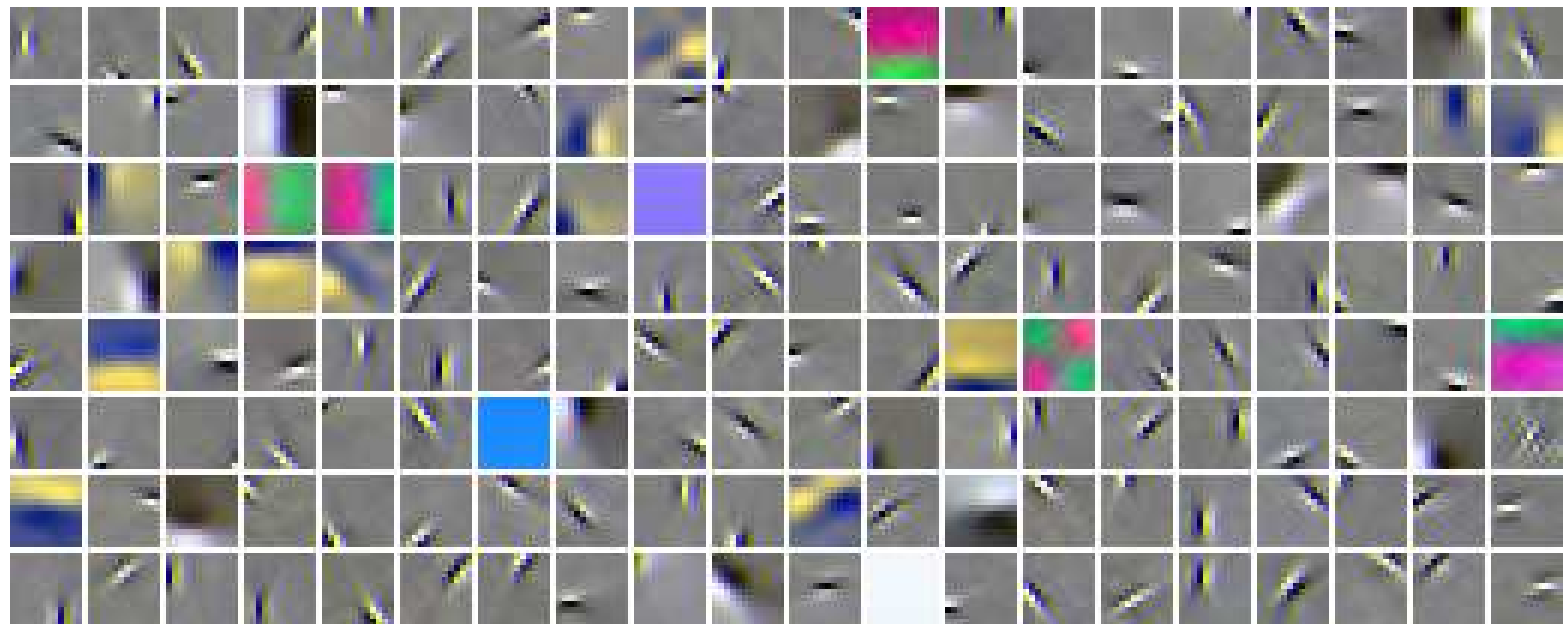
ICA / sparse coding of natural images

(Olshausen and Field, 1996; Bell and Sejnowski, 1997)



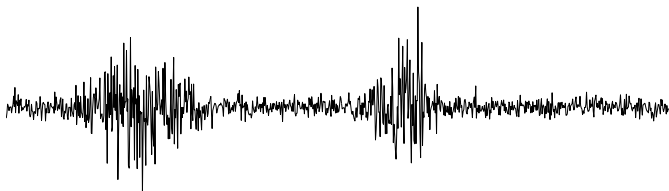
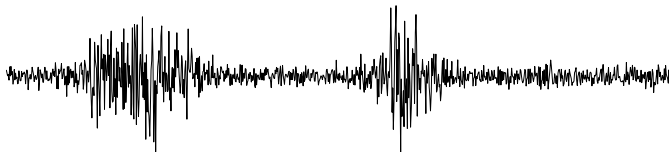
Using the FastICA algorithm (Hyvärinen, 1999)

ICA of natural images with colour (Hoyer and Hyvärinen, 2000)



Model II: Independent subspace analysis

- Components estimated from natural images are **not** really independent.
- The statistical structure much more complicated (of course!).
- Independent components cannot be found for most kinds of data: There are not enough free parameters.
- Dominant form of dependency after ICA is correlation of energies



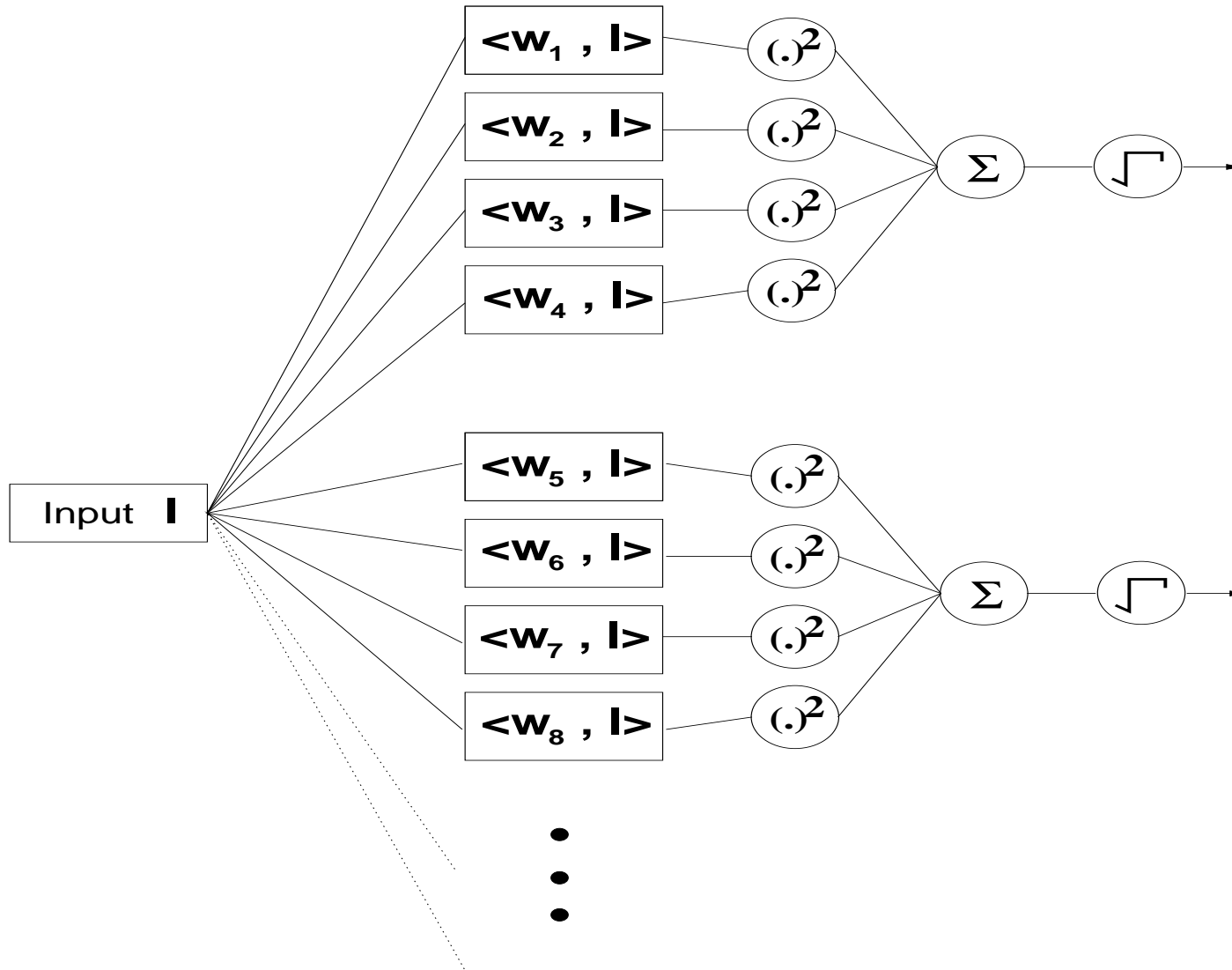
Signals which are uncorrelated but whose squares are correlated.

Using subspaces to model dependency

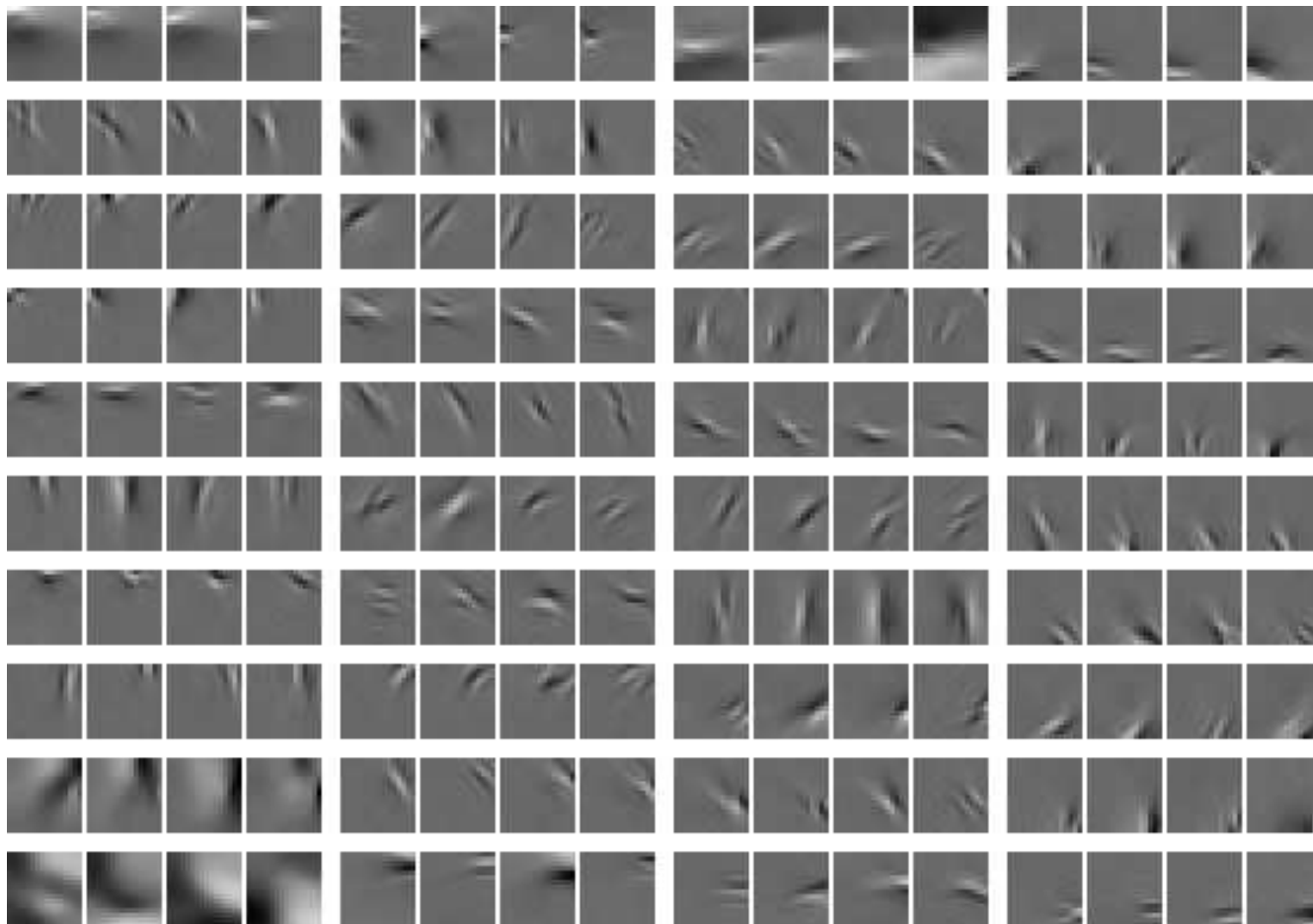
(Hyvärinen and Hoyer, 2000)

- Assumption: the s_i can be divided into groups or subspaces (Cardoso, 1998), such that
 - the s_i in the **same** group **are** dependent on each other
 - dependencies between **different** groups **are not** allowed.
- We also need to specify the distributions inside the groups
 - Use classic complex cell model, norm of projection in subspace
 - Leads to correlation of squares
 - Maximize independence / sparsity of complex cell output

Computation of features in independent subspace analysis



Independent subspaces of natural image patches

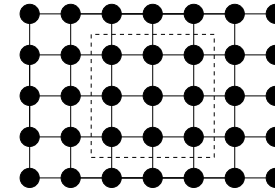


Each group of 4 basis vectors corresponds to one complex cell.

Model III: Spatial organization in V1

- In the brain, response properties mostly change continuously when moving on the cortical surface.
- We introduced Topographic ICA (Hyvärinen and Hoyer, 2001)

- Cells (components) are arranged on a **two-dimensional lattice**

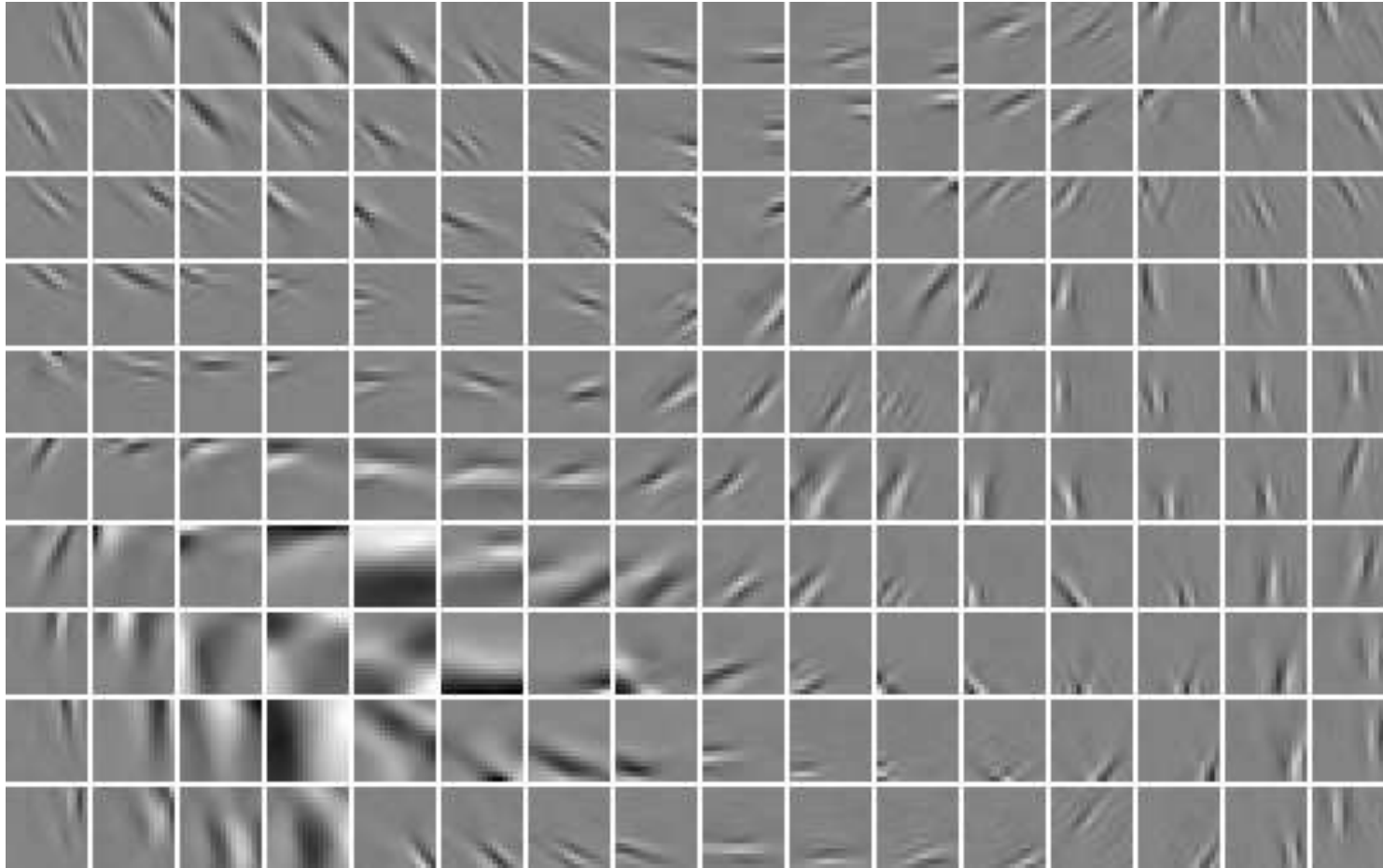


- Again, simple cell outputs are sparse, but not independent: Correlations of squares follows topography.
- Learn by maximizing likelihood which measures sparsity:

$$-\log p(\mathbf{x}|\mathbf{W}) = \sum_i \hat{E} \left\{ \sqrt{\sum_j h_{ij} (\mathbf{w}_i^T \mathbf{x})^2} \right\} \quad (5)$$

where h is distance on topographic grid.

Topographic ICA on natural image patches



Basic vectors (simple cell RF's) with spatial organization

Digression: Generalizations leading to unnormalized models

- Generalize ISA and topographic ICA by estimating two layers

$$p(\mathbf{x}; \mathbf{W}, \mathbf{M}) = \frac{1}{Z(\mathbf{W}, \mathbf{M})} \exp\left[\sum_i G\left(\sum_i m_{ij} (\mathbf{w}_j^T \mathbf{x})^2\right)\right] \quad (6)$$

- This is an unnormalized model:

Density function p is known only up to a multiplicative constant

$$Z(\mathbf{W}, \mathbf{M}) = \int \exp\left[\sum_i G\left(\sum_i m_{ij} (\mathbf{w}_j^T \mathbf{x})^2\right)\right] d\mathbf{x}$$

which **cannot be computed** with reasonable computing time

- Common problem in non-Gaussian models, e.g. ICA with more components than observed variables

$$p(\mathbf{x}; \mathbf{W}) = \frac{1}{Z(\mathbf{W})} \exp\left[-\sum_i |\mathbf{w}_i^T \mathbf{x}|\right] \quad (7)$$

Recent methods for unnormalized statistical models

- Score matching (Hyvärinen, JMLR, 2005)
 - Take derivative of model log-density w.r.t. \mathbf{x} , so partition function disappears
 - Fit this derivative to the same derivative of data density
 - Easy to compute due to an integration-by-parts trick
- Noise-contrastive estimation (Gutmann and Hyvärinen, JMLR, 2012)
 - Learn to distinguish data from artificially generated noise
 - Logistic regression learns ratios of pdf's of data and noise
 - For known noise pdf, we have in fact learnt data pdf

Another unnormalized model:

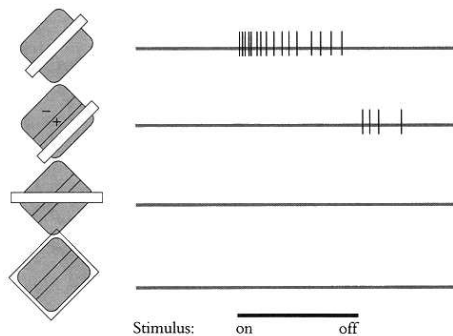
Linear correlations between components

- Many methods force components to be strictly uncorrelated
- So, any remaining linear correlations cannot be observed
- However, the “true” components could be correlated even linearly
- In ongoing work (Sasaki et al, 2013, 2014) we learn correlated components

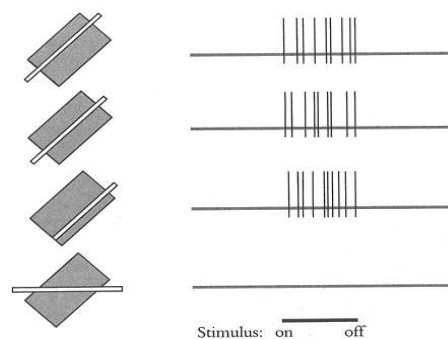
$$-\log p(\mathbf{x}|\mathbf{W}) = \sum_i \hat{E}\{|\mathbf{w}_i^T \mathbf{x}|\} + \sum_{i,j} \beta_{ij} \hat{E}\{|\mathbf{w}_i^T \mathbf{x} - \mathbf{w}_j^T \mathbf{x}|\} - \log Z(\beta_{ij}) \quad (8)$$

Model IV: Three-layer model

- Goal: Alternating selectivity and invariance in many layers
- Neurons are *selective* to certain properties of the stimulus:
 - response is strong when those properties take specific values
- Neurons are *tolerant* (invariant) to properties:
 - response does not change much when those properties change
- Example: Simple vs. complex cells in the primary visual cortex:



Simple cells:
Selective to orientation
and location of the bar

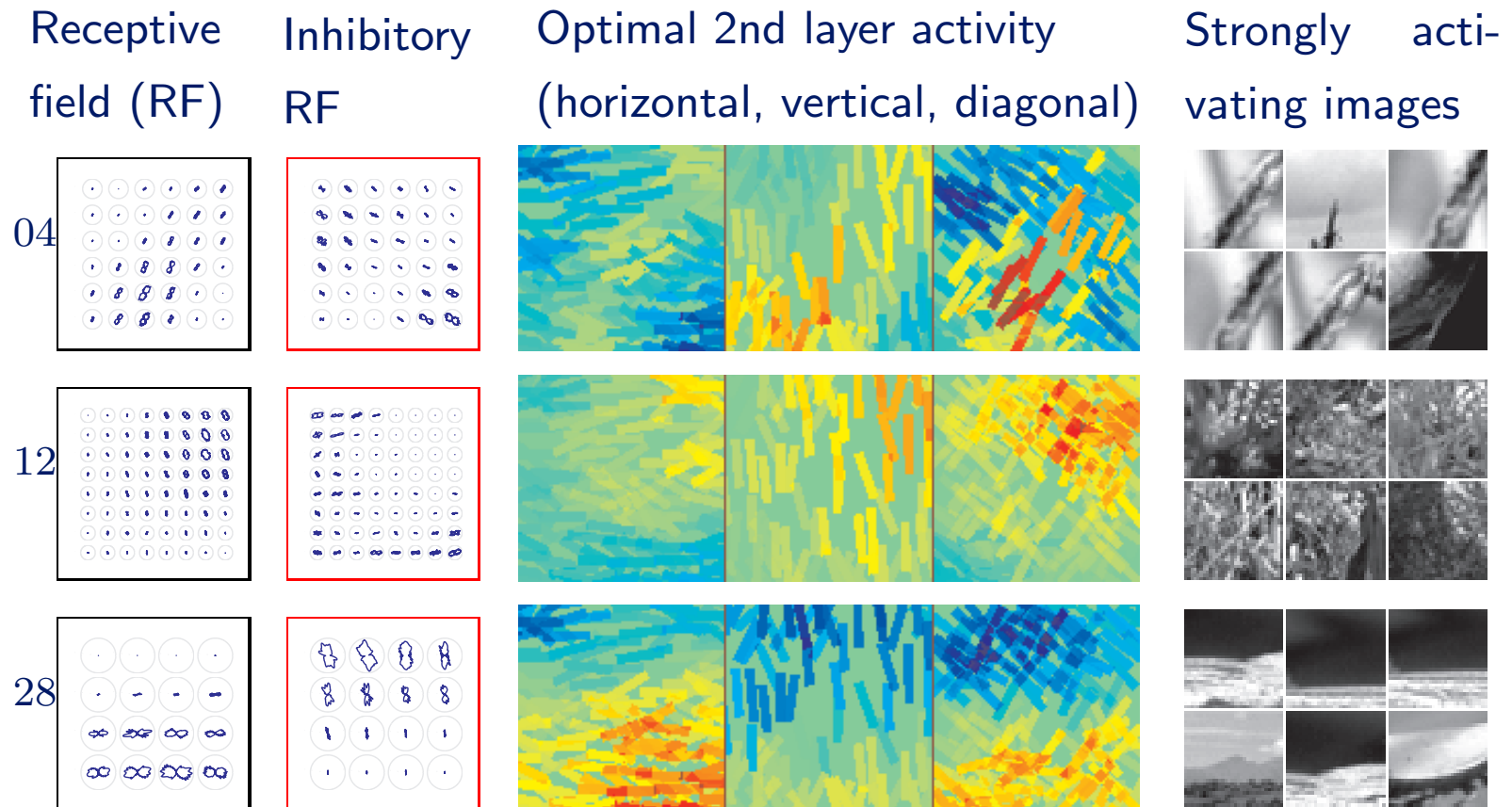


Complex cells:
Tolerant to exact location

Nonlinearities for neural selectivity and tolerance

- Selectivity has been modelled as
 - AND operation / MIN operation
- Tolerance (invariance) has been modelled as
 - OR operation / MAX operation
- Here, we use **convex** and **concave** nonlinearities
 - E.g. $(\sum_i x_i^4)^{1/4}$: similar to MAX
 - E.g. $(\sum_i x_i^{1/4})^4$: similar to MIN
- Two first layers similar to squaring + square root in complex cells
- Learning by maximization of sparsity in each layer

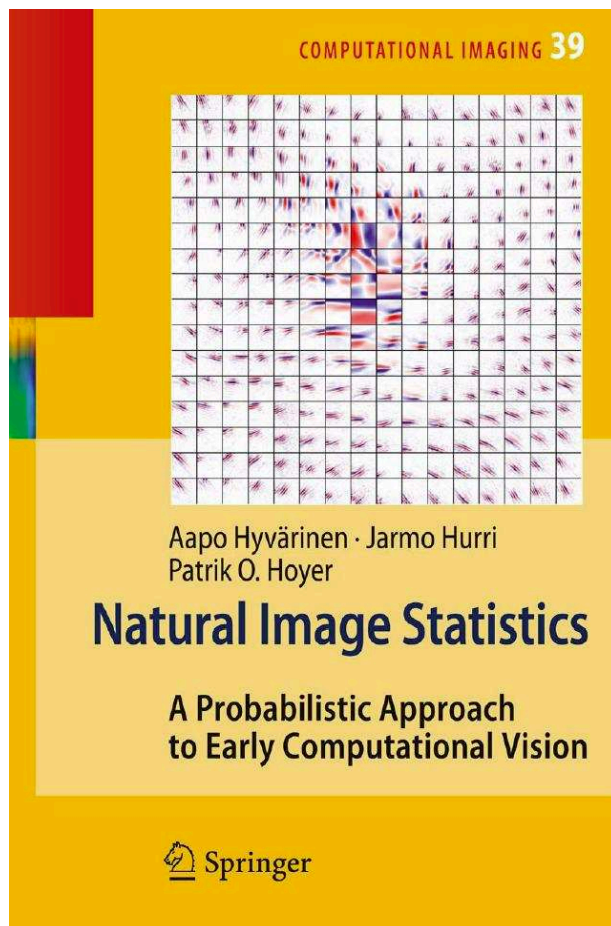
Layer three responses in three-layer model



Neurophysiological modelling vs. Deep learning

- Deep learning means neural network with many layers
- Result is often a black box: interpretation difficult
- For neurophysiological modelling, we would prefer a network where
 - The role of each unit is clear
 - All cell responses model biological responses
- Instead of blindly stacking many layers on top of each other, we must think about what each layer is doing

For more information on basic models



Conclusion

- Properties of visual neurons can be quantitatively modelled by statistical properties of natural images.
- Answers the **Why** question important in neuroscience
- Simple cell receptive fields can be learned by maximizing sparsity / independence of linear filters.
- By modelling dependencies between simple cell outputs we can model complex cells and topography
- Three-layer models can use alternating selectivity and invariance
- Theoretical development on estimation of unnormalized models
- Many applications in image processing and computer vision