On the choice of a sparse prior

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Abstract

An emerging paradigm analyses in what respect the properties of the nervous system reflect properties of natural scenes. It is hypothesized that neurons form sparse representations of natural stimuli: Each neuron should respond strongly to some stimuli while being inactive upon presentation of most others. For a given network, sparse representations need fewest spikes, and thus consume the least energy. To obtain optimally sparse responses the receptive fields of simulated neurons are optimized. Algorithmically this is identical to searching for basis functions that allow coding for the stimuli with sparse coefficients. The problem is thus identical to maximising the log likelihood of a generative model with prior knowledge of natural images. It is found that the resulting simulated neurons share most properties of simple cells found in primary visual cortex. Thus, forming optimally sparse representations is the most compact approach to described simple cell properties.

Many ways of defining sparse responses exist and it is widely believed that the particular choice of the sparse prior of the generative model does not significantly influence the estimated basis functions. Here we examine this assumption more closely. We include the constraint of unit variance of neuronal activity, used in most studies, into the objective functions. We then analyse learning on a database of natural (cat-camTM) visual stimuli. We show that the effective objective functions are largely dominated by the constraint, and are therefore very similar. The resulting receptive

fields show some similarities but also qualitative differences. Even in the region where the objective functions are dissimilar, the distributions of coefficients are similar and do not match the priors of the assumed generative model. In conclusion, the specific choice of the sparse prior is relevant, as is the choice of additional constraints, such as normalization of variance.

Introduction

It is important to analyze in what respect properties of sensory systems are matched to the properties of natural stimuli (Barlow 1961). Many recent studies analyse simulated neurons learning from natural scenes and compare their properties to the properties of real neurons in the visual system (Atick 1992; Dong and Atick 1995; Fyfe and Baddeley 1995; Bell 1996; Olshausen and Field 1996; Bell and Sejnowski 1997; Olshausen and Field 1997; Blais, Intrator et al. 1998; van Hateren and Ruderman 1998; Van Hateren and van der Schaaf 1998; Hyvärinen and Hoyer 2000; Lewicki and Sejnowski 2000; Kayser, Einhäuser et al. 2001; Olshausen 2001; Simoncelli and Olshausen 2001; Einhäuser, Kayser et al. 2002). Most of these studies follow the independent component analysis (ICA) paradigm: An explicit generative model is assumed where hidden, nongaussian generators are linearly combined to yield the image I(t) from the natural stimuli.

$$I(x, y, t) = \sum_{j} a_{j}(t) \phi_{j}(x, y)$$
(1)

The coefficients are assumed to occasionally be nonzero but about zero most of the time. This assumption is formulated probabilistically as $p(a_i)$ and referred to as a sparse prior is assumed for the coefficients $a_i(t)$. The log likelihood of the image given the model is very expensive to compute and is therefore typically approximated by an objective function Ψ . This objective Ψ is subsequently maximized using

standard optimisation algorithms. Frequently used options are scaled gradient descent, conjugate gradient descent and even faster methods like the fastICA (Hyvärinen 1999) method. The properties of the optimized neurons are subsequently compared to properties of real neurons in the visual system. It is found that these simulated neurons share selectivity to orientation, spatial frequency, localisation and motion with simple cells found in primary visual cortex. A number of further studies even directly addresses sparse coding in experiments and shows that the brain is indeed encoding stimuli sparsely (Baddeley, Abbott et al. 1997; Vinje and Gallant 2000; Willmore and Tolhurst 2001). Thus, sparse coding offers an approach that leads to simulated neurons with properties which compare well to those of real neurons.

An important property of the ICA paradigm was demonstrated in a seminal contribution by Hyvärinen and Oja (1998). Given a known and finite number of independent non-gaussian sources the resulting basis functions do not significantly depend on the chosen nonlinear objective. Applied to the problems considered here this would predict, that the specific definition of sparseness does not have any influence on the resulting basis functions.

However, do natural stimuli match the assumptions of this theorem? Firstly a finite number of generators in the real world is not obvious, and their number is definitely not known. Secondly, the generators in the real world are combined nonlinearly for obtaining the image. Occlusions, deformations and so forth make the generative process non-linear, making it intractable to directly invert it. Thirdly objects in natural scenes are not independent of each other, the real world is highly ordered and shows a high degree of dependence. The real world situation is therefore often different to the situation addressed in the paper of Hyvärinen and Oja (1998).

Nevertheless it is widely believed that the choice of the sparse objective does not significantly influence the estimated basis functions. Here we examine this assumption more closely. There are two factors that influence the objective to be optimised. The first part captures how well the assumed sparse prior on the coefficients is met. The second part captures how well the neurons collectively code for the image, i.e. how well the image can be reconstructed from their activities. Here we investigate the form of the combined effective objective functions. Furthermore, we analyse the distribution of activities (coefficient values) and in how far they match the specific priors. Finally, we compare the form of the basis functions obtained when optimizing different objectives.

Methods

Relation of generative models to objective function approaches

Here we relate probabilistic generative models of natural stimuli to the optimisation scheme used throughout this paper.

Let I be the image, ϕ a set of variables describing the model and a(t) a set of statistical variables, called coefficients, describing each image in terms of the model. The probability of the image given the data is calculated as:

$$p(I \mid \phi) = \int p(I \mid \phi, a) p(a) da$$
⁽²⁾

This integration however is typically infeasible. Olshausen and Field (1997) used the idea of maximizing an upper bound of this probability: If the probability of the image given ϕ and a is highly peaked at some maximal value we can rewrite (2) and obtain:

$$p(I \mid \phi) \approx p(I \mid \phi, a_{\max}) p(a_{\max})$$
(3)
The log-likelihood is thus:

$$\log p(I \mid \phi) \approx \log p(I \mid \phi, a_{\max}) + \log p(a_{\max})$$
(4)

If we assume Gaussian noise on the image the first term is proportional to the squared reconstruction error. The second term contains the (sparse) prior on the distribution of the generative model.

Typically the log likelihood of the data given the model is maximized. This leads to an optimisation problem on the following class of objective functions:

 $\Psi_{prob} = \Psi_{square\ error} + \Psi_{prior}$ (5) The latter term represents the a priori information about the coefficients a, the first term measures deviations from the models stimulus reconstruction.

The process sketched above can often be inverted. If Ψ consists of the standard Ψ_{error} ,

and Ψ_{prior} and is sufficiently peaked then we simply obtain $p(a) = e^{\Psi_{error}(a)}$.

The reconstruction error term

Here we simplify the minimum square error term so that different approaches can be

compared in a unified framework.

The reconstruction error is defined as

$$\Psi_{error} = -\left\langle \sum_{j} \left(I_{j} - \sum_{i} a_{i} \phi_{ij} \right)^{2} \right\rangle$$
(6)

Assuming unit variance of the input this can be simplified to:

$$\Psi_{error} = -\left\langle 1 - 2\sum_{i,j} a_{i,0} a_i + \sum_{i,j,k} (a_i a_j) (\phi_{ki} \phi_{kj}) \right\rangle$$
(7)

Where $a_{i,0} = \mathbf{I} \boldsymbol{\varphi}_{i}$ is the feedforward activity. If we furthermore assume whitened input $\langle I_i I_j \rangle = \delta_{ij}$ and linear activities then we can further simplify the last term:

$$\Psi_{error} = -1 + 2\sum_{i,j} \left\langle a_{i,0} a_i \right\rangle - \sum_{i,j} \left\langle a_i a_j \right\rangle \left\langle a_{i,0} a_{j,0} \right\rangle \tag{8}$$

The system is defined in a purely feedforward way since it does not directly depend on the input. In a linear system where each unit has unit variance this simplifies, after omitting constant terms, to:

$$\Psi_{error} \approx -\sum_{i,j} \left\langle a_i a_j \right\rangle^2 = \sum_{i,j} CC(a_i, a_j)^2$$
(9)

Here CC denotes the coefficient of covariation. Thus, if the system is linear, decorrelating the outputs is equivalent to minimizing the reconstruction error.

This formalism also captures methods that constrain the a to be uncorrelated which is identical to making the sparseness term small compared to the error term or having a noise free model (Olshausen and Field 1996). While most models effectively share the same $\Psi_{square\ error}$ there is a wide divergence for the sparseness objective $\Psi_{prior}(a) = \log(p(a))$.

Stimuli

Out of the videos described in (Kayser, Einhäuser et al. 2001) 40.000 30 by 30 patches are extracted from random positions and convolved with a Gaussian Kernel of standard deviation 15 pixels to minimize orientation artefacts. They are whitened to avoid effects of the second and lower order statistics that are prone to noise influences. Only the principal components 2 through 100 are used for learning since they contain more than 95% of the overall variance. Component 1 was removed since it contained the mean brightness.

The weights of the simulated neurons are randomly initialised with a uniform distribution in the whitened principal component space. For computational efficieny they are orthonormalized before starting the optimisation..

Decorrelation

As argued above, the considered models should allow the correct reconstruction of the image and thus minimize the squares of the coefficients of covariation CC between pairs of coefficients of different basis functions. The standard deviation can furthermore be biased to be 1. All optimisations are therefore done with the following objective function added.

$$\Psi_{decorr} = -\sum_{i,j} CC(a_i, a_j) - \sum_i (1 - std(a_i))^2$$
(10)

The first term biases the neurons to have distinct activity patterns while the latter term effectively normalizes the standard deviation. In the following we refer to this joint term as decorrelation term. If the variances of the coefficients are 1 then it is identical to the square error term. If this term is strong and the set of neurons is not overcomplete it removes all correlations between neurons.

Analysed objective functions

In the literature a large number of different definitions of objective functions can be found. Each results from a different way of defining sparseness. Here we analyze 6 different popular definitions:

$$\Psi_{Skewness} = 0.1* \sum_{i} \frac{\langle a_i^3 \rangle}{\langle a_i^2 \rangle^{3/2}}$$
(11)

$$\Psi_{Kurtosis} = 0.1* \sum_{i} \frac{\langle a_i^{\,4} \rangle}{\langle a_i^{\,2} \rangle^2} \tag{12}$$

$$\Psi_{Abs} = -\sum_{i} \left\langle abs(a_i) \right\rangle \tag{13}$$

$$\Psi_{ex2} = \sum_{i} \left\langle e^{-a_i^2} \right\rangle \tag{14}$$

$$\Psi_{Cauchy} = -\sum_{i} \left\langle \log(1+a_i^2) \right\rangle \tag{15}$$

$$\Psi_{CorrIJ} = -\sum_{i,j} \left(\left\langle a_i^2 a_j^2 \right\rangle - \left\langle a_i^2 \right\rangle \left\langle a_j^2 \right\rangle \right)^2$$
(16)

 $\Psi_{Skewness}$ and $\Psi_{Kurtosis}$ usually have higher values than the other objectives and are therefore divided by 10 to avoid them being overly strong compared to the decorrelation objective. The BCM learning rule (Bienenstock, Cooper et al. 1982) is another interesting algorithm that can be put into an objective function framework. It is largely identical to the $\Psi_{Skewness}$ objective (Intrator and Cooper 1992).

Optimization

The optimisation algorithm uses the above objective functions and their derivatives with respect to the weights. These derivatives are often complicated functions containing a large number of terms. We found it very useful to verify these numerically. The objective functions are maximised using 50 iterations of RPROP (resilient backprop Riedmiller and Braun 1993) with $\eta^+ = 1.2$ and $\eta^- = .5$, starting at a weightchange parameter of 0.01. We observe a significantly faster convergence of the objective function compared to scaled gradient descent. It is interesting to note that RPROP, where each synapse stores its weight and how fast it is supposed to change, is a local learning algorithm that could be implemented by neural hardware.

Results

The effective priors

Each of the analyzed objective functions is plotted in figure 1. They are divided into two groups. Ψ_{Abs} , $\Psi_{exp(-a^2)}$ and Ψ_{Cauchy} punish coefficients that differ from zero in a graded manner. $\Psi_{Skewness}$ and $\Psi_{Kurtosis}$ to the contrary reward high coefficients. The effective objective that is optimized by the system, however, consists of two terms. The first term represents the sparseness objective as plotted in Figure 1. At first sight it their wildly divergent properties could be expected to lead to differing basis functions, and it is counterintuitive to assume that all these objectives lead to similar basis functions. It however is necessary to take into account the term of the objective function that biases the neurons to avoid correlations and to have unitary variance (see methods). This term turns into a constraint when the neurons are directly required to be uncorrelated. This term should therefore be including into the prior. Figure 2 shows the objective function measured after convergence. The decorrelation term Ψ_{Decorr} has a strong influence on Ψ_{Abs} , the $\Psi_{exp(-a^2)}$ and the Ψ_{Cauchy} objective function. All these functions bias the coefficients to have small variance. The decorrelation term ensures that the variance does not approach zero. The definition of the decorrelation term results in a parabola shaped function that is added to the original objective functions. The resulting objective functions are almost identical for these three priors for higher coefficient values. After considering this effect they are also very similar to $\Psi_{Kurtosis}$ and the values of $\Psi_{Skewness}$ for positive activities. All these priors punish intermediate activities while preferring activities that are either small or very large. The priors are very similar for high values and only differ for smaller values.

The distribution of the coefficients

We want to investigate in how far the optimisation process leads to a distribution of coefficients which actually matches the log prior or the objective function. We furthermore want to know if the different objective functions lead to different distributions of coefficients.

Observing the histogram of the coefficients in response to the natural stimuli after convergence reveals high peaks at zero for all objective functions. They are thus very dissimilar to their priors. The coefficients a in response to different patches are not independent from each other. If any of the basis functions is changed then all the coefficients change. Since the number of coefficients is only a fraction of the number of stimuli used, it is impossible for the distribution to perfectly follow the prior. We will therefore consider the logarithm of the deviation of the distribution of coefficients divided by the distribution of coefficients before learning:

$$d_{rel} = \log \frac{\sum_{i} p\left(a_i \mid \phi_{after \, learning}\right)}{\sum_{i} p\left(a_i \mid \phi_{start}\right)}.$$
(17)

We cannot directly measure p and thus instead use the number of observations divided by the number of overall stimuli. Using the relative distribution instead of the original distribution automatically corrects for the distribution of contrasts in the natural scenes. It thus converts the highly peaked distribution of coefficients into a rather flat function. Before learning the basis functions are random and the distribution of the coefficients therefore is identical to the distribution of contrast in natural scenes. By dividing by the distribution before learning we correct for these effects. Similar behaviour could be achieved using nonlinear neurons that feature lateral divisive inhibition as in (Schwartz and Simoncelli 2001). Figure 3 shows that the relative distribution for large coefficients is well fit by the prior. It is interesting to note that the prior for larger coefficients is the decorrelation term that is not directly visible in most papers addressing sparse coding.

To analyze the differences between the different priors, we are interested in the range of small coefficients where the specific type of the prior matters most. Figure 4a shows that the relative distributions in this region cannot be fit by the prior. The peaks of the functions have conserved shape and only the position of their maximal value is determined by the objective function (Figure 4b). The distributions are largely identical even for coefficient values where the objective functions are considerably different.

Comparison of basis functions

The optimisation algorithm yields the basis functions that are the analogue of receptive fields of real neurons (Figure 5). A number of important similarities and differences can be observed. Some of the properties are identical for all the different

priors. All of them lead to receptive fields that are localised in orientation and spatial frequency. This can be understood from the fact that all analyzed objective functions reward high absolute values of the activity. High contrast regions are associated with defined orientation (Einhäuser, Kayser et al. 2002) explaining why all learned receptive fields are oriented. All of the considered objectives also lead to localisation in space if the decorrelation term is strong enough.

However also a large number of important differences exist between the resulting receptive fields: The differences between the functions punishing non-zero activities is small, they mostly exhibit small variations in the smoothness and the size of the receptive fields. This is not surprising knowing that their objectives only slightly differ for small coefficients. $\Psi_{_{Kurtosis}}$ is known to be prone to overfitting (Hyvärinen and Oja 1997). Due to its sensitivity to outliers, maximising $\Psi_{_{Kurtoxis}}$ leads to basis functions that can be expected to be specific to the real natural stimuli chosen in our study. Using less natural stimuli such as pictures from man made objects would be likely to significantly change the resulting receptive fields obtained from maximising $\Psi_{Kurtosis}$. While optimising the objective functions that punish non-zero coefficients lead to small localised, Gabor-type receptive fields, optimising $\Psi_{Kurtosis}$ leads to more elaborate filters. Optimising $\Psi_{Skewness}$ also leads to interesting basis functions. All of the basis functions show black lines on bright background. This is a very common feature in our dataset since many of the pictures show trees and branches in front of the bright sky (observation from the raw data). We attribute the possibility of learning from $\Psi_{{\it Skewness}}$ to these properties. It is an interesting violation of contrast reversal invariance. It would seem that many statistics of natural scenes are conserved if the contrast is reversed. If the statistics is invariant with respect to contrast reversal then all distributions of the coefficients a need to be symmetric and $\Psi_{Skewness}$ would not be a valid objective. The skewed receptive fields nevertheless share orientation and spatial frequency selectivity with the other objectives.

The methods analyzed in this paper all belong to the class of independent component analysis. They are however necessarily only an approximation to statistical independence. We therefore compare the receptive fields with another prior that more directly measures independence (Comon 1991). If the coefficients are independent then the coefficients as well as their squares should also be uncorrelated. We thus maximise Ψ_{CorrlJ} that punishes correlations between the squares. After a much slower convergence of 500 RPROP iterations the estimated basis functions are shown in Figure 5. They are also localised in orientation and spatial frequency. Their properties lie in between those obtained maximising $\Psi_{Kurtosis}$ and those obtained maximising the objective functions that punish non-zero activities. All variants of ICA analyzed here lead to basis functions that share some basic properties while having some individual characteristics.

Discussion

What general approach should be used in studying different systems of coding and learning? Above we shortly described the relation of generative model and objective function approaches. There are two obvious ways of comparing such learning systems. 1) It is possible to interpret the system as performing optimal regression with some a priori information. In this framework a generative model is fitted to the data.

2) It is possible to interpret that the system's task is to learn to extract variables of relevance from input data. In this interpretation the objective function is central since it measures the quality or importance of the extracted variables.

Both formulations come with inherent weaknesses and strengths.

1) The generative models used for describing the data are always so simple, for example linear, that they can not adequately describe the complexity of the real world. When fitting a generative model to data it is furthermore often infeasible to directly optimize the log likelihood of the data given the model. Instead it is typically necessary to simplify a sparse prior to an objective function that can be optimised efficiently (but see Olshausen and Millman 2000). This step can in fact lead to objective functions that seem counterintuitive. The objective function used in the model proposed by Hyvärinen and Hoyer (2000) for the emergence of complex cells derives from a sparse prior. The resulting objective function however can be interpreted as minimizing the average coefficient a and thus the number of spikes. This is equivalent to minimising the overall number of spikes fired by the neuron and thus their energy consumption. This property was not visible from analysing the prior. The choice of a prior furthermore is often arbitrary and as shown in this paper often far from the resulting distributions. The estimated generative model can however be a strong tool for analysing, improving and generating pictures. Image processing tools like super resolution (Hertzmann, Jacobs et al. 2001), denoising (Hyvärinen 1999) as well as sampling pictures from the learned distribution (Dayan, Hinton et al. 1995) is straightforward once such a model is learned.

2) When optimising an objective function the particular choice might often seem arbitrary because it needs to be indirectly deduced from evolutionary or design principles. Following these ideas the brains task is to extract relevant information from the real world while minimising its energy consumption (Barlow 1961). Objective functions can be interpreted as heuristics that measure the value of data and the price of computation in this framework. The energy consumption might be captured by a variant of the sparseness objective since each spike comes with an associated energy consumption (Attwell and Laughlin 2001). The Ψ_{Abs} objective for example punishes the average value of the coefficients that can be associated with the number of spikes and thus the energy consumption. One simple heuristic for measuring the usefulness of data might be temporal smoothness or stability (Földiak 1991; Kayser, Einhäuser et al. 2001). It derives from the observation that most variables that are important and that we have names for change on a timescale that is slow compared to for example the brightness changes of sensors of the retina. The advantage of the objective function approach is that hypotheses for the objectives of the system can sometimes directly be derived from evolutionary ideas while allowing comparison of the results to the properties of the animal's nervous system.

In the objective function approach also used in the present study, it is directly visible that the decorrelation term strongly influence learning. It ensures the normalization of the standard deviation of the coefficients. When designing systems of sparse learning it is thus also important to also take into account the way the system is normalized. Considering this normalisation makes it far easier to understand similarities and differences between objective functions, such as the similarities between the results of the Kurtosis and the Cauchy simulations.

Sparse coding and Independent component analysis are powerful methods that have many technical applications in dealing with real world data (c.f. Hyvärinen 1999). Their strength is that they do not merely depend on statistics of second order (as does PCA) that can easily be created by uninteresting noise sources. Its most impressive applications are blind deconvolution (Bell and Sejnowski 1995), blind source separation (Karhunen, Cichocki et al. 1997), the processing of EEG (Makeig, Westerfield et al. 1999; Vigario and Oja 2000) and fMRI (c.f. Quigley, Haughton et al. 2002) data as well as denoising (Hyvärinen 1999). Using heuristics that derive from the idea of data value might allow designing better objective functions for ICA. It could lead to algorithms that can better replicate physiological data (Kayser, Einhäuser et al. 2001) and potentially lead to outputs that are more useful as input to pattern recognition systems.

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Figure Captions

Figure 1:

The sparseness objectives are shown as a function of the value of the coefficient. The original forms of the objective functions as used in most papers are shown. For better comparison all of them are scaled to the same interval.

Figure 2:

The full objective functions, including the decorrelation term, are shown as a function of the coefficient. The objective function was only evaluated after convergence of the network, which is necessary since the decorrelation term is a function of the basis functions. For better comparison all objectives were scaled to the same interval. Figure 3:

The relative distribution of coefficients after convergence is shown as solid lines. The objective functions are shown as dotted lines.

Figure 4:

A) The same graphs as in figure 3 are shown, zoomed into the range -2.5...2.5 to depict the details for small coefficient values. The relative distribution of coefficients after convergence is shown as solid lines. The objective functions are also shown as dotted lines.

B) The relative distributions are shown, all aligned to their maximal value.

Figure 5:

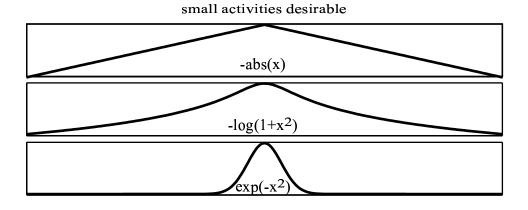
Typical examples of the resulting basis functions are depicted, each derived from the respective objective function .

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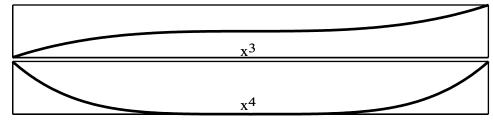
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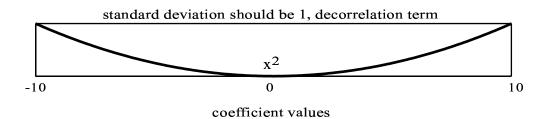
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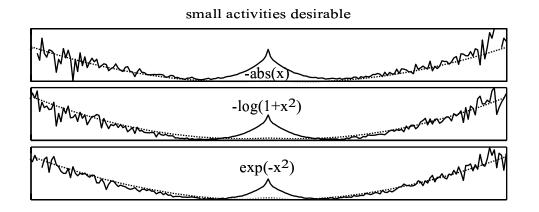
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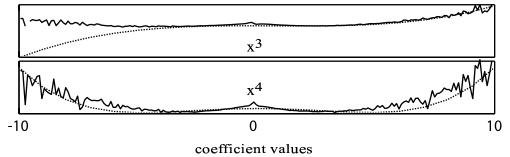
large activities desirable



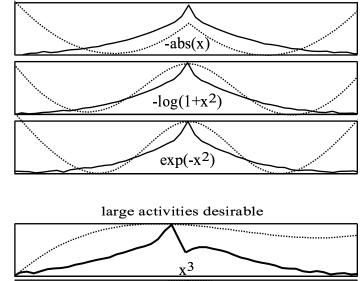


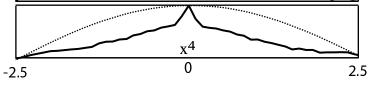


large activities desirable

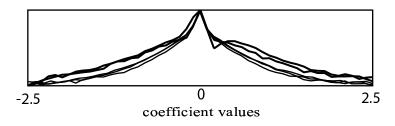


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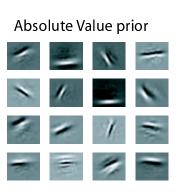




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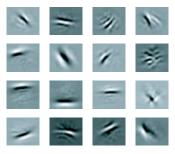
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Decorrelating the squares

