

Learning Wavelets Compositional Patterns in Sparse Representations of Natural Images

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Abstract

Sparsity is a fundamental principle in applied mathematics and statistics. It also plays a crucial role for representing natural images. In particular, a popular representational scheme for natural images is wavelets sparse coding, where each image is represented by a linear superposition of a small number of wavelets selected from a large dictionary of wavelets. The wavelets are usually localized, oriented and elongated basis functions organized in multiple scales. They are often called “atoms,” and the corresponding sparse coding is called “atomic decomposition.” In this article, we go beyond atomic decomposition by learning recurring compositional patterns formed by the wavelets in the sparse representations of natural images. If each wavelet is considered an atom, then a composition of a number of wavelets can be considered a “molecule.” We represent each compositional pattern by an active basis model, which is a composition of a small number of wavelets automatically selected from a given dictionary. The selected wavelets are allowed to perturb their locations and orientations so that the linear basis formed by the selected wavelets become active and the active basis forms a deformable template. For a given set of training images, our method learns a dictionary of wavelets compositional patterns in the form of active basis templates, so that each training image can be represented by a small number of templates that are translated, rotated, scaled and deformed versions of the active basis templates in the dictionary. Experiments show that our method is capable of learning dictionaries of meaningful wavelets compositional patterns from natural images.

Keywords: Compositionality, Group sparsity, Sparse coding, Unsupervised learning.

1 Introduction

Recent years have seen an explosion of research activities on sparsity as well as structured sparsity in applied mathematics, statistics and machine learning. See the recent books by Buhlmann and van de Geer [3] on the statistical side, and Elad [16] on the signal processing side, for timely reviews of the rapidly growing literature. Sparsity and structured sparsity also play a fundamental role in representing natural images, which contain bewildering varieties of patterns [46]. In a sparse representation scheme of natural images, there is usually a dictionary of representational elements, like a dictionary of “words,” so that each natural image can be represented by a small number of elements or words selected from the dictionary. Of course, for different images, different sets of elements or words will be selected for representation. Such sparse representations can be potentially useful for image processing and understanding.

The goal of this paper is to develop an unsupervised learning method to learn dictionaries of representational units from various types of natural images such as textures and objects. Our representational scheme is built upon wavelets representations of natural images, where our representational units are in the form of commonly occurring compositional patterns of wavelets. These compositional patterns are essentially shape templates. Intuitively, if we consider the dictionary of wavelets as an alphabet, then the dictionary of the wavelets compositional patterns can be considered a dictionary of words.

1.1 Wavelets representation, atomic decomposition, and sparsity

In wavelets representation of natural images, each image is represented by a linear superposition of wavelets in a dictionary. The number of wavelets in the dictionary may be the same as the dimension of the images, i.e., the number of pixels in the image domain, and the wavelets in the dictionary may even form an orthogonal basis. But more often than not, the number of wavelets in the dictionary is several folds larger than the dimension of the images. In this case, the dictionary is said to form an over-complete basis. An advantage of such an over-complete dictionary is that the elements in this rich dictionary can afford to be specific enough so that each image can be represented by only a small number of wavelets selected from the dictionary. An associated complication, however, is that in order to represent an image, it is necessary to select wavelets from the dictionary by some iterative algorithm. In statistics and machine learning, this is the variable selection problem in the

so-called $p > n$ linear regression setting [6], where n corresponds to the dimension of the image, and p corresponds to the number of wavelets in the dictionary. A popular method for wavelets selection is matching pursuit [35], which is a greedy algorithm that selects one wavelet in each iteration, which seeks the maximal reduction in the least squares reconstruction error. A related method is basis pursuit [8] or Lasso [48], which accomplishes variable selection by solving a penalized least squares problem where the penalty is in the form of the ℓ_1 norm of the wavelet coefficients. Such a norm has the effect of inducing sparsity in wavelet coefficients, while maintaining the convexity of the optimization problem. The ℓ_1 norm can be considered a convex relaxation of the ℓ_0 pseudo-norm that is a more direct measure of sparsity [15]. The ℓ_0 sparsity can be approximated more closely by non-convex penalties such as SCAD [18], although this leads to a potentially multi-modal objective function. The variable selection problem can also be treated in a Bayesian framework and solved by MCMC computation [25].

The dictionaries of wavelets can be designed under the guidance of sparsity. For instance, Donoho and co-authors designed various dictionaries of wavelets such as edgelets [14], wedgelets [13], ridgelets [4], curvelets [5], and beamlets [27] etc., and studied the sparsity of such dictionaries for representing various functional classes. These dictionaries of wavelets are all localized, elongated and oriented, and they are organized at multiple scales or resolutions. Such geometrical elements provide sparse representations of geometrical structures such as edges and region boundaries in natural images. The functional classes considered in this line of research reflect such geometric aspects of natural images.

The wavelets are often called “atoms” in the literature [12], meaning that they are basic representational elements that cannot be further decomposed. The sparse representation based on the atoms are called “atomic decomposition.” In this article, we use the words “wavelets,” “basis functions,” “basis elements” and “atoms” interchangeably.

The dictionary of wavelets can also be learned from natural images. The most striking feat in this endeavor was accomplished by Olshausen and Field (1996) [39], who proposed a method for learning an over-complete dictionary of basis functions from natural image patches. The scientific objective of their work is to explain the observed properties of the so-called “simple cells” in primary visual cortex or V1. Their objective function is similar to that of basis pursuit or Lasso, with a least squares error term and a sparsity inducing penalty term. However, in basis pursuit or Lasso,

the dictionary of wavelets is assumed given, and only the coefficients of the wavelets need to be computed by minimizing the objective function. Olshausen and Field proposed to minimize the objective function over both the coefficients and the dictionary of basis functions, which enabled them to learn a dictionary of localized, elongated and oriented wavelets that resemble the Gabor wavelets, which were proposed as a mathematical model for simple V1 cells [10].

In the terminology of linear regression, the goal of matching pursuit and basis pursuit/Lasso is to select the regressors or predictor vectors to explain a response vector, while assuming that the collection of all the regressors is already given. What Olshausen and Field accomplished is to learn the collection of regressors from multiple response vectors, so that each response vector can be explained by a small number of regressors. Of course, for different response vectors, they are explained by different sets of regressors.

1.2 Compositionality, molecular structures, and group sparsity

In atomic representations of images, the atoms (wavelets or basis functions) in a dictionary exist individually without any additional structures imposed on them, and they are selected individually to represent the images in matching pursuit or basis pursuit. We all know that the real atoms in physics and chemistry often form different molecules, it is also the case that the wavelets in the sparse representations of natural images often form commonly occurring compositional patterns or shape templates. If we consider the dictionary of wavelets as an alphabet, then the dictionary of compositional patterns can be considered a vocabulary or codebook of words. The goal of this article is to go beyond the atomic decomposition by learning the dictionaries of compositional patterns of the wavelets, so that when representing an image, the wavelets can be selected in groups instead of being selected individually. For each selected group of wavelets, we allow the coefficients of the group of wavelets to vary independently of each other. The number of selected groups is much smaller than the number of selected wavelets, resulting in sparser representations.

Such group structures have received considerable attention in statistics and machine learning in recent years, under the name of group sparsity or more generally structured sparsity. The most prominent example is the group Lasso [52], which replaces the ℓ_1 penalty of Lasso by a composite penalty according to the group structure. Just like in the Lasso where the dictionary of basis functions are assumed to be given, in the group Lasso, the dictionary of the groups of basis

functions are also assumed given. In this article, however, we attempt to learn dictionaries of the compositional patterns of the groups of wavelets, while assuming that the dictionary of wavelets is given, either by design or by learning. In linear regression language, we seek to learn from multiple response vectors the groups or group structures among the given collection of all the regressors, so that each response vector can be explained by a small number of groups of regressors selected from all possible groups. In other words, we seek to learn molecular structures formed by the atoms in atomic decompositions of natural images. This connects sparsity to another fundamental principle in vision, namely, compositionality [24, 56], which holds that patterns in natural images are composed of parts, which are themselves composed of sub-parts, and so on.

1.3 Image templates and dictionary learning

We represent each compositional pattern of wavelets by an active basis model [51], which is a composition of a small number of Gabor wavelets automatically selected from a dictionary of such wavelets. The selected wavelets are allowed to perturb their locations and orientations so that the linear basis formed by the selected wavelets become active and the active basis forms a deformable template. For a given set of training images, our method is to learn a dictionary of such active basis templates, so that each training image can be represented by a small number of templates that are translated, rotated, scaled and deformed versions of the learned templates in the dictionary. We assume that the dictionary of wavelets are given, and they are Gabor wavelets over a dense collection of locations, orientations and scales. Our learning scheme is unsupervised in the sense that the images are not annotated or labeled.

Figure (1) illustrates the basic idea. (a) displays the training image. (b) displays a mini-dictionary of 2 compositional patterns of wavelets learned from the training image. Here the number of training image is 1, although in general we may have many training images. The number of patterns in the dictionary is automatically determined by an adjusted BIC criterion. Each compositional pattern is represented by an active basis model, where each constituent Gabor wavelet is illustrated by a bar that has the same location, orientation and length as that Gabor wavelet. The 2 templates are displayed in different colors, so that it can be seen clearly how the translated, rotated, scaled and deformed versions of the 2 templates are used to represent the original image, as shown in (b). In (c), the templates are overlaid on the original image, where

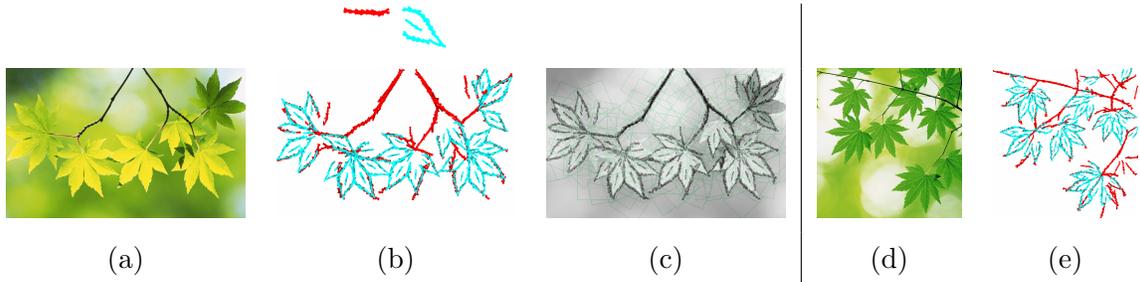


Figure 1: (a) Training image of 480×768 pixels. (b) Above: 2 compositional patterns of Gabor wavelets in the form of active basis templates learned from the training image. Each Gabor wavelet is illustrated by a bar with the same location, orientation and length as that wavelet. The bounding box of the templates is 100×100 pixels. The numbers of Gabor wavelets are respectively 22 in the red pattern and 40 in the green pattern. Below: Representing the training image by translated, rotated, scaled and deformed versions of the 2 templates. (c) Superpose deformed templates on the original image. Green squared boxes are bounding boxes of the templates. (d) Testing image. (e) Representation of the testing image by the 2 templates.

each green squared box is the bounding box of the template. In our current implementation, we allow some overlap between the bounding boxes of the templates.

The templates learned from the training image can be generalized to testing images, as shown in (d) and (e). Here the generalization is beyond what is commonly considered in machine learning literature. In machine learning, especially in supervised learning, the testing examples are assumed to be sampled from the same distribution as that of the training examples. In our situation, this is clearly not the case. We only assume that the testing examples consist of the same constituent templates as the training examples.



Figure 2: (a) Template of leaf learned in the first 7 iterations of the unsupervised learning algorithm. (b) In each iteration, the shared matching pursuit process selects wavelet elements sequentially to form each template. The sequence shows the process selecting 1, 3, 5, 10, 20, 30, 40 wavelet elements to form the leaf template in the last (10th) iteration.

The unsupervised learning algorithm starts from templates learned from randomly cropped image patches, so the initial templates are rather random. The algorithm then iterates the following two steps: (1) Image encoding: Encode each training image by translated, rotated, scaled and deformed versions of the current dictionary of templates, by a template matching pursuit process. (2) Dictionary learning: Re-learn each template from image patches currently encoded by this template, by a shared matching pursuit process. In Figure (2), (a) traces the template of leaf learned over the first 7 iterations of the learning algorithm. (b) shows the process of shared matching pursuit for learning this template in the last (10th) iteration, where the wavelet elements are sequentially added.

The rest of the paper is organized as follows. Section 2 reviews wavelets sparse coding and active basis model. Section 3 presents our representational scheme using active basis models as the composite representation units. This section then presents the unsupervised learning algorithm for learning dictionaries of active basis models from training images. Section 4 presents experimental results on image representation and classification. Section 5 concludes with a discussion, including comments on the limitations of our current work and comparisons with related work in the literature.

2 Wavelets sparse coding and active basis model

This section reviews wavelets sparse coding and the active basis model in order to fix the notation and set the stage for presenting our learning method.

2.1 Olshausen-Field model for sparse coding

Olshausen and Field [39] proposed that the role of simple V1 cells is to provide sparse representations of natural images. Let $\{\mathbf{I}_m, m = 1, \dots, M\}$ be a set of training image patches (e.g. 12×12), Olshausen-Field model seeks to represent these images by

$$\mathbf{I}_m = \sum_{i=1}^N c_{m,i} B_i + U_m, \quad (1)$$

where $(B_i, i = 1, \dots, N)$ is a dictionary of basis elements of the same dimension as \mathbf{I}_m , $c_{m,i}$ are the coefficients, and U_m is the unexplained residual image. N is often assumed to be greater than the

dimension of \mathbf{I}_m (e.g. $N = 2 \times 12 \times 12$), so the dictionary is said to be over-complete. On the other hand, the number of coefficients ($c_{m,i}, i = 1, \dots, N$) that are non-zero or significantly different from zero is assumed to be small for each image \mathbf{I}_m .

One may also assume that the basis elements in the dictionary are translated, rotated and dilated versions of one another, as in [41] (see also [47] and [53]), so that each B_i can be written as $B_{x,s,\alpha}$, where x is the location (a two-dimensional vector), s is the scale, and α is the orientation. We call such a dictionary self-similar, and we call (x, s, α) the geometric attribute of $B_{x,s,\alpha}$.

Model (1) then becomes

$$\mathbf{I}_m = \sum_{x,s,\alpha} c_{m,x,s,\alpha} B_{x,s,\alpha} + U_m, \quad (2)$$

where $B_{x,s,\alpha}$ are translated, rotated and dilated copies of a single basis element, e.g., $B = B_{x=0,s=1,\alpha=0}$, and (x, s, α) are properly discretized (default setting: α is discretized into 16 equally spaced orientations). B can be learned from training images $\{\mathbf{I}_m\}$ [41].

Assumption on wavelets in this paper. From now on, we assume that the dictionary of wavelets is self-similar, and $\{B_{x,s,\alpha}, \forall(x, s, \alpha)\}$ is already given. It can either be learned or designed. In the following, we assume that $B_{x,s,\alpha}$ is a Gabor wavelet, and we also assume that $B_{x,s,\alpha}$ is normalized to have unit ℓ_2 norm so that $|B_{x,y,\alpha}|^2 = 1$. $B_{x,s,\alpha}$ may also be a pair of Gabor sine and cosine wavelets, so that for each Gabor wavelet B , $B = (B_0, B_1)$. The corresponding coefficient $c = (c_0, c_1)$, and $cB = c_0B_0 + c_1B_1$. The projection $\langle \mathbf{I}, B \rangle = (\langle \mathbf{I}, B_0 \rangle, \langle \mathbf{I}, B_1 \rangle)$, and $|\langle \mathbf{I}, B \rangle|^2 = \langle \mathbf{I}, B_0 \rangle^2 + \langle \mathbf{I}, B_1 \rangle^2$.

Given the dictionary $(B_{x,s,\alpha}, \forall(x, s, \alpha))$, the encoding of an image \mathbf{I}_m amounts to inferring $(c_{m,x,s,\alpha}, \forall(x, s, \alpha))$ in (2) under the sparsity constraint, which means that only a small number of $(c_{m,x,s,\alpha})$ are non-zero. That is, we seek to encode \mathbf{I}_m by

$$\mathbf{I}_m = \sum_{i=1}^n c_{m,i} B_{x_{m,i}, s_{m,i}, \alpha_{m,i}} + U_m, \quad (3)$$

where $n \ll N$ is a small number, and $(x_{m,i}, s_{m,i}, \alpha_{m,i}, i = 1, \dots, n)$ are the geometric attributes of the selected wavelets whose coefficients $(c_{m,i})$ are non-zero. $(x_{m,i}, s_{m,i}, \alpha_{m,i}, i = 1, \dots, n)$ form a spatial point process (we continue to use i to index the wavelets, but here i only runs through the n selected wavelets instead of all the N wavelets as in (1)).

2.2 Active basis model for shared sparse coding of aligned image patches

The active basis model was proposed by Wu et al. [51] for modeling deformable compositional patterns of wavelet elements.

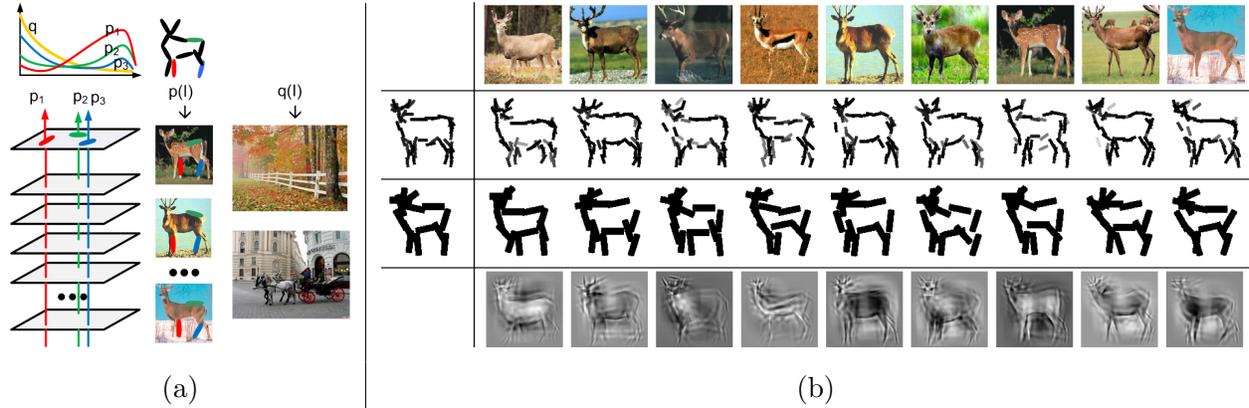


Figure 3: (a): The nominal template $\mathbf{B} = (B_{x_i, s, \alpha_i}, i = 1, \dots, n)$ of an active basis model, where each wavelet element B_{x_i, s, α_i} is illustrated by a bar at the same location and orientation, and with the same length. The elements (such as the colored bars) of \mathbf{B} are shared by all the training images $\{\mathbf{I}_m, m = 1, \dots, M\}$, subject to local perturbations $(\Delta x_{m,i}, \Delta \alpha_{m,i}, i = 1, \dots, n)$ that deform the template \mathbf{B} to match \mathbf{I}_m . The elements are selected based on the contrast between the foreground distribution p_i of the wavelet coefficients pooled from training images, and the background distribution q pooled from natural images. (b) The first row displays $\{\mathbf{I}_m, m = 1, \dots, M = 9\}$. The second row: the first plot is the nominal template $\mathbf{B} = (B_i = B_{x_i, s, \alpha_i}, i = 1, \dots, n = 50)$. The rest of the plots are the deformed templates $\mathbf{B}_m = (B_{m,i} = B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}})$. The third row: the same as the second row, except that the scale s is about twice as large, and $n = 14$. The last row displays the linear reconstruction $\mathbf{I}_m^{\text{syn}} = \sum_{i=1}^n c_{m,i} B_{m,i}$, where $n = 100$, and $(B_{m,i}, i = 1, \dots, n)$ contains Gabor wavelets and difference of Gaussian wavelets at multiple scales.

Suppose we have a set of training image patches $\{\mathbf{I}_m, m = 1, \dots, M\}$. This time we assume that they are defined on the same bounding box, and the objects in these images come from the same category. In addition, they appear at the same location, scale and orientation, and in the same pose. See Figure (3) for 9 image patches of deer. We call such image patches aligned.

The active basis model is of the following form

$$\mathbf{I}_m = \sum_{i=1}^n c_{m,i} B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}} + U_m, \quad (4)$$

where $\mathbf{B} = (B_{x_i, s, \alpha_i}, i = 1, \dots, n)$ form the nominal template of an active basis model (sometimes we simply call \mathbf{B} an active basis template). Here we assume that the scale s is fixed and given. $\mathbf{B}_m = (B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}}, i = 1, \dots, n)$ is the deformed version of the nominal template \mathbf{B} for encoding \mathbf{I}_m , where $(\Delta x_{m,i}, \Delta \alpha_{m,i})$ are the perturbations of the location and orientation from the nominal location x_i and the nominal orientation α_i respectively. The perturbations are introduced to account for shape deformation. See Figure (3) for illustration, where the wavelet elements that are depicted by colored bars can shift their locations and orientations in different training images. Both $\Delta x_{m,i}$ and $\Delta \alpha_{m,i}$ are assumed to vary within limited ranges (default setting: $\Delta x_{m,i} \in [-3, 3]$ pixels, and $\Delta \alpha_{m,i} \in \{-1, 0, 1\} \times \pi/16$).

2.3 Shared matching pursuit: local maximum pooling and arg-max explaining-away inhibition

Given the dictionary of Gabor wavelets $\{B_{x,s,\alpha}, \forall x, s, \alpha\}$, the learning of the active basis model from the aligned image patches $\{\mathbf{I}_m\}$ involves the sequential selection of B_{x_i, s, α_i} and the inference of its perturbed version $B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}}$ in each image \mathbf{I}_m . We call the learning as supervised, because the bounding boxes of the objects are given and the images are aligned. See Figure (3) for an illustration of the learning results.

In this subsection, we consider a prototype version of the shared matching pursuit algorithm, which is to be revised in the following subsections. The reason we start from this prototype algorithm is that it is simple and yet captures the key features of the learning algorithm.

The prototype shared matching pursuit algorithm is a greedy algorithm that seeks the maximal reduction of the following least squares reconstruction error

$$\sum_{m=1}^M \left| \mathbf{I}_m - \sum_{i=1}^n c_{m,i} B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}} \right|^2 \quad (5)$$

in each iteration (recall that the wavelet elements are normalized to have unit ℓ_2 norm).

[0] Initialize $i \leftarrow 0$. For $m = 1, \dots, M$, initialize the residual image $U_m \leftarrow \mathbf{I}_m$.

[1] $i \leftarrow i + 1$. Select the next element by

$$(x_i, \alpha_i) = \arg \max_{x, \alpha} \sum_{m=1}^M \max_{\Delta x, \Delta \alpha} |\langle U_m, B_{x+\Delta x, s, \alpha+\Delta \alpha} \rangle|^2,$$

where $\max_{\Delta x, \Delta \alpha}$ is a local maximum pooling within the small ranges of $\Delta x_{m,i}$ and $\Delta \alpha_{m,i}$.

[2] For $m = 1, \dots, M$, given (x_i, α_i) , infer the perturbations in location and orientation by retrieving the arg-max in the local maximum pooling of step [1]:

$$(\Delta x_{m,i}, \Delta \alpha_{m,i}) = \arg \max_{\Delta x, \Delta \alpha} |\langle U_m, B_{x_i+\Delta x, s, \alpha_i+\Delta \alpha} \rangle|^2.$$

Let $c_{m,i} \leftarrow \langle U_m, B_{x_i+\Delta x_{m,i}, s, \alpha_i+\Delta \alpha_{m,i}} \rangle$, and update the residual image

$$U_m \leftarrow U_m - c_{m,i} B_{x_i+\Delta x_{m,i}, s, \alpha_i+\Delta \alpha_{m,i}}. \quad (6)$$

[3] Stop if $i = n$, else go back to step [1].

The above algorithm is a generalization of the matching pursuit algorithm [35]. In step [1], the $\max_{\Delta x, \Delta \alpha}$ is the local maximum pooling, proposed by [43] as a function of V1 complex cells. The selected B_{x_i, s, α_i} is supposed to encode all the images $\{\mathbf{I}_m\}$ simultaneously, subject to local perturbations. That is, B_{x_i, s, α_i} is shared by all the images. After the selection of the shared wavelet B_{x_i, s, α_i} , we infer its perturbation on each image \mathbf{I}_m by retrieving the arg-max of the local maximum pooling on \mathbf{I}_m . This arg-max wavelet $B_{x_i+\Delta x_{m,i}, s, \alpha_i+\Delta \alpha_{m,i}}$ then explains away a small part from U_m according to (6), thereby implicitly inhibits nearby wavelets from being selected in the future. Note that the explain-away inhibition is image specific, because the perturbed versions of each selected wavelet are different for different images.

Assumption on orthogonality in this paper. Because of the arg-max explaining-away inhibition, the wavelet elements in each deformed template $\mathbf{B}_m = (B_{x_i+\Delta x_{m,i}, s, \alpha_i+\Delta \alpha_{m,i}}, i = 1, \dots, n)$ usually have little overlap with each other. So from now on, we shall assume that these wavelet elements are orthogonal to each other, so that the coefficient can be obtained by projection: $c_{m,i} = \langle \mathbf{I}_m, B_{x_i+\Delta x_{m,i}, s, \alpha_i+\Delta \alpha_{m,i}} \rangle$. We write $C_m = (c_{m,i}, i = 1, \dots, n)$. In practice, we allow small overlap between the elements of \mathbf{B}_m .

2.4 Statistical modeling: foreground pops out from natural image background

The above algorithm based on (5) implicitly assumes that the residual U_m is Gaussian white noise. This assumption can be problematic because the unexplained background in the image may contain

salient structures such as edges. This is why we need to revise the above algorithm which is based on the Gaussian white noise assumption. A better assumption is to assume that U_m follows the same distribution as that of natural images. More precisely, the distribution of \mathbf{I}_m given the deformed template $\mathbf{B}_m = (B_{x_i+\Delta x_{m,i},s,\alpha_i+\Delta\alpha_{m,i}}, i = 1, \dots, n)$, i.e., $p(\mathbf{I}_m | \mathbf{B}_m)$, is obtained by modifying the distribution of natural images $q(\mathbf{I}_m)$ in such a way that we only change the distribution of $C_m = (c_{m,i} = \langle \mathbf{I}_m, B_{x_i+\Delta x_{m,i},s,\alpha_i+\Delta\alpha_{m,i}} \rangle, i = 1, \dots, n)$ from $q(C_m)$ to $p(C_m)$, while leaving the conditional distribution of U_m given C_m unchanged. Here $p(C_m)$ and $q(C_m)$ are the distributions of C_m under $p(\mathbf{I}_m | \mathbf{B}_m)$ and $q(\mathbf{I}_m)$ respectively. We should find \mathbf{B} and perturb it to $\{\mathbf{B}_m\}$ so that $p(C_m)$ and $q(C_m)$ have the maximum contrast in terms of the Kullback-Leibler divergence. Thus the model is in the form of foreground $p(C_m)$ popping out from background $q(\mathbf{I}_m)$. Specifically, $p(\mathbf{I}_m | \mathbf{B}_m) = q(\mathbf{I}_m)p(C_m)/q(C_m)$. Such a density substitution scheme was first used in projection pursuit density estimation [23]. See Figure (3) for illustration.

For computational simplicity, we further assume that $(c_{m,i} = \langle \mathbf{I}_m, B_{x_i+\Delta x_{m,i},s,\alpha_i+\Delta\alpha_{m,i}} \rangle, i = 1, \dots, n)$ are independent given \mathbf{B}_m , under both p and q , so we have

$$p(\mathbf{I}_m | \mathbf{B}_m) = q(\mathbf{I}_m) \prod_{i=1}^n \frac{p_i(c_{m,i})}{q(c_{m,i})},$$

where $q(c)$ is assumed to be the same for $i = 1, \dots, n$ because $q(\mathbf{I}_m)$ is translation and rotation invariant. $q(c)$ can be pooled from natural images in the form of a heavy-tailed histogram of Gabor filter responses.

For parametric modeling, we assume the following exponential family model $p_i(c) = p(c; \lambda_i)$, where

$$p(c; \lambda) = \frac{1}{Z(\lambda)} \exp\{\lambda h(|c|^2)\} q(c). \quad (7)$$

$h(r)$ is a function of the response $r = |c|^2$ that saturates for large r . Specifically, we assume that $h(r) = \xi[2/(1 + e^{-2r/\xi}) - 1]$. $h(r)$ behaves like $h(r) \approx r$ for small r , but $h(r) \rightarrow \xi$ (default setting: $\xi = 6$) as $r \rightarrow \infty$. The reason we assume a saturation function is that in the natural image background, there are also occasional (albeit less frequent) edges that give equally large responses r as those in the foreground, so the probability ratio $p_i(c)/q(c)$ should go to a positive constant instead of 0 for large $|c|^2$. In (7),

$$Z(\lambda) = \int \exp\{\lambda h(r)\} q(c) dc = E_q[\exp\{\lambda h(r)\}]$$

is the normalizing constant. $\mu(\lambda) = \mathbb{E}_\lambda[h(r)] = \int h(r)p(c; \lambda)dc$ is the mean parameter. Both $Z(\lambda)$ and $\mu(\lambda)$ can be computed beforehand from natural images.

The log-likelihood is

$$l(\{\mathbf{I}_m\} | \mathbf{B}, \{\mathbf{B}_m\}) = \sum_{m=1}^M \sum_{i=1}^n \left[\lambda_i h(\langle \mathbf{I}_m, B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}} \rangle) - \log Z(\lambda_i) \right]. \quad (8)$$

2.5 Shared matching pursuit revised: saturation and hard inhibition

We can revise the shared matching pursuit in subsection (2.3) in order to maximize the log-likelihood (8) instead of minimizing the squared loss (5) as in subsection (2.3). The algorithm is as follows.

[0] Initialize $i \leftarrow 0$. For $m = 1, \dots, M$, initialize the response maps $R_m(x, \alpha) \leftarrow \langle \mathbf{I}_m, B_{x,s,\alpha} \rangle$ for all (x, α) .

[1] $i \leftarrow i + 1$. Select the next wavelet by finding

$$(x_i, \alpha_i) = \arg \max_{x, \alpha} \sum_{m=1}^M \max_{\Delta x, \Delta \alpha} h(|R_m(x + \Delta x, \alpha + \Delta \alpha)|^2),$$

where $\max_{\Delta x, \Delta \alpha}$ is again local maximum pooling.

[2] For $m = 1, \dots, M$, given (x_i, α_i) , infer the perturbations by retrieving the arg-max in the local maximum pooling of step [1]:

$$(\Delta x_{m,i}, \Delta \alpha_{m,i}) = \arg \max_{\Delta x, \Delta \alpha} |R_m(x_i + \Delta x, \alpha_i + \Delta \alpha)|^2.$$

Let $c_{m,i} \leftarrow R_m(x_i + \Delta x_{m,i}, \alpha_i + \Delta \alpha_{m,i})$, and update $R_m(x, \alpha) \leftarrow 0$ if $\text{corr}[B_{x,s,\alpha}, B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}}] > \epsilon$ (default setting: $\epsilon = .1$). Then compute λ_i by solving the maximum likelihood equation $\mu(\lambda_i) = \sum_{m=1}^M h(|c_{m,i}|^2)/M$.

[3] Stop if $i = n$, else go back to step [1].

The correlation is defined as the square of the inner product between the wavelets and can be stored beforehand.

There are two modifications to the original shared matching pursuit in subsection (2.3). (1) In step [1], we apply the saturation function $h(\cdot)$ to the response. This is justified by maximum likelihood based on the exponential family model (7). (2) In step [2], the arg-max element $B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}}$ directly inhibits nearby wavelets whose correlation with it is greater than a tolerance ϵ , instead of explaining away from U_m and indirectly inhibiting nearby wavelets as in (6).

This hard inhibition is to approximately enforce the orthogonality assumption in subsection (2.3) and can be viewed as an approximation to the linear subtraction (6). In step [2], the function $\mu()$ can be stored beforehand over a discrete list of values, so that the maximum likelihood equation can be solved by looking up this function with linear interpolation between adjacent discrete values.

2.6 Template matching for detection

After learning the template from training images $\{\mathbf{I}_m\}$, we can use the learned template to detect the object in the testing image \mathbf{I} . Assume that the testing image contains an instance of the object, we can detect the object by scanning the template over the whole image and compute the template matching score. At the location of the maximum template matching score, we can then deform the template to sketch the image. See Figure (4) for an illustration.



Figure 4: Inference by template matching. In each block, the left is the testing image \mathbf{I} , and the right is the translated and deformed template $(B_{\hat{X}+x_i+\Delta x_i, s, \alpha_i+\Delta \alpha_i}, i = 1, \dots, n)$.

[1] For every pixel X , compute the log-likelihood $l(X)$, which serves as the template matching score at putative location X :

$$l(X) = \sum_{i=1}^n [\lambda_i \max_{\Delta x, \Delta \alpha} h(|\langle \mathbf{I}, B_{X+x_i+\Delta x, s, \alpha_i+\Delta \alpha} \rangle|^2) - \log Z(\lambda_i)]. \quad (9)$$

[2] Find maximum likelihood $\hat{X} = \arg \max_X l(X)$. For $i = 1, \dots, n$, inferring perturbations by retrieving the arg-max in the local maximum pooling in step [1]:

$$(\Delta x_i, \Delta \alpha_i) = \arg \max_{\Delta x, \Delta \alpha} |\langle \mathbf{I}, B_{\hat{X}+x_i+\Delta x, s, \alpha_i+\Delta \alpha} \rangle|^2.$$

[3] Return the location \hat{X} , and the translated and deformed template $(B_{\hat{X}+x_i+\Delta x_i, s, \alpha_i+\Delta \alpha_i}, i = 1, \dots, n)$.

Rotation and multi-resolution. We can rotate the template and scan the template over multiple resolutions of the original image, to account for uncertainties about the orientation and scale of the object in the testing image.

3 Learning dictionaries of compositional patterns of wavelets

In Olshausen-Field model (1), the coefficients are assumed to be independent for simplicity [32, 40]. A natural question is how to correct this assumption, specifically, how to add another layer of model on the coefficients? In more plain words, what is beyond the Olshausen-Field model?

Since we argue that the Olshausen-Field model (1) with a dictionary of self-similar wavelets as in (2) is essentially a spatial point process as explicated in (3), the model on top of all the coefficients should focus on the geometric patterns formed by the wavelets with non-zero coefficients. In particular, we may search for recurring compositional patterns of the wavelets with non-zero coefficients. These patterns of spatial grouping can be modeled by active basis models.

In this section, we shall specify our representation where each representational unit is an active basis template. Then we shall describe the algorithm for learning dictionaries of active basis templates from training images.

3.1 Representation based on composite representational units

In this subsection, we strive to write down our representation in a form that is analogous to the Olshausen-Field model (3), by using compactified notation.

Compactified notation. As the first step of this exercise of compactification, let us slightly generalize the active basis model by assuming that the template may appear at location X_m in image \mathbf{I}_m , then we can write the representation in the following form:

$$\begin{aligned}\mathbf{I}_m &= \sum_{i=1}^n c_{m,i} B_{X_m+x_i+\Delta x_{m,i},s,\alpha_i+\Delta\alpha_{m,i}} + U_m \\ &= C_m \mathbf{B}_{X_m} + U_m,\end{aligned}\tag{10}$$

where $\mathbf{B}_{X_m} = (B_{X_m+x_i+\Delta x_{m,i},s,\alpha_i+\Delta\alpha_{m,i}}, i = 1, \dots, n)$ is the deformed template spatially translated to X_m , $C_m = (c_{m,i}, i = 1, \dots, n)$, and $C_m \mathbf{B}_{X_m}$ is defined to be $\sum_{i=1}^n c_{m,i} B_{X_m+x_i+\Delta x_{m,i},s,\alpha_i+\Delta\alpha_{m,i}}$. Here we no longer assume that the training images $\{\mathbf{I}_m\}$ are aligned.

\mathbf{B}_{X_m} explains the part of \mathbf{I}_m that is covered by \mathbf{B}_{X_m} . For each image \mathbf{I}_m and each X_m , we can define the log-likelihood ratio according to (9):

$$l(\mathbf{I}_m | \mathbf{B}_{X_m}) = \log \frac{p(\mathbf{I}_m | \mathbf{B}_{X_m})}{q(\mathbf{I}_m)}$$

$$= \sum_{i=1}^n \left[\lambda_i \max_{\Delta x, \Delta \alpha} h(|\langle \mathbf{I}_m, B_{X_m+x_i+\Delta x, s, \alpha_i+\Delta \alpha} \rangle|^2) - \log Z(\lambda_i) \right]. \quad (11)$$

As the next step of this compactification exercise, in addition to spatial translation and deformation, we can also rotate and scale the template. So a more general model than (10) is

$$\mathbf{I}_m = C_m \mathbf{B}_{X_m, S_m, A_m} + U_m,$$

where X_m is the location, S_m is the scale, and A_m is the orientation of the translated, rotated, scaled and deformed template. The scaling of the template is implemented by changing the resolution of the original image. We adopt the convention that whenever the notation \mathbf{B} appears in image representation, it always means the deformed template, where the perturbations of the basis elements can be inferred by local max pooling. The log-likelihood ratio $l(\mathbf{I}_m | \mathbf{B}_{X_m, S_m, A_m})$ can be similarly defined as in (11).

Compactified representation. Now suppose we have a dictionary of T types of templates, $\{\mathbf{B}^{(t)}, t = 1, \dots, T\}$, where each $\mathbf{B}^{(t)}$ is a compositional pattern of wavelets. Then we can represent an image \mathbf{I}_m by K_m templates that are spatially translated, rotated, scaled and deformed versions of these T templates in the dictionary:

$$\mathbf{I}_m = \sum_{k=1}^{K_m} C_{m,k} \mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})} + U_m, \quad (12)$$

where each $\mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_k)}$ is obtained by translating the template of type t_k , i.e., $\mathbf{B}^{(t_k)}$ in the dictionary, to location $X_{m,k}$, scale it to scale $S_{m,k}$, rotate it to orientation $A_{m,k}$, and deform it to match \mathbf{I}_m .

If the K_m templates do not overlap with each other, then the log-likelihood is

$$\sum_{m=1}^M \sum_{k=1}^{K_m} \left[l(\mathbf{I}_m | \mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}) \right]. \quad (13)$$

Packing and unpacking. The above representation is in analogy to Equation (3) in subsection (2.1), which we copy here: $\mathbf{I}_m = \sum_{i=1}^n c_{m,i} B_{x_{m,i}, s_{m,i}, \alpha_{m,i}} + U_m$. The difference is that each $\mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_k)}$ is a composite representational unit, which is itself a group of wavelets that follow a certain compositional pattern of type t_k . Because of such grouping or packing, the number of templates K_m needed to encode \mathbf{I}_m is expected to be much smaller than the total number of wavelets

needed to represent \mathbf{I}_m , thus resulting in sparser representation. Specifically, if each template is a group of g wavelet elements, then the number of wavelets in the representation (12) is $K_m g$. In fact, we can unpack model (12) into the wavelets representation (3). The reason that it is advantageous to pack the wavelets into groups is that these groups exhibit T types of recurring spatial grouping patterns, so that when we encode the image \mathbf{I}_m , for each selected group $\mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}$, we only need to code the overall location, scale, orientation and type of the group, instead of the locations, scales and orientations of the individual constituent wavelets.

Limited overlap assumption. It is desirable to allow some overlaps between the bounding boxes of the K_m templates that encode \mathbf{I}_m , so that no salient structures of \mathbf{I}_m fall through the cracks between the templates. Assume that the bounding boxes of all the templates are of the squared shape, we assume the following limited overlap constraint: For any template $\mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}$ centered at $X_{m,k}$, let D be the side length of its squared bounding box, then no other templates are allowed to be centered within a distance of ρD from $X_{m,k}$ (default setting: $\rho = .4$). Under such limited overlap condition, we may continue to use the log-likelihood (13). However, we need to re-scale it to account for the overlap between the K_m templates.

3.2 Model complexity

Before considering the learning algorithm that seeks to maximize the log-likelihood (13), we need to resolve two issues regarding model complexity. One is how to choose the number of wavelets $n^{(t)}$ in each template $\mathbf{B}^{(t)}$ in the dictionary. The other is how to choose the number of templates T in the dictionary $\{\mathbf{B}^{(t)}, t = 1, \dots, T\}$.

Determining the number of wavelets in a template in supervised learning. Suppose we are in the supervised learning setting where $\{\mathbf{I}_m, m = 1, \dots, M\}$ are aligned image patches, and we want to learn an active basis template $\mathbf{B} = \{B_{x_i, s, \alpha_i}, i = 1, \dots, n\}$. We employ the following penalized log-likelihood:

$$\sum_{m=1}^M \sum_{i=1}^n \left[\lambda_i h(\langle \mathbf{I}_m, B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}} \rangle) - \log Z(\lambda_i) - \gamma \right], \quad (14)$$

which is the sum of the log-likelihood (8) and a penalty term γ associated with each basis element. There are two interpretations of γ . One is from the minimum description length (MDL) [44] perspective, where γ can be interpreted as the cost of coding the perturbations $(\Delta x_{m,i}, \Delta \alpha_{m,i})$ in

the encoding of \mathbf{I}_m . The other interpretation is from the Bayesian information criterion (BIC) [45] perspective. From the Bayesian perspective, the perturbations $(\Delta x_{m,i}, \Delta \alpha_{m,i})$ should be integrated out according to their prior distributions, which are uniform distributions over the allowed ranges of perturbations. However, in our computation, the perturbations $(\Delta x_{m,i}, \Delta \alpha_{m,i})$ are inferred by local max pooling, i.e., they are actually maxed out instead of being integrated out. As a result, the resulting log-likelihood with perturbations maxed out actually over-estimates the log-likelihood with perturbations integrated out. The term γ may be considered an approximation to this over-estimation, which should be subtracted from the maxed out log-likelihood.

In order to maximize the penalized log-likelihood (14), we can continue to use the shared matching pursuit algorithm in subsection (2.5), except that we should stop the algorithm once the gain in the average log-likelihood is smaller than γ , i.e.,

$$\lambda_i \sum_{m=1}^M h(\langle \mathbf{I}_m, B_{x_i + \Delta x_{m,i}, s, \alpha_i + \Delta \alpha_{m,i}} \rangle) / M - \log Z(\lambda_i) < \gamma.$$

This determines n . Currently we set the tuning parameter γ empirically at the default value of 1.

We can apply the above idea to unsupervised learning of the dictionary $\{\mathbf{B}^{(t)}, t = 1, \dots, T\}$ from the non-aligned training images $\{\mathbf{I}_m\}$, by maximizing the penalized log-likelihood function

$$\sum_{m=1}^M \sum_{k=1}^{K_m} \left[l(\mathbf{I}_m | \mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}) - n^{(t_{m,k})} \gamma \right], \quad (15)$$

where $n^{(t)}$ is the number of wavelet elements in $\mathbf{B}^{(t)}$.

Determining the number of templates. The number of templates T in the dictionary can be selected by an adjusted BIC criterion. The BIC criterion has been advocated by Fraley and Raftery (2002) [22] for determining the number of clusters in mixture models. Our model is similar to mixture models or clustering in the sense that the image patches are clustered into T different clusters. The complication is that these image patches are not independent examples, and they are not randomly cropped from the training images either. Instead, they are cropped from the training images $\{\mathbf{I}_m\}$ by the template matching pursuit process under the limited overlap constraint. From the clustering perspective, our method also bears some similarity with bi-clustering [37] or co-clustering, where in each cluster, different basis elements are selected for representation. The complication is that these basis elements are not given coordinates (i.e., pixel values of the image patches). They need to be inferred by shared matching pursuit.

Recall that the BIC criterion is of the following form: maximized log-likelihood - number of parameters \times log (number of training examples)/2. We can take (15) as the log-likelihood term, however, we have to discount for the overlap between the templates. The number of parameters in our model is $\sum_{t=1}^T n^{(t)}$, which is the number of wavelets in the templates in the dictionary. The number of training examples can be taken as $\sum_{m=1}^M K_m$, which is the number of image patches that are explained by the templates. So we define our adjusted BIC criterion as:

$$\beta \sum_{m=1}^M \sum_{k=1}^{K_m} \left[l(\mathbf{I}_m \mid \mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}) - n^{(t_{m,k})} \gamma \right] - \frac{1}{2} \sum_{t=1}^T n^{(t)} \log \sum_{m=1}^M K_m, \quad (16)$$

where β is a ratio that discounts the overlap between the selected templates $\{\mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}\}$. We define $\beta = a/b$ where a is the total number of pixels that are actually covered by the bounding boxes of these templates, where the overlapping pixels are only counted once. b is the sum of the numbers of pixels within the bounding boxes of these templates, where a pixel is counted multiple times if it is covered by multiple templates.

3.3 Unsupervised learning algorithm

The learning algorithm seeks to maximize the penalized log-likelihood (15) subject to the limited overlap constraint. It is an iterative algorithm where each iteration seeks to maximally increase the penalized log-likelihood (15). Each iteration consists of two steps: (1) Image encoding. Given the current dictionary $\{\mathbf{B}^{(t)}, t = 1, \dots, T\}$, encode each training image \mathbf{I}_m by translated, rotated, scaled and deformed versions of the templates in the dictionary, i.e., $\{\mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}, k = 1, \dots, K_m\}$. (2) Dictionary learning. Given the current encoding of \mathbf{I}_m by $\{\mathbf{B}_{X_{m,k}, S_{m,k}, A_{m,k}}^{(t_{m,k})}, k = 1, \dots, K_m\}$, re-learn each $\mathbf{B}^{(t)}$ from the image patches covered by the translated, rotated, scaled and deformed versions of the current $\mathbf{B}^{(t)}$.

In the above learning process, we fix the total number of templates in the dictionary, T . We run the learning process for different values of T . Then we choose T that achieves the maximum value of the adjusted BIC (16).

The following are the details of the two steps:

Step (1): Image encoding by template matching pursuit. Suppose we are given the current dictionary $\{\mathbf{B}^{(t)}, t = 1, \dots, T\}$. Then for each \mathbf{I}_m , the template matching pursuit process seeks

to represent \mathbf{I}_m by sequentially selecting a small number of templates from the dictionary. Each selection seeks to maximally increase the penalized log-likelihood (15).

[0] Initialize the maps of template matching scores:

$$\mathbf{R}_m^{(t)}(X, S, A) \leftarrow l(\mathbf{I}_m \mid \mathbf{B}_{X,S,A}^{(t)}) - n^{(t)}\gamma,$$

for all (X, S, A, t) . This can be accomplished by first rotating the template $\mathbf{B}^{(t)}$ to orientation A , and then scanning the rotated template over the image zoomed to the resolution that corresponds to scale S . The larger the S is, the smaller the resolution is. Let $k \leftarrow 1$.

[1] Select the translated, rotated, scaled and deformed template by finding the global maximum of the response maps:

$$(X_{m,k}, S_{m,k}, A_{m,k}, t_{m,k}) = \arg \max_{X,S,A,t} \mathbf{R}_m^{(t)}(X, S, A).$$

[2] Let the selected arg-max template inhibit overlapping candidate templates to enforce limited overlap constraint. Let D be the side length of the bounding box of the selected template $\mathbf{B}_{X_{m,k},S_{m,k},A_{m,k}}^{(t_{m,k})}$, then for all (X, S, A, t) , if X is within a distance ρD from $X_{m,k}$, then set the response $\mathbf{R}_m^{(t)}(X, S, A) \leftarrow -\infty$ (default setting: $\rho = .4$).

[3] Stop if all $\mathbf{R}_m^{(t)}(X, S, A, t) \leq 0$. Otherwise let $k \leftarrow k + 1$, and go to [1].

The template matching pursuit algorithm implements a hard inhibition to enforce the limited overlap constraint. In a more rigorous implementation, we may update the residual image by $U_m \leftarrow U_m - C_m \mathbf{B}_{X_{m,k},S_{m,k},A_{m,k}}^{(t_{m,k})}$ as in the original version of matching pursuit. But the current simplified version is more efficient and works well enough.

Step (2): Dictionary re-learning by shared matching pursuit. For each $t = 1, \dots, T$, we re-learns $\mathbf{B}^{(t)}$ from all the image patches that are currently covered by $\mathbf{B}^{(t)}$. Each iteration of the shared matching pursuit process seeks to maximally increase the penalized log-likelihood (15), given the current encoding $(t_{m,k}, X_{m,k}, S_{m,k}, A_{m,k}, k = 1, \dots, K_m)$.

[0] Image patch cropping. For each \mathbf{I}_m , go through all the selected templates $\{\mathbf{B}_{X_{m,k},S_{m,k},A_{m,k}}^{(t_{m,k})}, \forall k\}$ that encode \mathbf{I}_m . If $t_{m,k} = t$, then crop the image patch of \mathbf{I}_m (at the resolution that corresponds to $S_{m,k}$) covered by the bounding box of the template $\mathbf{B}_{X_{m,k},S_{m,k},A_{m,k}}^{(t_{m,k})}$.

[1] Template re-learning. Re-learn template $\mathbf{B}^{(t)}$ from all the image patches covered by $\mathbf{B}^{(t)}$ that are cropped in [0], with their bounding boxes aligned. The learning is accomplished by the

shared matching pursuit algorithm of subsection (2.5). See Figure (2.b) for an illustration of the learning process in the last iteration.

In practice, we do not need to crop image patches directly. We only need to crop the maps of Gabor filter responses, and feed them into the shared matching pursuit algorithm.

Random initialization. The learning algorithm is initialized by learning each $\mathbf{B}^{(t)}$ from image patches that are randomly cropped from $\{\mathbf{I}_m\}$, so the initial templates are rather meaningless. Meaningful templates emerge very quickly after a few iterations. See Figure (2.a) for an illustration.

3.4 Notes on the learning algorithm

This subsection consists of some notes on various subtle aspects of the learning algorithm. More practically minded readers can jump to the next section for experimental results.

Polarization or specialization. Since the learning algorithm starts from a dictionary of templates learned from randomly cropped image patches, the differences among the initial templates are small. However, as the algorithm proceeds, the small differences among the initial templates quickly start a polarizing or specializing process, where the templates become more and more different, and they specialize in encoding different types of image patches. One may start the algorithm multiple times and select the one that achieves the maximum of the log-likelihood (15).

Stopping criterion. The regularization parameter γ plays an important role in determining when to stop the shared matching pursuit algorithm in the supervised learning in step (2). It also play an important role in determining when to stop the template matching pursuit algorithm in the image encoding step (1). The response map $\mathbf{R}_m^{(t)}(X, S, A)$ is initialized as $l(\mathbf{I}_m | \mathbf{B}_{X,S,A}^{(t)}) - n^{(t)}\gamma$. If $l(\mathbf{I}_m | \mathbf{B}_{X,S,A}^{(t)}) < n^{(t)}\gamma$, then the gain in terms of the log-likelihood ratio does not compensate the cost of coding the perturbations of the basis elements of the template. So we should stop the template matching pursuit if this is the case with all the remaining candidate templates.

Generalized matching pursuit in both steps. Both the image encoding step (1) and the dictionary re-learning step (2) are generalizations of the matching pursuit algorithm. In fact, the whole learning algorithm can be viewed as an encoding algorithm, which seeks to automatically discover the recurring compositional patterns of the selected wavelets that are otherwise overlooked by the plain matching pursuit algorithm. The re-learning of each template can be viewed as encoding multiple image patches by a single template, thus resulting in more efficient encoding than the

plain matching pursuit algorithm.

No early decision on wavelet representation. For learning the dictionary of compositional patterns of wavelets in the sparse representations of the training images $\{\mathbf{I}_m\}$, it is tempting to first apply the plain matching pursuit or basis pursuit/Lasso to each training image \mathbf{I}_m to obtain the sparse representation

$$\mathbf{I}_m = \sum_{i=1}^{n_m} c_{m,i} B_{x_{m,i}, s_{m,i}, \alpha_{m,i}} + U_m, \quad (17)$$

and then try to find the dictionary of commonly occurring spatial grouping patterns in the selected wavelets $\{B_{x_{m,i}, s_{m,i}, \alpha_{m,i}}, \forall i, m\}$. This was actually the strategy we adopted in our early work on textons [55]. The problem with such a two-step scheme is that the wavelets representation (17) produced by the matching pursuit or basis pursuit is an early decision or early commitment. Presumably, there may be many other wavelets representations that are no much less sparse than the representation (17), but they may be much more regular in terms of forming recurring compositional patterns. So we have to obtain the wavelets representation and discover the compositional patterns simultaneously or by an iterative scheme as in our learning algorithm. Not making early decision on wavelets sparse coding or edge detection is a key difference between our learning algorithm and those of [21] and [54].

Relationship with sparse component analysis and K-SVD. As our model is a recursion of the Olshausen-Field model, our learning algorithm can also be viewed as a recursion of the learning scheme of Olshausen and Field, which is sometimes called sparse component analysis in the literature. In the Olshausen-Field model $\mathbf{I}_m = \sum_{i=1}^N c_{m,i} B_i + U_m$, the dictionary of $(B_i, i = 1, \dots, N)$ is learned from the training image patches $\{\mathbf{I}_m\}$ by minimizing

$$\sum_{m=1}^M \left[\left\| \mathbf{I}_m - \sum_{i=1}^N c_{m,i} B_i \right\|^2 + \lambda \sum_{i=1}^N S(c_{m,i}) \right], \quad (18)$$

over both $(c_{m,i})$ and (B_i) , where $S()$ is a sparsity inducing penalty function, and λ is the regularization parameter. The learning algorithm iterates the following two steps. (1) Image encoding. For each \mathbf{I}_m , update $(c_{m,i}, \forall i)$ given $(B_i, \forall i)$. (2) Dictionary learning. Update (B_i) given $(c_{m,i}, \forall i, m)$. In Olshausen-Field learning algorithm, both steps are carried out by gradient descent. In a related learning algorithm called K-SVD [1], (1) can be accomplished by any pursuit algorithm such as matching pursuit or basis pursuit, and (2) is accomplished by SVD. Our algorithm is even more

similar to K-SVD than to the Olshausen-Field algorithm. It is interesting to notice that in both K-SVD and our algorithm, the updating of each representational unit in step (2) is performed on the image patches where this representational unit is currently active, i.e., the representational unit is re-learned from image patches that is currently encoded by this unit. Also similar to K-SVD, in our algorithm, the coefficients and the basis elements are updated together in dictionary re-learning in step (2).

Non-convex objective function. One complication about our learning method is that the log-likelihood (15) is not concave, or the negative log-likelihood is not convex, and our learning algorithm is a greedy algorithm that is similar to the EM algorithm [11]. In fact, this is also the case with the objective function (18) in Olshausen-Field sparse component analysis, which is non-convex in the joint domain of basis functions (wavelets) and their coefficients. It seems unlikely that a convex relaxations of the objective function can be obtained.

The sparse component analysis is related to independent component analysis (ICA) [2]. Both are related to factor analysis that is based on a linear additive structure, as is the case with other prominent unsupervised learning methods such as positive factor analysis [42], non-negative matrix factorization [30], and restricted Boltzmann machine [26]. In all these unsupervised learning methods, the objective functions are non-convex.

Linear subtraction versus occlusion. In both the template matching pursuit in step (1) and the shared matching pursuit in step (2), the explaining-away inhibition is carried out by hard inhibition that enforces limited overlap between templates in step (1) and the approximated non-overlap between basis elements in step (2). Such hard inhibition amounts to occlusion, where a selected template or basis element occludes nearby overlapping templates or elements. A more rigorous explaining-away mechanism in the context of linear additive structures (4) and (12) is by linear subtraction, where a selected template (group of wavelets) or wavelet is linearly subtracted from the training images, see Equation (6), so that other templates or wavelets continue to explain the residual images. This linear subtraction scheme is more computationally demanding than hard inhibition. We shall investigate the more rigorous subtraction scheme in future work.

Matching pursuit versus penalized least squares. Both steps (1) and (2) in our learning algorithm are generalizations of matching pursuit, and both can be replaced by generalizations of basis pursuit or Lasso. Step (1) can be replaced by group Lasso, where the groups are the groups of wavelets that

correspond to all possible translated, rotated, scaled and deformed versions of the templates in the current dictionary. Step (2) can be replaced by a different type of group Lasso, where the groups are formed across the aligned image patches from which we re-learn the template. Specifically, we group the coefficients of the same wavelet (up to local perturbations) across the aligned image patches, so that we always select the same set of wavelets for these aligned image patches. See Figure (3.a), where the vertical arrows indicate such a grouping scheme: we group the coefficients of the wavelets illustrated by the bars of the same color. This is sometimes called support union recovery in multivariate regression [38]. Such penalized least squares schemes can be much more expensive computationally than the corresponding versions of matching pursuit. We shall investigate them in future work.

4 Experiments

This section presents experiments based on the unsupervised learning algorithm in the previous section. The data and code for reproducing the experimental results reported in this paper can be downloaded at <http://www.stat.ucla.edu/~ywu/ABC/ABC.html>.

4.1 Image representation

In order to learn the dictionaries of compositional patterns from training images, we run the algorithm for 10 iterations. In the first iteration, we stop the template matching pursuit process for each \mathbf{I}_m until all $\mathbf{R}_m^{(t)}(X, S, A)$ become $-\infty$. That is because the initial templates in the dictionary are rather random, so we force them to explain the whole image of \mathbf{I}_m even if the templates do not match the image well. We fix $n^{(t)}$ to be the maximal value (default setting: 40) in the first 9 iterations. In the last iteration, we choose $n^{(t)}$ using the method described in subsection (3.2), and we use the templates with adaptively chosen $n^{(t)}$ to represent the training images.

Figure (5) shows an example of selecting the number of templates T in the dictionary. In each row, the first image is the training image. The remaining four blocks display the learned dictionaries of compositional patterns of wavelets in the form of active basis templates, as well as the representations of the training images using the learned dictionaries. The numbers of templates in the dictionaries are respectively 1, 2, 3, and 4. Just as in Figures (1) and (2), each wavelet

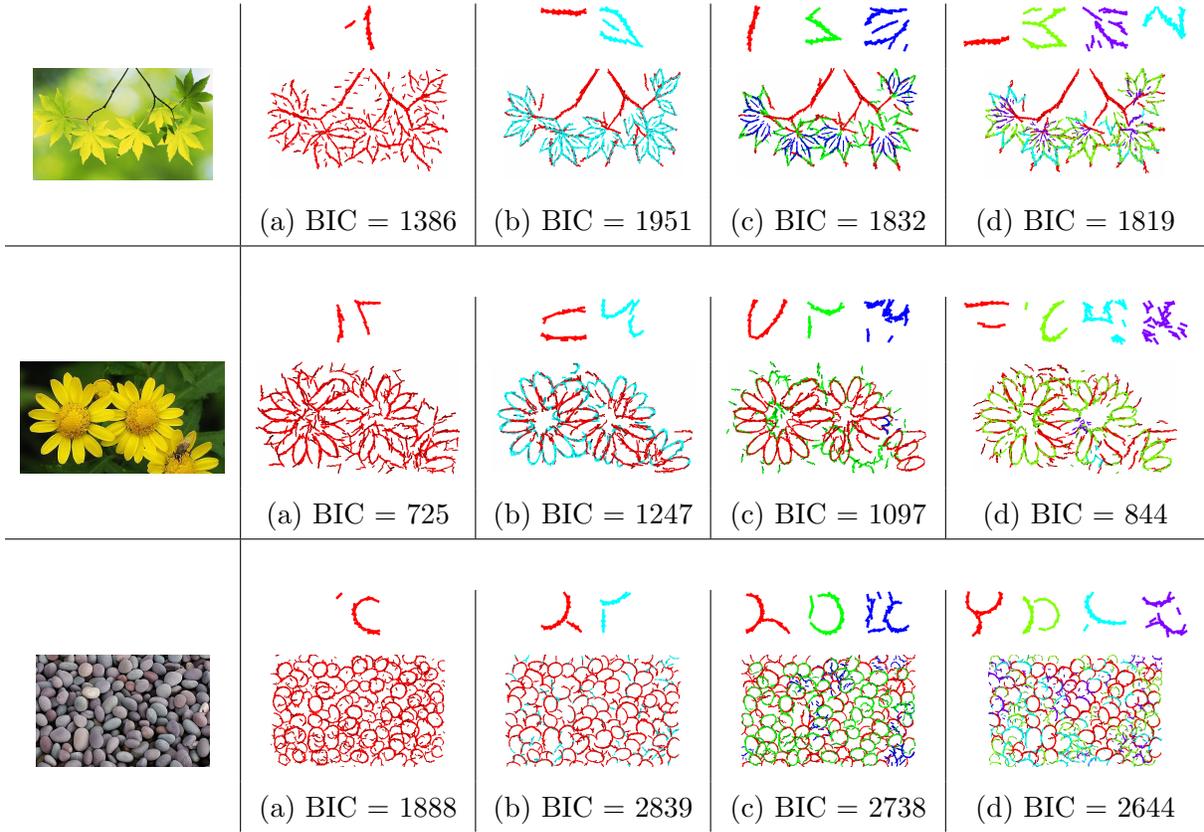


Figure 5: The adjusted BIC computed for different numbers of templates (1-4) in the dictionaries. The size of templates is 100×100 . The allowed range of scale change is $\{.8, 1, 1.2\}$ of the original scale. The templates are allowed full range of rotation. The maximal number of wavelet elements in each template is 40 and the actual number is automatically determined.

is illustrated by a bar at the same location and orientation, and with the same length as the wavelet. All the templates are of the size 100×100 . We also display the adjusted BIC criterion for each learned dictionary. The learned dictionaries are quite meaningful, and they give meaningful representations of the training images. It is interesting to observe how the dictionaries with only 1 template strive to represent the images.

As to the issue of selecting image resolution, for the example of maple leaves, we also learn dictionaries of templates at different resolutions of the training image. The resolution in Figure (5.b) achieves the maximum BIC per pixel. This is essentially equivalent to determining the size

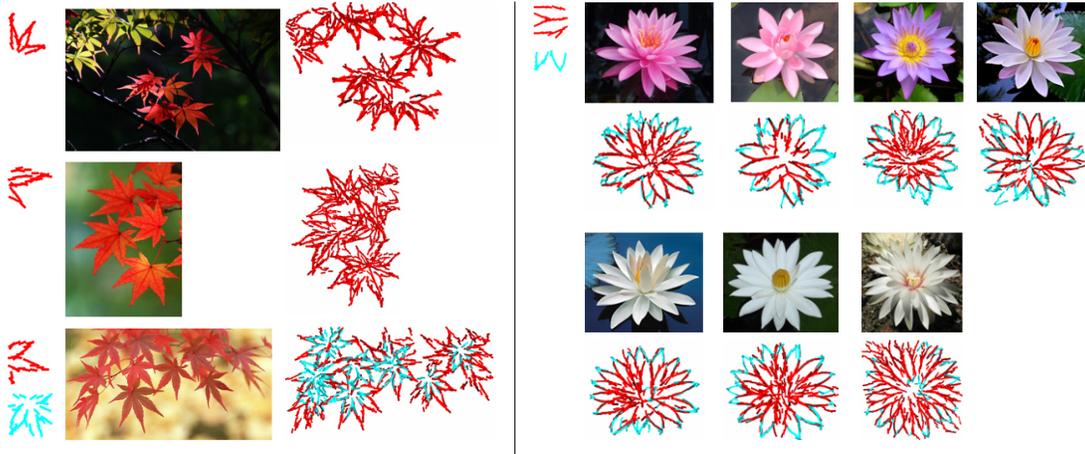


Figure 6: Maple leaves and lotus. Parameter setting is the same as in Figure (5).

of templates. However, we feel that instead of selecting the optimal resolution, it may be more appropriate to learn dictionaries at multiple resolutions or scales and combine them for image representation and understanding, just like the multiresolution analysis in wavelets theory.

The adjusted BIC is useful for determining the number of compositional patterns in the dictionary. However, it may be more appropriate to use it to determine the rough range of possible numbers of patterns, instead of using it to pinpoint the exact number. For instance, in the maple leaves example, the dictionary with 3 patterns also give a very meaningful representation of the training image. For applications such as image classification, the number of patterns in the dictionary may be determined by cross validation instead of BIC.

Figures (6) to (14) show more examples of representing natural images. In some of the images, such as flowers, leaves and brick wall, the composition patterns repeat themselves within the same images. For some other images, such as animal bodies and faces, the compositional patterns repeat across different images. For these images, the learning algorithm does not assume the images are aligned.

In the dictionary re-learning step (1), for each entry in the dictionary, we can learn a multi-scale template from the aligned image patches using wavelets at multiple scales. A multi-scale template has multiple component templates, each consisting of wavelets at a single scale. The learning of each component template can be done separately at each scale. Then in the image encoding step

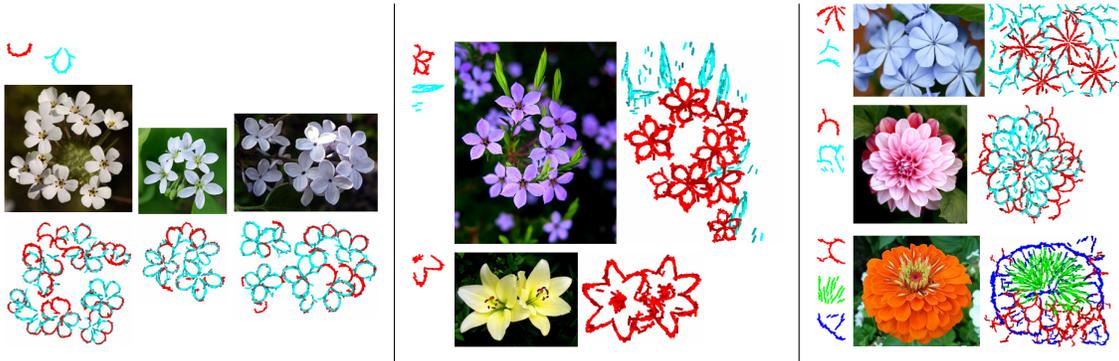


Figure 7: Flowers. Parameter setting is the same as in Figure (5). For the two examples in the middle, the templates consist of more than one petal. This is due to the fact that the templates are of a squared shape and are relatively large.

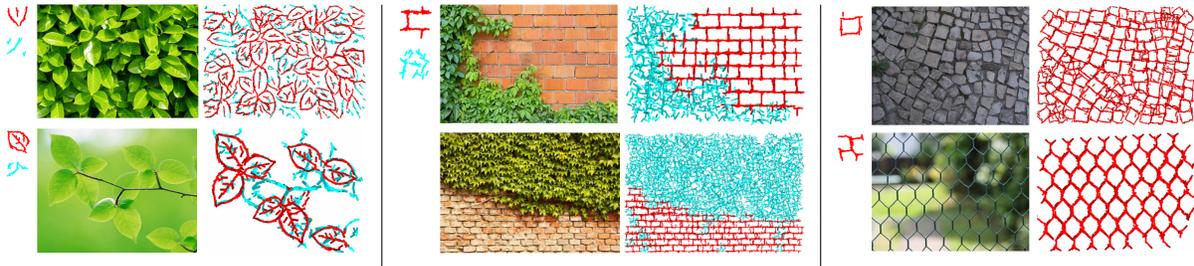


Figure 8: Ivy, leaves, ivy wall, pavement, and fence. Parameter setting is the same as in Figure (5). For the ivy wall example, the bottom row are testing image and its representation by the dictionary learned from the training image on the top row.

(2), for each entry in the dictionary, at each location, orientation and scale, we can combine the log-likelihood scores of the component templates at multiple scales in order to compute the overall template matching score at this location, orientation and scale. See the seagull example in Figure (11) for an illustration.

4.2 Image classification

The learned compositional patterns can be used as “words” in the bag-of-words method for image classification. Let $\{\mathbf{B}^{(t)}, t = 1, \dots, T\}$ be the templates learned from positive training images. For each image \mathbf{I}_m , let $\mathbf{R}_m^{(t)}(X, S, A) = l(\mathbf{I}_m | \mathbf{B}_{X,S,A}^{(t)})$ be the log-likelihood score of $\mathbf{B}^{(t)}$ at location

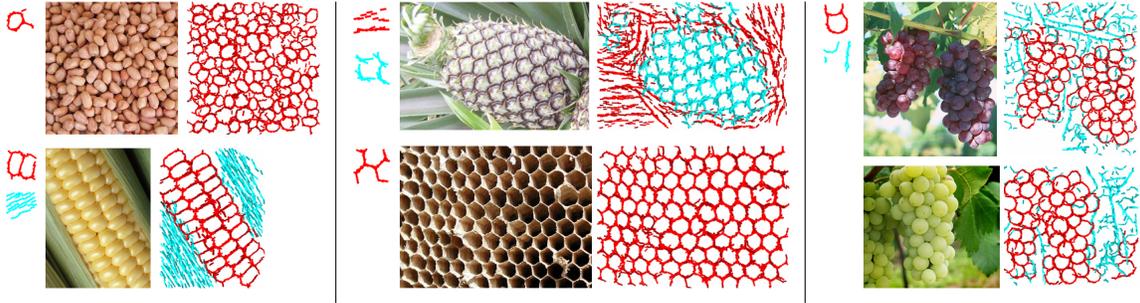


Figure 9: Peanuts, corn, pineapple, beehives, and grape. Parameter setting is the same as in Figure (5). For the grape example, the bottom row are testing image and its representation by the dictionary learned from the training image on the top row.

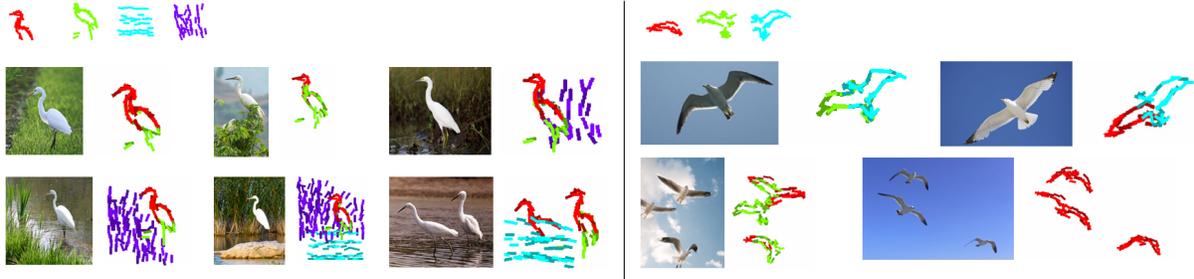


Figure 10: Egret (number of training images is 20) and seagull (number of training images is 11). Parameter setting is the same as in Figure (5), except that the allowed range of rotation for the templates is $[-2, 2] \times \pi/16$. In egret example, the templates for water waves and grasses are more texture-like than shape-like, because these patterns do not have well-defined shapes.

X , scale $S \in \{.8, 1, 1.2\}$ and orientation $A \in \{-1, 0, 1\} \times \pi/16$ (see subsection (3.1) and Equation (11)). Let $r_m^{(t)}(A) = \max(\max_{X,S} \mathbf{R}_m^{(t)}(X, S, A), 0)$ be the maximum score (lower-bounded by 0) at orientation A . Then for each \mathbf{I}_m , we have a $3T$ -dimensional feature vector $(r_m^{(t)}(A), t = 1, \dots, T, \forall A)$ (the factor 3 is due to the fact that we keep the scores of all three orientations). We then train a linear logistic regression on such $3T$ -dimensional feature vectors (regularized by ℓ_2 norm [17]) for image classification.

We evaluate this simple classifier on 16 categories from Caltech-101 [19], all ETHZ Shape [20] and all Graz-02 [36] data sets, where we test it on binary classification task. We resize all images to 150^2 pixels while maintaining their aspect ratios. We randomly sample 30 positive and negative

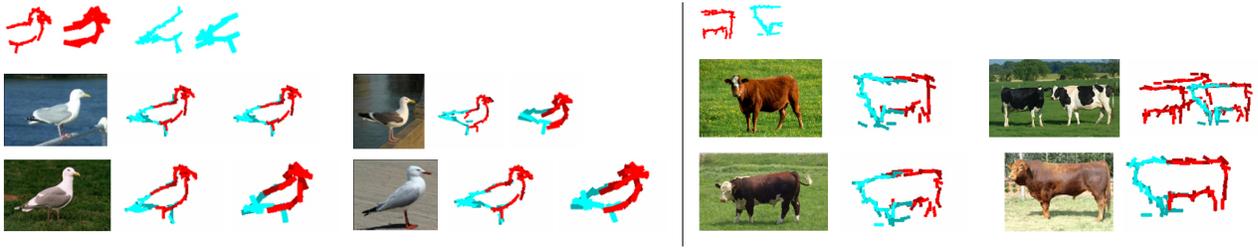


Figure 11: Seagull (number of training images is 10) and cattle (number of training images is 17). Parameter setting is the same as in Figure (10). For the seagull experiment, we learn multi-scale templates at two different scales (the scale parameters are .7 and 1.4 respectively). We sum the log-likelihood scores of the templates at the two scales in order to compute the template matching scores in the template matching pursuit process.

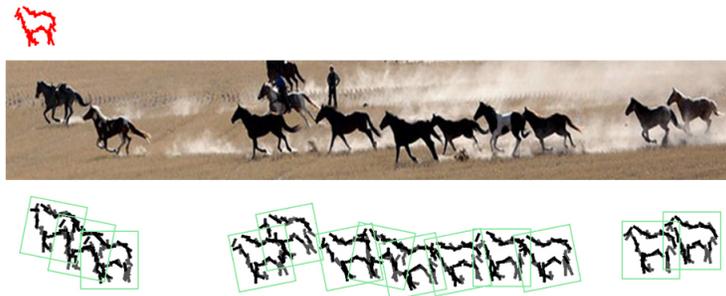


Figure 12: Horse. Parameter setting is the same as in Figure (5), except that the allowed range of rotation for the template is $[-1, 1] \times \pi/16$.

images respectively as training data, and leave the rest as testing data. For Caltech-101 and Graz-02, negative images are from background category. For ETHZ, negative examples are from images other than the target category. For each category, we learn a dictionary of $T = 10$ templates. Each is of the size 100×100 with $n = 30$ wavelet elements.

As a comparison, for each image, we densely extract SIFT features [33] with patch size 16×16 and step size 8, from both positive and negative images, quantize them into 50, 100 and 500 words respectively by k-means clustering [9] and feed them into SVM [49, 7] (linear and histogram intersection kernel [29, 34]). We take the best of these 6 results (3 numbers of words (50, 100, 500) \times 2 types of SVM (linear, kernel)) and compare it with our method. All experiments are carried

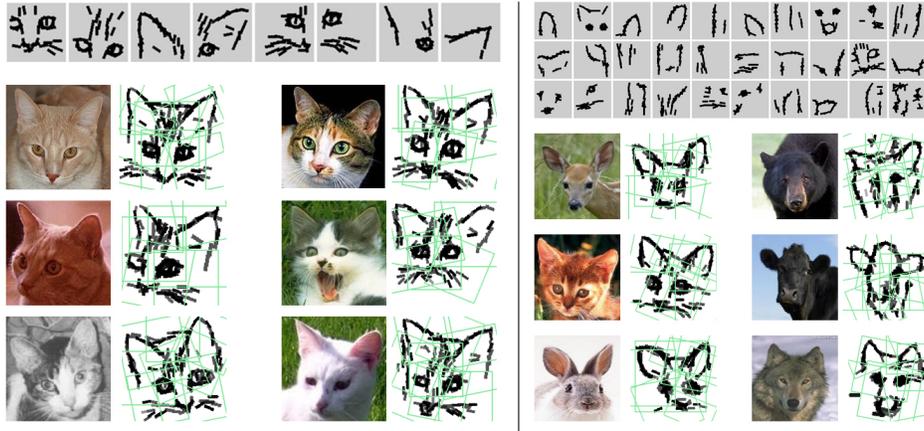


Figure 13: Cat faces (number of training images is 89) and animal faces (number of training images is 490). Parameter setting is the same as in Figure (10).



Figure 14: Cars. Parameter setting is the same as in Figure (10). Number of training images is 245.

out with 5 independent runs and the 95% confident intervals on accuracies are calculated. Table 1 presents the results. It shows that our method generally outperforms the popular SIFT + SVM method even though our method uses a much smaller dictionary of words (10 in our method versus 50, 100, or 500 in SIFT + SVM).

We also test our method on the whole Caltech-101 data set. We learn $T = 200$ templates from the training images from all the 101 categories together. In one set of experiments, 15 training images are randomly sampled from each category in each run. In another set of experiments, 30 training images are randomly taken from each category in each run. In each set of experiments, 5 runs are repeated.

Each template is of the size 64×64 with $n = 15$ wavelet elements. During the learning algorithm, if the number of image patches encoded by a template is less than a threshold, then this template

Table 1: Accuracies (%) on binary classification tasks for 24 categories from Caltech-101, ETHZ Shape and Graz-02 data sets.

Datasets	SIFT+SVM	Our method	Datasets	SIFT+SVM	Our method
Watch	90.1 ± 1.0	91.3 ± 2.0	Sunflower	76.0 ± 2.5	92.9 ± 2.5
Laptop	73.5 ± 5.3	87.9 ± 2.2	Chair	62.5 ± 5.0	89.1 ± 1.1
Piano	84.5 ± 4.2	93.4 ± 3.0	Lamp	61.5 ± 4.5	81.7 ± 3.7
Ketch	82.2 ± 0.8	89.2 ± 2.4	Dragonfly	66.0 ± 4.0	87.0 ± 4.1
Motorbike	93.9 ± 1.2	93.7 ± 0.9	Umbrella	73.4 ± 4.4	89.3 ± 2.5
Guitar	70.0 ± 2.4	80.9 ± 5.1	Cellphone	68.7 ± 5.1	87.9 ± 4.2
Schooner	64.3 ± 2.2	93.8 ± 2.7	Face	91.8 ± 2.3	95.8 ± 2.8
Ibis	67.8 ± 6.0	83.0 ± 1.9	Starfish	73.1 ± 6.7	85.3 ± 4.7
ETHZ-Bottle	68.6 ± 3.2	76.1 ± 3.3	ETHZ-Cup	66.0 ± 3.3	67.5 ± 4.4
ETHZ-Swans	64.2 ± 1.5	82.4 ± 0.5	ETHZ-Giraffes	61.5 ± 6.4	71.5 ± 3.5
ETHZ-Apple	55.0 ± 1.8	68.3 ± 5.2	Graz02-Person	70.4 ± 1.2	73.8 ± 2.3
Graz02-Car	64.0 ± 6.7	63.5 ± 5.1	Graz02-Bike	68.5 ± 2.8	77.6 ± 2.3

is eliminated from the dictionary. For 15 training images per category, the threshold is set at 5. For 30 training images per category, the threshold is set at 10. Figure (15) displays the learned dictionary of templates in one run of experiment with 30 training images per category. They seem to capture the mid-level structures such as lines, corners and circles etc.

For each image \mathbf{I}_m , and for each template $\mathbf{B}^{(t)}$ at each orientation A , besides the global maximum $r_m^{(t)}(A)$, we also divide \mathbf{I}_m equally into 2×2 sub-regions and take the maximum within each sub-region. In addition to each maximum, we also take the corresponding average. So each $\mathbf{B}^{(t)}$ extracts 30 features from \mathbf{I}_m . Thus in total, each \mathbf{I}_m produces a $30T$ -dimensional feature vector. We adopt standard evaluation protocol. For 15 training images per category, the accuracy of our method is $61.6 \pm 2.2\%$ (compared to $57.7 \pm 1.5\%$ in [31]). For 30 training images per category, our result is $68.5 \pm 0.9\%$ (compared to $65.4 \pm 0.5\%$ in [31]). While more recent papers such as [53] and the references therein report better performances based on spatial pyramid matching [29], we do not



Figure 15: Learned dictionary of templates from the Caltech-101 dataset in one run of an experiment, with 30 training images from each category. Template size is 64×64 . Number of wavelets in each template is 15.

use k-means to further cluster response maps into another layer of codewords, neither do we use any kernel.

5 Discussion

In this section, we first discuss the contributions and limitations of our current work. Then we shall compare our work with related work in the literature.

5.1 Contributions and limitations

The main contribution of this paper is conceptual: to explore what is beyond the wavelets sparse coding or atomic decomposition. We argue that a natural and necessary step to go beyond wavelets representation is to learn compositional patterns or shape templates formed by the wavelets. We propose a representational scheme based on composite representational units, which are groups of wavelets of recurring compositional patterns, and we have developed an unsupervised learning algorithm for learning dictionaries of compositional patterns from training images, where the compositional patterns arise from seeking commonly shared sparse coding of image patches. Our experiments on natural images of plants and animals etc. show that our method is capable of learning meaningful compositional patterns, which lead to meaningful representations of training

and testing images.



Figure 16: Image scaling and regimes of patterns: As we zoom out the images, the patterns undergo a transition from low-entropy regime of geometric structures to mid-entropy regime of object shapes to high-entropy regime of stochastic textures. Our current method targets the mid-entropy regime of shape patterns represented by compositions of wavelets.

While it is possible to design suitable dictionaries of wavelets for efficient representations of natural images, as in [14, 13, 4, 5], it may be impossible to design the rich varieties of compositional patterns or shape templates for natural images. Unsupervised learning methods such as ours are thus necessary for automatically discovering spatial grouping patterns of wavelets from natural images.

In terms of biological plausibility, the Olshausen-Field model is a model for simple V1 cells. The local max pooling of Riesenhuber and Poggio [43] and the arg-max retrieval and inhibition of the active basis model may be related to complex V1 cells. The dictionaries of active basis templates learned by our method may be related to V2 cells and beyond.

The following are limitations of our work. First, as illustrated by Figure (16), as we zoom out the images, the image patterns undergo a transition from low-entropy regime of geometric structures to mid-entropy regime of object shapes to high-entropy regime of stochastic textures [50] (patterns such as the brick wall are also textures, but they are structural textures instead of stochastic textures). Our current model mainly targets the mid-entropy regime of object shapes and textures. It does not account for stochastic texture patterns or appearance patterns that are ubiquitous in natural scenes. The current model does not account for flatness patterns that are prominent in the low-entropy regime either. Another limitation is that the model is still not a fully generative model. We have not modeled the spatial arrangements of the templates, nor have we considered further composing the templates into yet another layer of composite representational

units. A third limitation is that we assume that the dictionary of the wavelets are given as Gabor wavelets. It is desirable to learn these wavelets from training images by what may be called active component analysis. We shall study this issue in future work

The active basis models are currently learned by generative approach based on likelihood. It may be possible to learn the models discriminatively by regularized logistic regression after bringing in negative image patches, or to learn the models based on a combination of discriminative and generative loss functions.

To end this subsection on a positive note, despite all the above-mentioned limitations, we are pleased that our method can learn meaningful dictionaries and they prove to be useful for classification. At the minimum, we have proved the concept that it is possible to go beyond atomic decomposition and explore molecular representation, just as it is necessary to go beyond alphabet to have words to describe the real world.

5.2 Related work

AND-OR grammar. Our work connects the sparsity principle to another fundamental principle in vision, namely, compositionality [24, 56], which holds that the visual patterns are hierarchical compositions of constituent parts. In particular, in the language of AND-OR grammar of Zhu and Mumford (2007), the dictionary of active basis templates can be considered a big OR node, where each template is a child node of this OR node, and each template is itself an AND-OR structure, where AND means composition of the constituent wavelets, and OR means perturbations of the wavelets.

In terms of composing Gabor wavelets, our representation is similar to that of [21] as well as [54]. The difference is that, our method is based on a top-down generative model, where the compositional patterns are re-learned in each iteration of the learning algorithm from raw image patches by seeking to maximize the likelihood. This enables us to compose many wavelets into a large template in a single layer, instead of recursively building up the templates via multiple layers.

Deep learning. Our work is related to deep learning [26], especially deep learning with sparsity constraint [28, 31, 47, 53]. The difference is that our representational units are sparse compositions of automatically selected wavelet elements, where sparsity is explicitly built into the representational units by the shared matching pursuit process. The representational units are no longer linear basis

functions on top of the wavelets coefficients or filter responses at the lower layer. Also, our model is not built on a pre-processed sparse representation, which, as we have argued in subsection (3.3), amounts to early decision. In our learning algorithm, each iteration re-learns each template from raw image patches, where the sparse representations, their correspondences, and the template are obtained simultaneously instead of being obtained one after another.

HMAX model. Our work is related to the HMAX model of [43]. The local max pooling is employed for inferring the perturbations of the wavelet elements of the active basis model. In HMAX, the dictionary of the second layer consists of maps of local max pooling of Gabor responses. In our work, we explicitly learn the recurring compositional patterns of the wavelets guided by a generative model.

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