

# Efficient Inference of Spatial Hierarchical Models\*

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**Abstract:** The long term goal of artificial intelligence and computer vision is to be able to build models of the world automatically and to use them for interpretation of new situations. It is natural that such models are efficiently organized in a hierarchical manner; a model is build by sub-models, these sub-models are again build of another models, and so on. These building blocks are usually shareable; different objects may consist of the same components. In this paper, we describe a hierarchical probabilistic model for visual domain and propose a method for its efficient inference based on data partitioning and dynamic programming. We show the behaviour of the model, which is in this case made manually, and inference method on a controlled yet challenging dataset consisting of rotated, scaled and occluded letters. The experiments show that the proposed model is robust to all above-mentioned aspects.

## 1 INTRODUCTION

Efficient inference of complex hierarchical probabilistic models is an important milestone on the way to automatic, unsupervised learning of such structures. These structures are important because they represent a simple and efficient way of modelling real world objects, scenes and situations which almost always exhibit certain level of compositionality and hierarchical structure (Tsotsos, 1990; Bienenstock et al., 1996).

Hierarchical models are also very efficient in precisely describing very complex objects with completely natural ordering of semantic information on the given object. They model very local properties on lower layers and general ones on higher layers and they are also capable of describing very complex mutual relations, e.g. non-rigid shape deformations caused by rotations and stretching of certain model components. These advantages are unfortunately counterweighted by the fact that these models are computationally very demanding when classifying with them.

Although these models can work with any type of information like variously described image regions (Ommer and Buhmann, 2010; Ullman, 2007), the models describing shapes are of special interest

for us. The most common approaches to instantiating such shape models in unknown data are based on *dynamic programming* (Zhu and Mumford, 2006; Zhu et al., 2010; Zhu et al., 2011) with pruning, *coarse-to-fine* schemes (Kokkinos and Yuille, 2011) or *Bottom-up* building of instances and their sparsification<sup>2</sup> (Fidler and Leonardis, 2007). These works also deal with learning such models, either their complete structure or their parameters, which is also our long-term goal, but is outside the focus of this paper.

## 2 CONTRIBUTION

In this paper, a new simple yet effective probabilistic model is proposed. It is philosophically inspired by the *Bottom-up/Top-Down* system (Fidler et al., 2006; Fidler and Leonardis, 2007), but there are significant differences. Particularly novel is

- the compositional model itself, which is fully cast in probabilistic framework and can easily cope with missing or redundant data, though it is still reasonably simple and easily computable analytically (cf. (Kokkinos and Yuille, 2011)).

<sup>2</sup>Preferably globally optimal reduction of generated hypotheses as to avoid combinatorial explosion on higher layers. This basically means at least approximately solving a Set-Covering problem, known and proven to be NP-complete (Karp, 1972).

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- The inference system generates lower number of *Bottom-up* hypotheses, which means that sparsification is not necessary. This results in more efficient utilization of computational resources.
- The object structure is acyclic allowing for exact computing of model probability using simple *Belief propagation*<sup>3</sup>.
- As a side effect, the *Bottom-Up* process can be easily implemented and run in parallel, achieving another computational time savings.

### 3 CONCEPTS

The key concepts in this framework are an *instance* (of a composition) and an *area* which serves as a container for *instances*.

The notation is the following: the individual compositional models on the layer  $a$  are distinguished by subscript ( $a_1, a_2$ , etc.) and individual instances of a composition  $a_k$  are indexed by a superscript ( $a_k^1, a_k^2$ ).

An instance – always associated with a certain layer – is a hypothesis on presence of a realization of a composition from the library conditioned on the observed data  $D_i \in \mathcal{D}$ . This hypothesis is assigned probability, e.g.  $P(D_1, D_2, \dots, D_n | a_j^k)$  at the layer  $a$  or  $P(D_1, D_2, \dots, D_n | \sigma_j^k)$  at the root layer, etc.

At the beginning of processing, the initial set of areas to start with is the set of pixels. At each layer, an image gets partitioned into areas.

Having a partition set at the layer  $l$ , the partition set at the higher layer  $l + 1$  is obtained simply by merging spatially close sets from layer  $l$  together. That way, each area at the higher layer is composed of several areas in the lower layer. Optimally, we would like each area at the higher layer to have similar size and to contain similar number,  $N$ , of lower layer areas.

It is not possible to satisfy both constraints, so we use an algorithm which is only sub-optimal with this respect, however is simple and fast: It selects an area at the lower layer randomly, then merges it with  $k$  nearby areas ( $k$  approaches  $N$ , and the closer the better).

As a consequence, the collection of area sets over all layer forms a tree. Having formed the areas, instances are formed based on underlying children and every single instance has to contain all area children. That way each child is used exactly once within one

<sup>3</sup>To avoid the need of using the general loopy version which takes more computational time as well as has still some unsolved issues (Weiss, 2000; Mooij and Kappen, 2007).

instance and the tree structure (acyclicity) of instance is guaranteed.

Problems occur when there is a piece of data (an instance) in the area which is not modelled by a certain composition which on the other hand models the rest of the local data well. This is overcome by the choice that the compositions are represented by generative models, so they can be equipped by an ability of generating random patterns according to some probability distribution.

Another mechanism related to area is its so called *empty hypothesis* ( $\sigma_e$ ). This hypothesis states that the data within the area was created randomly without any known (explicit) model.

This *empty hypothesis* is necessary for correct inference of the model's presence. This necessity is justified by the fact that the library can not contain descriptions of all possible objects in the domain (world), because there is an infinite number of them, and therefore a mechanism for detecting unknown patterns (previously unseen) is required. The *empty hypothesis* does the job - if the probability  $P(\sigma_e | D_1, D_2, \dots, D_n)$  is higher than the probabilities of the other models, the underlying data are either part of a poorly-modelled object or real random noise and the learning algorithm can focus on this area.

#### 3.1 Example

An illustration of an area and a composition is shown in the Figure 1. The area contains three instances ( $b_1^1, b_2^1$  and  $b_3^1$ ) at the layer  $b$  and for the layer  $a$ , there is one composition  $a_1$  consisting of two components  $b_1$  and  $b_2$ . Because of the restriction that there can only be one instance in one area at a time, there are two hypotheses for such area possible. It can be either the hypothesis that the composition  $a_1$  generated the data within the area or that this data has been generated by the background process  $a_e$ .

The probability of the data conditioned on the presence of an instance of the composition  $a_1^1$  is given by

$$\begin{aligned}
 P(D_1, D_2, D_3 | a_1^1) &= P(D_1 | a_1^1) P(D_2 | a_1^1) P(D_3 | a_1^1) = \\
 &= \int_{b_1^1} P(D_1 | b_1^1) P(b_1^1 | a_1^1) db_1^1 \times \\
 &\times \int_{b_2^1} P(D_2 | b_2^1) P(b_2^1 | a_1^1) db_2^1 \times \\
 &\times \int_{\Omega} P(D_3 | b_e) P(b_e | a_1^1) db_e \quad (1)
 \end{aligned}$$

and  $P(D_3 | a_1^1)$  in this case represents the probability

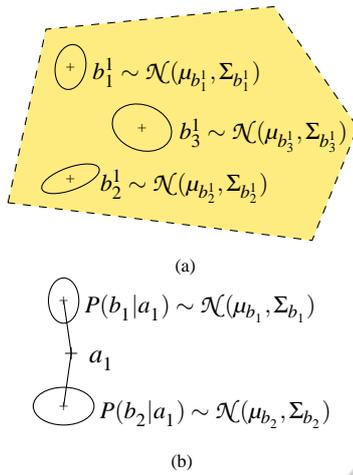


Figure 1: (a) An example of an area containing three instances  $b_1^1$ ,  $b_2^1$  and  $b_3^1$  on the layer  $b$ . (b) The model of the composition from the layer  $a$  consisting of two components whose configurations are specified by the conditional probabilities  $P(b_1|a_1)$  and  $P(b_2|a_1)$  respectively. The composition  $a_1$  models the area in the (a) with high probability and in that case the instance  $b_3^1$  is considered as a random noise.

that the  $D_3$  has been generated by the background process.  $\Omega$  represents the area size (the area of circle with radius of predefined constant related to the actual position in the hierarchy).

The probability that the whole of the content of the area has been generated accidentally (the probability of the *empty* hypothesis) is given by the formula

$$P(D_1, D_2, D_3 | a_e) = \prod_{i=1}^3 \left( \int_{\Omega} P(D_i | b_e) P(b_e | a_e) db_e \right), \quad (2)$$

where  $b_e$  and  $a_e$  stand for labels of *empty* hypotheses.

Having known prior probabilities  $P(a_1)$  and  $P(a_e)$ , the probability of presence of the object  $a_1$  can be easily computed using Bayes formula

$$P(a_1 | D_1, D_2, D_3) = \frac{P(D_1, D_2, D_3 | a_1) P(a_1)}{P(D_1, D_2, D_3 | a_1) P(a_1) + P(D_1, D_2, D_3 | a_e) P(a_e)} \quad (3)$$

and similarly for the probability of the *empty* hypothesis.

## 4 PROBABILISTIC MODEL

A detailed scheme of the probabilistic model of a composition is shown in the Figure 2. Nodes  $a$ ,  $b$  represent the basic compositional model as shown e.g. in the Figure 1(b). Apart from these compositional nodes, there is a hidden layer of nodes in this model.

These nodes, named as  $m_1$ ,  $m_2$  etc., are two-state and indicate if the corresponding data (or sub-component) is present or missing. The right-most node named  $e$  is a discrete-state node stating the number of unassigned (unexplainable) data. The individual conditional probabilities are

$$P(b_{11} | m_1 = 1) \propto \mathcal{N}(\mu, \sigma), \quad (4)$$

$$P(\{\} | m_1 = 0) = 1, \quad (5)$$

$$P(m_1 | a) = \begin{cases} p_1 & \text{if } m_1 = 1 \\ 1 - p_1 & \text{if } m_1 = 0 \end{cases}, \quad (6)$$

$$P(e | a) = \frac{\lambda^k}{k!} e^{-\lambda} \quad (7)$$

and

$$P(b_e | e) = \mathcal{N}(b_e; \mu_{\text{AREA}}, \Sigma_{\text{AREA}}) \quad (8)$$

with  $p_1$  and  $\lambda$  as new parameters. The symbol  $\{\}$  stands for a missing part.

The advantage of this model is that it can explicitly model every possible data configuration, regardless of the fact that the data can be completely alien to the structural model included. In that case, all data is assigned to the  $e$ -branch and all  $m$  nodes are set to the state *missing*. Naturally, the probability of this model would be in that case lower than the probability of the *empty* model, because besides the  $e$ -branch in the compositional model, there are also unassigned compositional branches which serve as a penalty term in such scenarios.

Even though the model is more complex than the schematically introduced one, it is still quite easy to evaluate (marginalize the hidden nodes) because all hidden nodes ( $m$  and  $e$ ) are constructed such that they have only one non-zero state at a time. An item  $D_i$  is always either present or missing, and the number of non-assigned items is also always known – it depends on the actual assignment of data to individual leaf nodes which is fixed – from all possible assignments, the one resulting in the maximal probability of instance is selected. This indeed implies that the probabilities are not precise but approximated, but simple experiments show that other assignments give significantly (in terms of magnitudes) lower probabilities and therefore even in the sum are negligible. Moreover, this choice keeps the resulting probability proportional to a normal distribution.

## 5 INFERENCE SCHEME

The inference works gradually from the bottom layer to higher layers exploiting principles of *Belief Propagation* (Bishop, 2006; Pearl, 1988) which means that first the compositions of the lowest model layer are

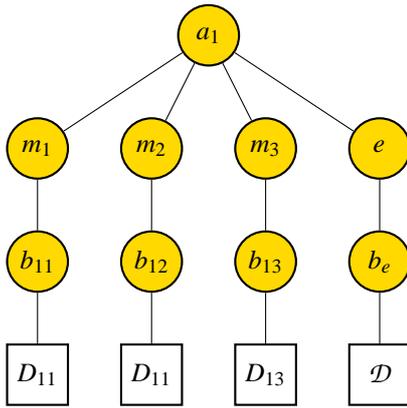


Figure 2: The structure of the probabilistic model of a composition, the rectangular nodes denoted with letter  $D$  are input data, the round nodes  $m$ ,  $e$  and  $b$  are internal nodes, Nodes  $b$  model spatial relations of children and parent,  $m$  model the decision if the child is present. Such compositions are organized hierarchically to model whole objects.

instantiated using a method similar to (Felzenszwalb and Huttenlocher, 2000) and all internal model nodes are marginalized (in the case of the example in the Section 3.1 nodes  $b$ ,  $m$  and  $e$ ) and instances are then taken as new 'input' data. This is done in each area and after the layer inference is done, new area partitioning on higher layer is made. This process goes iteratively until the top (root) layer is achieved.

What makes this inference scheme robust is a simultaneously running *Top-Down* adjustment process which takes non-optimally found instances and tries to improve them – either by focused search of missing underlying instance which might have been missed due to non-optimal area division, or by search of new underlying instances in order to improve instances' probability (when a sub-instance is e.g. too far from its expected position).

The inference can be summarized by the Algorithm 1.

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**Algorithm 1:** The sketch of inference algorithm with its steps.

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- 1) get layer 1 instances from an image **for**  $i \leftarrow 1$  **to** 5 **do**
    - 2) find random partitioning of instances of previous layer;
    - 3) infer instances of current layer (*bottom-up* process);
    - 4) merge the partitioning in previous layer;
    - 5) improve the instances - find missing parts (*top-down* process);
- end**
- 

In the step 3 of the Algorithm 1, the efficiency is

achieved by creating not all possible but only at most five most probable hypotheses. The step 4 discards the lower layer partitioning which allows to collect and complete the instances that have been split into multiple areas which affects their probabilities. This step is actually very simple, because these split instances have similar locations and therefore it is sufficient to track multiple compositions from different areas appearing close each other and fuse them together. This process cancels possible over-fitting of areas partitioning in favour of the highest-probable instance. The step 5 is also quite lightweight – again only a limited number of instances are subject to updating. The update itself works in the following scheme. The instance tree is analysed and missing and non-optimally detected sub-instances are detected. These candidates for updating are sorted according to their importance, which in this case means the highest layer has priority, because changes on this layer embodies the most significant changes of instance's probability. Furthermore, it does not make much sense to improve the model on a lower layer and then replace this branch completely. This *top-down* update of an instance ends when there are no defects detected or the maximal number of attempts is reached.

## 6 EXPERIMENTS

The aim of the experiment was to show that the probabilistic model and inference scheme are suitable for character recognition with possible rejection of decision when the model fails to model the data.

### 6.1 Character Recognition

The model (its structure) for two letters –  $a$  and  $b$  (see the Figure 3) has been built manually and consists of five layers of compositions. The whole library of compositions is shown in the Figure 4. The models are set in to represent the individual shapes with the goal to create share-able models and make the library as compact as possible. Nevertheless, the correspondence of the model to the dataset images is not perfect, especially the sub-models locations on higher layers were chosen by hand and no optimization with respect to the locations of individual composing components was done in order to come closer to the real-world situation when the unknown data always differ from learned models and to see if the model is not too sensitive to the choice of its parameters.

Probabilities  $p_1$  in the Eq. 6 were set equally for all library compositions as 0.9. Parameters  $\lambda$  for Poisson distribution used in the Eq. 7 were also set equally

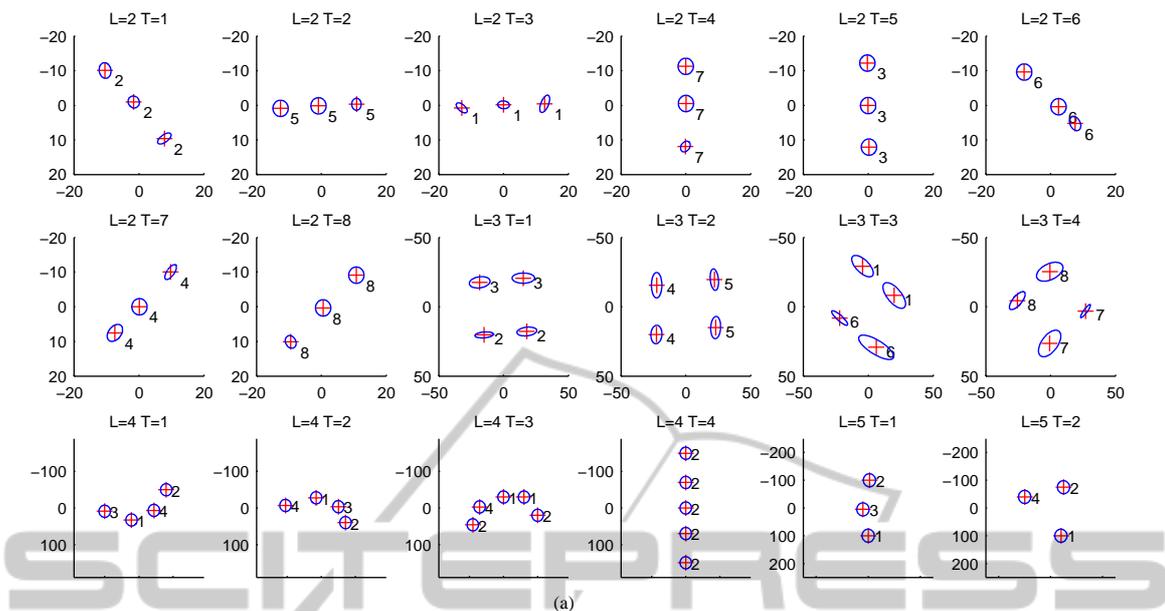


Figure 4: Complete model library. Letter  $L$  depicts the layer within the hierarchy, letter  $T$  its type (id) within the layer and numbers in the pictures encode the underlying component type (can be matched with  $T$  in preceding layer). The model library can be seen as a recursive structure, one would obtain the structures shown in the Figure 3 by taking the model from 5th layer and placing the appropriate models from the 4th layer at marked positions and repeating that until the second layer.

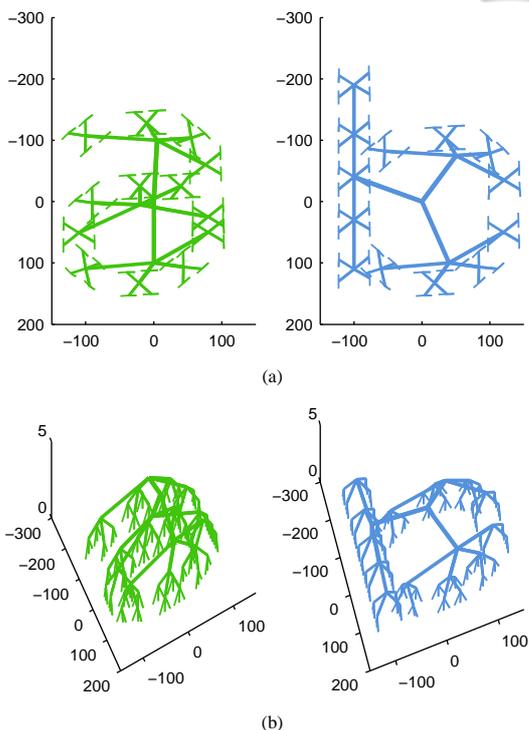


Figure 3: The complete structure models of  $a$ , resp.  $b$  in the (a) from top, in the (b). The  $z$  axis shows the depth of the models, layers are also emphasized by the thickness of the line.

for all compositions to the value 1 and for the case of *empty* hypothesis the value was set to be the *maximum likelihood* estimate over corresponding layer. This parameter reflects the average number of data points within an area on certain layer.

The dataset consists of images of letters  $a$ ,  $b$ ,  $c$  and  $d$  which were randomly rescaled by a factor ranging from 0.95 to 1.05 and also randomly rotated by a small angle, up to 5 degrees<sup>4</sup>.

These images were processed by a pair of orthogonal Gaussian filters to find edges and their orientation. The edges have been sorted according to their orientations into eight groups. The set of these edges represents the lowest (first) layer. To make the data more challenging and to show that the method is naturally insensitive to occlusions, a random square region covering 25% of that particular image was cut out in the first version of the experiment and converted into random noise in the second version of the experiment.

### 6.1.1 Results

The inference scheme was tested on 61 images from the described dataset, the numbers of each letter were set equally to 15 instances and the 61st image was generated randomly but with the same statistics of

<sup>4</sup>Within this range are e.g. the usual residual errors after preprocessing steps like rectification or rotation and scale compensation.

bottom layer instances<sup>5</sup>. This additional random pattern was intended to show the behaviour of the method on completely unknown data.

The selected letters exhibit a high level of similarity – especially in the case when a random 25% of an image is cut out. It can happen that the piece which has been cut out was the most discriminative feature between the letters. Consider the appearance of letter *a* with its right part cut and its similarity to letter *c* as an illustrative example. There is definitely a difference in the scale in this particular example, but this difference is cancelled (compensated) by scale invariance built in the model. Therefore, the success rate might not be as high as one would expect, especially on the second half of the dataset when poorly modelled but still compositional data is being dealt with.

In the first version of the experiment, the overall success rate on the first half of the data was 100% which means that all data that was explicitly represented in the model library were recognized properly, some examples are shown in the Figure 5(a), Figure 5(b) and Figure 5(c). However, in few images the models were not located precisely, as can be seen in the Figure 5(d), resulting in the overall location precision of 90% – 80% for the *a* category, 100% for the *b* category – correctly located models. These misplacements are caused mainly by the fact that both models are built of similar components – in the case of letter *a* the middle part and top part models differ only in two sub-models (see the Figure 4, models denoted  $L=4 T=2$  and  $L=4 T=3$ ) and when some of important sub-model is not found – due to occlusion or due to unfavourable partitioning – the winning hypothesis might be wrong. Nevertheless, this is a problem only on the top layer, lower layers get updated using the information from the top.

The results on the other half of the data show how the method behaves on a not-so-well modelled data. In the cases of letters *c* and *d*, it tends to instantiate any of the library models, because there is almost always a part of letter which can be interpreted using a model from library, especially on the lower layers. In the case of the last image – the random pattern – it shows the desired property that unknown pattern is quickly labelled as unknown (*empty* hypothesis wins).

The complete confusion matrix for the labelling is shown in the Table 1, the *empty* hypothesis is denoted as  $\emptyset$ .

In the second version of the experiment, when the occluded part of the image data was covered by ran-

<sup>5</sup>This random pattern was constructed by taking an image, processing it and then randomly permuting instances locations.

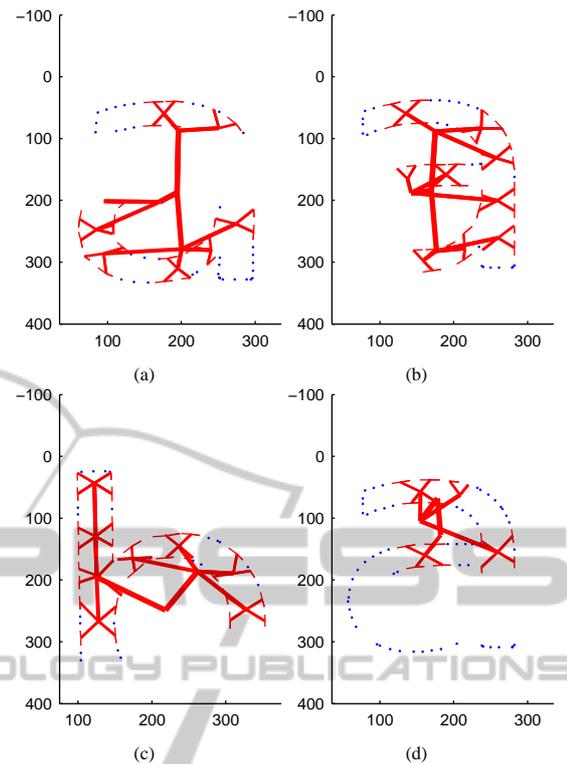


Figure 5: A few selected examples of winning model instantiation in the first version of the experiment (missing data), blue dots are unexplained edge segments. (d) shows a misplaced model (the only example in the results), (a), (b) and (c) show completely correct instantiations. It is also visible which parts of image were cut off.

Table 1: Confusion matrix for the letters dataset labelling for missing data, rows show the correct label and columns assigned label.

	a	b	$\emptyset$
a	1.00	0.00	0.00
b	0.00	1.00	0.00
$\emptyset$	0.52	0.32	0.16

Table 2: Confusion matrix for the letters dataset labelling (occlusion by random noise), rows show the correct label and columns assigned label.

	a	b	$\emptyset$
a	0.73	0.20	0.07
b	0.00	1.00	0.00
$\emptyset$	0.45	0.52	0.03

dom noise, the results are slightly worse, achieving 86.7% labelling accuracy on the *a* and *b* categories. On the rest of the data, the method behaved similarly as in the previous case and the completely random pattern – the image 61 – was again labelled correctly as unknown. Regarding the correctness of location, in

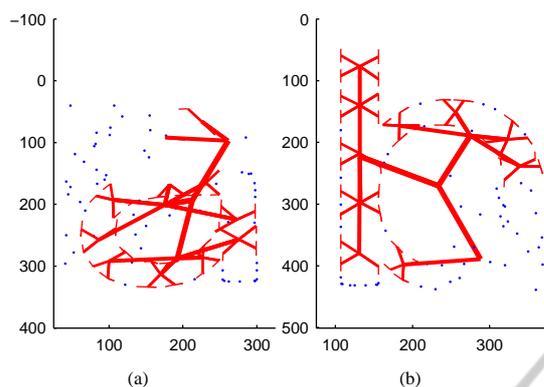


Figure 6: Two selected examples of winning model instantiation for the second version of the experiment (occlusion by random noise), the colour interpretation is the same as in the Figure 5.

all images that were correctly labelled were also the models correctly located, achieving 86.7% accuracy without any fine-tuning of the parameters. The complete confusion matrix can be found in the Table 2 and two illustrative results are shown in the Figure 6.

## 7 CONCLUSIONS

In this paper, a new hierarchical probabilistic model for modelling object appearance is introduced and an efficient method for its inference is described. Main and novel features of this approach are the following: first, the model is acyclic by definition, which allows for precise computation of probabilities using a very simple version of *Belief propagation*. Second, thanks to the partitioning of layers it is easy to compute the probabilities of the model conditioned on the observed data, which is very useful for *Maximum-likelihood* learning of parameters. This partitioning also prevents us from the combinatorial explosion in the *Bottom-up* generation of hypotheses and allows for parallel processing. The model and the method is also robust to occlusions.

The experiments on a controlled dataset of images of letters with a hand-made hierarchical model show that the proposed approach is generally usable for visual data. The dataset is characterized by small rotations and scale changes as well as occlusions (non-presence or replacing by random noise) of 25% of the image, none of which causes significant difficulties to the inference algorithm.

To briefly outline the future directions, the framework is planned to be used in a completely unsupervised structural learning method and is going to be compared to the state-of-the-art methods in structural learning and recognition.

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