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Introduction and motivation



3 Ranking task



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Introduction and motivation

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- How can we generalize in such a case?

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using a set of task representations or vectors (given a priori)

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• Why do this?

- this situation occurs in certain applications : drug discovery, song/movie recommandation, NLP tasks
- gives us a worst case scenario in an inductive transfer model
- hasn't been studied so much

Introduction and motivation



I will present some experiments in the case of a linear model



Introduction and motivation

In this talk

- I will present some experiments in the case of a linear model
- I will compare different training settings for the linear model in a classification task :

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- discriminative (with 2 different cost functions)
- generative

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- discriminative (with 2 different cost functions)
- generative
- I will present results for two problems :
 - classifi cation problem
 - ranking problem

Classification task

First problem : classifi cation

 Tasks' nature : character recognition in the context of license plates



Classification task

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- Tasks' nature : character recognition in the context of license plates
- **Training tasks** : we are given a set of class descriptors (vectors) with corresponding instances from these classes
- Test tasks : we are given two class descriptors and we need to discriminate between these two classes for a given set of untagged instances

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Classification task

First problem : classifi cation



Classification task

Model variations

• Question : how do we get a model (classifier) for a new task?

Classification task

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- Answer : by predicting it.

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- In general, we could model "models" like this :

$$f_{y}(x) = \left[g(d_{y})\right](x)$$

where x is the input, y is a task (class), d_y is it's descriptor and $g(d_y)$ predicts f_y **Classification task**

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 Here, we will have g(d_y) be a linear transformation A'd_y, hence

$$f_y(x) = d_y A x$$

or, in the energy-based framework

$$E(x,y)=-d_yAx$$

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Generalizing to a zero-data task: a linear model case study Classification task

Model variations

We will compare 2 training criteria

discriminative (1 vs all)

$$\mathcal{L}(x, y) = -\log(\operatorname{softmax}(-E(x, y)))$$

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 $D_y \sim \mathcal{U}(0, 1)^m$

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• We use standard regularisation techniques, such as early stopping, weight decay, etc.

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- For each of the 4 folds, we select two classes to isolate from others
- Validation is performed on a set of instances from the training classes
- We vary the number of classes present in the training set, but we keep the same test classes of the different folds

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Classification task

Results



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Classification task



Generative model does better when few task examples are available



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 Discriminative model does better when many task examples are available

Classification task



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- Discriminative model does better when many task examples are available
- More "examples" of tasks doesn't imply better generalisation

Classification task

Other model variation

Discriminative 0-1 multi-task classification

$$\mathcal{L}(x,y) = -\log(\operatorname{sigmoid}(-E(x,y))) \\ -\sum_{c \in \mathcal{Y}, c \neq y} \log(1 - \operatorname{sigmoid}(-E(x,c)))$$

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Classification task

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Ranking task

Second problem : ranking

• Tasks' nature : virtual screening for drug discovery

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We can then rank compounds in decreasing probability of activity, and makes suggestions of compounds to test.

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• Nature of descriptors d_y : TOP SECRET!

Experimental setup

 We have 7 proteins and a bank of compounds. All compounds has been previously tested for certain proteins. The target can be "1" for an "active" compound and "0" for an "inactive" compound.

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- The same model is used as in the first experiment. We use the cost of the sum of the cross-entropy, for the available activity target.
- The evaluation metric is the "lift" :

$$LIFT = 100 \cdot \frac{a_{s}}{a_{100}}$$

where a_s is the fraction of active compounds in the *s* first compounds, as ranked by the model. Here, we use s = 30, which is a standard size in computational chemistry.

Ranking task



LIFT results for the different proteins.

Protein	LIFT
А	167.90
D	152.89
F	97.07
Н	170.57
Ι	163.07
S	172.91
U	95.27

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For the training tasks, LIFT is around 220.

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Future work

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Extention to possibly get rid of task-overfitting, by having

$$E(x,y) = -d_yAx - w_yx$$

for training classes, but

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for test classes.

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Do more experiments, on other datasets (any suggestion ?)

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- Do more experiments, on other datasets (any suggestion?)
- Use a more complex model (Deep Belief Network?)