Interactive Learning

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Advanced Topics in Machine Learning & Optimization - 2023-24

Strategies

Extensions

Conclusion and Further Reading

"Imagine that you are the leader of a colonial expedition from Earth to an extrasolar planet. Luckily, this planet is habitable and has a fair amount of vegetation suitable for feeding your group. Impor- tantly, the most abundant source of food comes from a plant whose fruits are sometimes smooth and round, but sometimes bumpy and irregular."

Figure 1.1: Several alien fruits, which vary in shape from round to irregular.

"The physicians assure you that the shape of a fruit is the only feature that seems related to its safety. The problem, though, is that a wide variety of fruit shapes from these plants exist: almost a continuous range from round to irregular. Since the colony has essential uses for both safe and noxious fruits, you want to be able to classify them as accurately as possible." Credits: (Settles, 2012).

■ We know that *smoother* fruits are (monotonically) *safer*, but we don't know where to set the **threshold**.

We know that smoother fruits are (monotonically) safer, but we don't know where to set the threshold.

In other words, we want to learn a threshold function:

$$f_{ heta}(\mathbf{x}) = egin{cases} 1 & ext{if } x_3 < heta \ -1 & ext{otherwise} \end{cases}$$

where x are measurements of fruit features and x_3 captures its shape "irregularity".

Idea: use regular supervised learning

- Collect a large enough training set $\mathcal{L} = \{(\mathbf{x}, y)\}$, fit threshold classifier f_{θ} on \mathcal{L}
- If maximum % errors is ε ∈ (0, 1), enough to collect ≈ ¹/_ε examples (Shalev-Shwartz and Ben-David, 2014).
 For instance, if max error is 1%, then we need to collect 100 examples. Considering how simple this problem is, this is a lot!

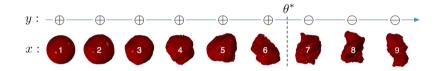


Figure 1.2: Supervised learning for the alien fruits example. Given a set of $\langle x, y \rangle$ instance-label pairs, we want to choose the threshold θ^* that classifies them most accurately.

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We want to find θ as quickly and as economically as possible, by requiring fewer tests.

Can we do better?

Key features:

- Fruits x are plentiful and easy to harvest and measure
- Obtaining y incurs a cost: person that eats the fruit may get sick

So we definitely want to minimize the number of needed labels.

Idea: gather large set of unlabeled fruits $U = \{x_i\}$ and arrange them by roughness.

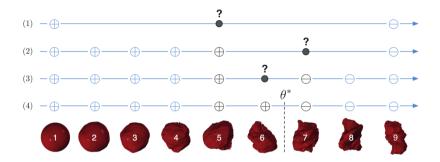


Figure 1.3: A binary search through the set of ordered, untested alien fruits. By only testing this subset of fruits, we can exponentially reduce costs while achieving the same result. The labels shown in light blue can be inferred, and therefore do not need to be tested.

Use binary search to find the threshold θ only takes $\approx \log_2 \frac{1}{\epsilon}$ tests! For $\epsilon = 1\%$, this amounts to ≈ 7 .

Idea: gather large set of unlabeled fruits $U = {x_i}$ and arrange them by roughness, then use binary search:

ϵ	$\frac{1}{\epsilon}$	$\log_2 \frac{1}{\epsilon}$
0.1	10	3.321
0.001	1000	9.966
0.00001	100000	16.610

In this (cleverly designed illustrative) scenario, there is an exponential improvement in sample complexity

"The key hypothesis is that if the learner is allowed to choose the data from which it learns — to be active, curious, or exploratory, if you will — it can perform better with less training." (Settles, 2012)

Preconditions:

- Collecting unlabelled instances x is cheap
- Obtaining their labels y is expensive

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Example: Citizen Science

There are tons of images of celestial bodies (think sky surveys). However, in order to undestand what's in an image (is it a spiral galaxy? is it a gravitational lensing effect?) you have to ask a human expert.

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Example: Recommendation

There are millions of products on online catalogues (think Amazon), but in order to discover what are the tastes of a user, you have to actually convince them to score the items. This information is personalized, so this is the only way to obtain supervision.

Adam, the "robot scientist" (King et al., 2009)

The Automation of Science

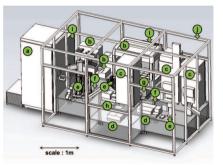
Ross D. King,¹* Jem Rowland,¹ Stephen G. Oliver,² Michael Young,³ Wayne Aubrey,¹ Emma Byrne,¹ Maria Liakata,¹ Magdalena Markham,¹ Pınar Pir,² Larisa N. Soldatova,¹ Andrew Sparkes,¹ Kenneth E. Whelan,¹ Amanda Clare¹

The basis of science is the hypothetico-deductive method and the recording of experiments in sufficient detail to enable reproducibility. We report the development of Robot Scientist "Adam," which advances the automation of both. Adam has autonomously generated functional genomics hypotheses about the yeast *Saccharomyces cerevisiae* and experimentally tested these hypotheses by using laboratory automation. We have confirmed Adam's conclusions through manual experiments. To describe Adam's research, we have developed an ontology and logical language. The resulting formalization involves over 10,000 different research units in a nested treelike structure, 10 levels deep, that relates the 6.6 million biomass measurements to their logical description. This formalization describes how a machine contributed to scientific knowledge.

The learner obtains labels by operating an automated testing machine.

tist Adam. The advances that distinguish Adam from other complex laboratory systems are the individual design of the experiments to test hypotheses and the utilization of complex internal cycles. Adam's basic operations are selection of specified yeast strains from a library held in a freezer. inoculation of these strains into microtiter plate wells containing rich medium, measurement of growth curves on rich medium. harvesting of a defined quantity of cells from each well, inoculation of these cells into wells containing defined media (minimal synthetic dextrose medium plus up to four added metabolites from a choice of six). and measurement of growth curves on the specified media. To achieve this functionality Adam has the following components: a, an automated -20°C freezer b, three liquid handlers (one

Fig. 1. The Robot Scien-



of which can separately control 96 fluid channels simultaneously); c, three strain and defined-growth-medium experiments each day (from a selection of automated +30°C incubators; d. two automated plate readers; e. three robot thousands of yeast strains), with each experiment lasting up to 5 days. The arms; f, two automated plate slides; g, an automated plate centrifuge; h, an design enables measurement of OD_server for each experiment at least once and y, the behavior of the second sec rigid transparent plastic enclosure. There are also two bar code readers, seven curate growth curves to be recorded (typically we take over a hundred meacameras, 20 environment sensors, and four personal computers, as well as the software. Adam is capable of designing and initiating over a thousand new online material for pictures and a video of Adam in action.

Similar strategies used in chemical engineering, material engineering, etc.

Notation

A summary of frequently used terms:

- Instances $\mathbf{x} \in \mathbb{R}^d$ are *unlabelled d*-dimensional vectors of observations
- Examples $z = (\mathbf{x}, y)$ are instances annotated by a label $y \in \{0, 1\}$ or $y \in \{1, \dots, c\}$
- A classifier $f: \mathbb{R}^d \to \{0,1\}$ maps instances to labels (e.g., a neural networks, ...)
- $\mathcal{F} = \{f_{\theta}\}$ is a family of classifiers parameterized by θ (e.g., all neural networks with a specified architecture)

The meaning of θ depends on the model class, e.g., for neural nets with a fixed architecture, θ represents their weights; for random forests, θ represents the structure and leaves of all trees.

Assumptions

We assume the data to be distributed according to a ground-truth distribution $p^*(Y, \mathbf{X})$, which combines a distribution over inputs ("how rare is this document/image?" and a distribution over labels given the input ("how likely is this document to be labeled as funny?")

$$p^*(Y, \mathbf{X}) \equiv p^*(Y \mid \mathbf{X}) \cdot p^*(\mathbf{X}) \tag{1}$$

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(1)

We focus on learning a probabilistic classifier, written as:

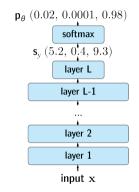
$$p_{\theta}(Y = y \mid \mathbf{X} = \mathbf{x}) \tag{2}$$

(3)

We always predict the most likely label, that is:

$$f_{\theta}(\mathbf{x}) = \operatorname*{argmax}_{y=1,...,c} p_{\theta}(Y = y \mid \mathbf{X} = \mathbf{x})$$

Possible models are anything from **logistic regression** to **neural nets** with a softmax activation (illustrated on the right).



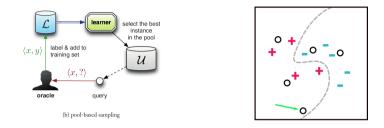
Structure of your average feed-forward neural network. Notice how the output consists of per-class probabilities. Here we write the vector p this using the notation $p_{\theta}(Y \mid \mathbf{x})$. Annotator modelled as an "oracle" that returns the correct label:

$$\operatorname{annot}(\mathbf{x}) := \operatorname{argmax}_{y \in \{0,1\}} p^* (Y = y \mid \mathbf{X} = \mathbf{x})$$
(4)

where p^* is the true (but unobserved) label distribution. In other words, we assume the annotator always answers correctly, i.e., they are knowledgeable and collaborative.

Invoking the oracle comes at a cost, which is unknown, but usually non-negligible, instance- and class-dependent.

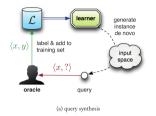
For simplicity, we assume the cost to be unitary: all questions cost the same.

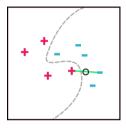


Active Learning (Pool-based). Given:

- a family of classifiers \mathcal{F} ,
- a set of unlabelled instances $U = \{\mathbf{x}_1, \dots, \mathbf{x}_m\} \subseteq \mathbb{R}^d$ sampled i.i.d. from $p^*(\mathbf{X})$,
- a (costly) labeling oracle $label: \mathbb{R}^d \to \{0, 1\},$

Find a classifier $\widehat{f} \in \mathcal{F}$ that achieves low risk on $p^*(\mathbf{X}, Y)$ while keeping annot. cost low

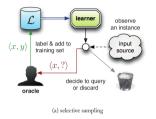


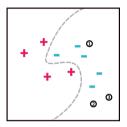


Active Learning (Query Synthesis). Given:

- a family of classifiers \mathcal{F} ,
- a generator of instances $\operatorname{synthesize}(\operatorname{region}) \mapsto x$,
- a (costly) labeling oracle $label: \mathbb{R}^d \to \{0, 1\},$

Find a classifier $\widehat{f} \in \mathcal{F}$ that achieves low risk on $p^*(\mathbf{X}, Y)$ while keeping annot. cost low



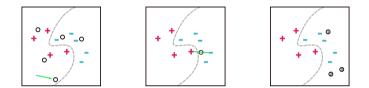


Active Learning (Selective Sampling). Given:

- a family of classifiers \mathcal{F} ,
- a sequence of unlabelled instances $x_1, x_2, x_3, \ldots,$
- a (costly) labeling oracle $label: \mathbb{R}^d \to \{0, 1\}$

Find a classifier $\widehat{f_t} \in \mathcal{F}$ that achieves low risk on future data x_{t+1}, x_{t+2}, \ldots while keeping annot. cost low

Query Sampling vs. Query Synthesis

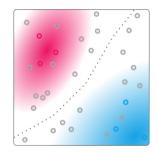


Left to right:

- Pool-based: moderate control over queries, requires memory to store U
- Query synthesis: maximum control over queries, can generate uninterpretable queries (Baum and Lang, 1992), although deep generative models can help somehow (Nguyen et al., 2016).
- Selective sampling: little control over the distribution of queries, often solved under tight memory constraints (online learning)
- We will focus on **pool-based AL**.

Strategies

Quiz Time!



Out of the many unlabeled points (in gray), which ones would you pick for a human annotator to label?

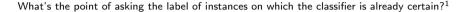
Template

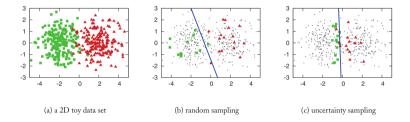
Input: models \mathcal{F} , examples \mathcal{L} , pool \mathcal{U} , query budget $T \geq 1$ **Output:** selected model $f \in \mathcal{F}$ 1: $f \leftarrow \operatorname{fit}(\mathcal{F}, \mathcal{L})$ \triangleright initialize the model 2: for t = 1, 2, ..., T do until the budget is exhausted $\mathbf{x} \leftarrow \operatorname{argmax}_{\mathbf{x} \in \mathcal{U}} \operatorname{acq}(f, \mathbf{x})$ \triangleright select a query instance 3: obtain label v of x from annotator 4. 5: $\mathcal{U} \leftarrow \mathcal{U} \setminus \{\mathbf{x}\}$ ▷ remove unlabeled instance from pool 6: $\mathcal{L} \leftarrow \mathcal{L} \cup \{(\mathbf{x}, \mathbf{v})\}$ ▷ update training set 7: $f \leftarrow \operatorname{fit}(\mathcal{F}, \mathcal{L})$ return f \triangleright update the model

I fit performs training (e.g., trains for a fixed # of epochs)

acq scores instances based on their "informativeness"

What instance $x \in U$ should be selected so to convey as much information as possible to f?





- **Left**: two Gaussians (40 points each)
- **Middle**: picking points completely at random (ignoring the class label!)
- **Right**: picking points based on **uncertainty**

¹There is a point to doing so, as we will see later.

Idea: pick $x \in U$ on which the classifier is most uncertain.

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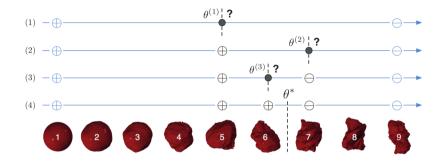


Figure 2.1: The binary search from Figure 1.3, re-interpreted as an uncertainty sampling approach. The best instance to query is deemed to be the one closest to the threshold θ .

Idea: pick $x \in U$ on which the classifier is most uncertain.

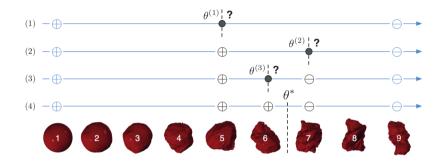


Figure 2.1: The binary search from Figure 1.3, re-interpreted as an uncertainty sampling approach. The best instance to query is deemed to be the one closest to the threshold θ .

How should uncertainty be defined?

Define uncertainty using the **confidence**, i.e., *distance from certainty*:

$$\operatorname{acq}(\theta, \mathbf{x}) := 1 - \rho_{\theta}(\hat{y} \mid \mathbf{x})$$
 (5)

where \hat{y} is the predicted label:

$$\hat{y} := f_{\theta}(\mathbf{x}) = \underset{y}{\operatorname{argmax}} p_{\theta}(y \mid \mathbf{x})$$
(6)

Define uncertainty using the margin, i.e., difference in (conditional) likelihood:

$$\operatorname{acq}(\theta, \mathbf{x}) := p_{\theta}(\hat{y}' \mid \mathbf{x}) - p_{\theta}(\hat{y} \mid \mathbf{x})$$
(7)

where \hat{y} is the predicted label and \hat{y}' is the **2nd best** label:

$$\hat{y} = \underset{v}{\operatorname{argmax}} p_{\theta}(y \mid \mathbf{x})$$
(8)

$$\hat{y}' := \underset{y \neq \hat{y}}{\operatorname{argmax}} p_{\theta}(y \mid \mathbf{x})$$
(9)

Define uncertainty using the Shannon **entropy** of the label:

$$\operatorname{acq}(\theta, \mathbf{x}) := H_{\theta}(Y \mid \mathbf{X} = \mathbf{x})$$
 (10)

where H_{θ} is defined as:

$$H_{\theta}(Y \mid \mathbf{X} = \mathbf{x}) := -\sum_{y \in [c]} p_{\theta}(y \mid \mathbf{x}) \log_2 p_{\theta}(y \mid \mathbf{x})$$
(11)

Remark: conventionally, $0 \times \log_2 0 = 0$.

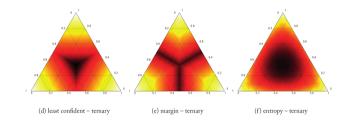
• It achieves a minimum on dead certain distributions:

$$p_{\theta}(Y \mid \mathbf{x}) = (0, 1, 0, \ldots, 0)$$

• and a maximum on the uniform distribution:

$$p_{\theta}(Y \mid \mathbf{x}) = (\frac{1}{c}, \cdots, \frac{1}{c})$$

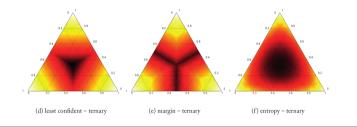
Confidence vs. Margin vs. Entropy



Left: confidence considers prob. of top class only
 Middle: margin considers prob. of top & runner up classes
 Right: entropy considers prob. of all classes

If c = 2, they are equivalent. If c > 2, no obvious best choice, it really depends on the task and loss (e.g., cross-entropy vs.accuracy)

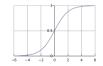
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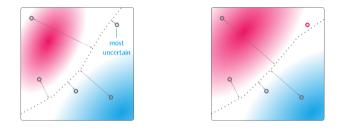
Example: for classifiers with a sigmoid top layer:



uncertainty depends on distance from separating hyperplane of predicted vs. top two vs. all classes

Example: Uncertainty Sampling

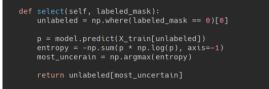
In a binary classification task (red vs. blue), when paired with a sigmoid-based classifier, uncertainty is inversely proportional to the distance from the separator between classes:



Left: gray points indicate unlabelled points, and their distance from the separation surface is indicated by an arrow. Uncertainty sampling picks the closest unlabelled point.

Right: that label of that point happens to be **red**, and the classifier is updated accordingly. Naturally, the distance of all other points from the separator (and hence their uncertainty) changes too.

Uncertainty Sampling



Uncertainty sampling is very easy to implement.

- Margin & Confidence can be defined even in terms of unnormalized scores.
- Usually performs reasonably well (though not optimally) in practice: a useful baseline/starting point.

Example: Structured Output

Consider an LSTM that takes a sequence of MNIST images $X = [x_1, \ldots, x_n]$ that composes a word and outputs the word itself $y = (y_1, \ldots, y_n)$.

- Computing the most likely output $\hat{\mathbf{y}}$ can be done efficiently.
- Computing the entropy amounts to:

$$H_{\theta}(\mathbf{Y} \mid \mathbf{X} = \mathbf{x}) := -\sum_{\mathbf{y} \in \{1, \dots, 26\}^n} p_{\theta}(\mathbf{y} \mid X) \log_2 p_{\theta}(\mathbf{y} \mid X)$$
(12)

This involves summing over 26^n possible outputs, which takes time exponential in *n*.

Computing the most likely output can be NP-hard. For instance, if y is molecular structure that mast satisfy specific hard constraints (chemical validity), then finding the best structure amounts to solving a hard combinatorial problem.

Hence, the **confidence** and **margin** can also be very hard.

When considering regression models with $Y \in \mathbb{R}$, uncertainty at x can be implemented as differential entropy:

$$H_{\theta}(Y \mid \mathbf{X} = \mathbf{x}) := \mathbb{E}[-\log_2 p_{\theta}(y \mid \mathbf{x}) \mid \mathbf{x}]$$
(13)

$$= -\int_{\mathbb{R}} p_{\theta}(y \mid \mathbf{x}) \log_2 p_{\theta}(y \mid \mathbf{x})$$
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As an alternative heuristic, use the variance:

$$\operatorname{Var}_{\theta}(Y \mid \mathbf{x}) := \mathbb{E}[(Y - \underbrace{\mathbb{E}[Y \mid \mathbf{x}]}_{\mu_{\theta}(Y \mid \mathbf{x}):=})^{2} \mid \mathbf{x}]$$
(15)

$$= \int_{\mathbb{R}} (y - \mu_{\theta} (\mathbf{Y} \mid \mathbf{x}))^2 p_{\theta} (y \mid \mathbf{x}) dy$$
(16)

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How to compute them?

Differential entropy and variance:

$$H_{\theta}(Y \mid \mathbf{X} = \mathbf{x}) = -\int_{\mathbb{R}} p_{\theta}(y \mid \mathbf{x}) \log_2 p_{\theta}(y \mid \mathbf{x}) \qquad \operatorname{Var}_{\theta}(Y \mid \mathbf{x}) = \int_{\mathbb{R}} (p_{\theta}(y \mid \mathbf{x}) - \mu_{\theta}(Y \mid \mathbf{x})) dy \qquad (18)$$

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Both expensive to compute for general models, can **approximate** via quadrature or sampling, but **closed-form solutions** exist for some models (e.g., Gaussian Processes and NNs with a Gaussian output)

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Example: 1-dimensional Gaussian Output

Consider one-dimensional output $y \in \mathbb{R}$ and a neural net:

$$nn: \mathbf{x} \mapsto (\mu, \sigma), \qquad \mathbf{y} \sim \mathcal{N}(\mu, \sigma)$$
 (19)

In this case, it is well known² that:

$$\operatorname{Var}_{\theta}(\boldsymbol{Y} \mid \mathbf{x}) = \sigma^{2}, \qquad H_{\theta}(\boldsymbol{Y} \mid \mathbf{x}) = \frac{1}{2}\log(2\pi\sigma^{2}) + \frac{1}{2}$$
(20)

Notice that $\operatorname{Var}_{\theta}(Y \mid x) \propto \exp H_{\theta}(Y \mid x)$, so they change monotonically.

²See https://en.wikipedia.org/wiki/Normal_distribution.

Differential entropy and variance:

$$H_{\theta}(Y \mid \mathbf{X} = \mathbf{x}) = -\int_{\mathbb{R}} p_{\theta}(y \mid \mathbf{x}) \log_2 p_{\theta}(y \mid \mathbf{x}) \qquad \operatorname{Var}_{\theta}(Y \mid \mathbf{x}) = \int_{\mathbb{R}} (p_{\theta}(y \mid \mathbf{x}) - \mu_{\theta}(Y \mid \mathbf{x})) dy \qquad (21)$$

Both expensive to compute for general models, can **approximate** via quadrature or sampling, but **closed-form solutions** exist for some models (e.g., Gaussian Processes and NNs with a Gaussian output)

Example: *k*-dimensional Gaussian Output

Consider one-dimensional output $y \in \mathbb{R}^k$ and a neural net:

$$nn: \mathbf{x} \mapsto (\boldsymbol{\mu}, \boldsymbol{S}), \qquad \boldsymbol{\Sigma} \leftarrow \boldsymbol{S} \boldsymbol{S}^{\mathsf{T}}, \qquad \mathbf{y} \sim \mathcal{N}(\boldsymbol{\mu}, \boldsymbol{\Sigma})$$
 (22)

with Σ PSD by construction. In this case, it is well known³ that:

$$\operatorname{Var}_{\theta}(Y \mid \mathbf{x}) \propto \operatorname{tr} \Sigma \qquad H_{\theta}(Y \mid \mathbf{x}) \propto \log \det \Sigma$$
(23)

where the trace is cheap to compute but the determinant is more challenging.

³See https://en.wikipedia.org/wiki/Multivariate_normal_distribution.

Illustration

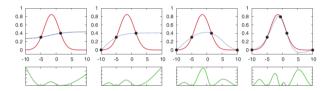
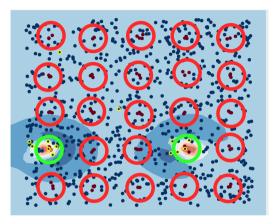
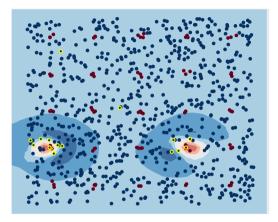


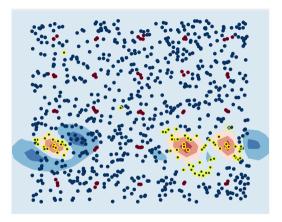
Figure 2.6: Variance-based uncertainty sampling for a toy 1D regression task. Each column represents an iteration of active learning. In the top row, solid lines show the target function to be learned, while dashed lines show a neural network approximation based on available training data (black dots). The bottom row plots the network's output variance across the input range, which is used to select the query for the next iteration.



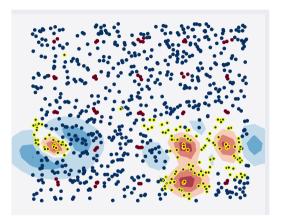
Synthetic dataset: 25 clusters of red points arranged in a 5×5 grid, surrounded by a sea of blue points



After **10** iterations of uncertainty sampling.



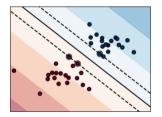
After **70** iterations of uncertainty sampling.



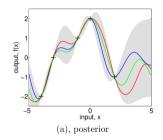
After **140** iterations of uncertainty sampling. Not nice!

Over-confidence

Discriminative models are **over-confident**:



Bayesian generative models not so much:



Uncertainty does $\ensuremath{\textbf{not}}$ decrease with distance from the training set.

Uncertainty **does** decrease with distance from the training set.

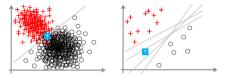


Figure 5: Left: Even with precise knowledge about the optimal hypothesis, the prediction at the query point (indicated by a question mark) is aleatorically uncertain, because the two classes are overlapping in that region. Right: A case of epistemic uncertainty due to a lack of knowledge about the right hypothesis, which is in turn caused by a lack of data.

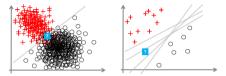


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Aleatoric uncertainty ("random") captures how much we can trust the supervision itself. It cannot be decreased. (left)

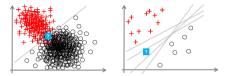


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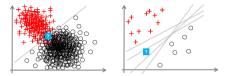


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There isn't much point in trying to reduce aleatoric uncertainty in AL (Sharma and Bilgic, 2017)

Uncertainty Sampling for Streaming Data

Input: models \mathcal{F} , bootstrap training set \mathcal{L} , threshold τ **Output:** selected model $f \in \mathcal{F}$ 1: $f \leftarrow \text{fit}(\mathcal{F}, \mathcal{L})$ ▷ initialize the model 2: for $t = 1, 2, 3, \ldots$, do receive instance x 3: if $unc(f, x) > \tau$ then \triangleright if f is uncertain about x 4: obtain label y of x from annotator 5: 6· $\mathcal{L} \leftarrow \mathcal{L} \cup \{(\mathbf{x}, \mathbf{y})\}$ ▷ update training set $f \leftarrow \operatorname{fit}(\mathcal{F}, \mathcal{L})$ \triangleright update the model 7. return f

The tricky bit is setting τ . Many algorithms update it dynamically by, e.g., starting from a large τ and lowering it as new data is received and the model improves

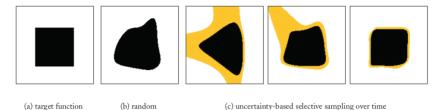


Figure 2.7: Stream-based uncertainty sampling for a simple toy classification task. (a) Positive instances lie inside the black box in 2D. (b) After 100 random samples, the function learned by a neural network is still somewhat amorphous. (c) Uncertainty-based selective sampling at 20, 60, and 100 queries. The highlighted areas represent the region of uncertainty, which gradually shrinks and becomes more focused as the network grows more confident. The output of the resulting network after 100 queries is much more square-like than (b).

For some problems, US **converges** to the right thing – because it is uncertain enough





Figure 2.8: An example of uncertainty sampling failure. (a) Positive instances lie inside the two black triangles. (b) An initial random sample fails to draw many training instances from the negative space in between the triangles. (c) The trained network becomes overly confident that instances in the center are positive. As a result, it avoids that region and begins to learn a different, more square-like shape.

If you are unluckly, US becomes **over-confident**: in this example, the model becomes confident that the regions inside the black blob cannot be white, so it does not sample them and converges to the wrong shape.

The Story So Far

■ In active learning the machine is allowed to ask questions to an oracle – and it should do so intelligently, so as to minimize the # of questions for obtaining a good model.

The most common query selection strategy is **uncertainty sampling**: the machine asks the oracle to label those (unlabelled) instances on which it is most unsure.

Issues:

- If the machine doesn't know that it doesn't know i.e., its self-assessed uncertainty is poorly calibrated – then uncertainty sampling can ignore informative instances altogether.
- Common uncertainty measures (e.g., entropy) mix together epistemic and aleatoric uncertainty, only the former of which we can reduce by acquiring more labels.

```
\begin{array}{l} \mbox{Input: models $\mathcal{F}$, examples $\mathcal{L}$, pool $\mathcal{U}$, query budget $T \geq 1$} \\ \mbox{Output: selected model $f \in $\mathcal{F}$} \\ \mbox{1: } f \leftarrow fit($\mathcal{F}$, $\mathcal{L}$) \\ \mbox{2: for $t=1,2,\ldots,T$ do} \\ \mbox{3: } \mathbf{x} \leftarrow \arg\max_{\mathbf{x} \in \mathcal{U}} \arg(f, \mathbf{x}) \\ \mbox{4: obtain labely of $x$ from annotator} \\ \mbox{5: } \mathcal{U} \leftarrow \mathcal{U} \setminus \{\mathbf{x}\} \\ \mbox{6: } \mathcal{L} \leftarrow \mathcal{U} \cup \{\mathbf{x}, y\} \\ \mbox{6: } \mathcal{L} \leftarrow \mathcal{L} \cup \{\{\mathbf{x}, y\}\} \\ \mbox{7: } \mathbf{return $f$} \quad \mbox{for $t\in \mathcal{I}$}, $\mathcal{L}$ \\ \mbox{return $f$} \quad \mbox{for $t\in \mathcal{I}$}, $\mathcal{L}$ \\ \end{tabular}
```

Uncertainty sampling is quite heuristic. Are there more principled approaches?

Version Space

Consider a hypothesis space $\mathcal{F} = \{f_{\theta} : \mathbf{x} \mapsto y\}$ and a data set $\mathcal{L} = \{(\mathbf{x}_i, y_i)\}$

Consistency

A hypothesis $f \in \mathcal{F}$ is consistent with \mathcal{L} , written $f \models \mathcal{L}$, iff it makes zero mistakes on it, that is:

$$(f \models \mathcal{L}) \iff \left(\sum_{(\mathbf{x}, y) \in \mathcal{L}} \mathbb{1}(f(\mathbf{x}) \neq y)\right) = 0$$
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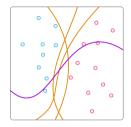
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Version Space

The version space $VS(\mathcal{L})$ of \mathcal{F} given \mathcal{L} is the set of hypotheses $f \in \mathcal{F}$ that are *consistent* with \mathcal{L} , that is:

$$\mathscr{IS}(\mathcal{L}) = \{ f \in \mathcal{F} : f \models \mathcal{L} \}$$

$$(25)$$



 $VS(\mathcal{L})$ contains those classifiers that are not ruled out by the examples \mathcal{L} (in orange). It does not include the purple classifier though!

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Example: \mathcal{L} contains the same instance twice but annotated with different labels – e.g., (x, 1) and (x, 3) – so no $f \in \mathcal{F}$ can classify both correctly.

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The Realizable Case

We assume the realizable case: $\exists f^* \in \mathcal{F}$ s.t. $y = f^*(\mathbf{x})$ for all \mathbf{x} and no noise.

This implies that $f^* \in VS(\mathcal{L})$ for all choices of labeled examples \mathcal{L} , because the supervision (x, y) is always consistent with f^* . Hence, the version space is never empty, regardless of what data we see!

Disagreement Region

Given \mathcal{F} and \mathcal{L} , the disagreement region is the set of points $\mathbf{x} \in \mathbb{R}^d$ such that there exist two classifiers f, f' in the version space $VS(\mathcal{L})$ that produce different predictions for them:

$$DIS(\mathcal{L}) = \{ \mathbf{x} \in \mathbb{R}^d : \exists f, f' \in VS(\mathcal{L}) : f(\mathbf{x}) \neq f'(\mathbf{x}) \}$$

$$(26)$$

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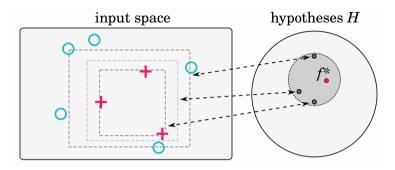
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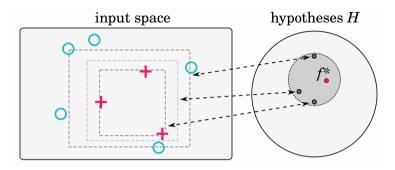
$$(26)$$

If $x \notin DIS(\mathcal{L})$, then all candidate classifiers f in the version space classify it the same: acquiring its label is pointless.

If $x \in DIS(\mathcal{L})$, then at least one f in the version space classifies it differently: acquiring its label is useful.

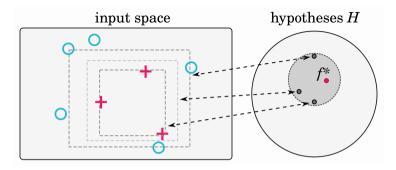


Left: input space \mathbb{R}^d , data set \mathcal{L} of red crosses vs blue circles. Right: hypothesis space \mathcal{F} , each f is a point; the ground-truth f^* is in red.



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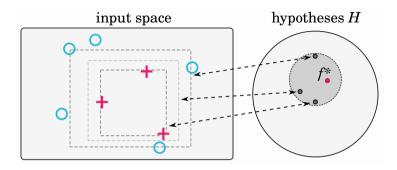
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The version space $VS(\mathcal{L})$ contains all the rectangles (pale gray) between inner & outer rectangles (darker gray) The disagreement region $DIS(\mathcal{L})$ is the space enclosed between these two rectangles.

Input: models \mathcal{F} Output: selected model $f \in \mathcal{F}$ 1: $\mathcal{L} \leftarrow \emptyset$ 2: $\mathcal{V} \leftarrow \mathcal{F}$ 3: for t = 1, 2, 3, ..., do4: receive instance x5: if $\mathbf{x} \in DIS(\mathcal{V})$ then6: obtain label y of x7: update $\mathcal{V} \leftarrow \{f \in \mathcal{V} : f(\mathbf{x}) = y\}$ 8: return any $f \in \mathcal{V}$

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If $x \in DIS(\mathcal{L})$, then there are at least two classifiers $f, f' \in VS(\mathcal{L})$ that disagree on how x should be labeled. Getting its label allows us to get rid of at least one of them, so $VS(\mathcal{L})$ and $DIS(\mathcal{L})$ both shrink.

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- **E** Recall that f^* is always compatible with examples (\mathbf{x}, y) , so it is always in $VS(\mathcal{L}) \rightarrow algorithm zooms$ into it!
- This algorithm makes **no useless queries**!

Question

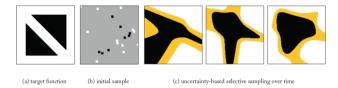


Figure 2.8: An example of uncertainty sampling failure. (a) Positive instances lie inside the two black triangles. (b) An initial random sample fails to draw many training instances from the negative space in between the triangles. (c) The trained network becomes overly confident that instances in the center are positive. As a result, it avoids that region and begins to learn a different, more square-like shape.

Does our streaming VS strategy fix this issue (assuming that the class of possible classifiers \mathcal{F} includes also the target shape)?

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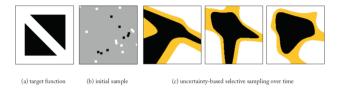


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Does our streaming VS strategy fix this issue (assuming that the class of possible classifiers \mathcal{F} includes also the target shape)? Yes! Incoming points x in the center region belong to the disagreement region (some classifiers in \mathcal{F} might believe they should be black, while f^* knows that they are white), so they are accepted and allow us to retrieve f^* .

Question

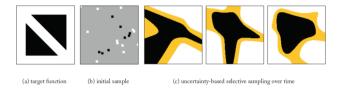


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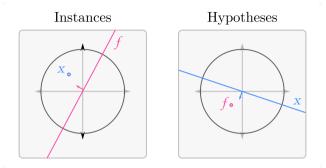
Can we do better if we can choose x?

Version Space for Pool-based AL

Input: models \mathcal{F} Output: selected model $f \in \mathcal{F}$ 1: $\mathcal{L} \leftarrow \emptyset$ 2: $\mathcal{V} \leftarrow \mathcal{F}$ > implements the version space $VS(\mathcal{L})$ 3: for t = 1, 2, ..., T do 4: $\mathbf{x} \leftarrow \operatorname{argmax}_{\mathbf{x} \in \mathcal{U}} \operatorname{acq}_{VS}(\mathcal{V}, \mathcal{F}, \mathbf{x})$ 5: obtain label y of \mathbf{x} 6: update $\mathcal{V} \leftarrow \{f \in \mathcal{V} : f(\mathbf{x}) = y\}$ > update version space 7: return any $f \in \mathcal{V}$

• We can always ensure that there is a point on which the classifiers in the version space disagree, unless the version space is empty or includes a single classifier. In this case we can simply terminate.

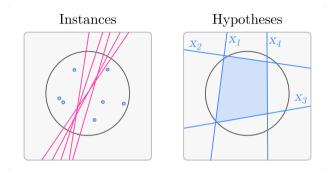
Problem: how do we define the acquisition function?



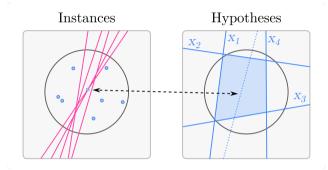
Consider the *linear classifiers*, i.e., \mathcal{F} is:

$$\left\{ f_{\theta}(\mathbf{x}) = \mathbb{1}\left(\boldsymbol{\theta}^{\top}\mathbf{x} > 0\right) : \boldsymbol{\theta} \in \mathbb{R}^{d}, \, \underbrace{\|\boldsymbol{\theta}\|_{2}}_{\text{length}} = 1 \right\}$$
(27)

The version space of $\mathcal L$ is essentially the set of direction vectors m heta that classify all points correctly.



Classifiers are hyperplanes in instance space and instances are hyperplanes in hypothesis space. In some sense, the two spaces are "dual" of one another.



Idea: pick the point $x \in U$ that (greedily) restricts the version space as much as possible. In this special case, x passes close to the **center** of $VS(\mathcal{L})$.

Idea: pick $x \in \mathcal{U}$ that reduces the volume of the version space $VS(\mathcal{L})$ as much as possible.

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$$\operatorname{Vol}\left(A\right)=\int_{A}d\theta$$

Idea: pick $x \in \mathcal{U}$ that reduces the *volume* of the version space $VS(\mathcal{L})$ as much as possible.

The volume of a region $A \subseteq \mathcal{F}$ is:

$$\operatorname{Vol}(A) = \int_{A} d\theta = \int_{\theta \in \mathbb{R}^{|\theta|}} \delta\{\theta \in A\} d\theta$$
(28)

So computing a volume in general requires integration.

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Pick instance x that minimizes the volume of the version space once it is added to the training set. Formally, the volume to be minimize is: $Vol(VS(\mathcal{L} \cup \{(x, y)\}))$.

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The best we can do is to compute the average volume based on the probability of the predicted labels given by the model:

$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmin}} \quad \frac{1}{c} \sum_{y=1}^{c} p_{\theta}(y \mid \mathbf{x}) \cdot \operatorname{Vol}\left(VS(\mathcal{L} \cup \{(\mathbf{x}, y)\})\right)$$
(29)

This tells us what the volume of the VS would be if we were to add x – with an unknown label y – to the data.

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(30)

• If \mathcal{F} is "simple" and/or \mathcal{L} is small, volume can be approximated cheaply using Monte Carlo techniques. For instance with *rejection sampling*, let $B \subseteq VS(\mathcal{L})$ of known volume:

$$\{\widetilde{\boldsymbol{ heta}}_i \sim \operatorname{Uniform}(\boldsymbol{B}) : i = 1, \dots, s\}, \quad \operatorname{Vol}\left(VS(\mathcal{L}')\right) \approx \frac{1}{\operatorname{Vol}\left(\boldsymbol{B}\right)} \cdot \frac{1}{s} \sum_{i=1}^{s} \mathbb{1}\left(\widetilde{\boldsymbol{ heta}}_i \in VS(\mathcal{L}')\right)$$
(31)

To check, $\mathbb{1}(\theta_i \in VS(\mathcal{L}'))$, check that f_{θ} classifies all examples in \mathcal{L} correctly.

If \mathcal{F} is finite, can explicitly store $f \models \mathcal{L}$. Bonus: computing expected volume is doable (integral becomes sum).

If \mathcal{F} is infinite, cannot store explicitly. However, we only need to compute its volume:

$$\frac{1}{c} \sum_{y \in [c]} p_{\theta}(y \mid \mathbf{x}) \cdot \underbrace{\operatorname{Vol}\left(VS(\mathcal{L} \cup \{(\mathbf{x}, y)\})\right)}_{\text{this is the difficult bit}}$$
(30)

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To check, $\mathbb{1}(\theta_i \in VS(\mathcal{L}'))$, check that f_{θ} classifies all examples in \mathcal{L} correctly.

Otherwise (think CNN on ImageNet), can be extremely challenging – we cannot use VS!

Not All Classifiers in VS Think Exactly The Same



Figure 3.5: Examples of committee and consensus distributions. $P_{\theta^{(j)}}$ refers the output distribution of the *i*th hypothesis, and P_C represents the consensus across all committee members.

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- These approaches make two **assumptions**:
 - Disagreement is measured using *all* hypotheses in the version space $VS(\mathcal{L})$.
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- Let's relax both of them \rightarrow speed-up!

Moreover, version space is only non-empty in the realizable case. How do we deal with this?

■ Idea: replace VS witha committee C:

- Select k representatives $C = \{c_1, \ldots, c_k\}$ from $VS(\mathcal{L})$, with k > 100.
- Then (efficiently) aggregate disagreement between them: no volume/integral is needed!

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In all cases, we end up having a set of classifiers that fit the data – assuming their accuracy is 100% – so they are all in the VS.

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In all cases, we end up having a set of classifiers that fit the data – assuming their accuracy is 100% – so they are all in the VS. In practice, less than perfect accuracy is allowed: members are "almost" in VS.

Measuring Disagreement of $\mathcal C$ on $\mathbf x\in \mathcal U$

"Hard" Voting + Entropy:

$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmax}} - \sum_{y} \frac{n(y,\mathbf{x})}{k} \log \frac{n(y,\mathbf{x})}{k}, \qquad n(y,\mathbf{x}) := \sum_{c\in\mathcal{C}} \mathbb{I}(c(\mathbf{x}) = y)$$
(32)

Each classifier votes either 0 or 1.

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Output probabilities of each $c \in C$ taken into account.

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Output probabilities of each $c \in C$ taken into account.

Average Kullback-Liebler divergence:

$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmax}} \quad \frac{1}{k} \sum_{c\in\mathcal{C}} \mathbb{KL}(p_c(Y \mid \mathbf{x}) \| p_{\mathcal{C}}(Y \mid \mathbf{x}))$$
(34)

$$\mathbb{KL}(p(Y \mid \mathbf{x}) \| q(Y \mid \mathbf{x})) := \sum_{y} p(y \mid \mathbf{x}) \log \frac{p(y \mid \mathbf{x})}{q(y \mid \mathbf{x})}$$
(35)

Very expressive, measures difference between whole distributions, i.e., prob. of all possible labels.

Idea: pick the point that gives the maximal improvement in model quality

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Useful Concepts

The loss of p_{θ} on example z = (x, y) is denoted $\ell(\theta, z)$. For instance, cross-entropy loss:

$$\ell(\theta, z) := -\sum_{j} \mathbb{1}(j = y) \log p_{\theta}(j \mid \mathbf{x}) = -\log p_{\theta}(y \mid \mathbf{x})$$
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The true risk \mathcal{L}^* of θ w.r.t. the ground-truth distribution $p^*(\mathbf{X}, Y)$ is:

$$\mathcal{L}^*(\theta) := \mathbb{E}_{z \sim p^*} \left[\ell(\theta, z) \right] \tag{37}$$

It measures the true quality of the model, unobserved.

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The empirical risk $\widehat{\mathcal{L}}_S$ of θ w.r.t. data set $S = \{z_1, \ldots, z_m\}$ sampled i.i.d. from p^* is:

$$\widehat{\mathcal{L}}_{\mathcal{S}}(\theta) := \frac{1}{|\mathcal{S}|} \sum_{z \in \mathcal{S}} \ell(\theta, z)$$
(38)

It estimates the quality of the model from a sample S, optimized during training.

Let $\widehat{\theta}$ be the parameters obtained by training on S and $\widehat{\theta}^{+z}$ those obtained by training on $S \cup \{z\}$, i.e.,

$$\widehat{\theta} := \underset{\theta}{\operatorname{argmin}} \ \widehat{\mathcal{L}}_{S}(\theta) \qquad \widehat{\theta}^{+z} := \underset{\theta}{\operatorname{argmin}} \ \widehat{\mathcal{L}}_{S \cup \{z\}}(\theta) \tag{39}$$

where optimization is possibly approximate, e.g., based on SGD.

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where optimization is possibly approximate, e.g., based on SGD.

Model Improvement

The model improvement (MI) given by a new example $z \notin S$ is the decrease in true risk:

$$\operatorname{acq}(\mathbf{x}) := \mathcal{L}^*(\widehat{\theta}) - \mathcal{L}^*(\widehat{\theta}^{+\mathbf{z}})$$
(40)

The higher, the better \longrightarrow pick the $x \in \mathcal{U}$ that maximizes the improvement.

Model Improvement as Greedy Optimization

MI amounts to solving:

$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmax}} \mathcal{L}^{*}(\widehat{\theta}) - \mathcal{L}^{*}(\widehat{\theta}^{+\mathbf{z}}) = \underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmin}} \mathcal{L}^{*}(\widehat{\theta}^{+\mathbf{z}})$$
(41)

It is guaranteed to find the best next candidate!

⁴Note: MI is *greedy*, *not* optimal! Non-greedy alternatives are conceptually better, but they also computationally infeasible and for this reason they are ignored in the AL literature.

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MI is essentially a greedy strategy for solving:⁴

$$\underset{S \subseteq \mathcal{U}}{\operatorname{argmin}} \quad \mathcal{L}^*(\widehat{\theta}) \tag{42}$$

s.t.
$$|S| \leq query budget$$
 (43)

In this view, AL is a subset optimization problem, and MI solves it directly.

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In this view, AL is a subset optimization problem, and MI solves it directly.

Compare this to uncertainty sampling, which is not as sound

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$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmin}} \ \mathcal{L}^*(\widehat{\theta}^{+z}) \tag{44}$$

 $^{^5 {\}rm The}$ unlabeled set ${\cal U}$ is ideally pretty large, so the approximation is reasonable.

We want to solve:

$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmin}} \ \mathcal{L}^*(\widehat{\theta}^{+z}) \tag{44}$$

Problem: $\mathcal{L}^*(\cdot)$ is an **integral over** $\mathbf{x}' \in \mathbb{R}^d$:

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) = \mathbb{E}_{z' \sim \rho^{*}}[\ell(\widehat{\theta}^{+z}, z')] = \int_{\mathbb{R}^{d}} \ell(\widehat{\theta}^{+z}, (\mathbf{x}', y')) d\mathbf{x}'$$
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which is intractable \rightarrow approximate using empirical average over $\mathcal{U}{:}^5$

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \approx \widehat{\mathcal{L}}_{\mathcal{U}}(\widehat{\theta}^{+z}) = \frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}' \in \mathcal{U}} \ell(\widehat{\theta}^{+z}, (\mathbf{x}', \mathbf{y}'))$$
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(46)

Example: if ℓ is the 0–1 loss, then this amounts to:

$$\frac{1}{\mathcal{U}|}\sum_{\mathbf{x}'\in\mathcal{U}}\mathbb{1}\left(f_{\hat{\theta}+z}(\mathbf{x}')\neq \mathbf{y}'\right)$$
(47)

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Problem: we don't have access to the ground-truth label $z = (x, y) \rightarrow$ marginalize w.r.t. ρ^* :

$$\mathbb{E}_{\boldsymbol{y}\sim\boldsymbol{\rho}^{*}(\boldsymbol{Y}|\boldsymbol{x})}\left[\frac{1}{|\mathcal{U}|}\sum_{\boldsymbol{x}'\in\mathcal{U}}\ell(\widehat{\theta}^{+(\boldsymbol{x},\boldsymbol{y})},(\boldsymbol{x}',\boldsymbol{y}'))\right]$$
(49)

This averages over alternative future models $\widehat{\theta}^{+(x,y)}$ obtained after retraining on $\mathcal{L} \cup (x,y)$.

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(50)

This averages over the **unknown labels** y' of the instances in $x' \in U$.

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \longrightarrow \mathbb{E}_{y \sim p^{*}(Y|\mathbf{x})} \left[\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}' \in \mathcal{U}} \mathbb{E}_{y' \sim p^{*}(Y|\mathbf{x}')} \left[\ell(\widehat{\theta}^{+}(\mathbf{x}, y), (\mathbf{x}', y')) \right] \right]$$
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Problem: we don't have access to p^* at all \rightarrow estimate using model's distribution:

$$\underbrace{\mathbb{E}_{y \sim \boldsymbol{p}_{\widehat{\theta}}(\boldsymbol{Y}|\mathbf{x})}\left[\underbrace{\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}' \in \mathcal{U}} \underbrace{\mathbb{E}_{y' \sim \boldsymbol{p}_{\widehat{\theta}^+}(\boldsymbol{Y}|\mathbf{x}')}\left[\ell(\widehat{\theta}^+, (\mathbf{x}', y'))\right]}_{(a)}}_{(b)}\right]}_{(c)}$$
(52)

where $\widehat{\theta}^+ := \widehat{\theta}^{+(\mathbf{x},y)}$. If p_{θ} is "good enough", then the approximation is good.

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \longrightarrow \mathbb{E}_{y \sim \rho^{*}(Y|x)} \left[\frac{1}{|\mathcal{U}|} \sum_{x' \in \mathcal{U}} \mathbb{E}_{y' \sim \rho^{*}(Y|x')} \left[\ell(\widehat{\theta}^{+}(x,y), (x', y')) \right] \right]$$
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where $\widehat{\theta}^+ := \widehat{\theta}^{+(\mathbf{x},y)}$. If p_{θ} is "good enough", then the approximation is good.

(a) Is the expected loss of the updated model on $x' \in \mathcal{U},$

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \longrightarrow \mathbb{E}_{y \sim \rho^{*}(Y|\mathbf{x})} \left[\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}' \in \mathcal{U}} \mathbb{E}_{y' \sim \rho^{*}(Y|\mathbf{x}')} \left[\ell(\widehat{\theta}^{+}(\mathbf{x}, y), (\mathbf{x}', y')) \right] \right]$$
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- (a) Is the expected loss of the updated model on $x' \in \mathcal{U},$
- (b) Is the **average** expected oss of the updated model on *all* of \mathcal{U} ,

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \longrightarrow \mathbb{E}_{y \sim \rho^{*}(Y|x)} \left[\frac{1}{|\mathcal{U}|} \sum_{x' \in \mathcal{U}} \mathbb{E}_{y' \sim \rho^{*}(Y|x')} \left[\ell(\widehat{\theta}^{+}(x,y), (x', y')) \right] \right]$$
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where $\widehat{\theta}^+ := \widehat{\theta}^{+(\mathbf{x},y)}$. If p_{θ} is "good enough", then the approximation is good.

- (a) Is the expected loss of the updated model on $x' \in \mathcal{U},$
- (b) Is the average expected oss of the updated model on all of \mathcal{U} ,
- (c) Is the above averaged over the possible updated models $\widehat{\theta}^{+(x,y)}.$

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \longrightarrow \mathbb{E}_{y \sim p_{\widehat{\theta}}(Y|\mathbf{x})} \Big[\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}' \in \mathcal{U}} \mathbb{E}_{y' \sim p_{\widehat{\theta}^{+}}(Y|\mathbf{x}')} \Big[\ell(\widehat{\theta}^{+}, (\mathbf{x}', y')) \Big] \Big]$$
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We pick $x \in \mathcal{U}$ that minimizes the above \rightarrow minimizes expected future confidence on \mathcal{U}

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \longrightarrow \mathbb{E}_{y \sim \rho_{\widehat{\theta}}(Y|\mathbf{x})} \Big[\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}' \in \mathcal{U}} \mathbb{E}_{y' \sim \rho_{\widehat{\theta}^{+}}(Y|\mathbf{x}')} \Big[\ell(\widehat{\theta}^{+}, (\mathbf{x}', y')) \Big] \Big]$$
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(60)

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We pick $x \in \mathcal{U}$ that minimizes the above \rightarrow minimizes expected future entropy on \mathcal{U}

In uncertainty sampling, we pick x that minimizes model's estimate of current uncertainty w.r.t. itself, this is myopic

■ In expected model improvement, we pick x that minimizes model's estimate of expected future uncertainty w.r.t. unlabeled set, this is less myopic

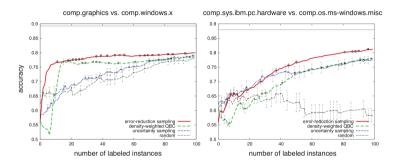


Figure 4.1: Learning curves showing that expected error reduction can outperform QBC and uncertainty sampling for two binary text classification tasks. *Source*: Adapted from Roy and McCallum (2001), with kind permission of the authors.

$$\mathcal{L}^{*}(\widehat{\theta}^{+z}) \longrightarrow \mathbb{E}_{y \sim p_{\widehat{\theta}}(Y|\mathbf{x})} \Big[\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}' \in \mathcal{U}} \mathbb{E}_{y' \sim p_{\widehat{\theta}^{+}}(Y|\mathbf{x}')} \Big[\ell(\widehat{\theta}^{+}, (\mathbf{x}', y')) \Big] \Big]$$
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Problem: this has to be done $|\mathcal{U}| \times [c]$ times.

Problem: this has to be done in each iteration of active learning.

Only practical for classes of models that support closed-form updates (e.g., Gaussian Processes) or stable incremental learning (e.g., perceptron-like learning algorithms).

Unless a candidate (x, y) induces a large change in the model $\hat{\theta}$ upon retraining, then it cannot reduce the model's risk by much: change is a prerequisite for improvement.

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Inituition:

$$\ell(\widehat{\theta}, z') - \ell(\widehat{\theta}^{+z}, z') \leq |\ell(\widehat{\theta}, z') - \ell(\widehat{\theta}^{+z}, z')| \leq c \cdot \|\widehat{\theta} - \widehat{\theta}^{+z}\|, \quad c > 0$$
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where $\|\cdot\|$ is, e.g., the Euclidean norm. This formally holds for all *c*-Lipshitz loss functions ℓ .

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Large change also occurs when the loss *increases* – hence the absolute value in the second step of Eq. 62. All in all, EMC looks for examples $x \in U$ that "make a difference" one way or the other.

But once (x, y) is acquired it is added to the training set \mathcal{L} on which $\hat{\theta}$ is fit, so loss is likely to go *down* rather than *up*.

The trick is that if $\hat{\theta}$ is obtained via gradient descent, the difference $\hat{\theta} - \hat{\theta}^{+z}$ is easy to compute:

$$\widehat{\theta} - \widehat{\theta}^{+z} = \eta \cdot \nabla_{\theta} \ell(\theta, z) \tag{63}$$

where η is the learning rate. This gives expected gradient length:

$$\operatorname{acq}_{\mathsf{EGL}}(\mathbf{x}) := \mathbb{E}_{\mathbf{y} \sim p_{\theta}(\mathbf{y}|\mathbf{x})} \left[\| \nabla_{\theta} \ell(\widehat{\theta}, (\mathbf{x}, \mathbf{y})) \|^{2} \right]$$
(64)

The square does not change ranking of examples & avoids computing a square root.

- Quite cheap to compute using automatic differentiation packages (using Jacobian to parallelize over \mathcal{U})
- Assuming η is constant across examples and GD, the computation is exact. For other optimizers, it is an approximation

Are Uncertain Points Representative?



Figure 5.1: An illustration of when uncertainty sampling can be a poor strategy. Shaded polygons represent labeled instances in \mathcal{L} , and circles represent unlabeled instances in \mathcal{U} . Since A is on the decision boundary, it would be queried as the most uncertain. However, B would probably provide more information about the input distribution as a whole.

Idea: pick instances $x \in U$ that are *both* locally informative and also similar to as many other unlabeled points as possible:

$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmax}} \operatorname{acq}(f,\mathbf{x}) \cdot \left(\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}'\in\mathcal{U}} \sin(\mathbf{x},\mathbf{x}')\right)^{\beta}$$
(65)

where:

- acq(f, x) is a "standard" acquisition function based on, e.g., pointwise uncertainty.
- sim(x, x') measures the similarity between x and x', e.g., a Gaussian kernel, Pearson's correlation coefficient, Spearman's rank correlation. Application specific.
- $\beta > 0$ is a hyper-parameter

Intuitively, x's label conveys information about the label on the other points in $\mathcal U$

Example

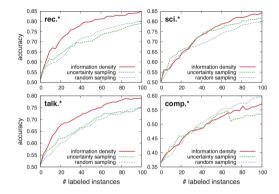


Figure 5.2: Learning curves showing that, by explicitly weighting queries by their representativeness among the input instances, information density can yield better results than the base uncertainty sampling heuristic by itself.

We optimize:

$$\underset{\mathbf{x}\in\mathcal{U}}{\operatorname{argmax}} \operatorname{acq}(f,\mathbf{x}) \cdot \left(\frac{1}{|\mathcal{U}|} \sum_{\mathbf{x}'\in\mathcal{U}} \operatorname{sim}(\mathbf{x},\mathbf{x}')\right)^{\beta}$$
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Properties:

- Tends to work better than pure more "local" acquisition functions (Settles, 2012)
- · Even when uncertainty sampling is worse than random, information density performs well

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Properties:

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- Similarity computation can be sped-up using caching: "simply" store similarity matrix $S_{ij} = [sim(x_i, x_j)]$ for all $x_i, x_i \in U$ (only needs to be done once)
- Approximate using clustering: cluster U so that points within cluster are similar and points across clusters are not → block-diagonal similarity matrix, lowers storage requirement from O(|U|²) to O(∑_i |cluster_i|²)

Do we gain anything by "summarizing" the data using **clustering**?

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Idea:

- Cluster unlabeled data set $\mathcal{U} \to \{C_i \subset \mathcal{U} : i \in [k]\}$
- Treat each C_i as a separate problem, e.g., query cluster centroids

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Problems:

- U may not have a good clustering structure or sim(·, ·) may not be able to capture it
- How many clusters and at what granularity?
- Clusters of x's may not correlate well with label y.

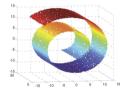


Figure: the swiss roll dataset has no obvious clustering structure.

Extensions

Consider a neural network $f_{\theta} : \mathbb{R}^d \to [c]$:

$$egin{array}{ll} f_{ heta}(\mathbf{x}) = rgmax_{y\in [c]} & p_{ heta}(y\mid \mathbf{x}) \ & y \in [c] & p_{ heta}(y\mid \mathbf{x}) = \operatorname{softmax}(W\phi_{\omega}(\mathbf{x}))_y \end{array}$$

where:

- $\theta = \{W, \omega\}$ are parameters
- $\phi_\omega: \mathbb{R}^d o \mathbb{R}^k$ is an embedding function (e.g., convolutions + pooling layers)
- $W \in \mathbb{R}^{c imes k}$ are the parameters of the top dense layer

Deep Architectures

 Very overconfident even away from the training set: their uncertainty cannot be trusted → strategies based on confidence, margin, entropy will underperform (including uncertainty sampling, model improvement, density-aware sampling, etc.)

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 ensure responsivity
- Quite insensitive to the addition of a single example \rightarrow what's the point of querying individual instances?
- Training is stochastic (i.e., not 100% stable) \rightarrow changes in performance can depend on factors other than new labeled examples, high variance

Overconfidence

Problem: Deep NNs tend to be very **overconfident** even away from the training set \rightarrow their *uncertainty* cannot be trusted

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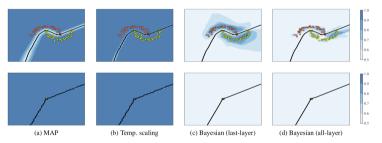


Figure 1. Binary classification on a toy dataset using a MAP estimate, temperature scaling, and both last-layer and all-layer Gaussian approximations over the weights which are obtained via Laplace approximations. Background color and black line represent confidence and decision boundary, respectively. Bottom row shows a zoomed-out view of the top row. The Bayesian approximations—even in the last-layer case—give desirable uncertainty estimates: confident close to the training data and uncertain otherwise. MAP and temperature scaling yield overconfident predictions. The optimal temperature is picked as in Guo et al. (2017).

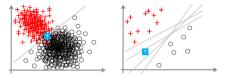


Figure 5: Left: Even with precise knowledge about the optimal hypothesis, the prediction at the query point (indicated by a question mark) is aleatorically uncertain, because the two classes are overlapping in that region. Right: A case of epistemic uncertainty due to a lack of knowledge about the right hypothesis, which is in turn caused by a lack of data.

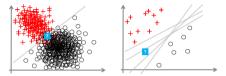


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Aleatoric uncertainty ("random") captures how much we can trust the supervision itself. It cannot be decreased. (left)

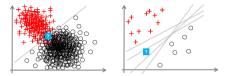


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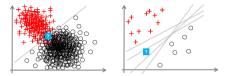


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There isn't much point in trying to reduce aleatoric uncertainty in AL (Sharma and Bilgic, 2017)

Bayesian NNs

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Idea of Bayesian NNs:

- Replace parameters θ with distribution over alternative parameters $p(\theta \mid \mathcal{L})$
- Compute predictions by marginalizing over θ :

$$p(y \mid \mathbf{x}) = \int \underbrace{p(y \mid \mathbf{x}, \theta)}_{\text{NN with params } \theta} \cdot \underbrace{p(\theta \mid \mathcal{L})}_{\text{posterior over params}} d\theta$$
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- Using Bayesian techniques i.e., maintaining a distribution over alternative NNs is also challenging.

Idea of Bayesian NNs:

- Replace parameters θ with distribution over alternative parameters $p(\theta \mid \mathcal{L})$
- Compute predictions by marginalizing over θ :

$$p(y \mid \mathbf{x}) = \int \underbrace{p(y \mid \mathbf{x}, \theta)}_{\text{NN with params } \theta} \cdot \underbrace{p(\theta \mid \mathcal{L})}_{\text{posterior over params}} d\theta$$
(67)

• Learn by updating distribution:

$$p(\theta \mid \mathcal{L}) \rightarrow p(\theta \mid \mathcal{L} \cup \{(\mathbf{x}, y)\})$$
 (68)

Not trivial! Is there an efficient approximation?

Dropout

Randomly set nodes to 0 with a fixed probability.

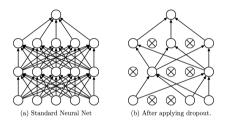


Figure 1: Dropout Neural Net Model. Left: A standard neural net with 2 hidden layers. Right: An example of a thinned net produced by applying dropout to the network on the left. Crossed units have been dropped.

Used as a regularization technique: by randomly removing neurons, prevents them from relying on each other "too much"

Computing class probabilities:

$$p(y \mid \mathbf{x}, \mathcal{L}) = \int p(y \mid \mathbf{x}, \theta) p(\theta \mid \mathcal{L}) d\theta$$
(69)

$$\approx \int p(y \mid \mathbf{x}, \theta) p_{\text{dropout}}(\theta) d\theta$$
(70)

$$\approx \frac{1}{R} \sum_{r=1}^{R} p(y \mid \mathbf{x}, \widehat{\boldsymbol{\theta}}_r), \qquad \widehat{\boldsymbol{\theta}}_r \sim p_{\text{dropout}}(\boldsymbol{\theta})$$
(71)

In other words, run NN R times with dropout enabled (during inference!) then average the R vectors of class probabilities.

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Immediately leads to more calibrated output probabilities!

Question: does dropout help with query selection too?

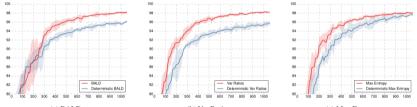
Question: does dropout help with query selection too? Yes.

Uncertainty sampling:

$$\operatorname{acq}_{UNC}(\mathbf{x}) = -\sum_{y \in [c]} p(Y = y \mid \mathbf{x}, \mathcal{L}) \log p(Y = y \mid \mathbf{x}, \mathcal{L})$$
(72)

Simply run the NN multiple times on your input x with different (random) dropout masks, then average the resulting probabilities.

Illustration



(a) BALD (b) Var Ratios (c) Max Entropy Figure 2. Test accuracy as a function of number of acquired images for various acquisition functions, using both a **Bayesian CNN (red)** and a deterministic CNN (blue).

For all choices of acquisition function, the dropout-based uncertainty helps!

Let us look at **batch-based** active learning.

Batch Selection

Given \mathcal{L} , \mathcal{U} and a classifier $f \in \mathcal{F}$ trained on \mathcal{L} , find a batch $B \subseteq \mathcal{U}$ of $b \gg 1$ unlabeled instances that brings maximal information to the model:

$$\operatorname{argmax}_{B \subseteq \mathcal{U}} \operatorname{acq}_{BALD}(f, B)$$
(73)
s.t. $|B| = b$ (74)

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$$\underset{B \subseteq \mathcal{U}}{\operatorname{argmax}} \operatorname{acq}_{BALD}(f, B) \tag{73}$$

s.t.
$$|B| = b$$
 (74)

Advantages:

- Only retrain the model after ever b examples, meaning that supervision has an effect.
- Retraining is less frequent, leading to faster overall execution (at the expense of possibly instance selection, because *b* examples depend on a fixed *f*).
- Supports parallel annotation for, e.g., crowd-sourcing scenarios.

Question: can regular acquisition function (like BALD) be extended to this setting?

Natural generalization of instance-level strategies:

$$\operatorname{acq}(f, B) = \sum_{\mathbf{x} \in B} \operatorname{acq}(f, \mathbf{x})$$
 (75)

How well does this work?

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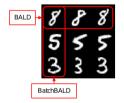
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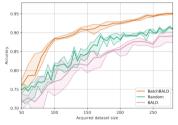
How well does this work?

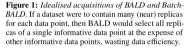
This ignores *correlation* between instances in x:

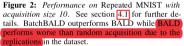
- Even if all of them are informative, they may carry the same information
- We want *B* to be informative as a whole!

Illustration









BatchBALD

The problem with the "natural generalization":

$$\operatorname{acq}(f, B) = \sum_{\mathbf{x} \in B} \operatorname{acq}(f, \mathbf{x})$$
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is that the sum doesn't consider the *overlap* between the information carried by different $x \in b$.

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Idea: don't break the acquisition function into a sum! For BALD, this means replacing:

$$\sum_{\mathbf{x}\in B} \left\{ \underbrace{\mathcal{H}(\mathbf{Y}\mid \mathbf{x}, \mathcal{L}) - \mathbb{E}_{\boldsymbol{\theta}\sim \boldsymbol{\rho}(\boldsymbol{\theta}\mid \mathcal{L})}[\mathcal{H}(\mathbf{Y}\mid \mathbf{x}, \boldsymbol{\theta})]}_{\mathcal{M}(\mathbf{Y}, \boldsymbol{\Theta}\mid \mathbf{x}, \mathcal{L})} \right\}$$
(77)

with

$$MI({Y_1,\ldots,Y_b},\Theta \mid {x_1,\ldots,x_b},\mathcal{L})$$
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with

$$MI(\{Y_1,\ldots,Y_b\},\Theta \mid \{x_1,\ldots,x_b\},\mathcal{L})$$
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In other words, don't assume independence between the elements of *B*!

Illustration

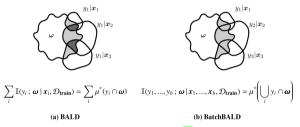


Figure 3: Intuition behind BALD and BatchBALD using I-diagrams [30]. BALD overestimates the joint mutual information. BatchBALD, however, takes the overlap between variables into account and will strive to acquire a better cover of ω . Areas contributing to the respective score are shown in grey, and areas that are double-counted in dark grey.

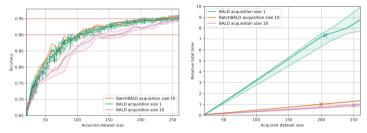


Figure 5: *Performance on* MNIST. BatchBALD outperforms BALD with acquisition size 10 and performs close to the optimum of acquisition size 1.

Figure 6: Relative total time on MNIST. Normalized to training BatchBALD with acquisition size 10 to 95% accuracy. The stars mark when 95% accuracy is reached for each method.

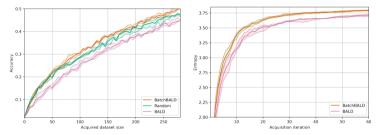


Figure 7: Performance on EMNIST. BatchBALD consistently outperforms both random acquisition and BALD while BALD is unable to beat random acquisi- a more diverse set of data points. tion.

Figure 8: Entropy of acquired class labels over acquisition steps on EMNIST. BatchBALD steadily acquires

Conclusion and Further Reading

\blacksquare AL useful when supervision is expensive high \rightarrow choose it wisely

Many variants: pool-based, streaming, and query synthesis

Many practical approaches: uncertainty-based (uncertainty sampling, QBC, expected gradient length), diversity-based (information density).

Some can be derived from version spaces and model improvement.

Deep variants select entire **batches** and often rely on Bayesian techniques

Critique & realistic annotators, costs, etc.: (Herde et al., 2021) (Settles, 2011)

Plenty of room for new research ;-)

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