

Tell Me What To Do: Prioritizing Data Labeling for NLP Systems with Active Learning

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Outline

- Background
- NLP systems for medical applications
- Active Learning
- Cancer pathology report classification
- Final thoughts

Little bit about myself

- Research Scientist within the Machine Learning group at LLNL since 2017
- Machine Learning interests:
 - Transfer and multitask learning
 - Probabilistic deep learning models
 - Uncertainty quantification in large ML models
 - Optimization for ML
- Recent projects:
 - ML for healthcare: cancer prognostics with MTL and synthetic EHR data generation
 - Deep Learning for climate (sub-)seasonal forecasting
 - Deep Reinforcement Learning for antibody design
 - ML for microbiome profile characterization



Background



Background

- Project developed with the National Cancer Institute (NCI) and other national laboratories:



- Development of natural language processing and deep learning algorithms to population-based cancer statistics collected by NCI's SEER program.
- Automate pathology reports classification and annotation process.

NLP Systems for Medical Applications



NLP systems for medical applications

- Many of the medical procedures, exams, and treatments are stored in written format.
- Notes contain the state and evolution of the patient throughout the treatment/medical procedure.
- High volumes of unstructured reports go to electronic health records (EHR) systems daily.
- 80% of healthcare-associated text data is unstructured and goes largely unutilized.¹



¹ <https://www.foreseemed.com/natural-language-processing-in-healthcare>

NLP systems for medical applications

- Physicians don't all "speak the same way" and there is not always a diagnosis consensus.
- This pile of data contains information that can dramatically improve current understandings of treatments and medical protocols.



NLP systems in Cancer Surveillance

- In the US, NCI's *Surveillance, Epidemiology, and End Results* (SEER) program is responsible for collecting, processing, and aggregating cancer data through local **Cancer Registries**.
- Surveillance data serve as a foundation for cancer research and are used to **plan and evaluate cancer prevention and control interventions**.

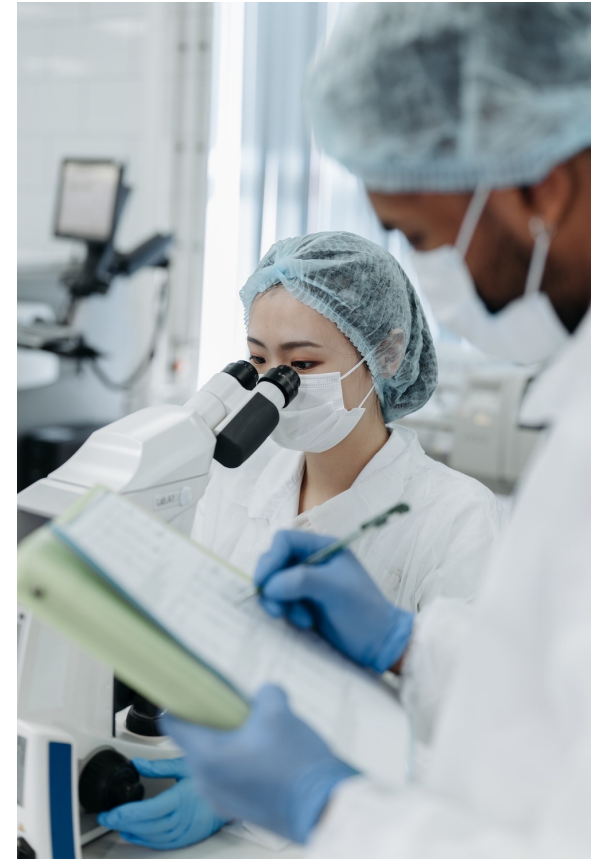


Source: [1]

[1] https://seer.cancer.gov/registries/cancer_registry/index.html

NLP systems in Cancer Surveillance

- Critical cancer data elements such as primary tumor site and tumor morphology reside in unstructured **cancer pathology reports** that are written during the time of diagnosis.
- Such reports are not only ungrammatical, fragmented, and marred with typos and abbreviations, but also exhibit linguistic variability across pathologists.



NLP systems in Cancer Surveillance

- Pathology reports:
 - “A pathology report is a document that contains the diagnosis determined by examining cells and tissues under a microscope. The report may also contain information about the size, shape, and appearance of a specimen as it looks to the naked eye.” ¹
- Pathologist is a doctor responsible for examining and writing a pathology report.
- Pathology reports are essential for cancer diagnosis and staging, therefore helping to determine treatment options.



¹ <https://www.cancer.gov/about-cancer/diagnosis-staging/diagnosis/pathology-reports-fact-sheet>

NLP systems in Cancer Surveillance

- Show pathology report examples:
 - [Example 1](#)
 - [Example 2](#)

NLP systems in Cancer Surveillance

- Cancer information is **manually** extracted from the pathology reports by **Certified Tumor Registrars (CTRs)**.
- Cancer registries face challenges scaling the manual effort to handle the increasing volumes of clinical reports they need to process and the amount of information they need to capture per report.
- NCI's effort to **automate** some of the involved processes **using NLP systems**.

Challenges in Pathology Report Classification

- Reports lack of universal structure, variation in linguistic patterns, and the presence of medical jargons.
- Reports are long documents (several pages) but only a few keywords may be relevant to a specific classification task.
- Long-distance linguistic dependencies across different sections of a document.
- Certain tasks may contain several classes (i.e., 525 histology types), and the number of documents per class tends to be highly unbalanced.

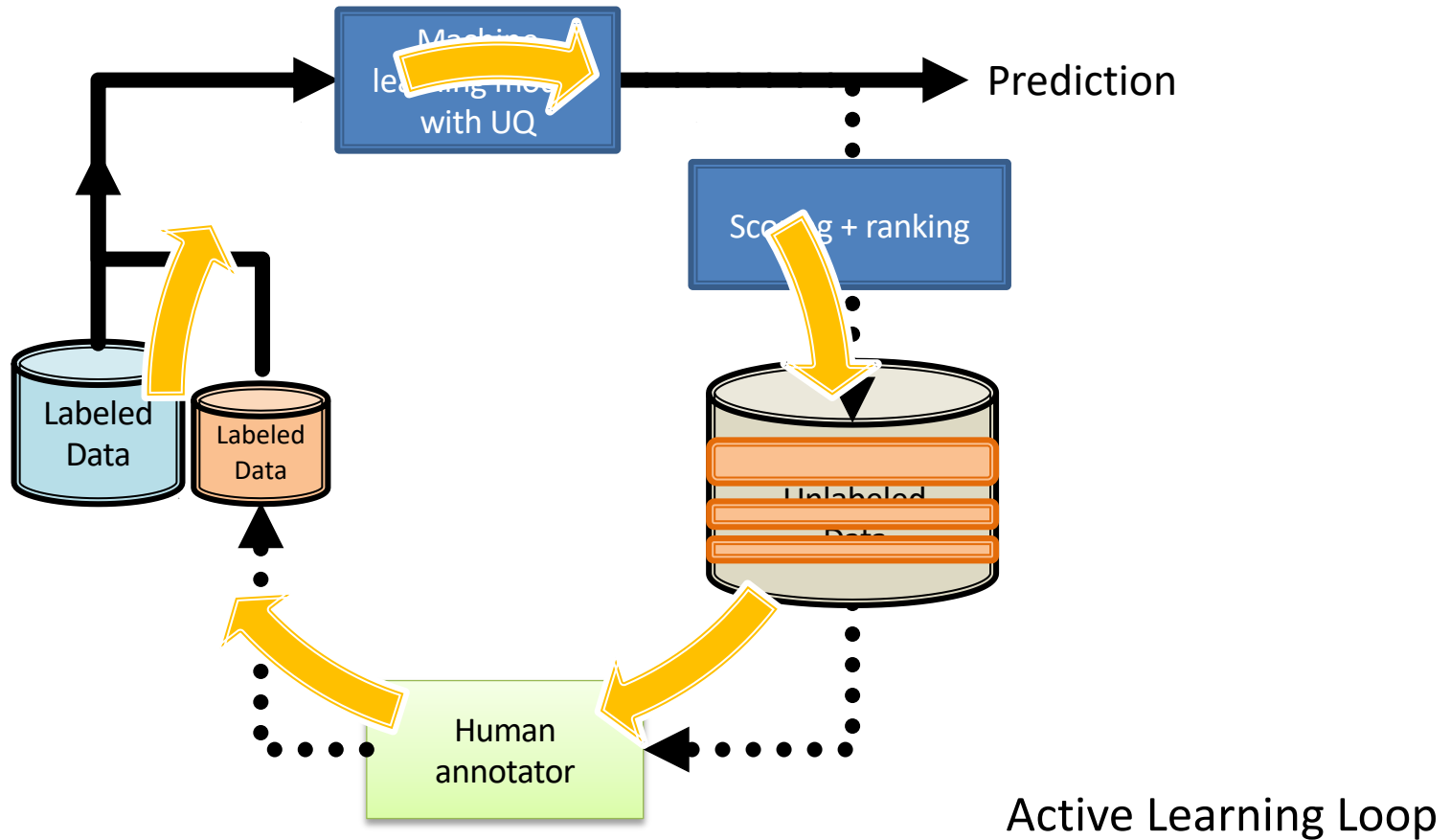
NLP for Pathology Report Classification

- State-of-the-art data-driven NLP systems require large sets of labeled data to reach reasonable accuracy.
- Few skilled Certified Tumor Registrars are available to provide labels to existing pathology reports.
- Solution: **use active learning to direct which pathology reports to label first, so the performance of NLP is maximized.**

Active Learning



Active Learning



Active Learning

- Active learning (AL) **seeks to maximize an ML model's performance gain while labeling the fewest number of samples possible** (reduce annotation cost).
- AL can be seen as a way to integrate humans into the machine learning process (**human-in-the-loop** paradigm).
- AL is a type of **semi-supervised learning**, meaning models are trained using both labeled and unlabeled data.

Active Learning

- Keys components in Active Learning setup:
 - Labeled data << unlabeled data
 - Acquisition function (*scoring function*):
 - Provides a measure of *informativeness* of a given sample to the ML model
 - Indicates how helpful that sample would be to the classifier/regressor if it were to be trained on that sample
- Active Learning loop can be executed indefinitely or until a target performance is reached or the budget is depleted.

AL categorization: Query mechanisms

- Membership query synthesis:
 - learner can request to query the label of any unlabeled sample in the input space, including the sample generated by the learner.
- Stream-based selective sampling:
 - makes an independent judgment on whether each sample in the data stream needs to query the labels of unlabeled samples.
- Pool-based selective sampling :
 - chooses the best query sample based on the evaluation and ranking of the entire dataset.

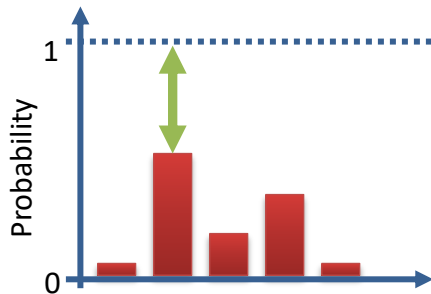
AL categorization: Scoring functions

- Uncertainty-based approach:
 - Select samples the current model are most uncertain about.
- Diversity-based approach:
 - Select samples following the underlying data distribution.
- Expected model change:
 - Calculate expected change in the model given the addition of a sample to the training data.
- Hybrid approaches:
 - Combinations of the approaches above.

Acquisition functions examples

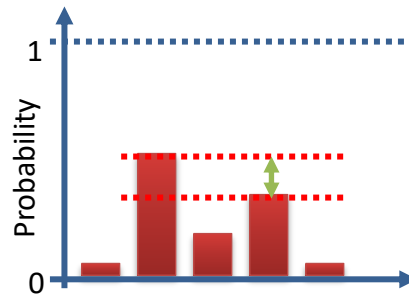
- Common uncertainty-based AF for classification:

Least Confidence



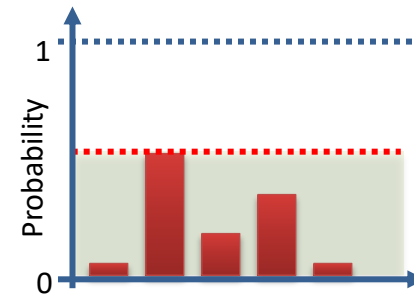
selects the instance for which it has the least confidence in its **most** likely label

Margin Sampling



selects the instance that has the smallest **difference** between the **first** and **second** most probable labels

Shannon's Entropy



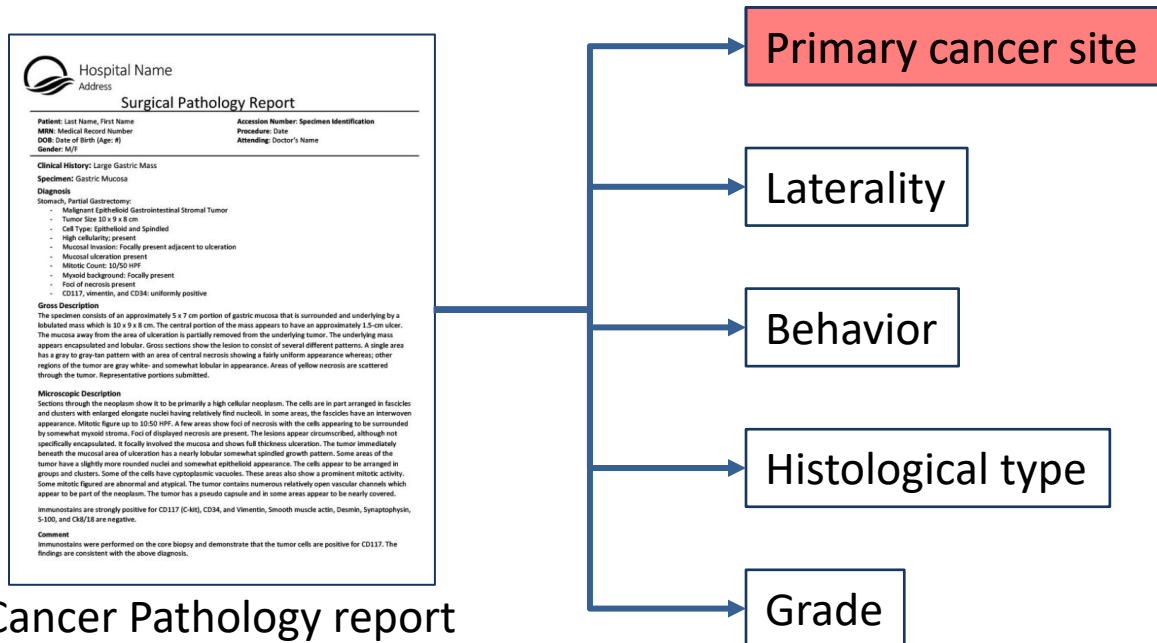
$$H(X) = - \sum_{i=1}^n p_i \log(p_i)$$

Experimental Study: Cancer Pathology Report Classification



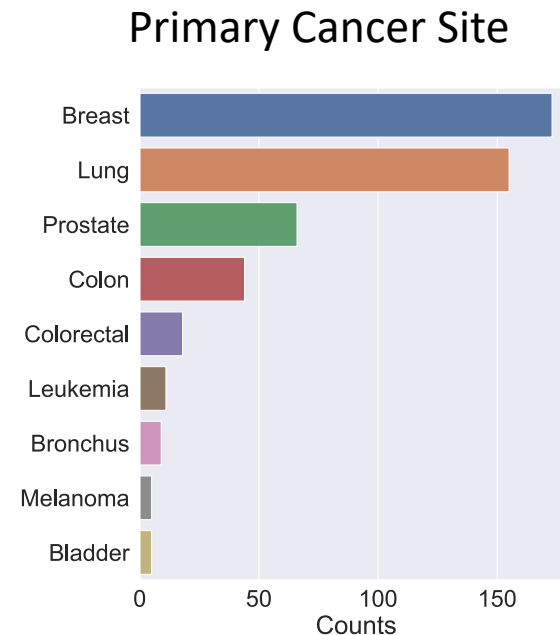
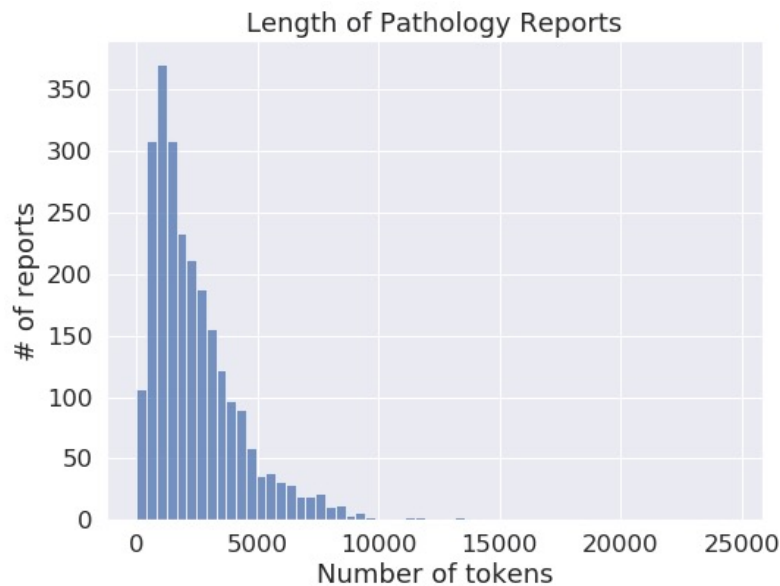
Cancer Pathology Reports Classification

■ Multi-label classification problem:



Pathology Reports Classification: Data

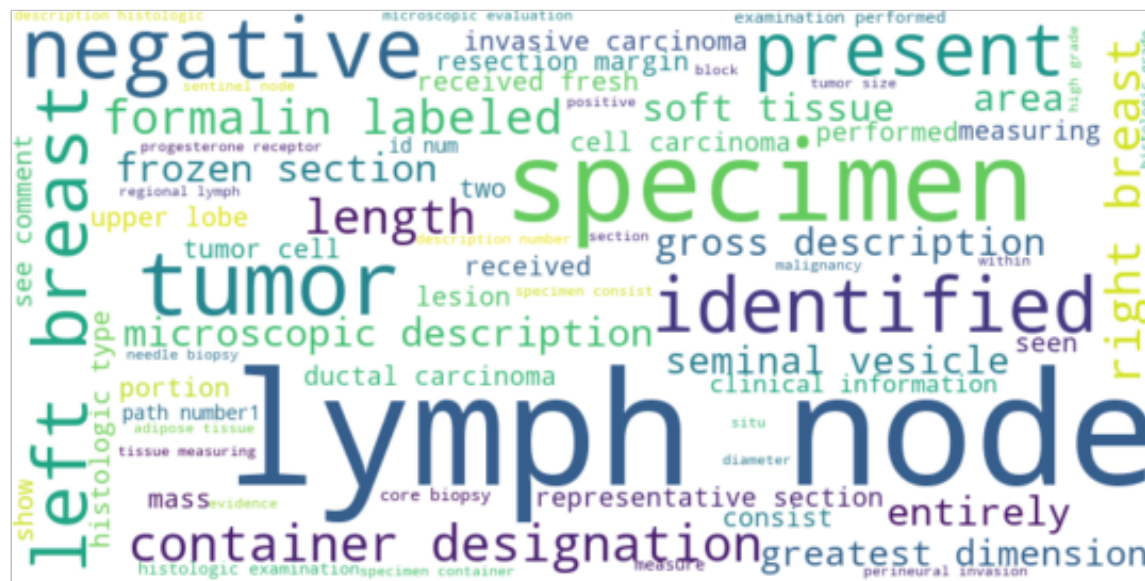
- Data description:
 - 2,500 de-identified cancer pathology reports:
 - 486 reports with **Primary Cancer Site** labelled



Pathology Reports Classification: Data cleaning

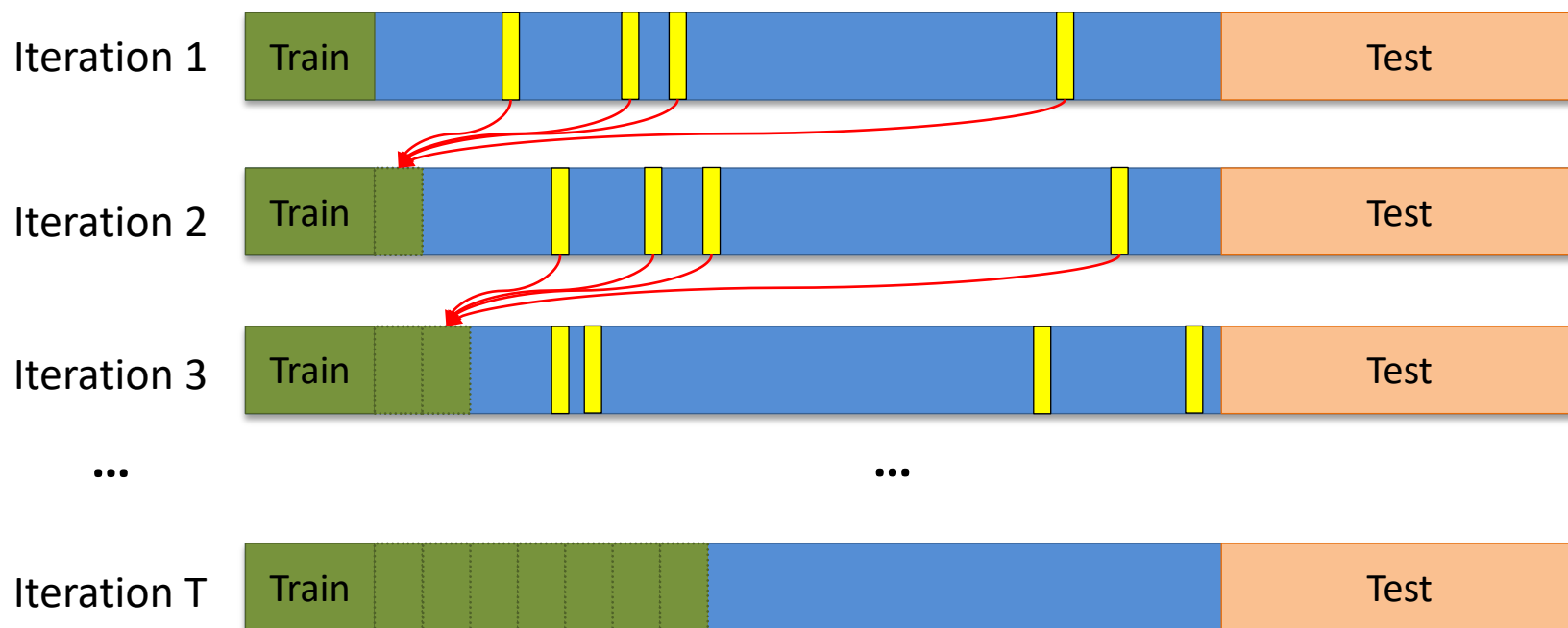
- Cleaning and normalization steps (nltk library):
 - Removed tags, section markers, headers, and footers
 - Removed punctuations
 - Removed non-ascii characters
 - Replaced integers to textual representations (1 → one, ...)
 - Converted to lowercase
 - Removed single character words
 - Removed stop words
 - Manually fixed some typos

- Most frequent words



Experimental Setup

- Stratified train/test/unlabeled split to account for class imbalance



Text Feature Extraction

- TF-IDF BoW with PCA (50 dimensions)
 - Bag-of-words (BoW) concept

$$\text{TF-IDF}(w, d) = \text{TF}(w, d) \times \text{IDF}_D(w)$$

Frequency of w in d

Rareness of the word w :
 $\log\left(\frac{N}{df_w}\right)$

The diagram illustrates the TF-IDF formula. The term $\text{TF}(w, d)$ is highlighted in a light blue box, with a callout box above it stating 'Frequency of w in d '. The term $\text{IDF}_D(w)$ is highlighted in a light green box, with a callout box below it stating 'Rareness of the word w : $\log\left(\frac{N}{df_w}\right)$ '. Arrows point from the callout boxes to their respective terms in the formula.

- Sklearn's TfidfVectorizer
- PCA is applied to the TF-IDF vectors

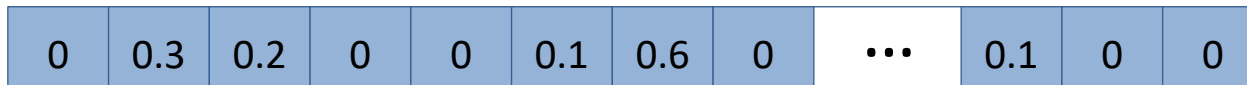
Text Feature Extraction

- TF-IDF BoW with PCA (50 dimensions)

“Microscopically there is a ragged ulcer penetrating to within 5 mm of the serosal surface. The ulcer is surrounded by dense connective response and chronic inflammatory infiltrate.”



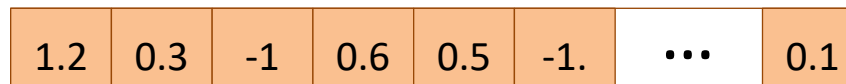
TF-IDF Vectorizer



D-dimensional vector



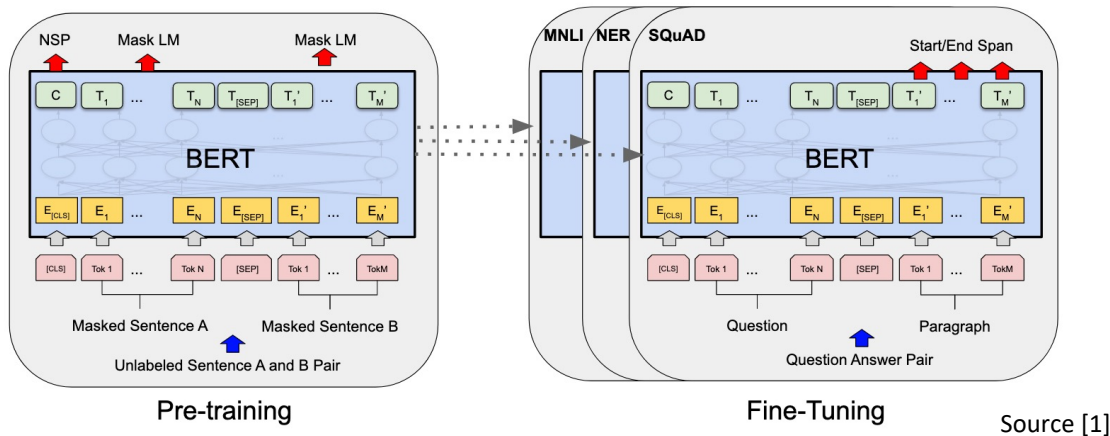
PCA



k-dimensional vector

Text Feature Extraction

- BERT (pre-trained Transformer-based model)



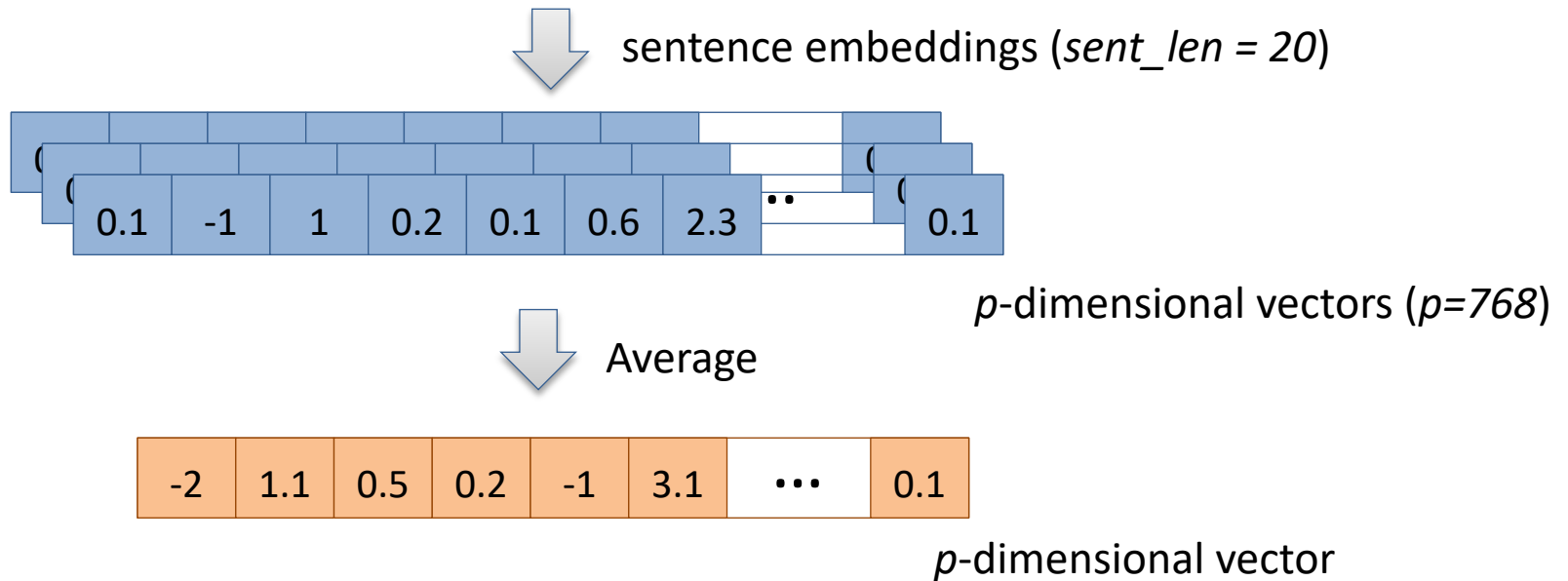
- Pathology report embedding is the mean value from all sentence's embeddings.
- Hugging-face's Transformer library (BertTokenizer, BertModel)

[1] Devlin, Jacob, Ming-Wei Chang, Kenton Lee, and Kristina Toutanova. "Bert: Pre-training of deep bidirectional transformers for language understanding." *arXiv preprint arXiv:1810.04805* (2018).

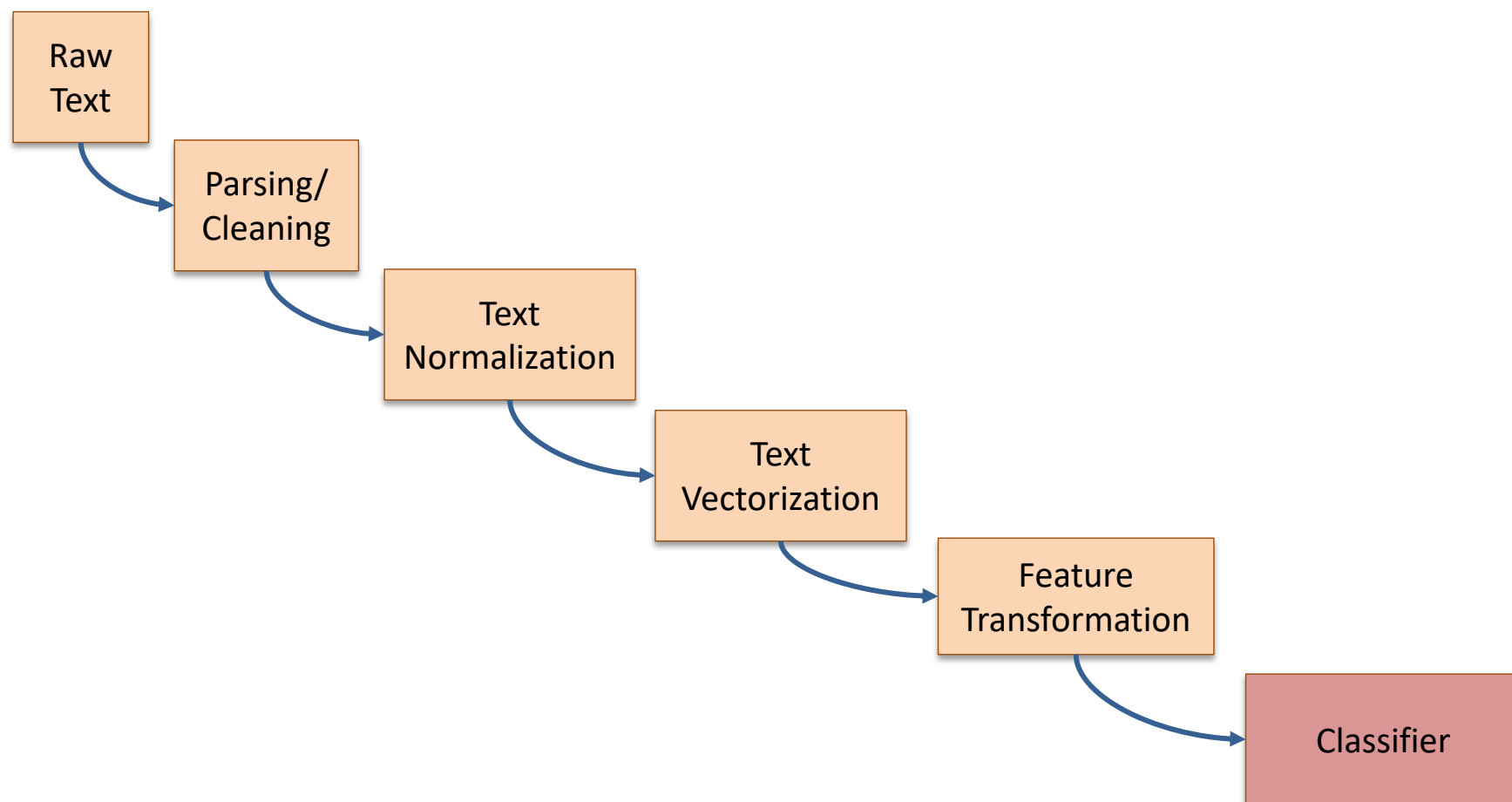
Text Feature Extraction

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Text Analysis Pipeline



Adapted from: *Applied Text Analysis with Python* by Benjamin Bengfort, Rebecca Bilbro, Tony Ojeda (2018)

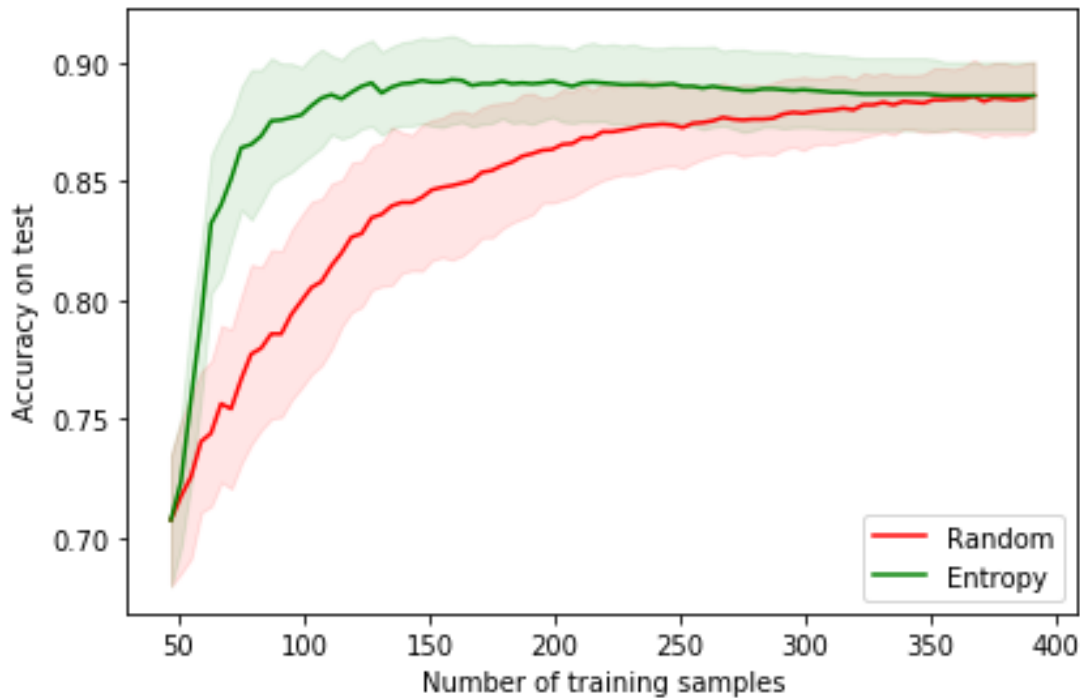
Classifiers (with UQ)

- Deterministic (use softmax probabilities)
 - Logistic regression
 - Deterministic (regular) Neural Network
- Bayesian Models
 - Gaussian Process Classifier
 - Bayesian Linear Regression
 - Bayes-by-Backprop NN model
- Bootstrapping
 - Logistic regression
 - Random Forest
 - Neural Networks

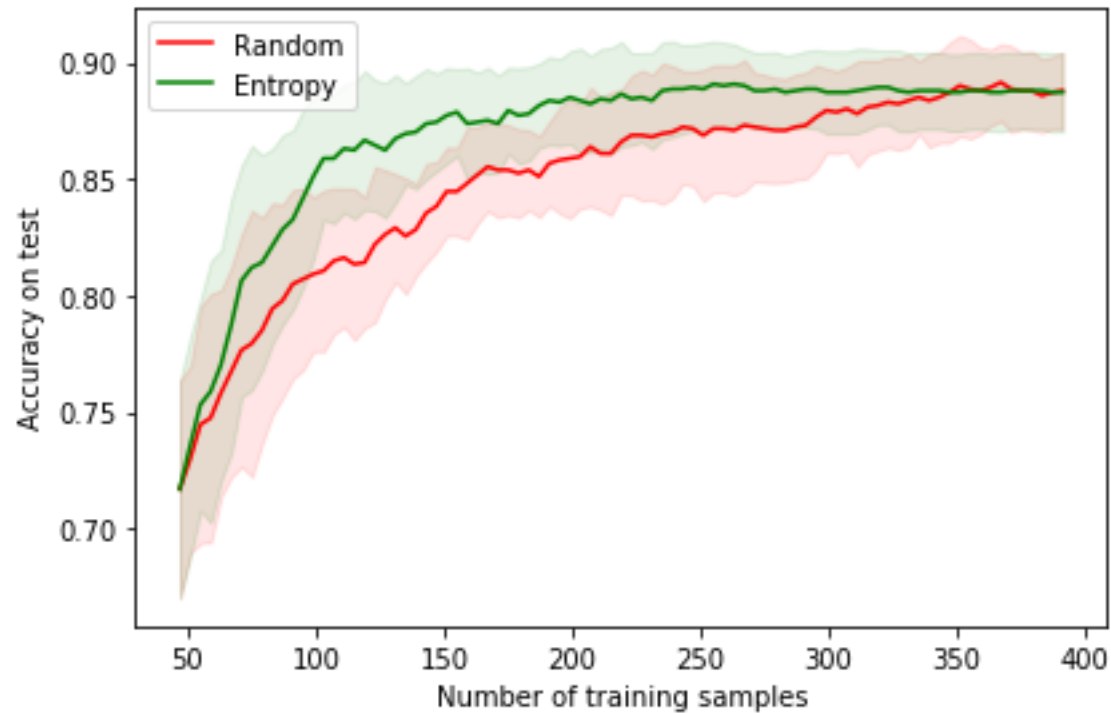
Acquisition Functions

- Random (baseline): randomly select the next set of samples to be labelled
- Entropy
- Least Confidence
- Margin Sampling

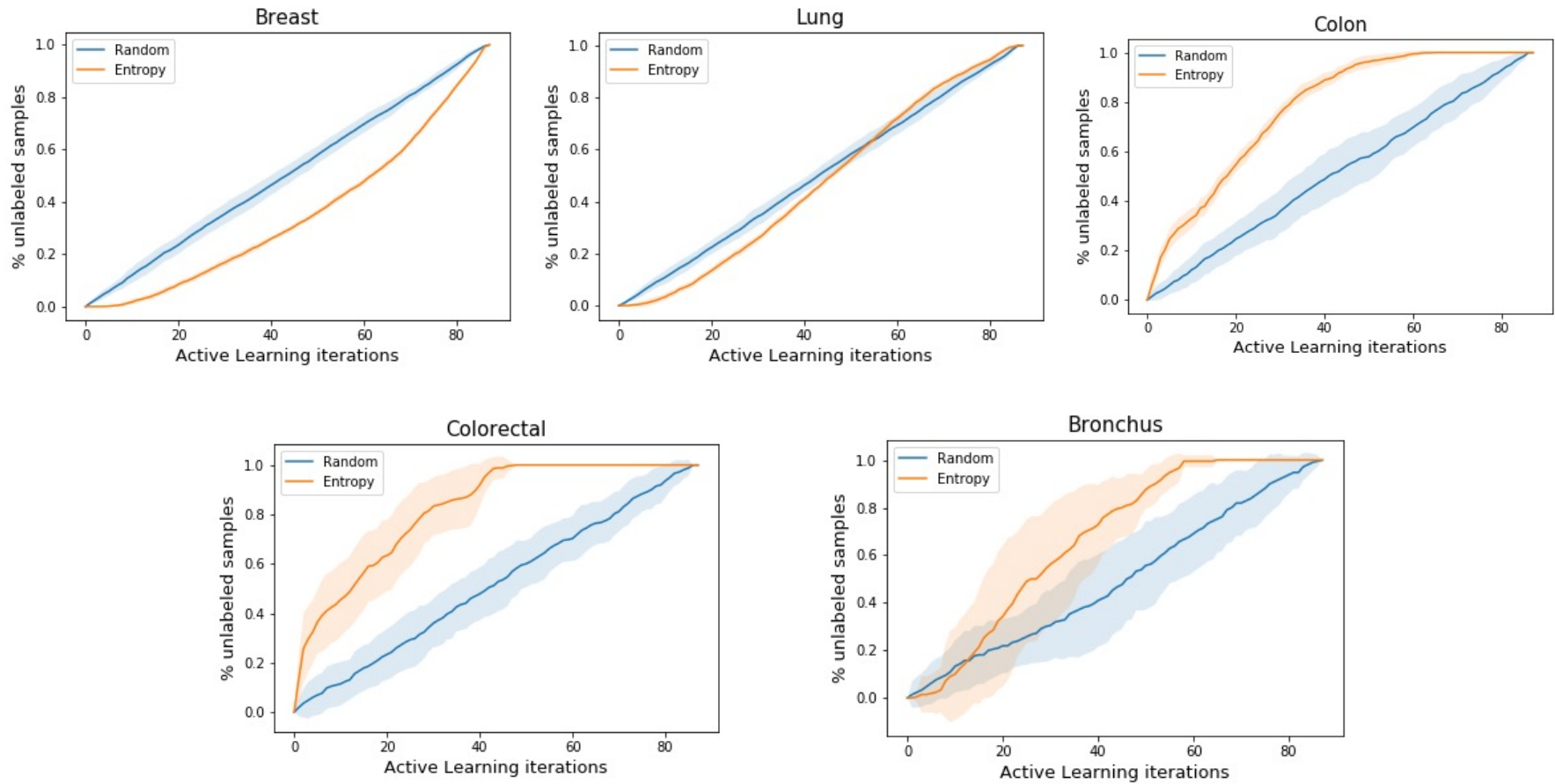
Primary Site: TF-IDF + PCA (50)



Primary Site: BERT model

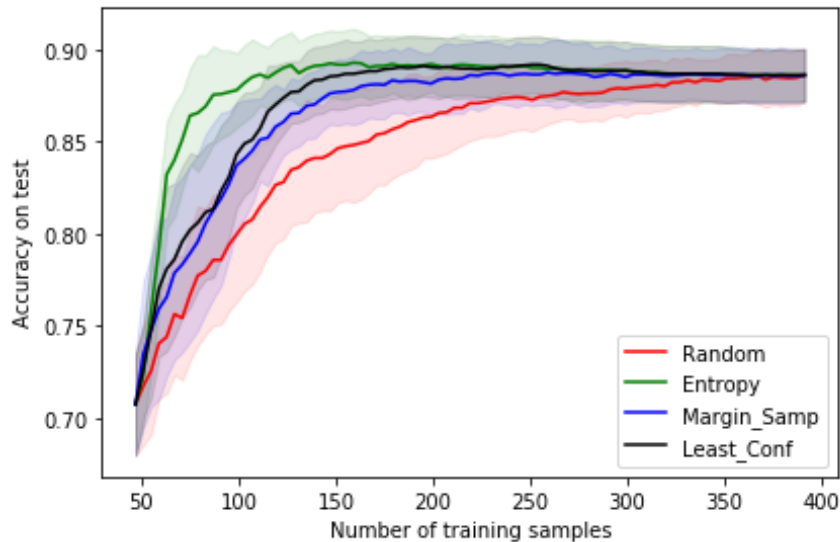


Primary Site: TF-IDF + PCA (50)

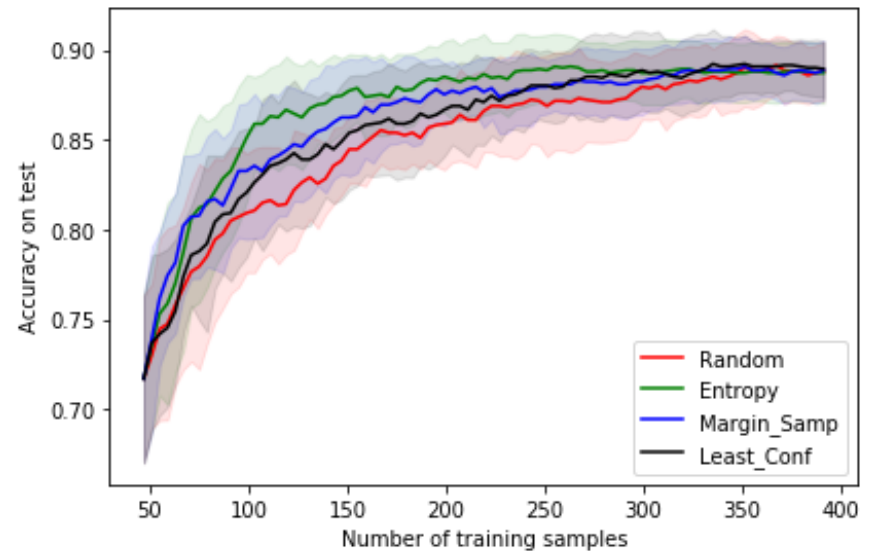


Primary Site: Other Acquisition Functions

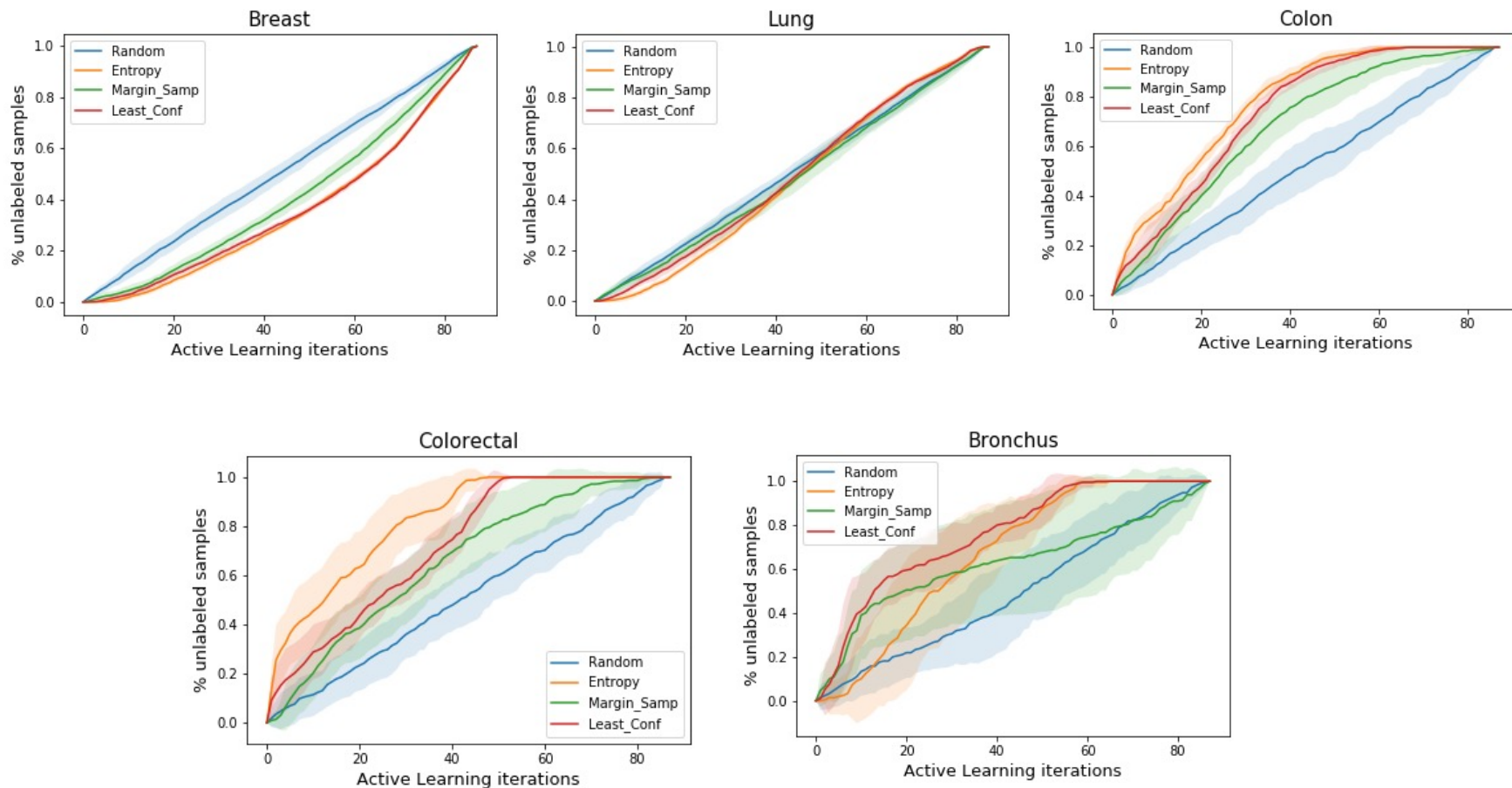
TF-IDF representation



BERT representation



Primary Site: Other Acquisition Functions



Final Thoughts



Final Thoughts

- Automated pathology report annotation can reduce the burden of manual labeling by CTRs at the cancer registries.
- It can accelerate data release and provide abundance of labeled cancer records for statistics/machine learning research.
- NLP is a powerful tool for automating manual labeling processes.
- Active Learning helps maximize a machine learning model's performance while reducing labeling costs.
- Active Learning is a general concept that can be used with many base ML models and setups.

Bibliography

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Thank you.

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