
An Interactive Visualization Tool for Understanding Active Learning

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Abstract

1 Despite recent progress in artificial intelligence and machine learning, many state-
2 of-the-art methods suffer from a lack of explainability and transparency. The
3 ability to interpret the predictions made by machine learning models and accurately
4 evaluate these models is crucially important. In this paper, we present an interactive
5 visualization tool to elucidate the training process of active learning. This tool
6 enables one to select a sample of interesting data points, view how their prediction
7 values change at different querying stages, and thus better understand when and
8 how active learning works. Additionally, users can utilize this tool to compare
9 different active learning strategies simultaneously and inspect why some strategies
10 outperform others in certain contexts. With some preliminary experiments, we
11 demonstrate that our visualization panel has a great potential to be used in various
12 active learning experiments and help users evaluate their models appropriately.

13 1 Introduction

14 Recent advances in machine learning (ML) and artificial intelligence (AI) have been mainly focused
15 on making models more accurate and efficient. However, there has been an increasing focus on
16 improving the explainability and interpretability of models as well as considering the human users of
17 these models [1]. It is crucially important for humans to be able to explain the predictions made by
18 ML models especially in some sensitive domains, such as automated financial investing, autonomous
19 driving, and healthcare management. There are still many ML applications where state-of-the-art
20 models can achieve strong predictive power, but the gain in accuracy comes at the cost of transparency,
21 and the decision reached lacks interpretability [2]. Meanwhile, several algorithms may achieve highly
22 similar performance based on some experimental setting, which makes it difficult to evaluate their
23 effectiveness and distinguish how they truly work.

24 Active Learning (AL) is an approach that has proven useful when labeled training data is scarce and
25 unlabeled data is plentiful, as is the case in many real-world settings [3]. AL works by selecting
26 unlabeled examples to be labeled by an oracle (usually a human annotator), in order to train an
27 accurate model with the least labeled examples [4]. The majority of theoretical work in AL has
28 taken place in classification problems, with many fundamental and successful algorithms developed
29 [5, 6]. To evaluate AL, the most common approach is to plot *accuracy* by number of queries, where
30 we expect accuracy to improve as more samples are queried. A few recent studies also adapt some
31 similar algorithms for regression [7, 8], and the usual performance measure is *Mean Squared Error*
32 (MSE). However, the information reflected from either the accuracy or MSE plots is both limited and
33 potentially biased to users. For example, many previous papers presented highly similar accuracy
34 curves [9, 10, 11], yet a number of contradictory findings have been recorded in past work [10].
35 Hence, users may not be able to well-distinguish different behaviors between diverse AL algorithms
36 singly from the accuracy plot.

37 In this paper, we propose a novel interactive visualization tool for people to better understand why
 38 and how AL works on certain classification and regression tasks. By using dimension reduction
 39 techniques to create a 2-D feature embedding for visualization, users are able to select a proportion
 40 of interesting test data points and view how their prediction values change according to more queries.
 41 The prediction values for selected points are arranged to a 2-D mesh-grid plot called **prediction-**
 42 **change** plots with each pixel showing the prediction differences. Compared to normal accuracy
 43 curves, our new tool illustrates how accuracy changes for a specific subgroup of data. We demonstrate
 44 that with our approach, users could compare different AL strategies simultaneously, get an intuition
 45 on why certain algorithms do not work on some test points, and evaluate them appropriately.

46 2 Related Work

47 **Active Learning and Interactive Visualization** To the best of our knowledge, there is no estab-
 48 lished Human-Centered or interactive visualization tool for understanding how and why AL works.
 49 The closest we can get is an online prototype "Active Learner" [12]. It serves as an exhibition and
 50 only gives settled results for three different datasets and four AL strategies. Other notable examples
 51 include work by Limberg et al. [13], in which they introduced a user interface for querying samples
 52 to be labeled, specifically for image recognition problems. In addition, Iwata et al. [14] proposed an
 53 AL framework for interactive visualization which selects objects for the user to re-locate so that they
 54 can obtain the desired visualization to their preference. Crucially, these studies did not investigate
 55 explainability or visualization for the evaluation of AL methods.

56 **Evaluation of Active Learning** Many successful AL querying methods have been developed
 57 and shown to outperform some baseline (often random sampling). Comparing accuracy/MSE over
 58 querying more samples is the most common approach for evaluation. Other metrics are sometimes
 59 used as well such as AUC and F1-score [15]. However, focusing on accuracy could lead to unexpected
 60 and unwarranted conclusions. For example, Ramirez-Loaiza et al. [15] claimed that AL often
 61 improves accuracy at the expense of recall, which means that the observed improvement in accuracy
 62 is not fully due to *effective* learning. To draw a proper evaluation of AL, accuracy plots are usually not
 63 sufficient and other facets should be considered, such as AL experimental setting, labeling efficiency,
 64 and redundancy [10]. Our proposed visualization tool can be helpful in many of them.

65 3 Prediction-change Plots

66 3.1 Definition of the prediction-change Check

67 We consider a general AL process, in which an oracle provides a fixed number of labels. The initial
 68 labeled dataset is $D_l = \{(x_1, y_1), \dots, (x_M, y_M)\}$ and the unlabeled data pool is $D_u = \{x_1, \dots, x_N\}$
 69 where each instance $x_i \in \mathbb{R}^d$ is a d -dimensional feature vector. y_i is the ground truth value of x_i . In
 70 each iteration, the model chooses $D' = \{(x', y')\}$ as the queried new sample.

71 Let f be the mapping function of our model (either a classifier or a regressor) and θ be its parameters
 72 (if any). For a test sample x_{test} from the test set $D_{\text{test}} = \{(x_{\text{test}}, y_{\text{test}})\}$, the prediction of the model
 73 trained on the original training set D_l is $\tilde{y}_{\text{test}} = f(x_{\text{test}}; \theta_{D_l})$. The prediction of our model trained
 74 with new added samples is $\tilde{y}'_{\text{test}} = f(x_{\text{test}}; \theta_{D_l+D'})$. Single \tilde{y}_{test} may or may not change.

75 **The first prediction-change check:** For some of the test samples, we expect the new prediction
 76 value of $\theta_{D_l+D'}$ is different from the old one of θ_{D_l} :

$$|f(x_{\text{test}}; \theta_{D_l+D'}) - f(x_{\text{test}}; \theta_{D_l})| \neq 0 \quad (1)$$

77 If there is no change for all test samples, this may suggest our new queried sample does not provide
 78 new information for the model.

79 **The second prediction-change check:** For some of the test samples x_{test} we selected, we expect the
 80 prediction of $\theta_{D_l+D'}$ is closer to the ground truth y_{test} , than to the prediction of θ_{D_l} :

$$|f(x_{\text{test}}; \theta_{D_l+D'}) - y_{\text{test}}| < |f(x_{\text{test}}; \theta_{D_l}) - y_{\text{test}}| \quad (2)$$

81 The prediction values can possibly be worse for some test samples. Overall, the sum of all prediction
 82 differences from our selected test samples compared to all ground truth values is expected to be
 83 smaller, which means the model is consistently performing better.

84 **3.2 Design of the Interactive Visualization Panel**

85 Regardless of how the AL experiment is conducted, this interactive visualization tool only requires
 86 two inputs: 1. the test dataset D_{test} ; 2. the prediction values on this set over a number of AL queries
 87 (could be stored in a matrix). We let users dynamically select "similar" and interesting test samples to
 88 look at how their prediction values change.

89 Firstly, Principal Component Analysis (PCA) is conducted and the first two PCs are extracted. Then,
 90 we plot these PCs to create a 2-D feature embedding for visualization and depict a general structure
 91 of all test samples, if there exists some similarity between the test samples. Finally, users can select a
 92 targeted region of points that they want to inspect. The interactive panel will automatically generate
 93 the *prediction-change* plots based on a number of (default is 20) nearest points. In detail, prediction
 94 values for selected points are arranged to a 2-D mesh-grid plot: **x-axis** represents the querying process
 95 (unit could be single or batch queries); **y-axis** represents the indices of selected test samples; **each**
 96 **pixel** shows prediction differences according to some criteria. Derived from definitions in 3.1, we
 97 propose three types of prediction-change plots based on the change of the prediction values between:

98 **1. current model vs. original model** (Equation (1)): If the model is progressively improving and
 99 learning from the queried instances, we expect the prediction values to be gradually more and more
 100 different. This can be reflected by more colorful mesh-grid graphs.

101 **2. current model vs. previous model** (Equation (1)): For some strategies, we expect later queried
 102 samples to bring less effect on changing the model. This can be reflected by lighter colors on the plot.

103 **3. current model vs. ground truth** (Equation (2)): As querying more samples, we expect the model
 104 to make better predictions that are closer to ground truth. This can also be reflected by a lighter plot.

105 Notice that for classification, pixels in the prediction-change plot have finite number of colors. For
 106 example, prediction-change values can only be $\{1, 0, -1\}$ for a binary classification with labels
 107 $\{0, 1\}$. In regression, these values are continuous, which suggests pixel colors to be gradient-based.

108 **4 Experiments**

109 **Settings** Because gradient colors are more recognizable in regression, we conducted our experiments
 110 on a regression problem with real world data: CASP [16], which contains 45,730 instances in
 111 total. We randomly selected 9,730 (%21) of them as the test set. For the AL process, we started with
 112 an empty training set, batch queried 500 at each iteration, and stopped after 15 batches. Fig. 1 (left)
 113 shows the MSE plot, and Fig. 1 (right) illustrates the PCA plot for all 9,730 test samples.

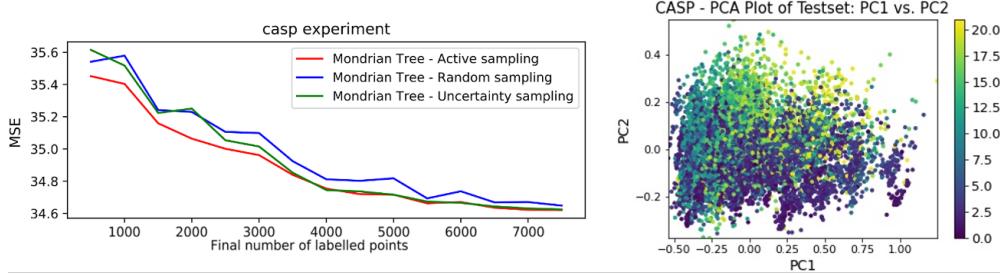


Figure 1: **Left:** MSE plot of the AL experiment. **Right:** PC1 vs. PC2 of all test samples

114 The regressor comes from Goetz et al. [7], which used purely random trees, specifically, Mondrian
 115 Trees. We compared the performance of three basic querying strategies: its tree-based active algorithm
 116 (al), a naive uncertainty sampling (uc) version of its active algorithm, and random sampling (rn).

117 **Results** There are some obvious clustering patterns in the PCA plot. To start off, we select the
 118 points at the bottom right corner with the smallest values. The red points in the PCA plot of Fig. 2
 119 mark the points we selected. Also, 3 * 3 panel plots are provided for the prediction-change plots of
 120 three strategies ("al", "uc", and "rn") in each row. A complete user-interface is in Appendix Fig. 3.

121 Looking at the *first* prediction-change plots across three querying algorithms, it is fairly clear that
 122 they perform differently in changing the regressor model, even though they have similar MSE curves

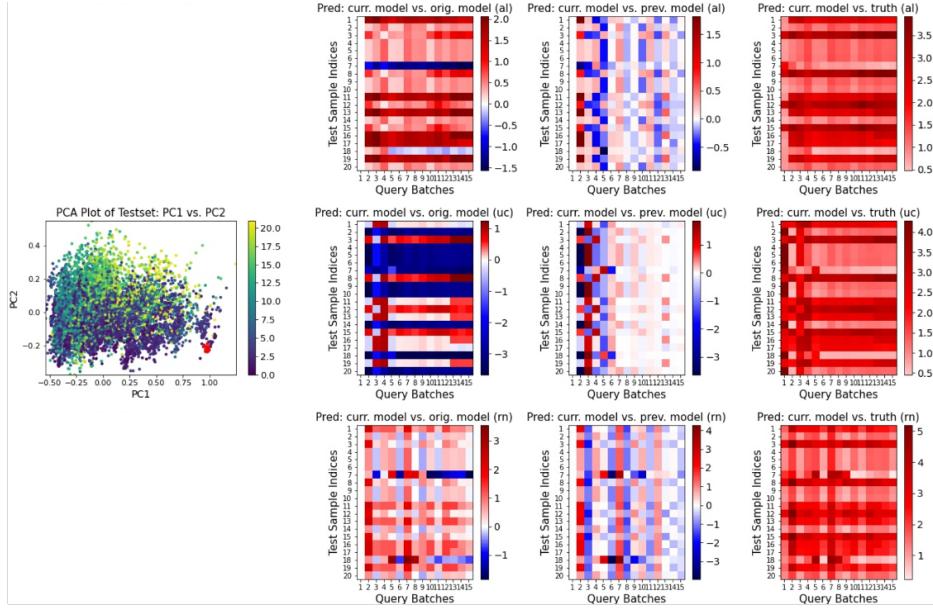


Figure 2: Prediction-change plots for three AL strategies on examples with small values

in Fig. 1. "al" performs more regularly than "uc" or "rn", with consistently larger values (18 out of 20 purely red rows) or smaller values (1 out of 20 purely blue rows) compared to the initial model. However, unlike this diverse performance on the low-value group, three algorithms show almost uniform performances for the large value group (plots shown in Appendix A.2 Fig. 4). To investigate potential reasons, we further check the batched queried samples and their true values (data shown in Appendix A.3 Fig. 5). Compared to the original CASP data distribution, all three algorithms show similar data distributions for the first 500 batch-queried samples. Although "al" and "uc" seem to query more large value points (near 20), unfortunately this does not help the Mondrian Tree model perform significantly better in making predictions. One inference we can draw is that the model does not learn well from the queried samples for fitting large value points or data points from a sparse area.

In the *second* prediction-change plot of "uc", we see dark pixels at first and quite light, nearly white color pixels after 5 query batches. This suggests that "uc" tends to change the model dramatically at the first stage and has less effect in later queries. The *third* prediction-change plots consistently showed red pixels, which intuitively makes sense because the dark points we selected have small truth values near 0. The tree regressor tends to fit the structure with larger predictions than the truth.

138 5 Conclusion

Human-Centered AI and ML are receiving increased attention from researchers, although there have not been many established Human-Centered techniques for AL. In this paper, we have proposed an interactive visualization tool for AL based on prediction values, inspired by the background that there is limited explainability potential from traditional accuracy/MSE plots. We introduce a simple definition of prediction-change checks, which is applicable in both classification and regression settings. Users are able to select a sample of data points and get a better intuition on how the predictions of those points change through AL. Our initial experiments on real-world regression tasks indicate several interesting insights and suggest that there are many more that this tool could reveal.

147 Limitations and Future Work It is easier for users to play around with this visualization panel
 148 and draw conclusions if there are clear clusters shown on the PCA plot. If test samples do not show
 149 any clear association or cluster groups, the AL task will be more complicated because of no clear
 150 decision boundaries, which further makes our prediction-change plots less distinguishable. There
 151 are many additional questions we can ask by running these panel plots in different experimental
 152 settings. Future work involves testing on different data sets, which leads to many interesting avenues
 153 for further exploration - e.g., do certain AL methods work well in certain data subgroups?

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192 **A Appendix**

193 **A.1 User Interface of the Prediction-change Plots**

194 This is a sample user interface of the visualization panel, using the example from Fig. 2.

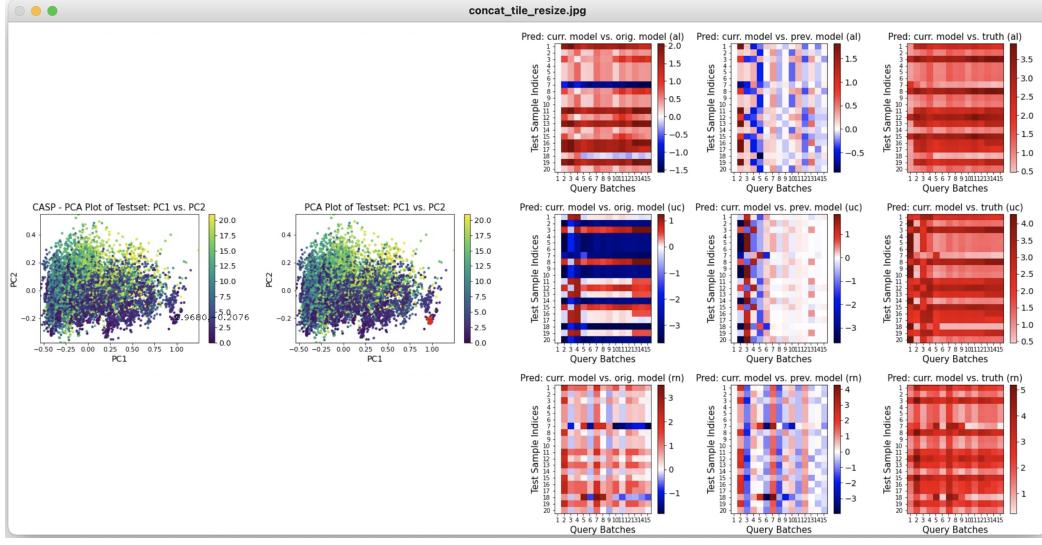


Figure 3: The actual user interface of Fig. 2

195 A.2 Prediction-change Plots for Large Value Group

196 In this example, the red points we selected contain large values (light yellow color in PCA plot;
 197 closed to 20). Three querying algorithms ("al", "uc", "rn") show highly similar performance for all
 198 prediction-change plots. The tree model might struggle to regress this large value group, regardless
 199 of what new samples being queried.

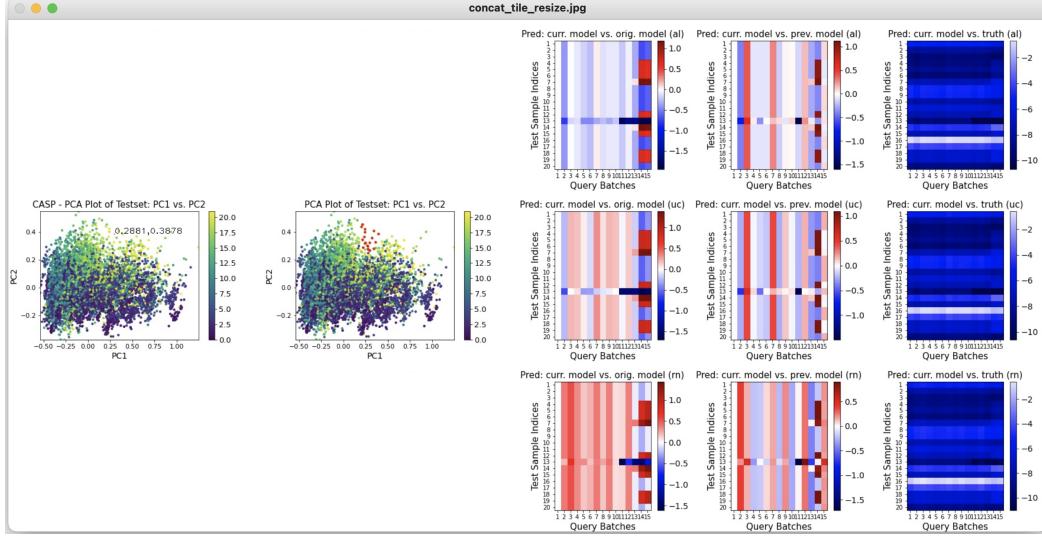


Figure 4: Prediction-change plots for three AL strategies on large value points

200 A.3 Distributions of All CASP Data and Queried Samples

201 By looking at the truth values of our batch-queried samples, we find that "al" and "uc" tend to query
 202 a bit more large value points, compared to the original distribution of all CASP data. However,
 203 unfortunately this does not make "al" or "uc" perform much better than "rn" in this large value group.

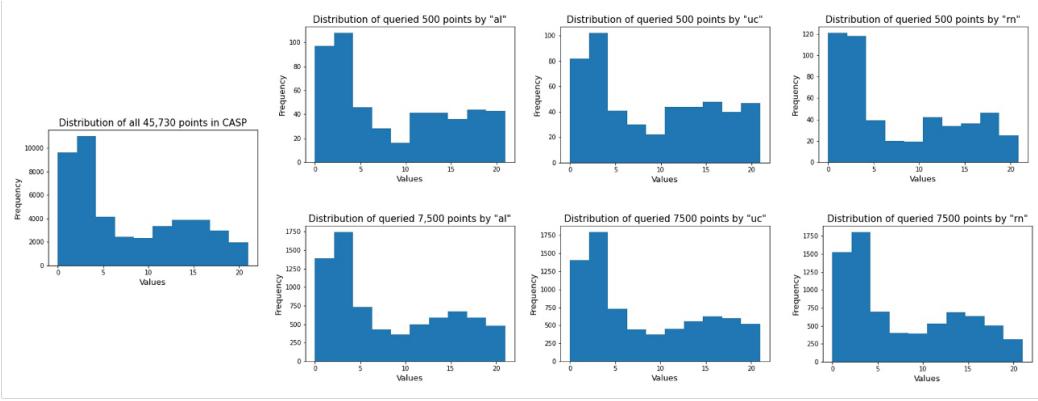


Figure 5: **Left:** Distribution of y-values for 45,730 points in CASP **Right-Top:** Distribution of 500 queries by three algorithms **Right-Down:** Distribution of 7500 queries by three algorithms

204 **A.4 Demo & Reproduce All Experiments**

205 A GitHub demo is available at: https://github.com/anxiousrabbit1/Active_Learning_Visualization_Demo

207 To reproduce all experimental results, all data and scripts are available at: https://github.com/anxiousrabbit1/Mondrian_Tree_AL