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ARTICLE TYPE

Visual Diagnostics of a Model Explainer – Tools for the Assessment of LIME Explanations

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Present Address

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Summary

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KEYWORDS:

LIME, black-box models, interpretability, diagnostics

To check on when editing writing:

- Generally: avoid 'can be'. Replace by 'is'.
- References like 'they', 'them' ... replace them by repeating the noun you are referring to to avoid any kind of ambiguity
- words to go back and make sure I am not using too often: ability, produce, understand

1 | INTRODUCTION

*** I added some subsections to the introduction, because it was getting so long. What do you think? Would it be better to make the introduction shorter and move the details to a separate section?

In the field of statistics, there are two main uses for models: inference and prediction. Machine learning models are often used for the latter purpose. While these models have been proven to perform well in a wide range of prediction problems, their accuracy comes at the cost of interpretability. They are often complex algorithms that are good at identifying patterns in the data but lack a functional form *** Is it okay for me to make this statement about the functional form?, so it not possible to

directly interpret them. As a result, they produce predictions that lack an explanation, which has earned them the reputation of being "black-box models".

1.1 | The Importance of Explanability

The ability to interpret a model serves multiple purposes. When a model is first fit, interpretations help to diagnose the model. Knowing which variables influence a prediction makes it possible to determine if the predictions are based on reasonable variables. If not, appropriate adjustments to the model can be made. After a model has been diagnosed, the explanations for the predictions are used to understand the underlying mechanism that produced the data and provide an argument for why the predictions should be trusted. In some areas of application, this ability to be able to explain the results from a model is critical.

The forensics sciences and the heath care industry are two areas that have benefited from machine learning but have an obvious need for explainable predictions. For example, Hare et al. [2] discuss the use of a random forest model in the forensics sciences to determine whether two bullets were fired from the same gun. Yu et al. [5] describe how healthcare has advanced through the use of machine learning in ways such as using neural networks to automate medical image diagnoses and implementing Bayesian networks to predict clinical outcomes. In both

of these fields, the decisions made based on the results of the machine learning models can greatly impact people's lives. If it is not possible to explain the output from a model, it becomes questionable whether to rely on the predictions to make important decisions.

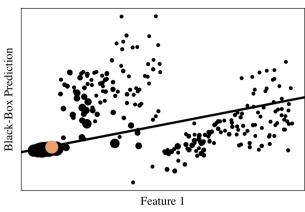
As an example, suppose that forensics examiners are trying to determine if a bullet used in a crime was fired from the gun that belongs to a suspect in the investigation. A random forest model could be used to produce a probability that the bullet was fired from the gun under question. Even if the model returns a high probability of a match, it will be difficult for a jury to trust the results from the model when deciding whether to convict the suspect without the forensics examiners having the ability to explain which factors in the model led to the high probability.

The European Union has taken the desire to provide explanations for machine learning model predictions a step further. In May 2018, the General Data Protection Regulation (GDPR) went into effect ¹. The regulation includes a policy that gives the right for individuals affected by decisions made via automated algorithms to understand the logic behind the decisions being made. As Goodman and Flaxman [1] point out, this regulation has a great effect on the way that machine learning algorithms are used to make decisions. The wording of the policy leaves room for interpretation, but the authors believe that, at a minimum, the regulations will require that an explanation of how the input variables relate to the predictions be provided.

1.2 | Explainer Models

As a result of the movement to produce explainable predictions, an area of research has emerged focusing on developing ways to explain output from machine learning algorithms. One approach that is being taken is to develop model explainers that provide insight into the performance of the complex model (reference). A model explainer is a method that is separate from the model but uses the model and output from the model to shed light on the process that the model goes through to produce predictions. LIME (local interpretable model-agnostic explanations) is one such model explainer [4].

While some model explainers are focused on understanding a model at the global level, LIME is designed provide a local explanations by focusing on a single prediction of interest. Additionally, LIME was designed to



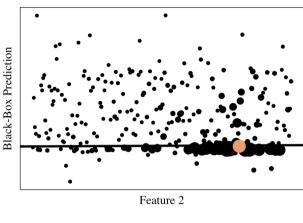


FIGURE 1 Conceptual depiction of LIME. The black lines represent a weighted linear regression model being used as the explainer model for a hypothetical black-box model fit with two features. The explainer model suggests that Feature 1 is driving the prediction made by the black-box model for the case of interest since it has captured the slope in the local region around the prediction of interest.

Prediction of Interest

work with any model and to produce easily understandable explanations. [4] XXX a citation might be good here for local and agnostic Is it okay to include the same reference to the original LIME paper again? Conceptually, LIME fits a simple interpretable model, referred to as the explainer, that is meant to capture the behavior of the complex "black-box model" in a local region around a prediction of interest. The interpretation of the simple model is then used to explain the variables that most influenced the prediction made by the complex model.

Figure 1 provides a visualization of this conceptual understanding of LIME. These figures show the predictions from a hypothetical black-box model plotted against

 $^{^{1} \}rm https://eugdpr.org/the-regulation/gdpr-faqs/$

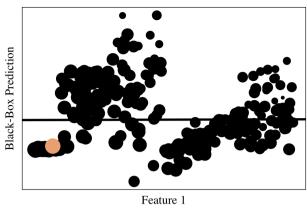
the two features used to fit the black-box model. The orange point represent one prediction that is of interest to explain. The size of the points represent the proximity to the prediction of interest measured using the Gower distance metric (reference and add exponent value?). The black lines represent the explainer model which is a weighted linear regression model in this case fit with the black-box predictions as the response variable and Feature 1 and Feature 2 as the covariates in the model where the proximity values used as the weights in the model. The top image shows that there is a complex relationship between black-box predictions and Feature 1. Here the explainer model is plotted with Feature 2 set to be the observed value of Feature 2 for the prediction of interest. The explainer model does a good job of capturing the relationship in the local region around the prediction of interest. The bottom shows that there is no relationship between the black-box predictions and Feature 2 in either the global sense or local region around the prediction of interest. Here the explainer model is plotted with Feature 1 set to be the observed value of Feature 1 for the prediction of interest, and it has a slope of approximately 0. This explainer model would indicate that Feature 1 is driving the prediction made by the black-box model for the prediction of interest.

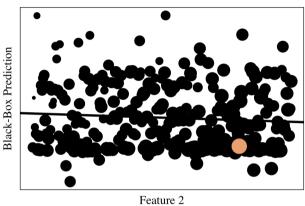
1.3 | Motivation for Diagnosing LIME

At a conceptual level, explainer models add another level of complexity to predictive models: in trying to explain the black-box model, a simpler explainer model is added. To be able to trust in the explanation it is imperative to check that the explainer model is reliable and does not over-simplify the black-box model.

The current implementations of LIME use a linear regression model as the explainer model (references to R and Python packages). These implementations are relying on the assumption that the relationship between the complex model predictions and the features is linear in a local region. It is important to assess this assumption in the local region of interest in order to trust the explanations produced by LIME. Additionally, the implementations offer various input settings, but little work has been done to provide advice on how to specify the settings in practice (look more into this). An assessment of the explanations could also help to determine which settings produce the explainer model with the best approximation of the complex model.

Figure 2 provides an example where the explainer model is doing a poor job of approximating the complex model. The plots were created using the same data as





Prediction of Interest

FIGURE 2 A second version of explainer model applied to the data from Figure 1 . The model explainer is refit using an adjusted distance measure. It leads to an explainer model that is doing a poor job of approximating the complex model in the local region around the prediction of interest for both features.

Figure 1 , but the Gower distance metric was adjusted (add exponent value), so that observations further away from the prediction of interest are given larger weights. In this case, the explainer model is doing a poor job of capturing the relationship between the black-box predictions and the feature for both Feature 1 and Feature 2. However, the magnitude of the slope of the line in the bottom figure appears to be larger than . As a result, this explainer model would return an explanation that Feature 2 is driving the prediction made by the black-box model for the case of interest, which is not an accurate explanation.

In the scenario depicted in Figure 1 , the distance metric was adjusted until the explainer models produced a good local approximation of the complex model. For

Figure 2 , the default setting for distance measures was used by the LIME R package. Without these images, it would not be obvious that the explainer model used in Figure 2 is a poor local approximation.

1.4 | Overview of Paper

In this paper, we will present some visualizations tools to assess the explanations from LIME. While predictive models are used in both regression and classification settings, we will focus on the classification setting for this paper. For additional simplicity, we will only discuss the case with a dichotomous response variable, but we believe that the work is adaptable for other situations.

Section 2 provides some background on the LIME algorithm and the current implementations of LIME. In Section 3, we discuss ways to assess the LIME explainer model and introduce our visualization tools. Section 4 demonstrates the use of our diagnostic tools with a random forest model fit to a forensics bullet matching dataset (***I went ahead and excluded the idea of including the iris data). To conclude, Section 5 reviews the importance of assessing the LIME explainer model, discusses how the results from the example provide possible insights into the workings of LIME, and suggests future research directions.

2 | BACKGROUND ON LIME

LIME was developed in 2016 by Ribeiro et al. [4]. The authors were interested in developing a model explainer method that produced easily understandable explanations for individual predictions made by any predictive model [4]. It was initially implemented in a Python package² by the original authors and was later adapted to an R package by Thomas Lin-Pedersen³.

The LIME algorithm is presented in the original paper in a general context that accounts for cases such as text classification and feature recognition. For this paper, we are only considering the tabular data situation with a binary categorical response variable and continuous feature variables. As a result, the following subsections are written in terms of this context. Furthermore, in this paper we are relying on the LIME implementation in R [3]. This implementation deviates at times from the original implementation by Ribeiro et al. [4]. Whenever the implementations differ, we will highlight these deviations.

2.1 | LIME Procedure in the Context of Tabular Data with a Dichotomous Response Variable

***Need to clean up the notation. Should I use the same notation as the original paper? Let \mathbf{X} be an n by p data matrix with p features and n observations, and let $x = \begin{pmatrix} x_1 & x_2 & \dots & x_p \end{pmatrix} \in \mathbb{R}^p$ be the observation of interest. Furthermore, let \mathbf{y} be a vector of length n of response values and $y \in \{0,1\}$ be the observed response value associated with x. Suppose that f is a classification model where $f: \mathbb{R}^p \to [0,1]$ that is applied to \mathbf{X} and \mathbf{y} . Let the predictions made by f applied to \mathbf{X} be denoted as $\hat{\mathbf{y}}$. It is of interest to explain the prediction made by f when f is applied to f. Note that f(f) is equal to the probability that f is a classification made by f when f is applied to f. Note that f is equal to the probability that f is a classification made by f when f is applied to f in the probability that f is equal to the probability that f is a classification made by f when f is applied to f. Note that f is equal to the probability that f is a classification model of f in the probability that f is a classification of f in the probability that f is a classification of f in the probability that f is a classification of f in the probability that f is a classification of f in the probability that f is a classification of f in the probability that f is a classification of f in the probability that f is a classification of f in the probability that f is a classification of f in the probability of f in the probability of f is a classification of f in the probability of f in the probability of f in the probability of f is an explicit f in the probability of f in the p

- 1. Generate a new dataset \mathbf{X}^* of size m by p using the observed values in \mathbf{X} . There are various ways to simulate the new dataset, and the methods currently used by the LIME R package will be described in more detail in Section 2.2.
- 2. Apply f to \mathbf{X}^* to obtain a vector of predictions $\hat{\mathbf{y}}'$ of length m. Let the prediction made by f when applied to the transformed case of interested be denoted as $f(x') = \hat{y}'$.
- 3. Apply a transformation T to \mathbf{X}^* that results in an interpretable representation. The transformation that is applied will depend on the simulation method used. Section 2.2 will discuss the transformations used by the LIME R package. Let $T(\mathbf{X}^*) = \mathbf{X}'$. Furthermore, apply T to x to obtain T(x) = x'.
- 4. Compute a proximity measure between x and \mathbf{x}'_i for each $i \in {1, ..., m}$ denoted by $\pi_x(\mathbf{x}'_i)$.
- 5. Identify a class of potentially interpretable models such as linear models or decision trees. Denote this class of models by $G: \mathbb{R}^p \to [0,1]$. Section 2.2 describes the class of interpretable models used by the R package.
- 6. Perform feature selection...
- 7. Fit explainer model
- 8. Interpret

Somehow I need to mesh the procedure performed by the R package and the description of LIME described in the original paper. The place where I left off in the procedure above is where the two start to disagree. Here

²https://github.com/marcotcr/lime

³https://github.com/thomasp85/lime

are my old "notes" on the LIME algorithm based on the paper:

- G: class of potentially interpretable models (e.g. linear models, decision trees, rule lists) in our case, we will use ridge regression
- g: explanation model where $g: \{0,1\}^{p'} \to \mathbb{R}$ and $g \in G$ do we need $\{0,1\}^{p'}$?
- $\Omega(g)$: measure of complexity of g (e.g. depth of a tree, number of non-zero coefficients in a linear model fit using LASSO)
- $\mathcal{L}(f, g, \Pi_x)$: the fidelity functions which is a measure of how unfaithful g is in approximating f in the locality defined by π_x
- $\xi(x)$: explanation produced by LIME where

$$\xi(x) = \underset{g \in G}{\arg\min} \ \mathcal{L}(f, g, \Pi_x) + \Omega(g)$$

(i.e. want to minimize $\mathcal{L}(f,g,\Pi_x)$ and keep $\Omega(g)$ low enough to be interpretable by humans)

XX references:)

2.2 | Implementation of LIME in R

make sure to include an output figure of a LIME explanation here

3 | METHODS

- 3.1 | Diagnostic Tool 1
- 3.2 | Diagnostic Tool 2
- 3.3 | etc...
- 4 | APPLICATION
- 4.1 | Bullet Matching Data
- 5 | DISCUSSION

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This is acknowledgment text. Provide text here.

Author contributions

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Financial disclosure

None reported.

Conflict of interest

The authors declare no potential conflict of interests.

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APPENDIX

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AUTHOR BIOGRAPHY

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