

Explaining Complex Machine Learning Models with LIME

Dr. Shirin Glander

shirin.glander@codecentric.de

2017-12-11

TwiML & AI

thunlai.com

Talk #7 Carlos Guestrin - Explaining the Predictions of Machine Learning

Marco Ribeiro,
Sameer Singh &
Carlos Guestrin:

"Why should I trust you?
Explaining the predictions of any classifier."
CORR 2016



models

LIME

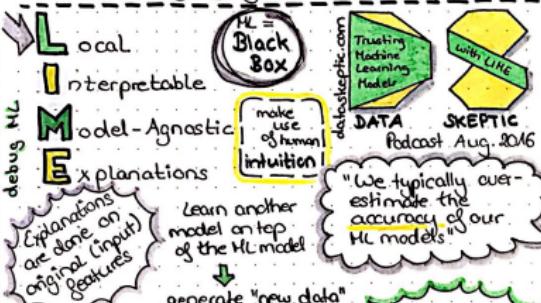
human interpretable explanations

- for neural networks, random forests, boosted decision trees, etc.



github.com/marcotcr/lime Python

recreated for R: github.com/thomasp85/lime



generate "new data" by permutation of the input data (test set)

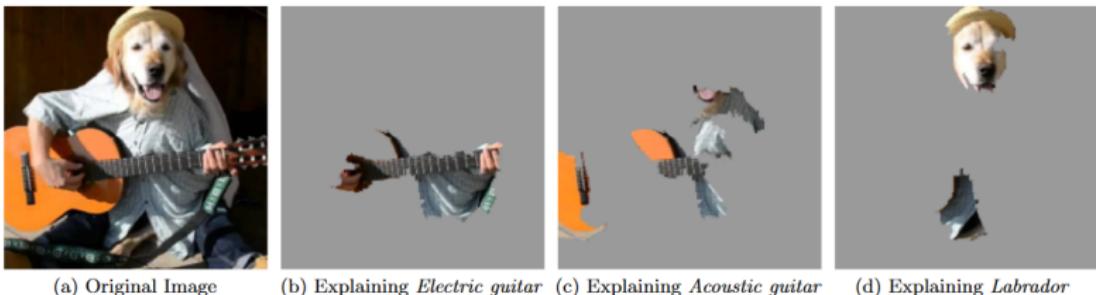
predictions on this new data with simple model. → Local fit (approximation)

more weight is being put on data that is similar to the original data



LIME can explain any classifier

- image recognition



(a) Original Image

(b) Explaining *Electric guitar*

(c) Explaining *Acoustic guitar*

(d) Explaining *Labrador*

Figure 4: Explaining an image classification prediction made by Google's Inception neural network. The top 3 classes predicted are "Electric Guitar" ($p = 0.32$), "Acoustic guitar" ($p = 0.24$) and "Labrador" ($p = 0.21$)

Ribeiro, Singh, and Guestrin (2016)



LIME can explain any classifier

- text classification

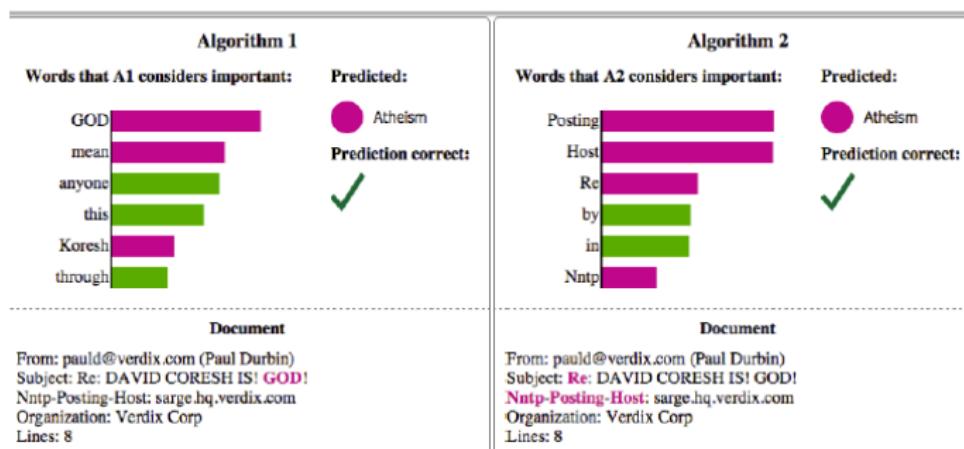
Example #3 of 6

True Class: Atheism

[Instructions](#)

[Previous](#)

[Next](#)



Ribeiro, Singh, and Guestrin (2016)



How LIME works

1. Permutation of each test case to explain
2. Complex model predicts all permuted test cases
3. Distance between permutations and original text case is calculated and converted to similarity scores
4. Subsetting features with highest importance in complex model for each permuted test case
5. Fitting a linear model with the subsetted features to the permuted data (weights represent similarity score)
6. Using simple model to explain test case prediction



An example in R

- Data: Chronic Kidney Disease
(http://archive.ics.uci.edu/ml/datasets/Chronic_Kidney_Disease)
- Nonparametric Missing Value Imputation using Random Forest
(library(missForest))
- Categorical features converted to dummy variables
(library(dummies))
- Scaled and centered

Predictor: ckd or notckd (class)

- Random Forest model with library(caret) (5x10 repeated CV)



The model

```
## Random Forest  
##  
## 360 samples  
## 48 predictor  
## 2 classes: 'ckd', 'notckd'  
##  
## No pre-processing  
## Resampling: Cross-Validated (10 fold, repeated 5 times)  
## Summary of sample sizes: 324, 324, 324, 324, 325, 324, ...  
## Resampling results across tuning parameters:  
##  
## mtry Accuracy Kappa  
## 2 0.9922647 0.9838466  
## 25 0.9917392 0.9826070  
## 48 0.9872930 0.9729881  
##  
## Accuracy was used to select the optimal model using the largest value.  
## The final value used for the model was mtry = 2.
```



Predictions

```
## Confusion Matrix and Statistics  
##  
##      Reference  
## Prediction ckd notckd  
##   ckd  23   2  
##   notckd 0  15  
##  
##          Accuracy : 0.95  
##          95% CI : (0.8308, 0.9939)  
##  No Information Rate : 0.575  
##  P-Value [Acc > NIR] : 1.113e-07  
##  
##          Kappa : 0.8961  
##  Mcnemar's Test P-Value : 0.4795  
##  
##          Sensitivity : 1.0000  
##          Specificity : 0.8824  
##          Pos Pred Value : 0.9200  
##          Neg Pred Value : 1.0000
```



Explaining the predictions

Explanation function:

- train_x is the training data
- model_rf is the complex model
- n_bins = 10 groups continuous variables into 10 bins
- quantile_bins = TRUE bases bins on quantiles (bins are not evenly spread across data range)
- dist_fun = "euclidean" sets distance function to calculate weights

```
library(lime)
explainer <- lime(train_x,
                    model_rf,
                    n_bins = 10,
                    quantile_bins = TRUE,
                    dist_fun = "euclidean")
```



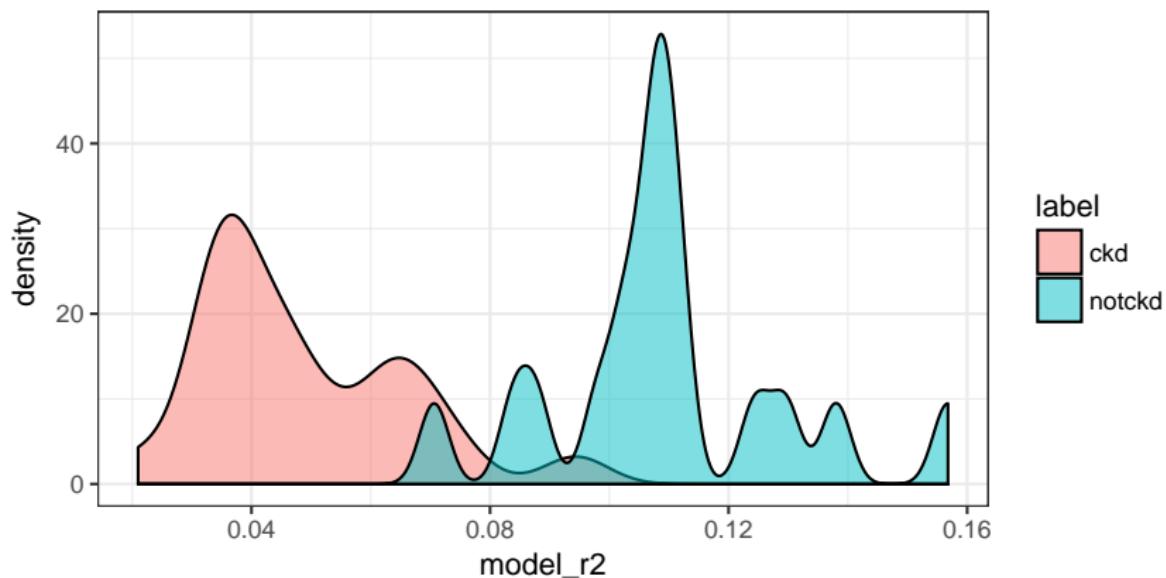
Explaining the predictions

- n_labels = 1 because we want to explain the most likely class predicted
 - n_features = 8 returns top 8 most important features for each test case
 - n_permutations = 1000 permutes test case 1000x
 - feature_select = "highest_weights":fits a ridge regression and selects the top features with highest absolute weight



Explanation quality

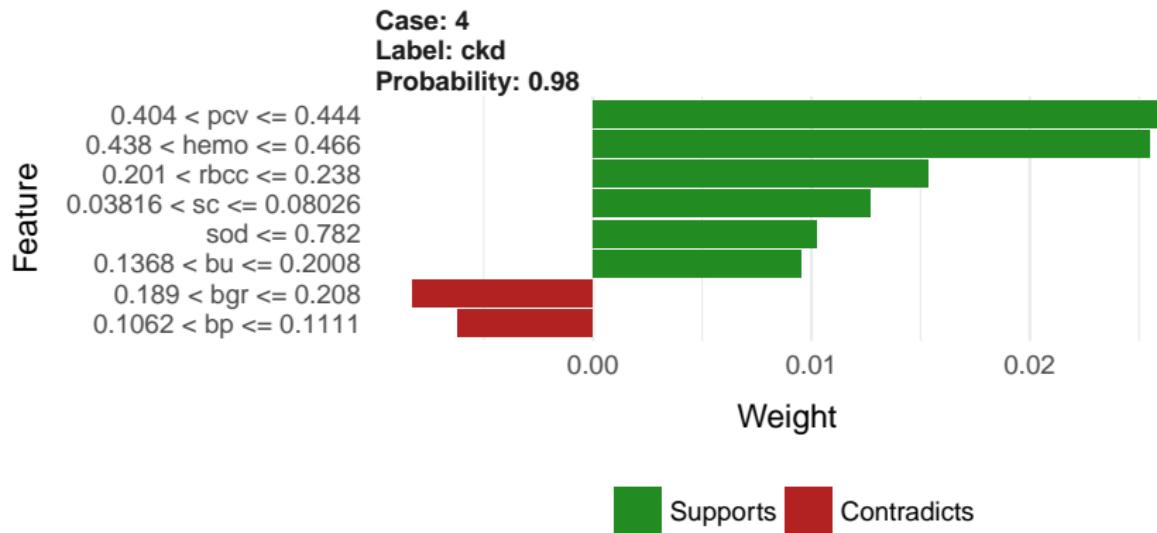
- model r^2





Plotting the explanations

```
plot_features(explanation_df[1:8, ])
```





Plotting the predictions

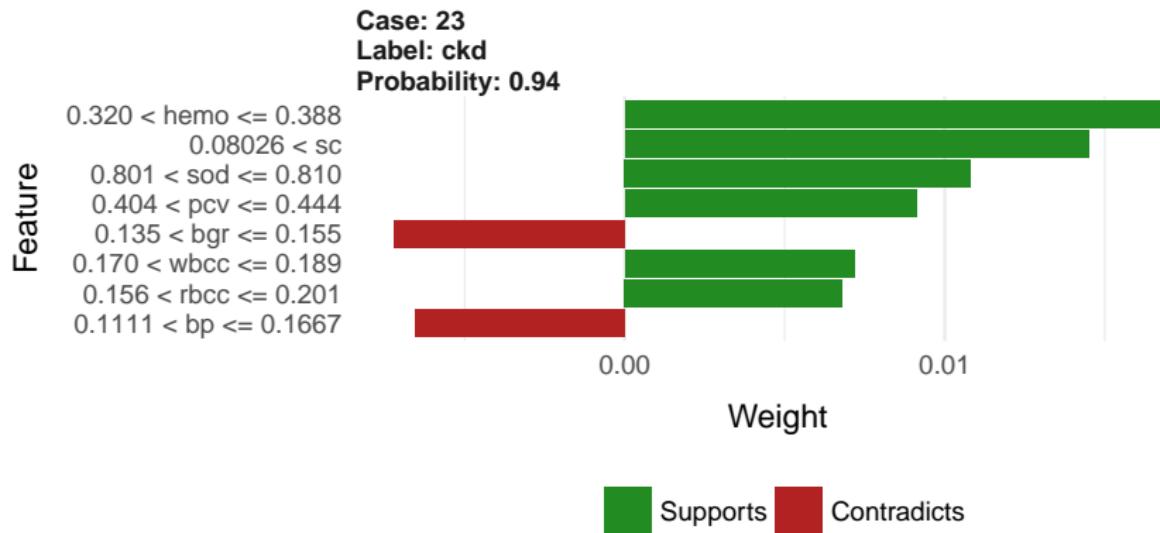
```
plot_features(explanation_df[9:16, ])
```





Plotting the predictions

```
plot_features(explanation_df[17:24, ])
```

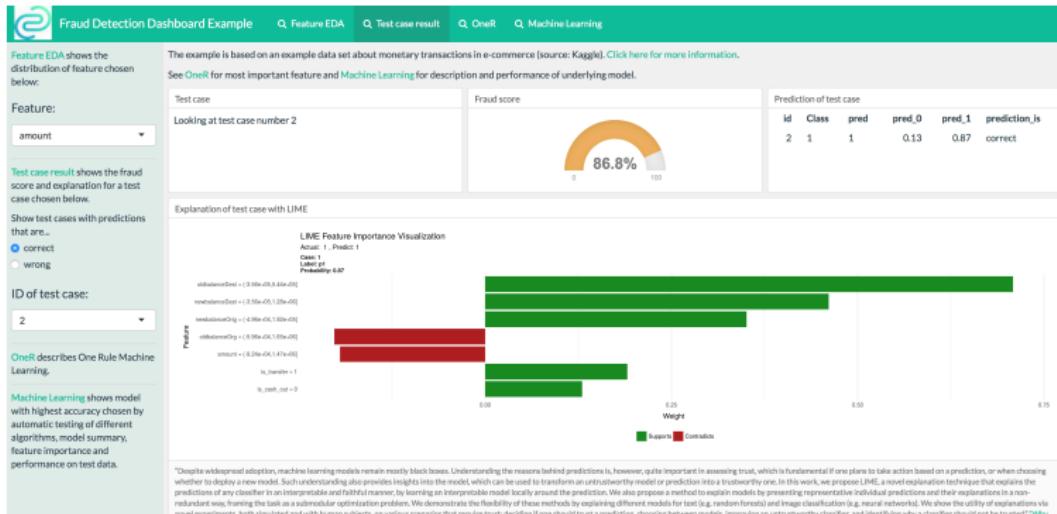




LIME in action

- Explaining fraud predictions:

https://shiring.shinyapps.io/fraud_example_dashboard/





More about LIME

Publication

- Ribeiro, Singh, and Guestrin (2016)

Contribute

- <https://github.com/marcotcr/lime>
- <https://github.com/thomasp85/lime>



Thank you!

...and stay connected...

You can find me on

- my blog: www.shirin-glander.de
- Twitter: <https://twitter.com/ShirinGlander>
- Github: <https://github.com/ShirinG>

Code and slides will go up on my blog!

MünsteR User group

- <https://www.meetup.com/Munster-R-Users-Group>

Ribeiro, Marco Túlio, Sameer Singh, and Carlos Guestrin. 2016. "Why Should I Trust You?': Explaining the Predictions of Any Classifier." CoRR abs/1602.04938. <http://arxiv.org/abs/1602.04938>.