

Efficient and simple prediction explanations with *groupShapley*

A practical perspective

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Prediction explanation

- ► Assume a model $f(x) \in \mathbb{R}$ that predicts some unknown outcome based on a set of features $x = (x_1, ..., x_M)$
- We apply the predictive model for a specific input $x = x^*$, reaching a certain prediction $f(x^*)$
- Individual prediction explanation
 - Want to understand how the different features, or types of features affect this specific prediction value $f(x^*)$
 - I.e. explain the predicted outcome in terms of the input $x = x^*$ (local explanation)

Frameworks…

- LIME
- Anchors

- Counterfactual explanations
- Explanation Vectors

- PredDiff
- Shapley values

Shapley values

- General concept
 - Stems from cooperative game theory (Shapley, 1953)
 - Used to distribute the total payoff to the players
 - Explicit formula for the "fair" payment to every player *j*:

$$\phi_j = \sum_{\text{all } S \text{ without } j} w(|S|) \left(v(S \cup \{j\}) - v(S) \right), w \text{ is a certain weight function,} \\ v(S) \text{ is the payoff with only players in subset } S$$

- Several mathematical optimality properties
- For prediction explanation
 - Players = features $(x_1, ..., x_M)$
 - Payoff = prediction outcome $(f(x^*))$
 - Contribution function: $v(S) = E[f(x)|x_S = x_S^*]$
 - Rough interpretation of ϕ_j
 - The prediction change caused by observing x_i



Bottlenecks



 $egin{aligned} M &= 5 \Rightarrow 2^M = 32 \ M &= 10 \Rightarrow 2^M = 1024 \ M &= 20 \Rightarrow 2^M = 1048676 \ M &= 40 \Rightarrow 2^M > 10^{12} \ M &= 100 \Rightarrow 2^M > 10^{30} \ M &= 1000 \Rightarrow 2^M > 10^{301} \end{aligned}$

1. The sum in the Shapley value formula is of size 2^{M} , growing exponentially in the number of features

2. How can we visualize, interpret and extract knowledge from 100s or 1000s of Shapley values?



- Typically: the sum of many small ϕ_i > sum of the few large ones
- Many highly dependent features complicates the interpretation

groupShapley

- Fundamentally very simple approach
 - Divide the *M* features into a small number of *G* disjoint groups $\{G_1, \dots, G_G\}$.
 - Replace the feature subsets *S* in the Shapley formula by group subsets *T*:

$$\phi_{G_i} = \sum_{\text{all } T \text{ without } G_i} w(|T|) \left(v(T \cup G_i) - v(T) \right)$$

• The scores are still Shapley values, so all mathematical properties are kept (on group level)

- What about the bottlenecks?
 - $2^G \ll 2^M \Rightarrow$ computationally tractable
 - G small \Rightarrow easy to visualize





How to group the features?

- Crucial to group features based on the desired explanation
- Grouping based on feature dependence
 - Highly dependent features grouped together, using e.g. a clustering method.
 - Easier to study theoretically
 - Often difficult to extract knowledge from in practice
- Grouping based on application/feature knowledge
 - Group features of similar type or general category
 - Gives directly meaningful interpretations of computed groupShapley values
 - May perform multiple explanations with different groupings for increased understanding
- We advocate grouping based on feature knowledge in practical applications

Practical example 1: Car insurance

- US Car insurance dataset
 - 10 302 customers with records of crash/no crash + 21 features
 - Fit a random forest model with 500 trees to predict crash based on the 21 features



- 3 feature groups based on type
 - Track record (4 features): # claims last 5 years, # licence record points, previous licence revokes, time as customer
 - Personal information (13 features): age of driver, education level, # children, job type, # driving children, marital status, gender, distance to work +++
 - Car information (4 features) value of car, age of car, type of car, whether car is red

Practical example 1: Car insurance

- ► We apply the model to 3 individuals
- 1 claim last 5 years, 3 licence record points. Single mother of 4 (2 driving). Driving a SUV, 27 miles to work.
- 2. Got licence revoked and 10 licence record points. 37 year old father of 2 (1 driving).
- 3 claims last 5 years, no licence record points
 60 year old married doctor with no children, with a PhD Red sports car.



Practical example 2: Gene data

- Disease classification with high dimensional gene data
 - 127 patients where 85 are diseased with either Crohn's disease (CD) or Ulcerative colitis (UC) + 42 healthy controls.
 - 4 834 genes (after pre-processing)
 - Using 100 random individuals, we fit a Lasso penalized linear regression model to predict P(diseased with either CD or UC) based on the patient's genes



Feature groups

- Use the so-called Hallmark gene set to group the features (genes) into 23 different groups commonly used in gene set enrichment analysis
- The Hallmark gene set "conveys a specific biological state or process" (Liberzon et al., 2015)

Practical example 2: Gene data

- Compute groupShapley values for the remaining 27 patients
- Make separate groupShapley boxplots for UC, CD and controls
- Can we identify genetical similarities and differences for UC and CD?
- Note: Model not trained to separate UC and CD



Concluding remarks

Other use cases

- Classification with time series data (see paper)
- Explain original categorical features by grouping one-hotencoded features
- Explain image classification by grouping pixels into superpixels
- Explain models with large number of feature-engineered variables based on original base features

Implementation

- Easy to apply in practice with the shapr R-package*
- Code snippet for Car insurance example

```
2 • #### "model" are pre-defined ####
4 library(shapr)
5 - #### 1 Define aroups ####
  group_Names = list(Personal_Info = c("AGE", "EDUCATION", "HOMEKIDS", "HOME_VAL",
                                         "OCCUPATION", "TRAVTIME", "KIDSDRIV", "MSTATUS",
                                         "PARENT1", "GENDER", "URBANICITY", "YOJ"),
                      Car_Info = c("BLUEBOOK", "CAR_AGE", "CAR_TYPE", "RED_CAR"),
                      Track Record = c("CLM FREQ", "MVR PTS", "REVOKED", "TIF"))
  explainer_group <- shapr::shapr(x_train,</pre>
                                    model,
                                    group = group Names)
   # Compute groupShapley values
  explanation group <- shapr::explain(x_test,</pre>
                                        approach = "ctree",
                                        explainer = explainer_group,
                                        prediction_zero = mean(x_train$CLAIM_FLAG))
   plot(explanation_group)
```

*groupShapley is currently only available in the GitHub version of *shapr*. <u>https://github.com/NorskRegnesentral/shapr</u> Will be included in the next CRAN release