Example of how to use XGBoost optimized with SERA

Use case

In this markdown two use cases are provided where three variants are taken into consideration:

- XGBoost with SERA optimization where weights are obtained automatically;
- XGBoost with SERA optimization where weights are obtained with domain knowledge;
- XGBoost with Squared Error as objective.

The first use case uses a number of intervals I = 1000, while the second I = 10.

Let us start.

Load required packages

```
library(ModelOptimizationIR)
library(xgboost)
library(dplyr)
library(IRon)
library(scam)
```

Load data and perform random partition

```
data("NO2Emissions")
n <- nrow(NO2Emissions)
s <- sample(1:n, size = n*0.8)

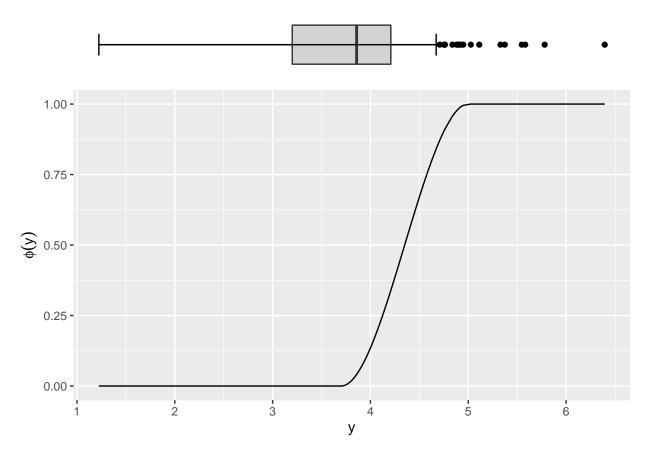
formula <- LNO2 ~ .
train <- NO2Emissions %>% dplyr::slice(s)
test <- NO2Emissions %>% dplyr::slice(-s)
```

Extract weights using domain knowledge

```
y <- train$LN02
control.points <- matrix(c(1.1, 0, 0, 3.7, 0, 0, 5, 1, 0), byrow = TRUE, ncol=3) # Extreme points above
ph.ctrl <- phi.control(y = y, method="range", control.pts = control.points)
phi <- phi(y = y, phi.parms = ph.ctrl)</pre>
```

Plot with the relevance function

```
phiPlot(ds=y, phi.parms = ph.ctrl)
```



Add some hyper-parameters for modeling

```
params <- list(max_depth=7, eta=10^{-1}, gamma=10^{-2})
```

First use case I = 1000

Define steps to discretize SERA and calculate weights

```
I = 1000 # Default value
steps <- seq(0, 1, 1/I)
sigma <- sigma(phis = phi, steps = steps)</pre>
```

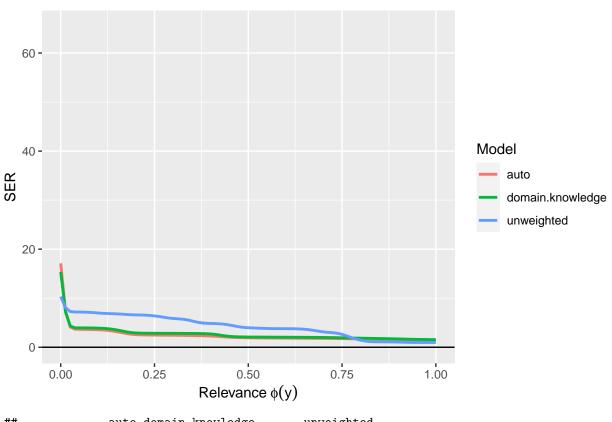
Train the models and save results

)

Evaluate SERA

```
sera(trues=df$trues, preds=dplyr::select(df, -trues), ph = ph.ctrl, pl=TRUE)
```

SERA



```
## auto domain.knowledge unweighted
## 2.331837 2.563095 4.234552
```

Evaluate MSE

Second use case I = 10

Define steps to discretize SERA and calculate weights

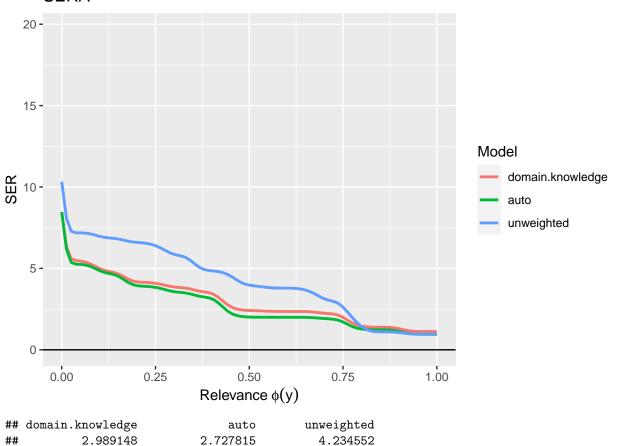
```
I = 10 # Default value
steps <- seq(0, 1, 1/I)
sigma <- sigma(phis = phi, steps = steps)</pre>
```

Train the models and save results

Evaluate SERA

```
sera(trues=df$trues, preds=dplyr::select(df, -trues), ph = ph.ctrl, pl=TRUE)
```

SERA



Evaluate MSE

```
df %>%
summarise(across(!matches("trues"), ~mean((trues - .x)^2)))
```

Better generalization was achieved, but at the cost of worst predictive focus on extreme values.