Pipelines

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Materials from Modeling in R and Python

- tidymodels: meta-package for 'tidy' modeling in R
- scikit-learn: modeling in Python

parsnip

- parsnip package (CART \rightarrow caret \rightarrow parsnip)
- unify modeling interfaces (lm, glmnet, randomForest, etc etc etc)
- model (algorithm), mode (classification/regression), engine (implementation/package)

rsample

- resampling, cross-validation, bootstrapping, holdout sets ...
- $train/test split (initial_split()/training()/testing())$
- cross-validation (vfold_cv()), bootstrap (bootstrap)
- blocked/grouped methods! group_vfold_cv, group_bootstraps()

recipes

- feature engineering
- preprocessing (centering/scaling, imputation, dimension reduction, etc.)

more

- workflows: bundle preprocessing/modeling/post-processing
- tune: hyperparameter tuning
- yardstick: assessment

example

```
library(tidyverse)
library(tidymodels)
historical <- (read_csv("../code/historical_baseball.csv")</pre>
    |> mutate(across(inducted, ~fct_rev(factor(.))))
    |> filter(ab > 250)
data_split <- initial_split(historical, prop = 2/3, strata = inducted)</pre>
train_data <- training(data_split)</pre>
testing_data <- testing(data_split)</pre>
b_recipe <- (
    recipe(inducted ~ ., data = train_data)
    |> step_rm("last_year")
    |> update_role(player_id, new_role = "ID") ## not predictor or outcome
    |> step_center(all_numeric())
    |> step_scale(all_numeric())
    |> step_nzv(all_numeric())
    |> prep()
)
data(iris)
svm.model <- e1071::svm(Species ~ ., data = iris, probability = TRUE)</pre>
```

```
pred <- predict(svm.model, iris, probability = TRUE)</pre>
  prob <- attr(pred, "probabilities")</pre>
  spmat <- matrix(0, ncol = length(levels(iris$Species)),</pre>
                   nrow = length(iris$Species))
  spmat[cbind(1:nrow(prob), as.numeric(iris$Species))] <- 1</pre>
  -1 * mean(sapply(1:nrow(prob),
                    \(i) dmultinom(x = spmat[i,], size = 1, prob[i,], log = TRUE)))
[1] 0.07936639
  yardstick::mn_log_loss_vec(truth = iris$Species, estimate = prob)
[1] 0.07936639
  b_recipe |> bake(train_data) |> rsample::vfold_cv()
# 10-fold cross-validation
# A tibble: 10 x 2
   splits
                       id
   t>
                       <chr>
1 <split [1598/178] > Fold01
 2 <split [1598/178] > Fold02
 3 <split [1598/178] > Fold03
4 <split [1598/178] > Fold04
 5 <split [1598/178] > Fold05
 6 <split [1598/178] > Fold06
7 <split [1599/177] > Fold07
8 <split [1599/177] > Fold08
 9 <split [1599/177] > Fold09
10 <split [1599/177] > Fold10
  lr_mod <-</pre>
      logistic_reg(mode = "classification", penalty = tune(), mixture = 1) %>%
      set_engine(engine = "glmnet")
```

```
tt <- tune_grid(
    object = lr_mod,
    preprocessor = b_recipe,
    resamples = vfold_cv(train_data),
    metrics = metric_set(mn_log_loss)
)</pre>
```