

# Tidymodels

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Math 243: Stat Learning

November 22nd, 2021

## Outline

In today's class, we will...

- Discuss the `tidymodels` packages for model building in the `tidyverse` framework

## Section 1

### Intro to tidymodels

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glm	stats	<code>predict(object, type = "response")</code>
gbm	gbm	<code>predict(object, type = "response", n.trees)</code>
rpart	rpart	<code>predict(object, type = "prob")</code>
kknn	kknn	<code>kknn(...)\$prob</code>

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- Each method has significantly different methods for making class probability predictions
- Additionally, each model takes in different types of data arguments (vectors, model matrices, data frames, model formulas)

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Additionally, `tidymodels` fits in the broader `tidyverse` framework:

- Packages and functions should be accessible and easily interpreted
- Outputs should be data frames (or tibbles) whenever possible
- Functions should be compatible with the `%>%` operator and functional programming
- Model objects should be compatible with `ggplot2`

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- Model objects should be compatible with `ggplot2`

`tidymodels` takes the mechanics from each individual model package (`mass`, `tree`, `glm` etc.) and unifies the input and output

# The tidymodels framework

- ① Preprocess data using the `recipes` package
- ② Create training-test data splits using the `rsample` package
- ③ Give a model a functional form and specify fitting method using the `parsnip` package
- ④ Fit the model, tidy the results, and make predictions using the `fit`, `tidy`, and `predict` functions
- ⑤ Estimate model performance using cross-validation from the `rsample` package
- ⑥ Tune model parameters by adding model specifications

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We'll investigate each of these in-depth (although slightly out of order)

## Section 2

### Build a Model

# The Data

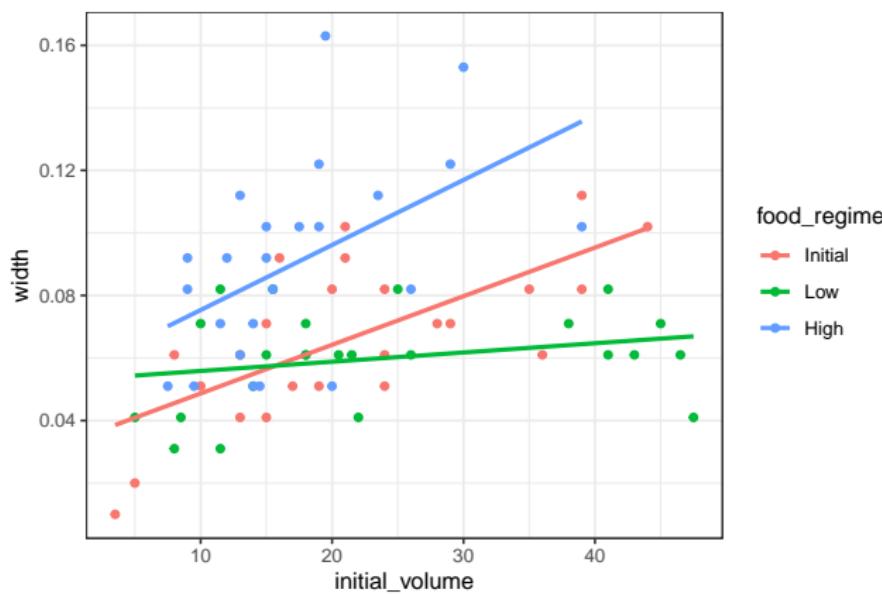
The sea\_urchins data set explores the relationship between feeding regimes and size of sea urchins over time:

```
sea_urchins<-read_csv("https://tidymodels.org/start/models/urchins.csv") %>%  
  setNames(c("food_regime", "initial_volume", "width")) %>%  
  mutate(food_regime = factor(food_regime, levels = c("Initial", "Low", "High")))  
head(sea_urchins)
```

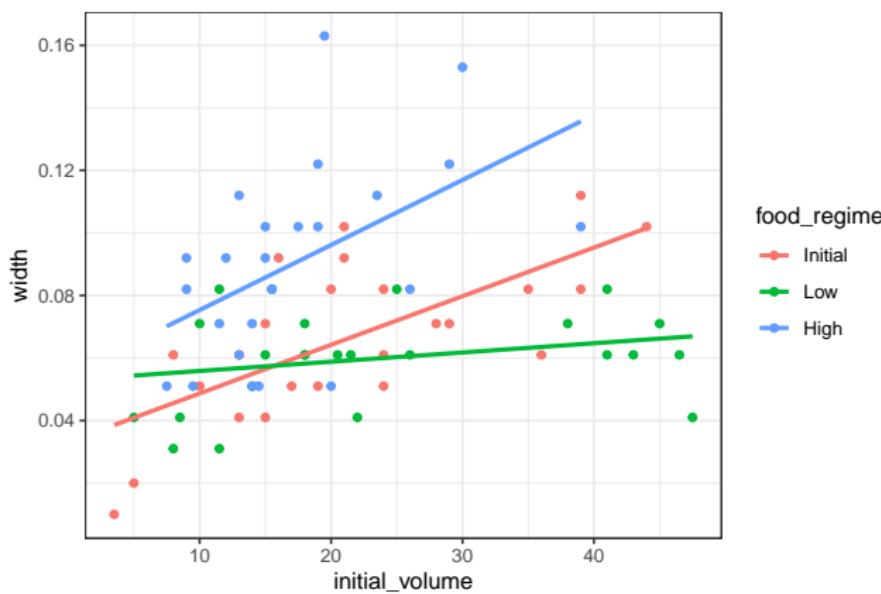
  

```
## # A tibble: 6 x 3  
##   food_regime initial_volume width  
##   <fct>          <dbl>  <dbl>  
## 1 Initial         3.5    0.01  
## 2 Initial         5      0.02  
## 3 Initial         8      0.061  
## 4 Initial        10     0.051  
## 5 Initial        13     0.041  
## 6 Initial        13     0.061
```

# Scatterplot

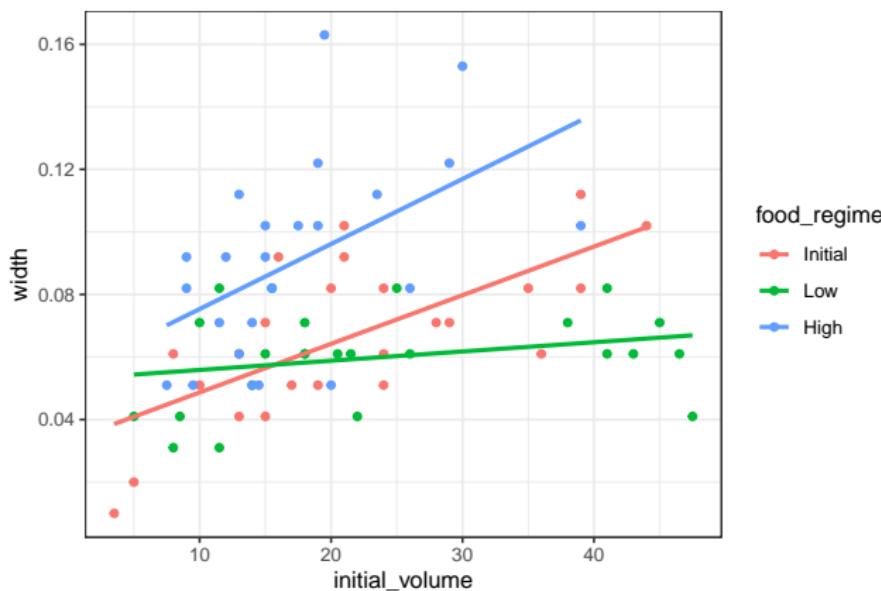


## Scatterplot



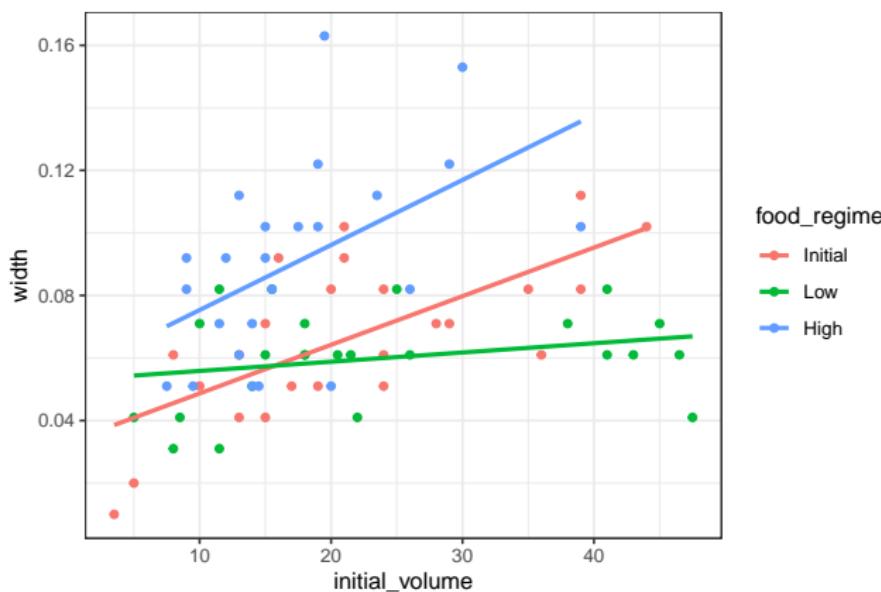
- Goal: Predict width as a function of `food_regime` and `initial_volume`.

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  - Does an additive model seem appropriate?

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- Goal: Predict `width` as a function of `food_regime` and `initial_volume`.
  - Does an additive model seem appropriate?
  - One option might be a linear model with interaction terms.

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Our model formula takes the form:

```
width ~ initial_volume + food_regime + initial_volume:food_regime
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```
library(parsnip)
linear_reg() %>%
  set_engine("lm")
```

```
## Linear Regression Model Specification (regression)
##
## Computational engine: lm
```

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## Linear Regression Model Specification (regression)
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- Other engines are possible for `linear_reg()`: `glmnet`, `stan`, and more

Now we create the model based on data using the `fit` function:

```
lm_mod<-linear_reg() %>%
  set_engine("lm")

lm_fit<- lm_mod %>%
  fit(width ~ initial_volume*food_regime, data = sea_urchins)
```

# Results

The output of our lm\_fit object:

```
lm_fit
```

```
## parsnip model object
##
## Fit time:  0ms
##
## Call:
## stats::lm(formula = width ~ initial_volume * food_regime, data = data)
##
## Coefficients:
##             (Intercept)          initial_volume
##                   0.0331216           0.0015546
##             food_regimeLow        food_regimeHigh
##                   0.0197824           0.0214111
##   initial_volume:food_regimeLow  initial_volume:food_regimeHigh
##                   -0.0012594           0.0005254
```

## Summary Table

To get the traditional **summary table**:

```
tidy(lm_fit) %>% kable()
```

term	estimate	std.error	statistic	p.value
(Intercept)	0.0331216	0.0096186	3.4434873	0.0010020
initial_volume	0.0015546	0.0003978	3.9077643	0.0002220
food_regimeLow	0.0197824	0.0129883	1.5230864	0.1325145
food_regimeHigh	0.0214111	0.0145318	1.4733993	0.1453970
initial_volume:food_regimeLow	-0.0012594	0.0005102	-2.4685525	0.0161638
initial_volume:food_regimeHigh	0.0005254	0.0007020	0.7484702	0.4568356

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Note that the output is a data frame with standard column names

## New Data

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- First, we generate data:

```
new_urchins <- expand.grid(initial_volume = c(5,30),  
                           food_regime = c("Initial", "Low", "High"))  
new_urchins %>% kable()
```

initial_volume	food_regime
5	Initial
30	Initial
5	Low
30	Low
5	High
30	High

# Make predictions

Then we make predictions

```
new_preds <- predict(lm_fit, new_data = new_urchins)
conf_int_preds<-predict(lm_fit, new_data = new_urchins, type = "conf_int")
new_preds %>% kable()
```

.pred
0.0408948
0.0797608
0.0543803
0.0617621
0.0649329
0.1169338

```
conf_int_preds %>% kable()
```

.pred_lower	.pred_upper
0.0251382	0.0566514
0.0688612	0.0906605
0.0396403	0.0691204
0.0522641	0.0712601
0.0483265	0.0815393
0.0999144	0.1339532

## Combining Data and Predictions

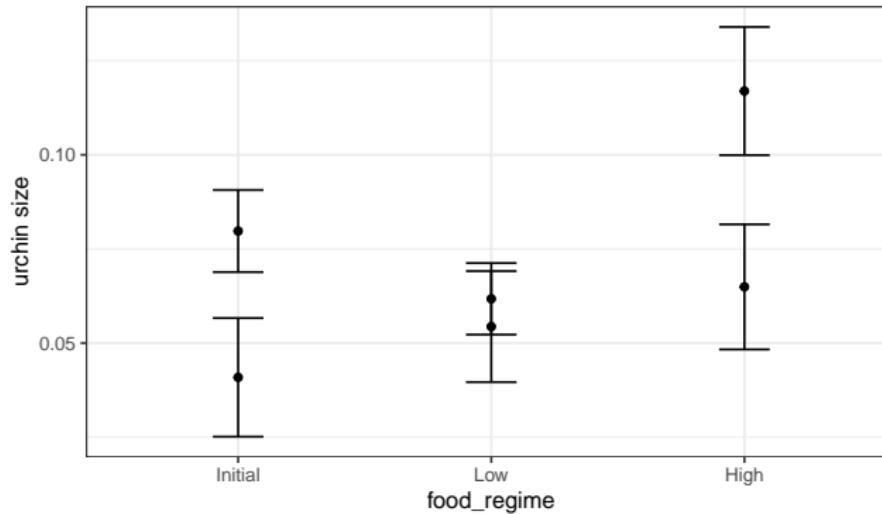
Because the result of `predict()` is tidy, we can easily combine it with the original data:

```
combined_data <- new_urchins %>% cbind(new_preds) %>% cbind(conf_int_preds)  
combined_data %>% kable()
```

initial_volume	food_regime	.pred	.pred_lower	.pred_upper
5	Initial	0.0408948	0.0251382	0.0566514
30	Initial	0.0797608	0.0688612	0.0906605
5	Low	0.0543803	0.0396403	0.0691204
30	Low	0.0617621	0.0522641	0.0712601
5	High	0.0649329	0.0483265	0.0815393
30	High	0.1169338	0.0999144	0.1339532

## Predictions Plot

```
ggplot(combined_data, aes(x = food_regime)) +  
  geom_point(aes(y = .pred)) +  
  geom_errorbar(aes(ymin = .pred_lower, ymax = .pred_upper), width = .2) +  
  labs(y = "urchin size") + theme_bw()
```



# Using a different engine

## LASSO?

- With only 3 predictors (`food_regime`, `initial_width` and the interaction term), its unlikely our model will be improved by Penalized Regression. But let's try anyway:

```
glmnet_mod <- linear_reg(penalty = 0.01, mixture = 1) %>% set_engine("glmnet")
```

- `mixture = 1` indicates LASSO (`mixture = 0` is used for Ridge Regression)
- `glmnet` requires us to indicate a value of penalty parameter  $\lambda$  to make predictions.
  - Here, we choose `penalty = 0.01` somewhat arbitrarily (we'll tune later); in any case, `glmnet` will still create models for all  $\lambda$

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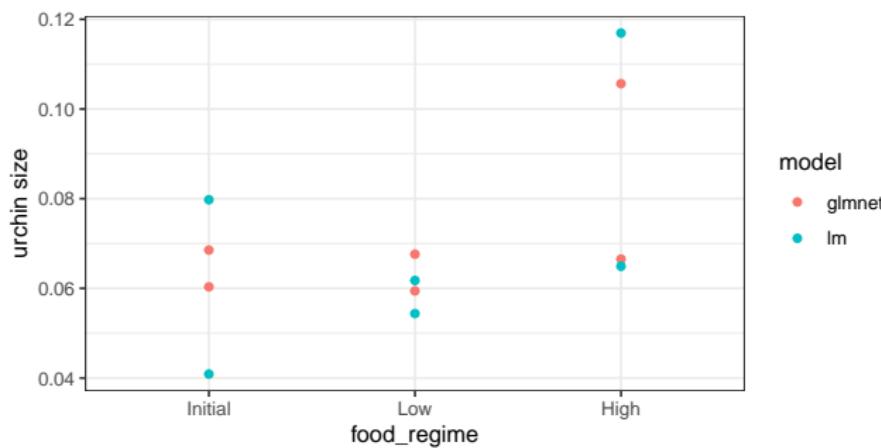
```
glmnet_fit <- glmnet_mod %>% fit(width ~ initial_volume*food_regime, data = sea_urchins)
tidy(glmnet_fit, penalty = .004) #penalty selects particular value of lambda
```

```
## # A tibble: 6 x 3
##   term                  estimate  penalty
##   <chr>                 <dbl>    <dbl>
## 1 (Intercept)            0.0587   0.004
## 2 initial_volume         0.000328  0.004
## 3 food_regimeLow        -0.000918  0.004
## 4 food_regimeHigh        0          0.004
## 5 initial_volume:food_regimeLow  0          0.004
## 6 initial_volume:food_regimeHigh  0.00124   0.004
```

## Results from glmnet

```
new_glmnet_preds <- predict(glmnet_fit, new_data = new_urchins, penalty = 0.004)
combined_glmnet_data <- new_urchins %>% cbind(new_glmnet_preds)
two_models <- rbind(combined_glmnet_data,
                      combined_data %>% select(-.pred_lower, -.pred_upper )) %>%
  mutate(model = rep(c("glmnet", "lm"), each = 6))

ggplot(two_models, aes(x = food_regime)) +
  geom_point(aes(y = .pred, color = model) ) +
  labs(y = "urchin size") + theme_bw()
```



## Section 3

### Preprocessing with recipes

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  - Transforms data to be on a different scale
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- The `recipes` package assists with preprocessing before a model is trained
  - Converts qualitative predictors to dummy variables
  - Transforms data to be on a different scale
  - Transforms several predictors at the same time
  - Extracts features from variable
- The main advance of `recipes` is that it allows us combine several steps at once, in a reproducible fashion

# House Prices

- The house data contains information on 30 predictors for 200 houses in Ames, Iowa  
names(house)

```
## [1] "SalePrice"      "Id"           "Functional"     "BldgType"  
## [5] "Foundation"    "LotShape"       "LandSlope"      "SaleCondition"  
## [9] "RoofMatl"       "ScreenPorch"    "MSSubClass"     "GarageCars"  
## [13] "BedroomAbvGr"   "TotalBsmtSF"   "LotArea"        "OpenPorchSF"  
## [17] "BsmtFullBath"   "WoodDeckSF"    "OverallCond"    "YrSold"  
## [21] "GrLivArea"      "MoSold"        "TotRmsAbvGrd"   "PoolArea"  
## [25] "YearBuilt"       "GarageArea"    "OverallQual"    "Fireplaces"  
## [29] "EnclosedPorch"  "FullBath"      "HalfBath"
```

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## [25] "YearBuilt"       "GarageArea"    "OverallQual"    "Fireplaces"  
## [29] "EnclosedPorch"   "FullBath"      "HalfBath"
```

- Note that the variable Id is not useful as a predictor, but is useful for referring to houses in the data set.

## Investigate Predictors

- Additionally, note that several of the variables are factors, so should be converted to a collection of dummy variables.

# Investigate Predictors

- Additionally, note that several of the variables are factors, so should be converted to a collection of dummy variables.
- Moreover, for a few variables, some levels are very underrepresented.

```
house %>% count(RoofMat1)
```

```
##   RoofMat1    n
## 1 CompShg 195
## 2 Membran    1
## 3 Tar&Grv    2
## 4 WdShake    1
## 5 WdShngl    1
```

# Data Splitting

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```
library(rsample)
set.seed(1221)
data_split <- initial_split(house , prop = 3/4)
train_data <- training(data_split)
test_data <- testing(data_split)
```

## Create a recipe and update roles

- We now create a recipe for some data pre-processing

```
library(recipes)
house_rec <-
  recipe(SalePrice ~ ., data = train_data) %>%
  update_role(Id, new_role = "ID")
```

## Create a recipe and update roles

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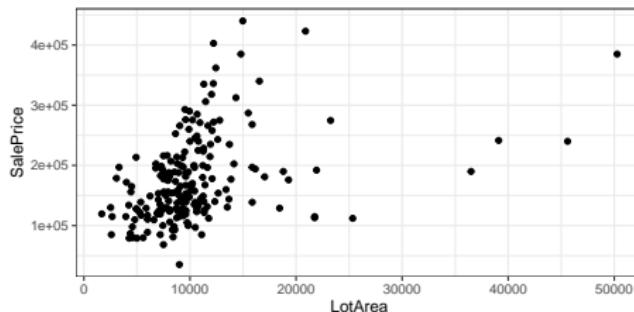
```
library(recipes)
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```

```
summary(house_rec)
```

```
## # A tibble: 31 x 4
##   variable     type    role    source
##   <chr>        <chr>   <chr>   <chr>
## 1 Id          numeric ID      original
## 2 Functional  nominal predictor original
## 3 BldgType    nominal predictor original
## 4 Foundation  nominal predictor original
## 5 LotShape    nominal predictor original
## 6 LandSlope   nominal predictor original
## 7 SaleCondition nominal predictor original
## 8 RoofMatl   nominal predictor original
## 9 ScreenPorch numeric predictor original
## 10 MSSubClass  numeric predictor original
## # ... with 21 more rows
```

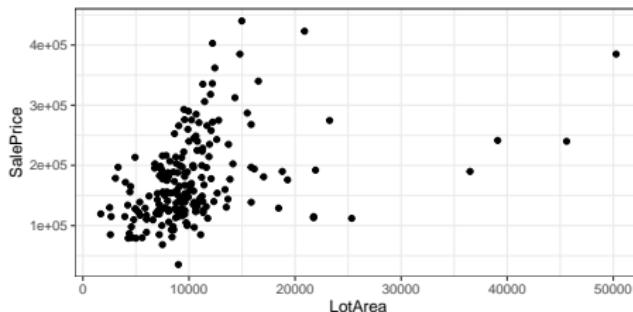
## Add steps to recipes

- Consider the relationship between sale price and lot area:

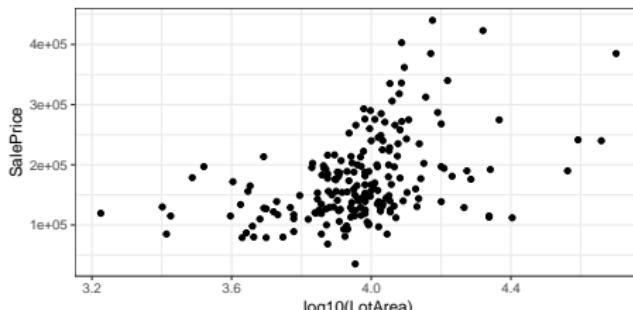


## Add steps to recipes

- Consider the relationship between sale price and lot area:



- Accuracy of a linear model may improve by performing log transformation on LotArea:



## Adding steps to recipes

- Let's update our recipe:

```
house_rec <- house_rec %>%
  step_log(LotArea, base = 10)
```

```
house_rec
```

```
## Recipe
##
## Inputs:
##
##       role #variables
##       ID             1
##     outcome            1
##   predictor           29
##
## Operations:
##
## Log transformation on LotArea
```

## Create New Variables from Old

- The original data set contains variables `FullBath` and `HalfBath`. But we want a measure of total number of baths:

$$\text{TotalBath} = \text{FullBath} + \frac{1}{2}\text{HalfBath}$$

## Create New Variables from Old

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$$\text{TotalBath} = \text{FullBath} + \frac{1}{2}\text{HalfBath}$$

- We can also add a `mutate` step in our recipe to do just this:

```
house_rec <- house_rec %>%
  step_mutate(TotalBath = FullBath+0.5*HalfBath) %>%
  step_rm(FullBath, HalfBath)
```

```
house_rec
```

```
## Recipe
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##   predictor            29
##
## Operations:
##
## Log transformation on LotArea
```

## Create Dummy Variables

- Recall that 7 of our variables are factors (Functional, BldgType, Foundation, LotShape, LandSlope, SaleCondition, RoofMatl). To create appropriate dummy variables:

```
house_rec <- house_rec %>% step_dummy(all_nominal(), -all_outcomes())  
house_rec
```

```
## Recipe  
##  
## Inputs:  
##  
##       role #variables  
##       ID      1  
##     outcome      1  
##   predictor      29  
##  
## Operations:  
##  
## Log transformation on LotArea  
## Variable mutation  
## Delete terms FullBath, HalfBath  
## Dummy variables from all_nominal(), -all_outcomes()
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- The first argument `all_nominal` selects all variables that are either factors or characters
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## Remove Problematic Predictors

- Finally, to avoid the situation where an infrequently occurring level doesn't exist in the training or test sets:

```
house_rec <- house_rec %>% step_zv(all_predictors())
house_rec
```

```
## Recipe
##
## Inputs:
##
##       role #variables
##           ID          1
##       outcome         1
##   predictor        29
##
## Operations:
##
## Log transformation on LotArea
## Variable mutation
## Delete terms FullBath, HalfBath
## Dummy variables from all_nominal(), -all_outcomes()
## Zero variance filter on all_predictors()
```

## Remove Problematic Predictors

- Finally, to avoid the situation where an infrequently occurring level doesn't exist in the training or test sets:

```
house_rec <- house_rec %>% step_zv(all_predictors())
house_rec
```

```
## Recipe
##
## Inputs:
##
##       role #variables
##           ID          1
##       outcome         1
##   predictor        29
##
## Operations:
##
## Log transformation on LotArea
## Variable mutation
## Delete terms FullBath, HalfBath
## Dummy variables from all_nominal(), -all_outcomes()
## Zero variance filter on all_predictors()
```

- The `step_zv` verb removes columns from the training data which have a single value

# Workflows

- Why create a recipe when we could just as easily perform the pre-processing steps using `dplyr`?

# Workflows

- Why create a recipe when we could just as easily perform the pre-processing steps using `dplyr`?
  - ① The recipe allows us to apply the same procedures to both test and training data.
  - ② The recipe gives instructions for processing the data **without actually performing that action**

# Workflows

- Why create a recipe when we could just as easily perform the pre-processing steps using `dplyr`?
- ① The recipe allows us to apply the same procedures to both test and training data.
  - ② The recipe gives instructions for processing the data **without actually performing that action**

To use our recipe across several steps, we will use a *workflow*, which will

- ① Process the recipe using the training set
- ② Apply the recipe to the training set
- ③ Apply the recipe to the test set

## Create the workflow

```
house_mod <- linear_reg() %>% set_engine("lm")

house_wf <- workflow() %>%
  add_model(house_mod) %>%
  add_recipe(house_rec)

house_wf

## == Workflow =====
## Preprocessor: Recipe
## Model: linear_reg()
##
## -- Preprocessor -----
## 5 Recipe Steps
##
## * step_log()
## * step_mutate()
## * step_rm()
## * step_dummy()
## * step_zv()
##
## -- Model -----
## Linear Regression Model Specification (regression)
##
## Computational engine: lm
```

# Fitting Models with Workflows

```
house_fit <- house_wflow %>% fit(data = train_data)

house_fit %>% pull_workflow_fit() %>% tidy()

## # A tibble: 47 x 5
##   term      estimate std.error statistic p.value
##   <chr>     <dbl>     <dbl>     <dbl>    <dbl>
## 1 (Intercept) 1920452.  3087160.    0.622  0.535
## 2 ScreenPorch    59.7     69.1     0.863  0.390
## 3 MSSubClass     -323.    129.     -2.50   0.0138
## 4 GarageCars     3602.    7179.     0.502  0.617
## 5 BedroomAbvGr    1509.    3933.     0.384  0.702
## 6 TotalBsmtSF     12.4     8.82     1.41   0.162
## 7 LotArea        20971.   20280.     1.03   0.304
## 8 OpenPorchSF     -17.0     37.0     -0.461  0.646
## 9 BsmtFullBath    16888.   5087.     3.32   0.00124
## 10 WoodDeckSF      18.5     17.9     1.03   0.305
## # ... with 37 more rows
```

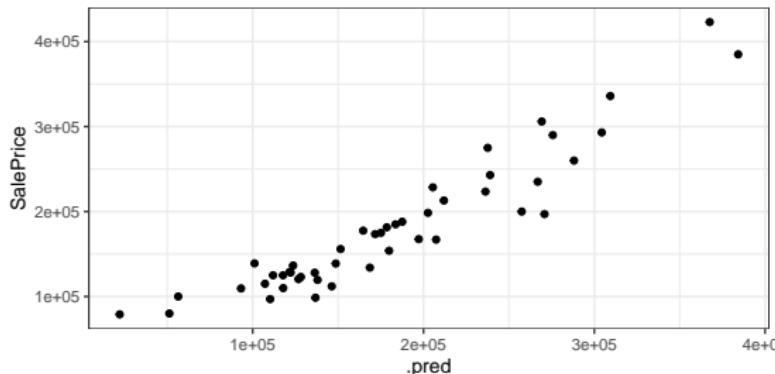
## Making predictions with workflow

```
house_preds<- predict(house_fit, test_data)
house_preds

## # A tibble: 50 x 1
##       .pred
##   <dbl>
## 1 178531.
## 2 236275.
## 3 257502.
## 4 269208.
## 5 51212.
## 6 123668.
## 7 136742.
## 8 136343.
## 9 238991.
## 10     NA
## # ... with 40 more rows
```

## Evaluate performance

```
house_results <- house_preds %>% cbind(test_data)
```



```
rbind(  
  rmse(house_results, truth = SalePrice, estimate = .pred),  
  rsq(house_results, truth = SalePrice, estimate = .pred)  
)
```

```
## # A tibble: 2 x 3  
##   .metric .estimator .estimate  
##   <chr>    <chr>        <dbl>  
## 1 rmse     standard    27049.  
## 2 rsq      standard     0.885
```

## Section 4

### Resampling

## Resampling with rsample

- We previously built a linear model for SalePrice as a function of predictors in the house data and found the following accuracy measures on **test** data:

```
## # A tibble: 2 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 rmse    standard    27049.
## 2 rsq     standard      0.885
```

## Resampling with rsample

- We previously built a linear model for SalePrice as a function of predictors in the house data and found the following accuracy measures on **test** data:

```
## # A tibble: 2 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>     <dbl>
## 1 rmse    standard    27049.
## 2 rsq     standard     0.885
```

- But how typical are these estimates? Let's perform cross-validation.

```
set.seed(271)
library(rsample)
folds <- vfold_cv(train_data, v = 10, statra = RoofMat1)
```

# Delving Deeper

- Which observations are in each fold?

```
folds$splits[[1]]
```

```
## <Analysis/Assess/Total>
## <135/15/150>
folds$splits[[1]] %>% analysis() %>% head() %>% select(1:5)
```

```
##      SalePrice   Id Functional BldgType Foundation
##  58       81000  387        Typ     1Fam    PConc
##  37      113000  240        Typ     1Fam    CBlock
##  85      124000  631        Typ     1Fam    BrkTil
## 108      183000  821        Typ     1Fam    PConc
## 192      340000 1418        Typ     1Fam    PConc
## 165      93000  1180       Min2     1Fam    Slab
```

```
folds$splits[[1]] %>% assessment() %>% head() %>% select(1:5)
```

```
##      SalePrice   Id Functional BldgType Foundation
## 193      271000 1427        Typ     1Fam    PConc
##  74      130000  499        Typ     1Fam    PConc
## 131      136500  997        Typ     1Fam    CBlock
## 123      169990  923        Typ     1Fam    PConc
##  82      240000  622        Typ     1Fam    CBlock
## 194      119000 1429        Typ     1Fam    CBlock
```

## Adding resampling to workflow

```
house_fit_resamples <- house_wflow %>% fit_resamples(folds)  
house_fit_resamples
```

```
## # Resampling results  
## # 10-fold cross-validation  
## # A tibble: 10 x 4  
##   splits      id    .metrics      .notes  
##   <list>      <chr> <list>      <list>  
## 1 <split [135/15]> Fold01 <tibble [2 x 4]> <tibble [1 x 1]>  
## 2 <split [135/15]> Fold02 <tibble [2 x 4]> <tibble [1 x 1]>  
## 3 <split [135/15]> Fold03 <tibble [2 x 4]> <tibble [1 x 1]>  
## 4 <split [135/15]> Fold04 <tibble [2 x 4]> <tibble [1 x 1]>  
## 5 <split [135/15]> Fold05 <tibble [2 x 4]> <tibble [1 x 1]>  
## 6 <split [135/15]> Fold06 <tibble [2 x 4]> <tibble [1 x 1]>  
## 7 <split [135/15]> Fold07 <tibble [2 x 4]> <tibble [1 x 1]>  
## 8 <split [135/15]> Fold08 <tibble [2 x 4]> <tibble [1 x 1]>  
## 9 <split [135/15]> Fold09 <tibble [2 x 4]> <tibble [1 x 1]>  
## 10 <split [135/15]> Fold10 <tibble [2 x 4]> <tibble [1 x 1]>
```

# Metrics

- Let's look at the results:

```
house_fit_resamples$.metrics[[1]]
```

```
## # A tibble: 2 x 4
##   .metric .estimator .estimate .config
##   <chr>   <chr>       <dbl> <chr>
## 1 rmse    standard    28540.  Preprocessor1_Model1
## 2 rsq     standard      0.874 Preprocessor1_Model1
```

```
house_fit_resamples$.metrics[[2]]
```

```
## # A tibble: 2 x 4
##   .metric .estimator .estimate .config
##   <chr>   <chr>       <dbl> <chr>
## 1 rmse    standard    22605.  Preprocessor1_Model1
## 2 rsq     standard      0.884 Preprocessor1_Model1
```

```
house_fit_resamples$.metrics[[3]]
```

```
## # A tibble: 2 x 4
##   .metric .estimator .estimate .config
##   <chr>   <chr>       <dbl> <chr>
## 1 rmse    standard    22019.  Preprocessor1_Model1
## 2 rsq     standard      0.876 Preprocessor1_Model1
```

# CV Performance

- How do the models do overall?

```
#Baseline
rbind(
  rmse(house_results, truth = SalePrice, estimate = .pred),
  rsq(house_results, truth = SalePrice, estimate = .pred)
)

## # A tibble: 2 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 rmse    standard   27049.
## 2 rsq     standard    0.885
```

# CV Performance

- How do the models do overall?

```
#Baseline
rbind(
  rmse(house_results, truth = SalePrice, estimate = .pred),
  rsq(house_results, truth = SalePrice, estimate = .pred)
)

## # A tibble: 2 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 rmse    standard    27049.
## 2 rsq     standard     0.885
```

- Cross-validation:

```
collect_metrics(house_fit_resamples)

## # A tibble: 2 x 6
##   .metric .estimator      mean      n   std_err .config
##   <chr>   <chr>       <dbl> <int>    <dbl> <chr>
## 1 rmse    standard    24994.      10  1939.  Preprocessor1_Model1
## 2 rsq     standard     0.862      10   0.0237 Preprocessor1_Model1
```

## Section 5

### Tuning Hyperparameters

# Building a LASSO model

- The linear model did fine. But can we improve our results using penalized regression?

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  - If we wanted a LASSO model with particular penalty (say  $\lambda = 4$ ) we could use

```
house_lasso_mod <- linear_reg(penalty =4 ) %>% set_engine("glmnet")
```

## Building a LASSO model

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```
house_lasso_mod <- linear_reg(penalty = 4) %>% set_engine("glmnet")
```

- But we are really interested in finding the **BEST** value of  $\lambda$ . So instead

```
house_lasso_mod <- linear_reg(penalty = tune()) %>% set_engine("glmnet")
```

## Building a LASSO model

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```
house_lasso_mod <- linear_reg(penalty = 4) %>% set_engine("glmnet")
```

- But we are really interested in finding the **BEST** value of  $\lambda$ . So instead

```
house_lasso_mod <- linear_reg(penalty = tune()) %>% set_engine("glmnet")
```

- Let's fit the model and tune

```
lasso_grid <- grid_regular(penalty() %>% range_set(c(-5,5)), levels = 10)
lasso_wf <- workflow() %>% add_model(house_lasso_mod) %>% add_recipe(house_rec)
lasso_res <- lasso_wf %>% tune_grid(grid = lasso_grid, resamples = folds)
```

# Results

```
collect_metrics(lasso_res)
```

```
## # A tibble: 20 x 7
##       penalty .metric .estimator     mean      n   std_err .config
##       <dbl> <chr>   <chr>     <dbl> <int>    <dbl> <chr>
## 1     0.00001 rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 2     0.00001 rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 3     0.000129 rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 4     0.000129 rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 5     0.00167 rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 6     0.00167 rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 7     0.0215  rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 8     0.0215  rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 9     0.278   rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 10    0.278   rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 11    3.59   rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 12    3.59   rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 13    46.4   rmse standard  24499.     10  1981. Preprocessor1_Mod-
## 14    46.4   rsq  standard    0.868     10   0.0209 Preprocessor1_Mod-
## 15    599.   rmse standard  23612.     10  2259. Preprocessor1_Mod-
## 16    599.   rsq  standard    0.878     10   0.0111 Preprocessor1_Mod-
## 17    7743.  rmse standard  28677.     10  3090. Preprocessor1_Mod-
## 18    7743.  rsq  standard    0.844     10   0.0281 Preprocessor1_Mod-
## 19  100000  rmse standard  67654.     10  5580. Preprocessor1_Mod-
## 20  100000  rsq  standard    NaN       0     NA  Preprocessor1_Mod-
```

# Results

```
collect_metrics(lasso_res)
```

```
## # A tibble: 20 x 7
##       penalty .metric .estimator     mean      n   std_err .config
##       <dbl> <chr>   <chr>     <dbl> <int>    <dbl> <chr>
## 1     0.00001 rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 2     0.00001 rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 3     0.000129 rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 4     0.000129 rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 5     0.00167 rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 6     0.00167 rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 7     0.0215  rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 8     0.0215  rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 9     0.278   rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 10    0.278   rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 11    3.59   rmse standard  24577.     10  1966. Preprocessor1_Mod-
## 12    3.59   rsq  standard    0.867     10   0.0209 Preprocessor1_Mod-
## 13    46.4   rmse standard  24499.     10  1981. Preprocessor1_Mod-
## 14    46.4   rsq  standard    0.868     10   0.0209 Preprocessor1_Mod-
## 15    599.   rmse standard  23612.     10  2259. Preprocessor1_Mod-
## 16    599.   rsq  standard    0.878     10   0.0111 Preprocessor1_Mod-
## 17    7743.  rmse standard  28677.     10  3090. Preprocessor1_Mod-
## 18    7743.  rsq  standard    0.844     10   0.0281 Preprocessor1_Mod-
## 19  100000  rmse standard  67654.     10  5580. Preprocessor1_Mod-
## 20  100000  rsq  standard    NaN       0     NA  Preprocessor1_Mod-
```

# Which penalties?

- Focus just on optimal penalties for rmse:

```
lasso_res %>%  
  show_best("rmse")
```

```
## # A tibble: 5 x 7  
##   penalty metric estimator  mean     n std_err .config  
##   <dbl> <chr>    <chr>     <dbl> <int>   <dbl> <chr>  
## 1 599.   rmse     standard  23612.    10    2259. Preprocessor1_Model08  
## 2 46.4    rmse     standard  24499.    10    1981. Preprocessor1_Model07  
## 3 0.00001 rmse     standard  24577.    10    1966. Preprocessor1_Model01  
## 4 0.000129 rmse     standard  24577.    10    1966. Preprocessor1_Model02  
## 5 0.00167  rmse     standard  24577.    10    1966. Preprocessor1_Model03
```

# Which penalties?

- Focus just on optimal penalties for rmse:

```
lasso_res %>%  
  show_best("rmse")
```

```
## # A tibble: 5 x 7  
##   penalty .metric .estimator  mean     n std_err .config  
##   <dbl> <chr>   <chr>    <dbl> <int>   <dbl> <chr>  
## 1 599.   rmse    standard  23612.    10    2259. Preprocessor1_Model08  
## 2 46.4    rmse    standard  24499.    10    1981. Preprocessor1_Model07  
## 3 0.00001 rmse    standard  24577.    10    1966. Preprocessor1_Model01  
## 4 0.000129 rmse    standard  24577.    10    1966. Preprocessor1_Model02  
## 5 0.00167  rmse    standard  24577.    10    1966. Preprocessor1_Model03
```

- Let's collect the best model:

```
best_lasso <- lasso_res %>% select_best(metric = "rmse")  
best_lasso
```

```
## # A tibble: 1 x 2  
##   penalty .config  
##   <dbl> <chr>  
## 1 599.  Preprocessor1_Model08
```

## Finalize the model

- We update or finalize our workflow with the values from `select_best`:

```
final_lasso_wf <- lasso_wf %>% finalize_workflow(best_lasso)
final_lasso_wf
```

```
## == Workflow =====
## Preprocessor: Recipe
## Model: linear_reg()
##
## -- Preprocessor -----
## 5 Recipe Steps
##
## * step_log()
## * step_mutate()
## * step_rm()
## * step_dummy()
## * step_zv()
##
## -- Model -----
## Linear Regression Model Specification (regression)
##
## Main Arguments:
##   penalty = 599.484250318942
##
## Computational engine: glmnet
```

## Fit the Best Model

- Thus far, we've just focused on finding the best parameter. But we haven't actually built a LASSO model on training data. Let's do that:

## Fit the Best Model

- Thus far, we've just focused on finding the best parameter. But we haven't actually built a LASSO model on training data. Let's do that:

```
final_lasso_fit <- final_lasso_wf %>% last_fit(data_split)
```

```
final_lasso_fit$.metrics
```

```
## [[1]]
## # A tibble: 2 x 4
##   .metric .estimator .estimate .config
##   <chr>   <chr>       <dbl> <chr>
## 1 rmse    standard    26889. Preprocessor1_Model1
## 2 rsq     standard      0.883 Preprocessor1_Model1
final_lasso_fit$.predictions
```

```
## [[1]]
## # A tibble: 50 x 4
##   .pred   .row SalePrice .config
##   <dbl> <int>    <int> <chr>
## 1 178628.     1    181500 Preprocessor1_Model1
## 2 231755.     2    223500 Preprocessor1_Model1
## 3 247905.     3    200000 Preprocessor1_Model1
## 4 265349.     7    306000 Preprocessor1_Model1
## 5 58165.     12    80000 Preprocessor1_Model1
## 6 129233.    14   136500 Preprocessor1_Model1
## 7 135097.    16    98600 Preprocessor1_Model1
```