Notes on Ch8: Feature Engineering with Recipes

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2025-08-20

Prerequisites

```
library(tidymodels)
## -- Attaching packages ------ tidymodels 1.3.0 --
               1.0.8
## v broom
                         v recipes
                                      1.3.1
## v dials
               1.4.1 v rsample
                                      1.3.1
## v dplyr
               1.1.4 v tibble
                                     3.3.0
## v ggplot2 3.5.2 v tidyr
## v infer 1.0.9 v tune
                                      1.3.1
                                      1.3.0
## v modeldata 1.5.0 v workflows 1.2.0
## v parsnip 1.3.2 v workflowsets 1.1.1
               1.1.0 v yardstick 1.3.2
## v purrr
## -- Conflicts ----- tidymodels_conflicts() --
## x purrr::discard() masks scales::discard()
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                 masks stats::lag()
## x recipes::step() masks stats::step()
tidymodels_prefer()
# Data prep from previous chapter
data(ames)
ames <- mutate(ames, Sale_Price = log10(Sale_Price))</pre>
# Data split
set.seed(502)
ames_split <- initial_split(ames, prop = 0.80, strata = Sale_Price)</pre>
ames_train <- training(ames_split)</pre>
ames_test <- testing(ames_split)</pre>
```

A simple recipe() for the Ames Housing Data

Sample recipe:

```
simple_ames <-
  recipe(
    Sale_Price ~ Neighborhood + Gr_Liv_Area + Year_Built + Bldg_Type,
    data = ames_train
) |>
  step_log(Gr_Liv_Area, base = 10) |>
```

```
step_dummy(all_nominal_predictors())
simple_ames

##

## -- Recipe ------
##

## -- Inputs

## Number of variables by role

## outcome: 1

## predictor: 4

##

## -- Operations

## * Log transformation on: Gr_Liv_Area

## * Dummy variables from: all_nominal_predictors()
```

Advantages of using recipes for modeling:

- computations can be recycled across models since they are not coupled to the modeling function.
- enables a broader set of data processing choices than what formulas can offer.
- the syntax can be very compact (like when using all_nominal_predictors(), etc).
- all data preprocessing can be captured in a single R object, instead of being in scripts that are scattered across different files.

Using recipes

Attaching the recipe object to the workflow:

```
lm_model <- linear_reg() |> set_engine("lm")
lm_wflow <-</pre>
 workflow() |>
 add_model(lm_model) |>
 add_recipe(simple_ames)
lm_wflow
## Preprocessor: Recipe
## Model: linear_reg()
## -- Preprocessor ------
## 2 Recipe Steps
##
## * step_log()
## * step_dummy()
## -- Model -----
## Linear Regression Model Specification (regression)
##
## Computational engine: lm
```

```
Fitting the training data to the model and recipe:
```

```
lm_fit <- fit(lm_wflow, ames_train)</pre>
Making a prediction using the model and recipe:
predict(lm_fit, ames_test |> slice(1:3))
## Warning in predict.lm(object = object$fit, newdata = new_data, type =
## "response", : prediction from rank-deficient fit; consider predict(.,
## rankdeficient="NA")
## # A tibble: 3 x 1
##
    .pred
##
    <dbl>
## 1 5.08
## 2 5.32
## 3 5.28
Extracting details from the bare model object or recipe:
lm_fit |> extract_recipe(estimated = TRUE)
##
## -- Recipe ------
##
## -- Inputs
## Number of variables by role
## outcome:
## predictor: 4
##
## -- Training information
## Training data contained 2342 data points and no incomplete rows.
##
## -- Operations
## * Log transformation on: Gr_Liv_Area | Trained
## * Dummy variables from: Neighborhood Bldg_Type | Trained
Extract model details in tidy format:
lm_fit |>
 # This returns the parsnip object:
 extract_fit_parsnip() |>
 # Now tidy the linear model object:
 tidy() |>
 slice(1:5)
## # A tibble: 5 x 5
##
    term
                               estimate std.error statistic
                                                              p.value
##
    <chr>
                                  <dbl>
                                            <dbl> <dbl>
                                                                <dbl>
                                                      -2.90 3.80e- 3
## 1 (Intercept)
                               -0.669
                                         0.231
## 2 Gr_Liv_Area
                                0.620
                                         0.0143
                                                     43.2 2.63e-299
```

```
## 3 Year_Built 0.00200 0.000117 17.1 6.16e- 62
## 4 Neighborhood_College_Creek 0.0178 0.00819 2.17 3.02e- 2
## 5 Neighborhood Old Town -0.0330 0.00838 -3.93 8.66e- 5
```

How data are used by the recipe()

- 1. The recipe(..., data) is called, the dataset is used to determine the data types of each column so that selectors like all_numeric() or all_numeric_predictors() can be used.
- 2. When fit(workflow, data) to prepare the data, the training data are used for all estimation operations.
- 3. When predict(workflow, new_data) is used, no model or preprocessor parameters are re-estimated using the values in new_data.

Examples of recipe steps

Encoding qualitative data in a numeric format

- step_unkown(): used to change missing values to a dedicated factor level.
- step_novel(): used to anticipate a new factor level that may be encountered in future data.
- step_other(): used to analyze the frequencies of the factor levels in the training set and convert infrequently occurring variables to a catch-all level or "other". Thresholds can be specified by the user.

Example:

```
simple_ames <-
   recipe(
     Sale_Price ~ Neighborhood + Gr_Liv_Area + Year_Built + Bldg_Type,
     data = ames_train
) |>
   step_log(Gr_Liv_Area, base = 10) |>
   step_other(Neighborhood, threshold = 0.01) |>
   step_dummy(all_nominal_predictors())
```

Interaction terms

Interaction effects occur when one predictor has an effect on the outcome that is contingent on one or more other predictors.

Example: When trying to predict how much traffic there will be during your commute using the specific time of day and the weather, the relationship between the amount of traffic and bad weather can be different depending on the time of day.

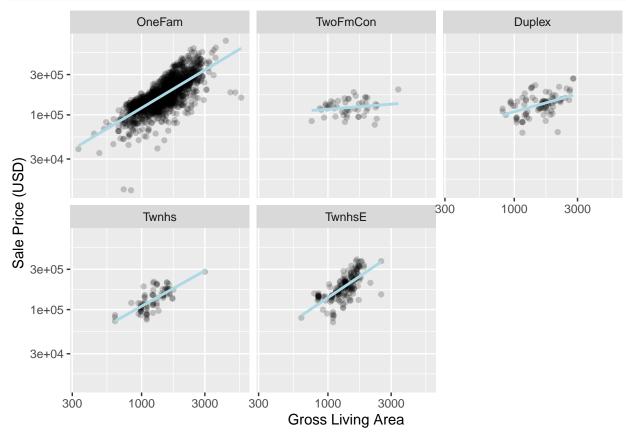
In this case, one could add to the model an interaction term between the two predictors, along with the original two predictors (which are also called "the main effects").

Numerically, the interaction term between predictors is encoded as their product.

Exploring the Ames training set (EDA):

```
ames_train |>
  ggplot(aes(x = Gr_Liv_Area, y = 10^Sale_Price)) +
  geom_point(alpha = 0.2) +
  facet_wrap(~ Bldg_Type) +
  geom_smooth(method = lm, formula = y ~ x, se = FALSE, color = "lightblue") +
  scale_x_log10() +
```

```
scale_y_log10() +
labs(x = "Gross Living Area", y = "Sale Price (USD)")
```



In base R formula, the interaction term is specified using a :.

```
Sale_Price ~ Neighborhood + log10(Gr_Liv_Area) + Bldg_Type + log10(Gr_Liv_Area):Bldg_Type # shorhand form of the formula above:
Sale_Price ~ Neighborhood + log10(Gr_Liv_Area) * Bldg_Type
```

In recipes, interaction terms are specified in a more explicit manner (using step_interact():

```
simple_ames <-
  recipe(
    Sale_Price ~ Neighborhood + Gr_Liv_Area + Year_Built + Bldg_Type,
    data = ames_train
) |>
  step_log(Gr_Liv_Area, base = 10) |>
  step_other(Neighborhood, threshold = 0.01) |>
  step_dummy(all_nominal_predictors()) |>
  # Gr_Liv_Area is on the log scale already!
  step_interact(~ Gr_Liv_Area:starts_with("Bldg_Type_"))
```

The function starts_with() was used here because step_dummy() uses an underscore as a separator between the column name and the level.

Additional interactions can be specified by separating them with +.

Any interaction term that interacts with itself (var_1:var_1), it will be ignored.

The interaction terms will be named according to the syntax var_1_x_var_2 in order for the column name to be a valid data frame column name.

Spline functions

These functions are used to represent nonlinear relationship between the predictors and the outcome.

The function geom_smooth() in ggplot, is a spline representation of the data.

Example:

```
library(patchwork)
library(splines)
plot_smoother <- function(deg_free) {</pre>
  ggplot(ames_train, aes(x = Latitude, y = 10^Sale_Price)) +
    geom_point(alpha = 0.2) +
    scale_y_log10() +
    geom_smooth(
       method = lm,
       formula = y ~ ns(x, df = deg_free),
       color = "lightblue",
       se = FALSE
    ) +
    labs(title = paste(deg_free, "Spline Terms"),
          y = "Sale Price (USD)")
}
( plot_smoother(2) + plot_smoother(5) ) / ( plot_smoother(20) + plot_smoother(100) )
          2 Spline Terms
                                                            5 Spline Terms
Sale Price (USD)
                                                  Sale Price (USD)
   3e+05
                                                     3e+05
                                                     1e+05
   3e+04
                                                     3e+04
                                          42.06
               42.00
                        42.02
                                 42.04
                                                                 42.00
                                                                          42.02
                                                                                   42.04
                                                                                            42.06
                        Latitude
                                                                          Latitude
          20 Spline Terms
                                                            100 Spline Terms
Sale Arice (USD) 3e+05 -
                                                  Sale Price (USD)
                                                     3e+05
                                                     1e+05
                                                     3e+04 -
```

42.00

42.02

Latitude

42.04

42.06

42.06

42.00

42.02

Latitude

42.04

The plot shows that two spline terms underfit the data, while 100 terms overfit. Having spline terms between 5 and 20 seem reasonable (i.e. the splines are fairly smooth and catch the main patterns of the data).

Adding a natural spline (ns) representation in recipes:

```
recipe(
    Sale_Price ~ Neighborhood + Gr_Liv_Area + Year_Built + Bldg_Type + Latitude,
    data = ames_train
) |>
    step_log(Gr_Liv_Area, base = 10) |>
    step_other(Neighborhood, threshold = 0.01) |>
    step_dummy(all_nominal_predictors()) |>
    step_interact( ~ Gr_Liv_Area:starts_with("Bldg_Type_")) |>
    step_ns(Latitude, deg_free = 20)
```

The number of splines to specify could be considered as a tuning parameter for the model.

Feature extraction

This is a method for representing multiple features at once. Most methods falling under this category create new features (aka columns) that capture the information in the broader set as a whole. Common example: PCA or Principal component analysis. This method tries to extract as much of the original information in the predictor set as possible using a smaller number of features.

In the Ames dataset, several predictors are used to measure the size of the property – the Total_Bsmt_SF, First_Flr_SF, and Gr_Liv_Area (total basement size, size of first floor, and gross living area, respectively). PCA can be used to represent these potentially redundant variables as a smaller feature set.

Possible recipe step for PCA for the use case explained above:

```
step_pca(matches("(SF$)|(Gr_Liv)"))
```

Note: step_pca() assumes that all of the predictors are on the same scale. For this reason, this step is often preceded by step_normalize(), which centers and scales each column.

Other recipe steps for feature extraction include independent component analysis (ICA), non-negative matrix factorization (NNMF), multidimensional scaling (MDS), uniform manifold approximation and projection (UMAP), etc.

Row sampling steps

Subsampling techniques are popularly used to deal with class imabalance. Common subsampling approaches:

- Downsampling: the data keeps the minority class and takes a random sample of the majority class. Aim is to make the class frequencies balance.
- Upsampling: replicates samples from the minority class to balance the classes.
- Hybrid: combination of both.

Subsampling syntax using the **themis** package:

```
step_downsample(outcome_column_name)
```

General transformations

The step_mutate() can be used to conduct a variety of basic operations to the data, similar to the dplyr mutate().

Natural language processing

Functions from the **textrecipes** package can apply NLP methods to the data.

Skipping steps for new data

Sometimes, there will be a step in the modeling process that should not be applied to new data. Example: the subsampling step performed to handle class imbalances during training - this step should not be performed when predicting on new data.

Solution: Each function in the step_*() family has a skip argument that when set to TRUE, will be ignored by the predict() function.

tidy() method for recipes

Sample recipe using the steps discussed above:

```
ames_rec <-
recipe(
    Sale_Price ~ Neighborhood + Gr_Liv_Area + Year_Built + Bldg_Type + Latitude + Longitude,
    data = ames_train
) |>
step_log(Gr_Liv_Area, base = 10) |>
step_other(Neighborhood, threshold = 0.01) |>
step_dummy(all_nominal_predictors()) |>
step_interact( ~ Gr_Liv_Area:starts_with("Bldg_Type_")) |>
step_ns(Latitude, Longitude, deg_free = 20)
```

Calling the tidy() method on the recipe object above:

```
tidy(ames_rec)
```

```
## # A tibble: 5 x 6
##
     number operation type
                                trained skip id
      <int> <chr>
##
                      <chr>
                                <lgl>
                                        <lgl> <chr>
## 1
          1 step
                      log
                                FALSE
                                        FALSE log_1iDg1
## 2
          2 step
                      other
                               FALSE
                                        FALSE other_anRnU
## 3
          3 step
                      dummy
                               FALSE
                                        FALSE dummy_Xe89S
## 4
                      interact FALSE
                                        FALSE interact_yj20u
          4 step
## 5
                                FALSE
                                        FALSE ns_1akCq
          5 step
                      ns
```

This result can be very helpful for verifying the individual steps.

Note: the entries under the id column contains some random suffix which are not really helpful. Specifying the id ahead of time for step_other() will improve this situation:

```
ames_rec <-
recipe(
    Sale_Price ~ Neighborhood + Gr_Liv_Area + Year_Built + Bldg_Type + Latitude + Longitude,
    data = ames_train
) |>
step_log(Gr_Liv_Area, base = 10) |>
step_other(Neighborhood, threshold = 0.01, id = "my_id") |>
step_dummy(all_nominal_predictors()) |>
step_interact( ~ Gr_Liv_Area:starts_with("Bldg_Type_")) |>
step_ns(Latitude, Longitude, deg_free = 20)

tidy(ames_rec)
```

```
## # A tibble: 5 x 6
##
    number operation type
                                trained skip id
      <int> <chr>
##
                      <chr>>
                                <lgl>
                                        <lgl> <chr>
                                FALSE
## 1
                                        FALSE log_OCwGV
          1 step
                      log
## 2
          2 step
                      other
                                FALSE
                                        FALSE my id
## 3
                                FALSE
                                        FALSE dummy_NkzTj
          3 step
                      dummy
## 4
                      interact FALSE
                                        FALSE interact AmBz1
          4 step
## 5
                                        FALSE ns_8R5Nn
          5 step
                      ns
                                FALSE
Refitting the workflow with the new recipe:
```

```
lm_wflow <-</pre>
  workflow() |>
  add_model(lm_model) |>
  add_recipe(ames_rec)
lm_fit <- fit(lm_wflow, ames_train)</pre>
```

Calling the tidy() method again, along with the id identifier specified for step_other():

```
estimated_recipe <-</pre>
  lm_fit |>
  extract_recipe(estimated = TRUE)
tidy(estimated_recipe, id = "my_id")
```

```
## # A tibble: 22 x 3
##
      terms
                                      id
                  retained
##
      <chr>
                   <chr>
                                      <chr>>
## 1 Neighborhood North_Ames
                                      my_id
## 2 Neighborhood College_Creek
                                      my_id
## 3 Neighborhood Old_Town
                                      my_id
## 4 Neighborhood Edwards
                                      my_id
## 5 Neighborhood Somerset
                                      my_id
## 6 Neighborhood Northridge_Heights my_id
## 7 Neighborhood Gilbert
                                      my_id
## 8 Neighborhood Sawyer
                                      my_id
## 9 Neighborhood Northwest_Ames
                                      my_id
## 10 Neighborhood Sawyer_West
                                      my_id
## # i 12 more rows
```

If we know the number identifier from the result of calling tidy() on the recipe (i.e., tide(ames_rec)), we can use that instead of id when generating the tidied up estimated_recipe:

```
tidy(estimated_recipe, number = 2)
```

```
## # A tibble: 22 x 3
##
      terms
                  retained
                                      id
                   <chr>
##
      <chr>
                                      <chr>
## 1 Neighborhood North_Ames
                                      my_id
## 2 Neighborhood College_Creek
                                      my id
## 3 Neighborhood Old_Town
                                      my_id
## 4 Neighborhood Edwards
                                      my_id
## 5 Neighborhood Somerset
                                      my_id
## 6 Neighborhood Northridge_Heights my_id
## 7 Neighborhood Gilbert
                                      my_id
## 8 Neighborhood Sawyer
                                      my_id
```

```
## 9 Neighborhood Northwest_Ames my_id
## 10 Neighborhood Sawyer_West my_id
## # i 12 more rows
```

Each tidy() call for a step_*() function returns relevant information about that step.

Column roles

The formula used with the initial call to recipe() assigns *roles* to each of the columns, depending on which side of the tilde (~) they are on. The roles are either predictor or outcome.

Sometimes, it may be necessary to keep a column even though it is neither a predictor nor an outcome. (Example: the address column in the Ames dataset).

The function *_role() can be used to add, remove, or update roles. Example: updating the role of address in the recipe:

```
ames_rec |>
  update_role(address, new_role = "street address")
```

After which the address column will cease to be a predictor, but instead will be a "street address" in the recipe.

All functions in the step_*() family have a role field that we can tweak, depending on what is necessary for each step.

Summary

Standard code for this chapter:

```
library(tidymodels)
tidymodels_prefer()
data(ames)
ames <- mutate(ames, Sale_Price = log10(Sale_Price))</pre>
set.seed(502)
ames_split <- initial_split(ames, prop = 0.80, strata = Sale_Price)</pre>
ames_train <- training(ames_split)</pre>
ames_test <- testing(ames_split)</pre>
ames rec <-
 recipe(
    Sale_Price ~ Neighborhood + Gr_Liv_Area + Year_Built + Bldg_Type + Longitude + Latitude,
    data = ames_train
  ) |>
  step_log(Gr_Liv_Area, base = 10) |>
  step other(Neighborhood, threshold = 0.01) |>
  step_dummy(all_nominal_predictors()) |>
  step_interact( ~ Gr_Liv_Area:starts_with("Bldg_Type_")) |>
  step_ns(Latitude, Longitude, deg_free = 20)
lm_model <- linear_reg() |> set_engine("lm")
lm_wflow <-</pre>
  workflow() |>
  add_model(lm_model) |>
```

```
add_recipe(ames_rec)

lm_fit <- fit(lm_wflow, ames_train)</pre>
```