

### Simple Linear Regression in R

Regression models allow you explore relationships between a response and explanatory variables.

You can use the model to make predictions.

It is always a good idea to visualize a dataset such as scatterplots.

The intercept is the y value when x is zero. In some cases, its interpretations might make sense. For instance, on average a house with zero convenience stores nearby had a price of 8.2242 TWD per square meter.

The slope is the amount y increases by if you increase x by one.

If your sole explanatory variable is categorical, the intercept is the response variable's mean of the omitted category. The coefficients of each category are means relative to the intercept. You can change this if you like so that the coefficients are the means of each category.

### Simple regression code

```
# Assume you have a real estate dataset and want to build a model predicting prices using n_convenience
stores nearby within walking distance.
# Visualize the two variables. What is the relationship?
ggplot(taiwan_real_estate, aes(n_convenience, price_twd_msq)) +
  geom_point(alpha = 0.5) +
  geom_smooth(method='lm', se=FALSE)
# Run a linear regression of price_twd_msq vs. n_convenience
lm(price_twd_msq ~ n_convenience, data = taiwan_real_estate)
# Visualize prices v age_category using a histogram. What do you see?
ggplot(taiwan_real_estate, aes(price_twd_msq)) +
  # Make it a histogram with 10 bins
  geom_histogram(bins=10) +
  # Facet the plot so each house age group gets its own panel
  facet_wrap(vars(house_age_years))
# calculate means by each age category
taiwan_real_estate %>%
  # Group by house age
  group_by(house_age_years) %>%
  # Summarize to calculate the mean house price/area
  summarize(mean_by_group = mean(price_twd_msq))
# Run a linear regression of price_twd_msq vs. house_age_years
mdl_price_vs_age <- lm(data=taiwan_real_estate, price_twd_msq~house_age_years) #add +0 after house_age-
_years to get each category's mean.

# See the result
mdl_price_vs_age
```



### Predictions and Model objects

Extrapolating means making predictions outside the range of observed data. Even if you use nonsense explanatory data to make predictions, the model won't throw an error and give you a prediction. Understand your data to determine if a prediction is nonsense or not.

`coefficients(model_object)` returns a named, numeric vector of the coefficients.

`residuals(model_object)` returns the difference between actual response values minus predictions. They are a measure of inaccuracy.

`broom::augment(model_object)` returns observation level detail such as residuals, fitted values, etc.

Residuals exist due to problems in the model and fundamental randomness. And extreme cases are also often due to randomness.

Due to regression to the mean, a baseball player does not hit as many home runs this year as he did the year before.

If you transformed the response variable, you must "back-transform" your predictions.

It is useful to have values you want to use to make predictions (test data) in a tibble. Thus, you can store your predictions in the same tibble and make plots.

`fitted_values(model_object)` returns predictions on the original data.

`broom::tidy(model_object)` returns coefficients and its detail.

`broom::glance(model_object)` returns model-level results (performance metrics).

Eventually, extreme cases will more look like average cases (b/c they don't persist over time). This is called regression to the mean.

If there is no straight line relationship between the response variable and the explanatory variable, it is sometimes possible to create one by transforming one or both of the variables.

### How to make predictions and view model objects?

```
# Model prices and n_convenience
mdl_price_vs_conv <- lm(formula = price_twd_msq ~ n_convenience, data = taiwan_real_estate)
# Create a tibble of integer values from 0 to 10.
explanatory_data <- tibble(n_convenience = 0:10)
# Make predictions and store them in prediction_data
prediction_data <- explanatory_data %>%
  mutate(price_twd_msq = predict(mdl_price_vs_conv, explanatory_data))
# Plot the predictions along with all points
ggplot(taiwan_real_estate, aes(n_convenience, price_twd_msq)) +
  geom_point() +
  geom_smooth(method = "lm", se = FALSE) +
  # Add a point layer of prediction data, colored yellow
  geom_point(color='yellow', data=prediction_data)
# ----- Regression to the mean example-----
# Suppose you have data on annual returns from investing in companies in the SP500 index and you're
interested in knowing if the invested performance stays the same from 2018 to 2019.
# Using sp500_yearly_returns, plot return_2019 vs. return_2018
ggplot(data=sp500_yearly_returns, aes(x=return_2018, y=return_2019)) +
  # Make it a scatter plot
  geom_point() +
```



### How to make predictions and view model objects? (cont)

```
# Add a line at y = x, colored green, size 1
geom_abline(slope=1, color='green', size=1) +
# Add a linear regression trend line, no std. error ribbon
geom_smooth(method='lm', se=FALSE) +
# Fix the coordinate ratio
coord_fixed()

# Transforming variables and back-transforming the response-----
# Assume you've facebook advertising data; how many people see the adds and how many people click on them.
mdl_click_vs_impression <- lm(
  I(n_clicks^0.25) ~ I(n_impressions^0.25),
  data = ad_conversion
)
explanatory_data <- tibble(
  n_impressions = seq(0, 3e6, 5e5)
)
prediction_data <- explanatory_data %>%
  mutate(
    n_clicks_025 = predict(mdl_click_vs_impression, explanatory_data),
    n_clicks = n_clicks_025 ^ 4
  )
ggplot(ad_conversion, aes(n_impressions ^ 0.25, n_clicks ^ 0.25)) +
  geom_point() +
  geom_smooth(method = "lm", se = FALSE) +
  # Add points from prediction_data, colored green
  geom_point(data=prediction_data, color='green')
```

### Quantifying Model Fit

Coefficient of determination is the proportion of variance in the response variable that is predictable from the explanatory variable. 1 means a perfect fit and 0 means the worst possible fit.

For simple linear regression, coeff of determination is correlation between the response and explanatory squared.

Residual standard error (or sum of squared residuals) is a typical difference between prediction and an observed response. This is sigma in broom::glance(model)

Root mean squared error (RMSE) also works but the denominator is number of observations and not degrees of freedom.

If the linear regression model is a good fit, then the residuals are normally distributed and their mean is zero. This assumption can be checked using the residual v fitted values plot. The blue trend line should closely follow the y=0 line.

The Q-Q plot whether the residuals follow a normal distribution. If the points track along the diagonal line, they are normally distributed.

The scale-location plot shows whether the size of residuals get bigger or smaller as the fitted values change.

Leverage quantifies how extreme your explanatory variables are. These values are stored under .hat in augment().

Influence measures how much model the model would change if you left the observation out of the dataset when modeling. Contained in .cooksd column in augment().



### Code to quantify model's fit.

```
# Plot three diagnostics for mdl_price_vs_conv
library(ggplot2)
library(ggfortify)
autoplot(mdl_price_vs_conv, which=1:3, nrow=3, ncol=1)
# Plot the three outlier diagnostics for mdl_price_vs_conv
autoplot(mdl_price_vs_dist, which=4:6, nrow=3, ncol=1)
```

### Simple Logistic regression in R

Build this model when the response is binary. Predictions are probabilities and not amounts.

The responses follow a logistic (s-shaped) curve. You can have the model return probabilities by specifying the response type in predict().

Odds ratio is the probability of something happening divided by the probability that it doesn't.

This is sometimes easier to reason about than probabilities, particularly when you want to make decisions about choices. For example, if a customer has a 20% chance of churning, it may be more intuitive to say "the chance of them not churning is four times higher than the chance of them churning".

One downside to probabilities and odds ratios for logistic regression predictions is that the prediction lines for each are curved.

A nice property of logistic regression odds ratio is that on a log-scale they change linearly with the explanatory variable.

This makes it harder to reason about what happens to the prediction when you make a change to the explanatory variable. The logarithm of the odds ratio (the "log odds ratio") does have a linear relationship between predicted response and explanatory variable.

We use confusion matrices to quantify the fit of logistic regression.

Accuracy is the proportion of correct predictions.

Sensitivity is the proportion of true positives.  $TP/(FN+TP)$ . Proportion of observations where the actual response was true where the model also predicted were true.

Specificity is the proportion of true negatives.  $TN/(TN+FP)$ . Proportion of observations where the actual response was false where the model also predicted that they were false.

### Code for Logistic regression in R

```
plt_churn_vs_relationship ggplot(churn, aes(time_since_first_purchase, has_churned)) +
  geom_point() +
  geom_smooth(method = "lm", se = FALSE, color = "red") +
  # Add a glm trend line, no std error ribbon, binomial family
  geom_smooth(method='glm', se=FALSE, method.args=list(family=binomial))
# Fit a logistic regression of churn vs.
# length of relationship using the churn dataset
mdl_churn_vs_relationship <- glm(has_churned ~ time_since_first_purchase, data=churn, family='binomial')
# See the result
mdl_churn_vs_relationship
# Make predictions. "response" type returns probabilities of churning.
prediction_data <- explanatory_data %>%
  mutate(
```



### Code for Logistic regression in R (cont)

```

has_churned = predict(mdl_churn_vs_relationship, explanatory_data, type = "response"),
most_likely_outcome = round(has_churned) # easier to interpret.
)
# Update the plot
plt_churn_vs_relationship +
  # Add most likely outcome points from prediction_data, colored yellow, size 2
  geom_point(data=prediction_data, size=2, color='yellow', aes(y=most_likely_outcome))
# Odds ratio-----
# From previous step
prediction_data <- explanatory_data %>%
  mutate(
    has_churned = predict(mdl_churn_vs_relationship, explanatory_data, type = "response"),
    odds_ratio = has_churned / (1 - has_churned),
    log_odds_ratio = log(odds_ratio)
  )
# Using prediction_data, plot odds_ratio vs. time_since_first_purchase
ggplot(data=prediction_data, aes(x=time_since_first_purchase, y=odds_ratio)) +
  # Make it a line plot
  geom_line() +
  # Add a dotted horizontal line at y = 1. Indicates where churning is just as likely as not churning.
  geom_hline(yintercept=1, linetype='dotted')

```

### Quantifying Logistic's Model Fit

```

# Get the confusion matrix.
library(yardstick)
# Get the actual and most likely responses from the dataset
actual_response <- churn$has_churned
predicted_response <- round(fitted(mdl_churn_vs_relationship))
# Create a table of counts
outcomes <- table(predicted_response, actual_response)
# Convert outcomes to a yardstick confusion matrix
confusion <- conf_mat(outcomes)
# Get performance metrics for the confusion matrix
summary(confusion, event_level = 'second')

```

