

Libraries to Import	Basic Codes (cont)	Data Information
library(readr) library(dplyr)	<b>factor(x, levels=c("wt", "mutant"))</b> [Create factor specifying level order]	is.na(x) is.null(x)
library(ggplot2) library(broom)	<b>relevel(x, ref="wildtype")</b> [Re-level a factor variable]	is.nan(x) is.array(x)
library(caret) library(rpart)	<b>t.test(y~grp, data=df)</b> [T-test mean y across grp in data df]	is.data.frame(x) is.numeric(x)
library(splines) library(party)	<b>lmfit &lt;- lm(y~x1+x2, data=df)</b> [Fit linear model y against two x's]	is.complex(x) is.character(x)
library(leaps) library(glmnet)	<b>anova(lmfit)</b> [Print ANOVA table on object returned from lm()]	head(x) tail(x)
library(MASS) library(class)	<b>summary(lmfit)</b> [Get summary information about a model fit with lm()]	summary(x) str(x)
<b>Data Conversion</b>	<b>TukeyHSD(aov(lmfit))</b> [ANOVA Post-hoc pairwise contrasts]	length(x) dim(x)
as.array(x) as.character(x)	<b>wilcox.test(y~grp, data=df)</b> [Wilcoxon rank sum / Mann-Whitney U test]	dimnames(x) attr(x,which)
as.data.frame(x) as.factor(x)	<b>xt &lt;- xtabs(~x1+x2, data=df)</b> [Cross-tabulate a contingency table]	nrow(x) ncol(x)
as.logical(x) as.numeric(x)	<b>addmargins(xt)</b> [Adds summary margin to a contingency table xt]	NROW(x) NCOL(x)
<b>K-Fold</b>	<b>addmargins(xt)</b> [Adds summary margin to a contingency table xt]	class(x) unclass(x)
<b>folds &lt;- crossv_kfold(data, k = 5)</b> [k is the number of folds]	<b>chisq.test(xt)</b> [Chi-square test on a contingency table xt]	<b>Data Splitting</b>
<b>Regularization - Ridge &amp; Lasso</b>	<b>fisher.test(xt)</b> [Fisher's exact test on a contingency table xt]	<b>createDataPartition(y,p=0.8)</b> [createDalt splits a vector 'y' with 80 percent data in one part and 20 percent in other parttaPartition(y,p=0.8)]
lambda_to_try <- 10^seq(-3, 5, length.out = 100)^{1/2}	<b>mosaicplot(xt)</b> [Mosaic plot for a contingency table xt]	<b>trainControl( summaryFunction=&lt;Rfunction&gt;,classProbs=&lt;logical&gt;)</b> [It is used for controlling training parameters like resampling, number of folds, iteration etc.]
<b>ridge_cv &lt;- cv.glmnet(X, y, alpha = 0, lambda = lambda_to_try, standardize = TRUE, nfolds = 10)</b> [Setting alpha = 0 implements ridge regression]	<b>power.t.test(n, power, sd, delta)</b> [T-test power calculations]	<b>densityplot.rfe(x,data,...)</b> [Lattice functions for plotting resampling results of recursive feature selection]
<b>lasso &lt;- glmnet(data_x, data_y, alpha = 1.0)</b>	<b>power.prop.test(n, power, p1, p2)</b> [Proportions test power calculations]	<b>featureplot(x,y,plot...)</b> [A shortcut to produce lattice plots]
<b>Random Forest</b>	<b>tidy() augment() glance()</b> [Model tidying functions in the broom package]	
<b>randomForest(formula, data)</b> [formula is a formula describing the predictor and response variables. data is the name of the data set used]		
<b>Basic Codes</b>		
<b>read_csv("path/nhanes.csv")</b> [Read nhanes.csv in the path/ folder (readr)]		
<b>View(df)</b> [View tabular data frame df in a graphical viewer]		
<b>mean, median, range</b> [Descriptive stats. Remember na.rm=TRUE if desired]		
<b>filter(df, ...)</b> [ Filters data frame according to condition ... (dplyr)]		

