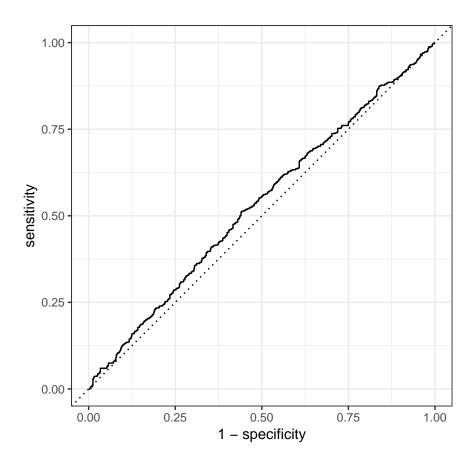
```
library(ISLR)
head(Smarket)
     Year
           Lag1
                 Lag2 Lag3 Lag4
                                      Lag5 Volume Today Direction
## 1 2001  0.381  -0.192  -2.624  -1.055  5.010  1.1913  0.959
## 2 2001 0.959 0.381 -0.192 -2.624 -1.055 1.2965 1.032
                                                                 Uр
## 3 2001 1.032 0.959 0.381 -0.192 -2.624 1.4112 -0.623
                                                               Down
## 4 2001 -0.623 1.032 0.959 0.381 -0.192 1.2760 0.614
                                                                 Uр
## 5 2001 0.614 -0.623 1.032 0.959 0.381 1.2057 0.213
                                                                 Uр
## 6 2001 0.213 0.614 -0.623 1.032 0.959 1.3491 1.392
                                                                 Uр
Logistic Regression
glm_fit <- glm(Direction ~ Lag1 + Lag2,</pre>
   data = Smarket,
   family = binomial
)
glm_fit
## Call: glm(formula = Direction ~ Lag1 + Lag2, family = binomial, data = Smarket)
##
## Coefficients:
## (Intercept)
                      Lag1
                                   Lag2
##
      0.07425
                  -0.07151
                               -0.04450
##
## Degrees of Freedom: 1249 Total (i.e. Null); 1247 Residual
## Null Deviance:
                       1731
## Residual Deviance: 1728 AIC: 1734
library(tidyverse)
## -- Attaching packages -----
## v ggplot2 3.1.0
                        v purrr
                                  0.2.5
## v tibble 2.0.1
                        v dplyr 0.8.0.1
## v tidyr
            0.8.1
                        v stringr 1.4.0
## v readr
            1.1.1
                        v forcats 0.3.0
## -- Conflicts -----
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
## x dplyr::select() masks MASS::select()
library(modelr)
new_data <- read_csv("Lag1, Lag2
                     0.5, 0.3
                     0.4, 0.3")
new_data %>% add_predictions(glm_fit) %>% mutate(prob = exp(pred)/ (1 + exp(pred)))
```

library(MASS)

```
## # A tibble: 2 x 4
## Lag1 Lag2 pred prob
## <dbl> <dbl> <dbl> <dbl> <dbl> <0.5 0.3 0.0251 0.506
## 2 0.4 0.3 0.0323 0.508</pre>
```

ROC curve

```
Smarket2 <- Smarket %>%
   add_predictions(glm_fit) %>%
   mutate(prob = exp(pred)/ (1 + exp(pred)), EstDir = ifelse(prob > 0.5, "Up", "Down"))
Smarket2 %>% count(Direction, EstDir) %>% spread(Direction, n)
## # A tibble: 2 x 3
   EstDir Down
   <chr> <int> <int>
## 1 Down
           114
                  102
## 2 Up
            488
                  546
library(tidymodels)
## -- Attaching packages ------
## v broom 0.5.0 v recipes 0.1.4
## v dials 0.0.2 v rsample 0.0.4
## v infer 0.4.0
                      v yardstick 0.0.2
## v parsnip
            0.0.1
## -- Conflicts ------
## x broom::bootstrap() masks modelr::bootstrap()
## x scales::discard() masks purrr::discard()
## x dplyr::filter() masks stats::filter()
## x recipes::fixed() masks stringr::fixed()
## x dplyr::lag() masks stats::lag()
## x yardstick::mae() masks modelr::mae()
## x yardstick::mape() masks modelr::mape()
## x yardstick::rmse() masks modelr::rmse()
                      masks MASS::select()
## x dplyr::select()
## x yardstick::spec() masks readr::spec()
                      masks stats::step()
## x recipes::step()
## x yardstick::tidy() masks rsample::tidy(), recipes::tidy(), broom::tidy()
autoplot(roc_curve(Smarket2, Direction, prob))
```



roc_auc(Smarket2, Direction, prob)

Multinomial logistic

variable name	type	about the variable
id	scale	student id
female	nominal	(0/1)
race	nominal	ethnicity (1=hispanic 2=asian 3=african-amer 4=white)
ses	ordinal	socio economic status (1=low 2=middle 3=high)
schtyp	nominal	type of school (1=public 2=private)
prog	nominal	type of program (1=general 2=academic 3=vocational)
read	scale	standardized reading score
write	scale	standardized writing score
math	scale	standardized math score
science	scale	standardized science score
socst	scale	standardized social studies score
hon	nominal	honors english $(0/1)$

```
ml <- read_csv("hsb2.csv")</pre>
## Parsed with column specification:
## cols(
##
    id = col_integer(),
##
    female = col_integer(),
##
    race = col_integer(),
##
    ses = col_integer(),
    schtyp = col_integer(),
    prog = col_integer(),
##
##
    read = col_integer(),
## write = col_integer(),
## math = col_integer(),
##
   science = col_integer(),
##
   socst = col_integer()
## )
hsb2 <- ml %>% mutate(
   prog = recode_factor(prog, `1` = "general", `2` = "academic", `3` = "vocational"),
   ses = recode_factor(ses, `1` = "low", `2` = "middle", `3` = "high"))
hsb2 %>% count(prog, ses) %>% spread(prog, n)
## # A tibble: 3 x 4
##
   ses general academic vocational
   <fct> <int> <int> <int>
## 1 low 16 19
                                 12
## 2 middle
              20
                       44
                                   31
               9
                                    7
## 3 high
                       42
hsb2 %>% group_by(prog) %>% summarize(mwrite = mean(write))
## # A tibble: 3 x 2
   prog
          mwrite
                <dbl>
   <fct>
## 1 general 51.3
## 2 academic 56.3
## 3 vocational 46.8
library(nnet)
multi_fit <- multinom(prog ~ ses + write, data = hsb2)</pre>
## # weights: 15 (8 variable)
## initial value 219.722458
## iter 10 value 179.985215
## final value 179.981726
## converged
```

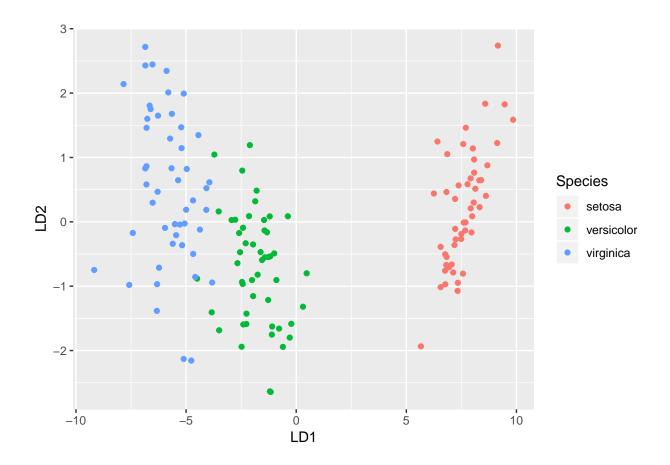
```
new_data <- tibble(ses = "middle", write = 56)</pre>
predict(multi_fit, new_data, type = "probs")
      general
               academic vocational
##
  0.2174957 0.5485545 0.2339499
# tidyverse
new_data %>% add_predictions(multi_fit)
## # A tibble: 1 x 3
         write pred
    ses
     <chr> <dbl> <fct>
##
## 1 middle
              56 academic
predict(multi_fit, new_data)
## [1] academic
## Levels: general academic vocational
# a few month later
# new_data %>% add_predictions(multi_fit, type = "probs")
```

LDA

```
lda_fit <- lda(Direction ~ Lag1 + Lag2, data = Smarket)</pre>
Smarket3 <- Smarket %>% mutate(EstDir = predict(lda_fit, newdata = Smarket)$class)
Smarket3 %>% count(Direction, EstDir) %>% spread(Direction, n)
## # A tibble: 2 x 3
## EstDir Down
    <fct> <int> <int>
## 1 Down
            114 102
## 2 Up
              488 546
new_data <- read_csv("Lag1, Lag2</pre>
                      0.5, 0.3
                      0.4, 0.3")
predict(lda_fit, new_data)
## $class
## [1] Up Up
## Levels: Down Up
##
## $posterior
##
         Down
                      Uр
```

An exmple with more than one class

```
head(iris)
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
                                                0.2 setosa
## 1
             5.1
                         3.5
                                     1.4
## 2
             4.9
                         3.0
                                     1.4
                                                 0.2 setosa
## 3
             4.7
                         3.2
                                     1.3
                                                 0.2 setosa
## 4
                                                 0.2 setosa
             4.6
                         3.1
                                      1.5
                                                 0.2 setosa
## 5
                         3.6
             5.0
                                      1.4
## 6
             5.4
                         3.9
                                      1.7
                                                 0.4 setosa
iris_fit <- lda(Species ~ ., data = iris)</pre>
new_data <- tibble(Sepal.Length = 5.906, Sepal.Width = 2.77, Petal.Length = 3, Petal.Width = 0.246)
predict(iris_fit, new_data)$posterior
##
        setosa
                versicolor
                              virginica
## 1 0.9996068 0.0003931945 7.917338e-19
iris2 <- iris %>% bind_cols(as_tibble(predict(iris_fit, newdata = iris)$x))
ggplot(iris2) + geom_point(aes(LD1, LD2, color = Species))
```



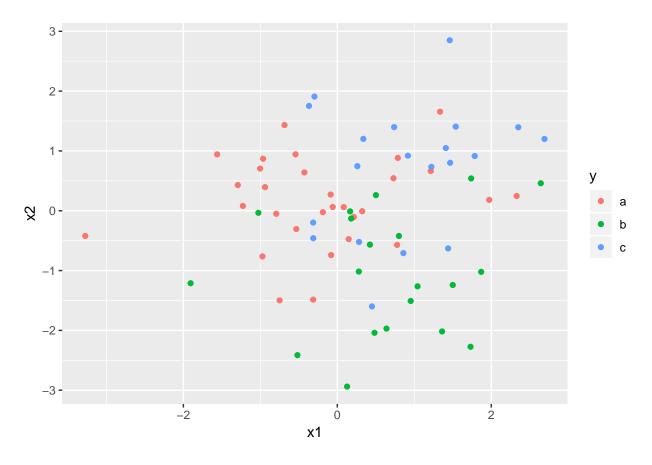
Reduced rank LDA

```
# only use LD1
predict(iris_fit, new_data, dimen = 1)
## $class
## [1] setosa
## Levels: setosa versicolor virginica
##
## $posterior
                               virginica
##
        setosa
                 versicolor
## 1 0.9999577 4.230918e-05 1.925451e-18
##
## $x
          LD1
## 1 3.958893
```

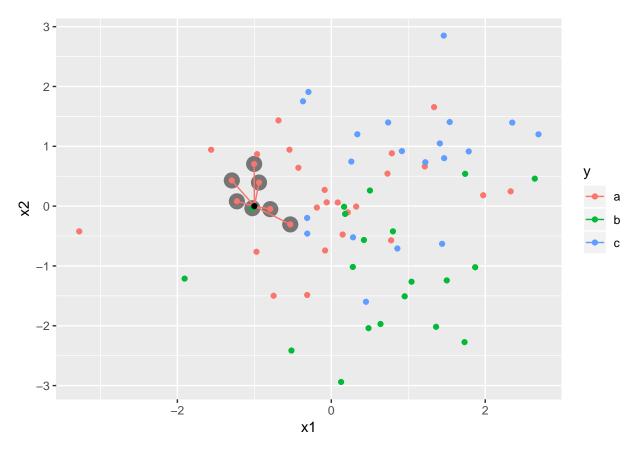
Explaining kNN

```
library(mvtnorm)
g1 <- rmvnorm(30, mean=c(-0.5,0))
g2 <- rmvnorm(20, mean=c(1,-1))
g3 <- rmvnorm(20, mean=c(1,1))
x <- rbind(g1, g2, g3)
colnames(x) <- c("x1", "x2")
x <- as_tibble(x)
y <- rep(c("a", "b", "c"), c(30, 20, 20))
knn_example <- bind_cols(x, y = y)</pre>
```

```
ggplot(knn_example) + geom_point(aes(x1, x2, color = y))
```



```
k = 7
point = c(-1, 0)
knn_neighor <- knn_example %>%
  mutate(dist = sqrt((x1 - point[1])^2 + (x2 - point[2])^2)) %>%
  filter(row_number(dist) <= k)
ggplot(knn_example) +
  geom_point(data = knn_neighor, aes(x1, x2), alpha = 0.5, size = 5) +
  geom_segment(data = knn_neighor, aes(x = x1, y = x2, xend = point[1], yend = point[2], color = y)) +
  geom_point(aes(x1, x2, color = y)) +
  annotate("point", x = point[1], y = point[2])</pre>
```



1

2

0.2

0.6

1 c

-1 b