

Lab 4 MATH 342W

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Create a dataset D which we call Xy such that the linear model has R^2 about 0% but x, y are clearly associated.

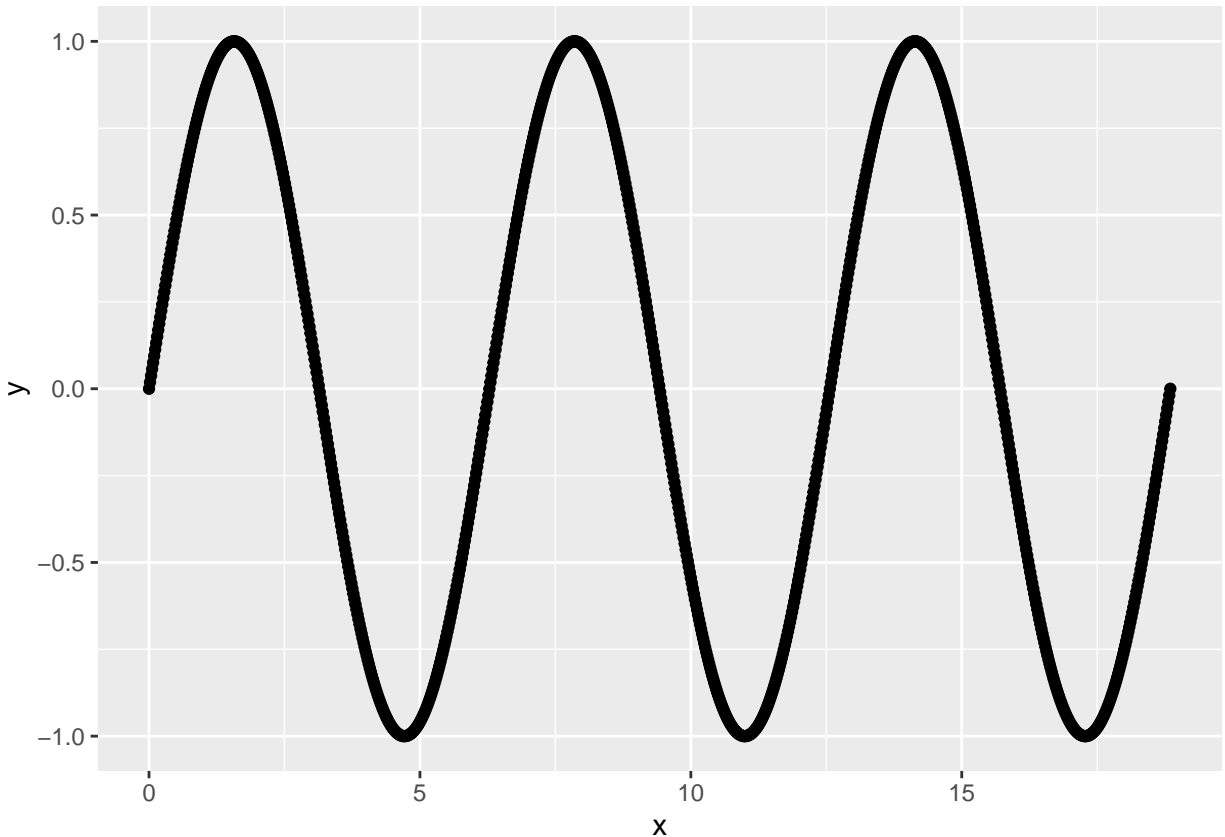
```
x = seq(0, 6 * pi, length.out=1000) # 1000 output
y = sin(x)
```

```
pacman::p_load(ggplot2)
```

```
#first check that Rsq is around zero
summary(lm(y ~ x))$r.squared
```

```
## [1] 0.06734109
```

```
#now check association visually
ggplot(data.frame(x = x, y = y)) + geom_point(aes(x = x, y = y))
```



Write a function `my_ols` that takes in `X`, a matrix with `p` columns representing the feature measurements for each of the `n` units, a vector of `n` responses `y` and returns a list that contains the `b`, the `p+1`-sized column vector of OLS coefficients, `yhat` (the vector of `n` predictions), `e` (the vector of `n` residuals), `df` for degrees of freedom of the model, `SSE`, `SST`, `MSE`, `RMSE` and `Rsqr` (for the R-squared metric). Internally, you cannot use `lm` or any other package; it must be done manually. You should throw errors if the inputs are non-numeric or not the same length. Or if `X` is not otherwise suitable. You should also name the class of the return value `my_ols` by using the `class` function as a setter. No need to create ROxygen documentation here.

$df = \text{degrees of freedom} = p + 1 / \text{number of dimensions} / \# \text{ of parameters}$

```
# X -> Columns / features
my_ols = function(X, y){
  # Step 1 concatenate 1 column
  X = cbind(1, X)

  # t(X) -> Transpose
  # Get best weights using OLS
  b = solve(t(X) %*% X) %*% t(X) %*% y

  # Get predictions
  y_hat = X %*% b

  # Get residuals
  e = y - y_hat

  # GET METRICS
  SSE = sum(e^2)
```

```

SST = sum((y - mean(y))^2)

df = ncol(X)
n = nrow(X)
MSE = SSE / (n - df)
RMSE = sqrt(MSE)
RSQ = (SST - SSE) / SST

# RETURN
lmobj = list(
  b = b,
  y_hat = y_hat,
  e = e,
  SSE = SSE,
  SST = SST,
  df = df,
  MSE = MSE,
  RSQ = RSQ
)

class(lmobj) = "my_ols"
lmobj
}

```

Verify that the OLS coefficients for the Type of cars in the cars dataset gives you the same results as we did in class (i.e. the ybar's within group).

```

#TO-DO
cars = MASS::Cars93
colnames(cars)

```

```

## [1] "Manufacturer"      "Model"              "Type"
## [4] "Min.Price"         "Price"              "Max.Price"
## [7] "MPG.city"          "MPG.highway"        "AirBags"
## [10] "DriveTrain"        "Cylinders"          "EngineSize"
## [13] "Horsepower"        "RPM"                "Rev.per.mile"
## [16] "Man.trans.avail"   "Fuel.tank.capacity" "Passengers"
## [19] "Length"            "Wheelbase"          "Width"
## [22] "Turn.circle"       "Rear.seat.room"     "Luggage.room"
## [25] "Weight"            "Origin"              "Make"

```

```
head(cars)
```

```

##   Manufacturer  Model   Type Min.Price Price Max.Price MPG.city MPG.highway
## 1      Acura Integra  Small    12.9   15.9    18.8     25         31
## 2      Acura Legend Midsize    29.2   33.9    38.7     18         25
## 3       Audi   90 Compact    25.9   29.1    32.3     20         26
## 4       Audi  100 Midsize    30.8   37.7    44.6     19         26
## 5        BMW  535i Midsize    23.7   30.0    36.2     22         30
## 6      Buick Century Midsize    14.2   15.7    17.3     22         31

```

```
##           AirBags DriveTrain Cylinders EngineSize Horsepower  RPM
## 1           None      Front         4         1.8         140 6300
## 2 Driver & Passenger      Front         6         3.2         200 5500
## 3      Driver only      Front         6         2.8         172 5500
## 4 Driver & Passenger      Front         6         2.8         172 5500
## 5      Driver only      Rear          4         3.5         208 5700
## 6      Driver only      Front         4         2.2         110 5200
##  Rev.per.mile Man.trans.avail Fuel.tank.capacity Passengers Length Wheelbase
## 1          2890              Yes          13.2           5     177     102
## 2          2335              Yes          18.0           5     195     115
## 3          2280              Yes          16.9           5     180     102
## 4          2535              Yes          21.1           6     193     106
## 5          2545              Yes          21.1           4     186     109
## 6          2565              No           16.4           6     189     105
##  Width Turn.circle Rear.seat.room Luggage.room Weight  Origin      Make
## 1     68          37          26.5          11   2705 non-USA Acura Integra
## 2     71          38          30.0          15   3560 non-USA Acura Legend
## 3     67          37          28.0          14   3375 non-USA Audi 90
## 4     70          37          31.0          17   3405 non-USA Audi 100
## 5     69          39          27.0          13   3640 non-USA BMW 535i
## 6     69          41          28.0          16   2880  USA Buick Century
```

```
# model.matrix => augments 1 and dummifies levels.
cars_X = model.matrix(~Type, cars)
head(cars_X)
```

```
## (Intercept) TypeLarge TypeMidsize TypeSmall TypeSporty TypeVan
## 1           1           0           0           1           0           0
## 2           1           0           1           0           0           0
## 3           1           0           0           0           0           0
## 4           1           0           1           0           0           0
## 5           1           0           1           0           0           0
## 6           1           0           1           0           0           0
```

```
# There are 6 types of cars, 1 became the intercept
head(cars_X)
```

```
## (Intercept) TypeLarge TypeMidsize TypeSmall TypeSporty TypeVan
## 1           1           0           0           1           0           0
## 2           1           0           1           0           0           0
## 3           1           0           0           0           0           0
## 4           1           0           1           0           0           0
## 5           1           0           1           0           0           0
## 6           1           0           1           0           0           0
```

```
# TRYING TO MODEL THE PRICE OF THE CAR FROM FEATURES
```

```
cars_y = cars$Price # Our label, y => Price, what we're guessing
```

```
# Initially it will fail because they the extra 1 column augmented makes the X not symmetrical
# -1 removes the intercept column or our 1 vector
my_ols(cars_X[, -1], cars_y)
```

```

## $b
##           [,1]
##      18.212500
## TypeLarge    6.087500
## TypeMidsize  9.005682
## TypeSmall   -8.045833
## TypeSporty   1.180357
## TypeVan      0.887500
##
## $y_hat
##           [,1]
## 1  10.16667
## 2  27.21818
## 3  18.21250
## 4  27.21818
## 5  27.21818
## 6  27.21818
## 7  24.30000
## 8  24.30000
## 9  27.21818
## 10 24.30000
## 11 27.21818
## 12 18.21250
## 13 18.21250
## 14 19.39286
## 15 27.21818
## 16 19.10000
## 17 19.10000
## 18 24.30000
## 19 19.39286
## 20 24.30000
## 21 18.21250
## 22 24.30000
## 23 10.16667
## 24 10.16667
## 25 18.21250
## 26 19.10000
## 27 27.21818
## 28 19.39286
## 29 10.16667
## 30 24.30000
## 31 10.16667
## 32 10.16667
## 33 18.21250
## 34 19.39286
## 35 19.39286
## 36 19.10000
## 37 27.21818
## 38 24.30000
## 39 10.16667
## 40 19.39286
## 41 19.39286
## 42 10.16667
## 43 18.21250

```

```

## 44 10.16667
## 45 10.16667
## 46 19.39286
## 47 27.21818
## 48 27.21818
## 49 27.21818
## 50 27.21818
## 51 27.21818
## 52 24.30000
## 53 10.16667
## 54 10.16667
## 55 18.21250
## 56 19.10000
## 57 19.39286
## 58 18.21250
## 59 27.21818
## 60 19.39286
## 61 27.21818
## 62 10.16667
## 63 27.21818
## 64 10.16667
## 65 18.21250
## 66 19.10000
## 67 27.21818
## 68 18.21250
## 69 27.21818
## 70 19.10000
## 71 24.30000
## 72 19.39286
## 73 10.16667
## 74 18.21250
## 75 19.39286
## 76 27.21818
## 77 24.30000
## 78 18.21250
## 79 10.16667
## 80 10.16667
## 81 10.16667
## 82 18.21250
## 83 10.16667
## 84 10.16667
## 85 19.39286
## 86 27.21818
## 87 19.10000
## 88 10.16667
## 89 19.10000
## 90 18.21250
## 91 19.39286
## 92 18.21250
## 93 27.21818
##
## $e
##           [,1]
## 1  5.733333e+00

```

```
## 2 6.681818e+00
## 3 1.088750e+01
## 4 1.048182e+01
## 5 2.781818e+00
## 6 -1.151818e+01
## 7 -3.500000e+00
## 8 -6.000000e-01
## 9 -9.181818e-01
## 10 1.040000e+01
## 11 1.288182e+01
## 12 -4.812500e+00
## 13 -6.812500e+00
## 14 -4.292857e+00
## 15 -1.131818e+01
## 16 -2.800000e+00
## 17 -2.500000e+00
## 18 -5.500000e+00
## 19 1.860714e+01
## 20 -5.900000e+00
## 21 -2.412500e+00
## 22 5.200000e+00
## 23 -9.666667e-01
## 24 1.133333e+00
## 25 -4.912500e+00
## 26 -1.000000e-01
## 27 -1.161818e+01
## 28 6.407143e+00
## 29 2.033333e+00
## 30 -5.000000e+00
## 31 -2.766667e+00
## 32 -6.666667e-02
## 33 -6.912500e+00
## 34 -3.492857e+00
## 35 -5.392857e+00
## 36 8.000000e-01
## 37 -7.018182e+00
## 38 -3.400000e+00
## 39 -1.766667e+00
## 40 -6.892857e+00
## 41 4.071429e-01
## 42 1.933333e+00
## 43 -7.125000e-01
## 44 -2.166667e+00
## 45 -1.666667e-01
## 46 -9.392857e+00
## 47 -1.331818e+01
## 48 2.068182e+01
## 49 7.818182e-01
## 50 7.981818e+00
## 51 7.081818e+00
## 52 1.180000e+01
## 53 -1.866667e+00
## 54 1.433333e+00
## 55 -1.712500e+00
```

```

## 56 -3.552714e-15
## 57  1.310714e+01
## 58  1.368750e+01
## 59  3.468182e+01
## 60 -5.292857e+00
## 61 -1.231818e+01
## 62  1.333333e-01
## 63 -1.118182e+00
## 64  1.633333e+00
## 65 -2.512500e+00
## 66 -3.552714e-15
## 67 -5.718182e+00
## 68 -4.712500e+00
## 69 -1.091818e+01
## 70  4.000000e-01
## 71 -3.600000e+00
## 72 -4.992857e+00
## 73 -1.166667e+00
## 74 -7.112500e+00
## 75 -1.692857e+00
## 76 -8.718182e+00
## 77  1.000000e-01
## 78  1.048750e+01
## 79  9.333333e-01
## 80 -1.766667e+00
## 81  7.333333e-01
## 82  1.287500e+00
## 83 -1.566667e+00
## 84 -3.666667e-01
## 85 -9.928571e-01
## 86 -9.018182e+00
## 87  3.600000e+00
## 88 -1.066667e+00
## 89  6.000000e-01
## 90  1.787500e+00
## 91  3.907143e+00
## 92  4.487500e+00
## 93 -5.181818e-01
##
## $SSE
## [1] 5162.586
##
## $SST
## [1] 8584.021
##
## $df
## [1] 6
##
## $MSE
## [1] 59.34007
##
## $RSQ
## [1] 0.3985819
##

```



```
## attr("class")
## [1] "my_ols"
```

```
print("...")
```

```
## [1] "..."
```

Create a prediction method `g` that takes in a vector `x_star` and the dataset `D` i.e. `X` and `y` and returns the OLS predictions. Let `X` be a matrix with `p` columns representing the feature measurements for each of the `n` units

```
g = function(x_star, X, y){
  #TO-DO
  # c1 to x_star so it will be of same size to b since b = length(p+1)
  #x_star is our unseen data / unseen features
  # ols weights * x_star == y_hat
  c(1,x_star) %*% my_ols(X,y)$b
}
```

Load up the famous iris dataset. We are going to do a different prediction problem. Imagine the only input `x` is Species and you are trying to predict `y` which is Petal.Length. A reasonable prediction is the average petal length within each Species. Prove that this is the OLS model by fitting an appropriate `lm` and then using the predict function to verify.

```
data(iris)
#TO-DO
# We're trying tho show that this gives y_
coef(lm(Petal.Length ~ Species, iris))
```

```
##          (Intercept) Speciesversicolor Speciesvirginica
##             1.462             2.798             4.090
```

```
# We try to pull petal length for all species
mean(iris$Petal.Length[iris$Species == "setosa"]) # PULLS OUT ALL THE PETAL.LENGTHS OF SETOSA -> Then w
```

```
## [1] 1.462
```

```
mean(iris$Petal.Length[iris$Species == "versicolor"])
```

```
## [1] 4.26
```

```
mean(iris$Petal.Length[iris$Species == "virginica"])
```

```
## [1] 5.552
```

```
# 5.552 = coefficient of Virginica -> 1.462 + 4.090
```

Construct the design matrix with an intercept, `X` without using `model.matrix`.

```
## design matrix == x matrix
cbind(1, ifelse(iris$Species == "versicolor", 1, 0), ifelse(iris$Species == "virginica", 1, 0))
```

```
##      [,1] [,2] [,3]
## [1,]    1    0    0
## [2,]    1    0    0
## [3,]    1    0    0
## [4,]    1    0    0
## [5,]    1    0    0
## [6,]    1    0    0
## [7,]    1    0    0
## [8,]    1    0    0
## [9,]    1    0    0
## [10,]   1    0    0
## [11,]   1    0    0
## [12,]   1    0    0
## [13,]   1    0    0
## [14,]   1    0    0
## [15,]   1    0    0
## [16,]   1    0    0
## [17,]   1    0    0
## [18,]   1    0    0
## [19,]   1    0    0
## [20,]   1    0    0
## [21,]   1    0    0
## [22,]   1    0    0
## [23,]   1    0    0
## [24,]   1    0    0
## [25,]   1    0    0
## [26,]   1    0    0
## [27,]   1    0    0
## [28,]   1    0    0
## [29,]   1    0    0
## [30,]   1    0    0
## [31,]   1    0    0
## [32,]   1    0    0
## [33,]   1    0    0
## [34,]   1    0    0
## [35,]   1    0    0
## [36,]   1    0    0
## [37,]   1    0    0
## [38,]   1    0    0
## [39,]   1    0    0
## [40,]   1    0    0
## [41,]   1    0    0
## [42,]   1    0    0
## [43,]   1    0    0
## [44,]   1    0    0
## [45,]   1    0    0
## [46,]   1    0    0
## [47,]   1    0    0
## [48,]   1    0    0
## [49,]   1    0    0
```

##	[50,]	1	0	0
##	[51,]	1	1	0
##	[52,]	1	1	0
##	[53,]	1	1	0
##	[54,]	1	1	0
##	[55,]	1	1	0
##	[56,]	1	1	0
##	[57,]	1	1	0
##	[58,]	1	1	0
##	[59,]	1	1	0
##	[60,]	1	1	0
##	[61,]	1	1	0
##	[62,]	1	1	0
##	[63,]	1	1	0
##	[64,]	1	1	0
##	[65,]	1	1	0
##	[66,]	1	1	0
##	[67,]	1	1	0
##	[68,]	1	1	0
##	[69,]	1	1	0
##	[70,]	1	1	0
##	[71,]	1	1	0
##	[72,]	1	1	0
##	[73,]	1	1	0
##	[74,]	1	1	0
##	[75,]	1	1	0
##	[76,]	1	1	0
##	[77,]	1	1	0
##	[78,]	1	1	0
##	[79,]	1	1	0
##	[80,]	1	1	0
##	[81,]	1	1	0
##	[82,]	1	1	0
##	[83,]	1	1	0
##	[84,]	1	1	0
##	[85,]	1	1	0
##	[86,]	1	1	0
##	[87,]	1	1	0
##	[88,]	1	1	0
##	[89,]	1	1	0
##	[90,]	1	1	0
##	[91,]	1	1	0
##	[92,]	1	1	0
##	[93,]	1	1	0
##	[94,]	1	1	0
##	[95,]	1	1	0
##	[96,]	1	1	0
##	[97,]	1	1	0
##	[98,]	1	1	0
##	[99,]	1	1	0
##	[100,]	1	1	0
##	[101,]	1	0	1
##	[102,]	1	0	1
##	[103,]	1	0	1

```

## [104,] 1 0 1
## [105,] 1 0 1
## [106,] 1 0 1
## [107,] 1 0 1
## [108,] 1 0 1
## [109,] 1 0 1
## [110,] 1 0 1
## [111,] 1 0 1
## [112,] 1 0 1
## [113,] 1 0 1
## [114,] 1 0 1
## [115,] 1 0 1
## [116,] 1 0 1
## [117,] 1 0 1
## [118,] 1 0 1
## [119,] 1 0 1
## [120,] 1 0 1
## [121,] 1 0 1
## [122,] 1 0 1
## [123,] 1 0 1
## [124,] 1 0 1
## [125,] 1 0 1
## [126,] 1 0 1
## [127,] 1 0 1
## [128,] 1 0 1
## [129,] 1 0 1
## [130,] 1 0 1
## [131,] 1 0 1
## [132,] 1 0 1
## [133,] 1 0 1
## [134,] 1 0 1
## [135,] 1 0 1
## [136,] 1 0 1
## [137,] 1 0 1
## [138,] 1 0 1
## [139,] 1 0 1
## [140,] 1 0 1
## [141,] 1 0 1
## [142,] 1 0 1
## [143,] 1 0 1
## [144,] 1 0 1
## [145,] 1 0 1
## [146,] 1 0 1
## [147,] 1 0 1
## [148,] 1 0 1
## [149,] 1 0 1
## [150,] 1 0 1

```

We now load the diamonds dataset. Skim the dataset using skimr or summary. What is the datatype of the color feature? : ORDERED FACTOR originally before we turn it into an integer after one-hot-encoding the different levels.

```
rm(list = ls())
pacman::p_load(ggplot2, skim)
```

```
## Installing package into 'C:/Users/lenovo/AppData/Local/R/win-library/4.3'
## (as 'lib' is unspecified)
```

```
## Warning: package 'skim' is not available for this version of R
##
## A version of this package for your version of R might be available elsewhere,
## see the ideas at
## https://cran.r-project.org/doc/manuals/r-patched/R-admin.html#Installing-packages
```

```
## Warning: unable to access index for repository http://www.stats.ox.ac.uk/pub/RWin/bin/windows/contrib/4.3/PACKAGES
## cannot open URL 'http://www.stats.ox.ac.uk/pub/RWin/bin/windows/contrib/4.3/PACKAGES'
```

```
## Warning: 'BiocManager' not available. Could not check Bioconductor.
##
## Please use 'install.packages('BiocManager')' and then retry.
```

```
## Warning in p_install(package, character.only = TRUE, ...):
```

```
## Warning in library(package, lib.loc = lib.loc, character.only = TRUE,
## logical.return = TRUE, : there is no package called 'skim'
```

```
## Warning in pacman::p_load(ggplot2, skim): Failed to install/load:
## skim
```

```
pacman::p_load(skim)
```

```
## Installing package into 'C:/Users/lenovo/AppData/Local/R/win-library/4.3'
## (as 'lib' is unspecified)
```

```
## Warning: package 'skim' is not available for this version of R
##
## A version of this package for your version of R might be available elsewhere,
## see the ideas at
## https://cran.r-project.org/doc/manuals/r-patched/R-admin.html#Installing-packages
```

```
## Warning: unable to access index for repository http://www.stats.ox.ac.uk/pub/RWin/bin/windows/contrib/4.3/PACKAGES
## cannot open URL 'http://www.stats.ox.ac.uk/pub/RWin/bin/windows/contrib/4.3/PACKAGES'
```

```
## Warning: 'BiocManager' not available. Could not check Bioconductor.
##
## Please use 'install.packages('BiocManager')' and then retry.
```

```
## Warning in p_install(package, character.only = TRUE, ...):
```

```
## Warning in library(package, lib.loc = lib.loc, character.only = TRUE,
## logical.return = TRUE, : there is no package called 'skim'
```

```
## Warning in pacman::p_load(skim): Failed to install/load:
## skim
```

```
diamonds = ggplot2::diamonds
#TO-DO
summary(diamonds)
```

```
##      carat      cut      color      clarity      depth
## Min.   :0.2000   Fair      : 1610   D: 6775   SI1      :13065   Min.    :43.00
## 1st Qu.:0.4000   Good      : 4906   E: 9797   VS2      :12258   1st Qu.:61.00
## Median :0.7000   Very Good:12082   F: 9542   SI2      : 9194   Median :61.80
## Mean   :0.7979   Premium  :13791   G:11292   VS1      : 8171   Mean    :61.75
## 3rd Qu.:1.0400   Ideal     :21551   H: 8304   VVS2     : 5066   3rd Qu.:62.50
## Max.   :5.0100                      I: 5422   VVS1     : 3655   Max.    :79.00
##                                J: 2808   (Other): 2531
##
##      table      price      x      y
## Min.   :43.00   Min.    : 326   Min.    : 0.000   Min.    : 0.000
## 1st Qu.:56.00   1st Qu.: 950   1st Qu.: 4.710   1st Qu.: 4.720
## Median :57.00   Median : 2401   Median : 5.700   Median : 5.710
## Mean   :57.46   Mean    : 3933   Mean    : 5.731   Mean    : 5.735
## 3rd Qu.:59.00   3rd Qu.: 5324   3rd Qu.: 6.540   3rd Qu.: 6.540
## Max.   :95.00   Max.    :18823   Max.    :10.740   Max.    :58.900
##
##      z
## Min.   : 0.000
## 1st Qu.: 2.910
## Median : 3.530
## Mean    : 3.539
## 3rd Qu.: 4.040
## Max.    :31.800
##
```

```
colnames(diamonds)
```

```
## [1] "carat" "cut" "color" "clarity" "depth" "table" "price"
## [8] "x" "y" "z"
```

```
typeof(diamonds$color)
```

```
## [1] "integer"
```

Find the levels of the color feature.

```
levels(diamonds$color)
```

```
## [1] "D" "E" "F" "G" "H" "I" "J"
```

```
# Different entries in color feature
```

Create new feature in the diamonds dataset, `color_as_numeric`, which is color expressed as a continuous interval value.

```
#TO-DO
# NUMERIC
# one hot encoding color
diamonds$color_as_numeric = as.numeric(diamonds$color)
head(diamonds)
```

```
## # A tibble: 6 x 11
##   carat cut    color clarity depth table price     x     y     z color_as_numeric
##   <dbl> <ord> <ord> <ord>   <dbl> <dbl> <int> <dbl> <dbl> <dbl>         <dbl>
## 1  0.23 Ideal E      SI2     61.5    55   326   3.95   3.98   2.43             2
## 2  0.21 Prem~ E      SI1     59.8    61   326   3.89   3.84   2.31             2
## 3  0.23 Good  E      VS1     56.9    65   327   4.05   4.07   2.31             2
## 4  0.29 Prem~ I      VS2     62.4    58   334   4.2    4.23   2.63             6
## 5  0.31 Good  J      SI2     63.3    58   335   4.34   4.35   2.75             7
## 6  0.24 Very~ J      VVS2     62.8    57   336   3.94   3.96   2.48             7
```

```
# NOMINAL
diamonds$color_as_nominal = factor(diamonds$color)
head(diamonds)
```

```
## # A tibble: 6 x 12
##   carat cut    color clarity depth table price     x     y     z color_as_numeric
##   <dbl> <ord> <ord> <ord>   <dbl> <dbl> <int> <dbl> <dbl> <dbl>         <dbl>
## 1  0.23 Ideal E      SI2     61.5    55   326   3.95   3.98   2.43             2
## 2  0.21 Prem~ E      SI1     59.8    61   326   3.89   3.84   2.31             2
## 3  0.23 Good  E      VS1     56.9    65   327   4.05   4.07   2.31             2
## 4  0.29 Prem~ I      VS2     62.4    58   334   4.2    4.23   2.63             6
## 5  0.31 Good  J      SI2     63.3    58   335   4.34   4.35   2.75             7
## 6  0.24 Very~ J      VVS2     62.8    57   336   3.94   3.96   2.48             7
## # i 1 more variable: color_as_nominal <ord>
```

Use that converted feature as the one predictor in a regression. How well does this regression do as measured by RMSE?

```
#TO-DO
# Trying to fit a linear line using price as a function of color_as_numeric
# Different colors of diamonds cost different amounts
diamonds_coeff = lm(price ~ color_as_numeric, diamonds) # gets w_0, w_1
diamonds_coeff
```

```
##
## Call:
## lm(formula = price ~ color_as_numeric, data = diamonds)
##
## Coefficients:
##      (Intercept)  color_as_numeric
##           2478.7             404.6
```

```
summary(diamonds_coeff)$sigma # RSQ
```

```
## [1] 3929.665
```

Create new feature in the diamonds dataset, `color_as_nominal`, which is color expressed as a nominal categorical variable.

```
#TO-DO
diamonds_coeff_nominal = lm(price ~ color_as_nominal, diamonds)
summary(diamonds_coeff_nominal)$sigma
```

```
## [1] 3926.777
```

Use that converted feature as the one predictor in a regression. How well does this regression do as measured by RMSE?

```
#TO-DO

# Assuming the linear model has been fit and is named diamonds_coeff
predicted_prices = predict(diamonds_coeff, newdata = diamonds)

# Calculate residuals (differences between actual and predicted prices)
residuals = diamonds$price - predicted_prices

# Calculate MSE and then RMSE
mse = mean(residuals^2)
rmse = sqrt(mse)

# Print RMSE
print(rmse)
```

```
## [1] 3929.592
```

```
# We are off 3,929.59 in regards to price.
```

Which regression does better - `color_as_numeric` or `color_as_nominal`? Why?

#TO-DO

Now regress both `color_as_numeric` and `color_as_nominal` in a regression. Does this regression do any better (as gauged by RMSE) than either `color_as_numeric` or `color_as_nominal` alone?

```
#TO-DO

numeric_nominal_model = lm(diamonds$price ~ diamonds$color_as_numeric + diamonds$color_as_nominal, newdata = diamonds)

## Warning: In lm.fit(x, y, offset = offset, singular.ok = singular.ok, ...) :
## extra argument 'newdata' will be disregarded

numeric_nominal_predictions = predict(numeric_nominal_model, newdata = diamonds)
numeric_nominal_residuals = diamonds$price - numeric_nominal_predictions
nn_mse = mean(numeric_nominal_residuals^2)
nn_rmse = sqrt(nn_mse)
cat("Color as Nominal only RMSE:", rmse, "Both Numeric and Nominal RMSE:", nn_rmse, "\n")
```

```
## Color as Nominal only RMSE: 3929.592 Both Numeric and Nominal RMSE: 3926.522
```


What are the coefficients (the b vector)?

```
# EXTRACT AND Y
X = subset(diamonds, select = -c(color_as_nominal, price))
X = as.matrix(X)
X = cbind(1, X)
y = diamonds$price

# Compute the coefficients vector 'b_vector' using the OLS formula
#b_vector <- solve(t(X) %*% X) %*% t(X) %*% y

# Print the coefficients vector
#print(b_vector)
```

Something appears to be anomalous in the coefficients. What is it? Why? 1. Intercept is really high, meaning if everything is 0 the diamond will cost insane amounts? 2. x has a negative coefficient which does not make sense since x is a measurement of the diamond. it should be that the larger the x the larger the price

#TO-DO

Return to the iris dataset. Find the hat matrix H for this regression of diamond price on diamond color. Use only the first 1,00 observations in the diamond dataset.

```
rm(list = ls())
diamonds1000 = ggplot2::diamonds[1:1000,]
# WE NEED TO FIND THE X MATRIX =>  $H = X(X^T X)^{-1} X^T$ 
X_diamonds = model.matrix(price ~ color, diamonds1000) #Regress price to color
H = X_diamonds %*% solve(t(X_diamonds) %*% X_diamonds) %*% t(X_diamonds)
```

Verify this hat matrix is symmetric using the `expect_equal` function in the package `testthat`.

```
#TO-DO
pacman::p_load(testthat)
expect_equal(H, t(H)) #Test for symmetry ; NO RESPONSE HENCE IT IS SYMMETRICAL
```

Verify this hat matrix is idempotent using the `expect_equal` function in the package `testthat`.

```
pacman::p_load(testthat)
#TO-DO
expect_equal(H %*% H, H) # NO RESPONSE MEANING IT IS IDEMPOTENT
```

Using the `diag` function, find the trace of the hat matrix.

```
#TO-DO
sum(diag(H))
```

```
## [1] 7
```

```
# Trace of an orthogonal matrix is its rank
```

It turns out the trace of a hat matrix is the same as its rank! But we don't have time to prove these interesting and useful facts..

Using the hat matrix, compute the \hat{y} vector and using the projection onto the residual space, compute the e vector and verify they are orthogonal to each other.

#TO-DO

```
y_diamond = diamonds1000$price
yhat_diamond = H%*%y_diamond #
yhat_diamond
```

```
##          [,1]
## 1    2542.571
## 2    2542.571
## 3    2542.571
## 4    1947.695
## 5    1990.217
## 6    1990.217
## 7    1947.695
## 8    2307.808
## 9    2542.571
## 10   2307.808
## 11   1990.217
## 12   1990.217
## 13   2692.438
## 14   1990.217
## 15   2542.571
## 16   2542.571
## 17   1947.695
## 18   1990.217
## 19   1990.217
## 20   1990.217
## 21   1947.695
## 22   2542.571
## 23   2307.808
## 24   1990.217
## 25   1990.217
## 26   2576.259
## 27   1947.695
## 28   1990.217
## 29   2594.380
## 30   2692.438
## 31   2692.438
## 32   2692.438
## 33   2542.571
## 34   2542.571
## 35   2594.380
## 36   2692.438
## 37   2542.571
## 38   2307.808
## 39   2594.380
## 40   1947.695
## 41   1947.695
```

##	42	1990.217
##	43	2594.380
##	44	2594.380
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##	84	1947.695
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```

```
I = diag(nrow(H))
e = (I - H) %*% y_diamond
```

```
t(e) %*% yhat_diamond # 0 -> but since the inversion the bits are off and giving us an error
```

```
##           [,1]
## [1,] 1.063338e-06
```

```
print("...")
```

```
## [1] "..."
```

Compute SST, SSR and SSE and R^2 and then show that $SST = SSR + SSE$.

```
SST = sum((y_diamond - mean(y_diamond))^2)
SSE = sum(e^2)
SSR = sum((yhat_diamond - mean(y_diamond))^2)

SST - sum(SSR + SSE)
```

```
## [1] 2.384186e-07
```

```
RSQ = SSR / SST
RSQ
```

```
## [1] 0.07918666
```

Find the angle theta between $y - \bar{y}$ and $\hat{y} - \bar{y}$ and then verify that its cosine squared is the same as the R^2 from the previous problem.

```
#TO-DO

numer = sqrt(sum((yhat_diamond - mean(y_diamond))^2))
denom = sqrt(sum((y_diamond - mean(y_diamond))^2))
theta = acos(numer/denom)
cos(theta)^2
```

```
## [1] 0.07918666
```

Project the y vector onto each column of the X matrix and test if the sum of these projections is the same as \hat{y} .

```
n = nrow(X_diamonds)
sum_proj_y = matrix(0, nrow=n, ncol=1)

for (j in 1:ncol(X_diamonds)) {
  X_j = X_diamonds[, j, drop=FALSE]
  sum_proj_y = sum_proj_y + (X_j %*% t(X_j)/sum(X_j^2)) %*% y_diamond
}

sum_proj_y
```

```
##           [,1]
## 1    4476.1881
## 2    4476.1881
## 3    4476.1881
## 4    1062.8515
## 5     137.4839
## 6     137.4839
## 7    1062.8515
## 8    2020.1621
## 9    4476.1881
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##	22	4476.1881
##	23	2020.1621
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```

```
print("...")
```

```
## [1] "..."
```

Convert this design matrix into Q, an orthonormal matrix.

```
#TO-DO
Q = matrix(NA, nrow=nrow(X_diamonds), ncol=ncol(X_diamonds))

Q[,1] = X_diamonds[,1]

for(j in 2:ncol(X_diamonds)){
  Q[,j] = X_diamonds[,j]
  for (k in 1:(j-1)){
    q_k = Q[,k , drop=FALSE]
    Q[,j] = X_diamonds[,j] - (q_k %*% t(q_k)/sum(q_k^2)) %*% X_diamonds[,j]
  }
}

# NORMALIZE
for (j in 1:ncol(X_diamonds)){
  Q[,j] = Q[,j] / sqrt(sum(Q[,j]^2))
}
```

Project the y vector onto each column of the Q matrix and test if the sum of these projections is the same as yhat.

```
#TO-DO
n = nrow(X_diamonds)
sum_proj_y = matrix(0, nrow=n, ncol=1)

for (j in 1:ncol(Q)) {
  q_j = Q[, j, drop=FALSE]
```



```

    sum_proj_y = sum_proj_y + (q_j %*% t(q_j)/sum(q_j^2)) %*% y_diamond
}

yhat_diamond = sum_proj_y

```

```

##           [,1]
## 1    -670.15854
## 2    -670.15854
## 3    -670.15854
## 4      72.18379
## 5    1134.59896
## 6    1134.59896
## 7      72.18379
## 8   -202.61394
## 9    -670.15854
## 10   -202.61394
## 11    1134.59896
## 12    1134.59896
## 13  -1003.83845
## 14    1134.59896
## 15    -670.15854
## 16    -670.15854
## 17      72.18379
## 18    1134.59896
## 19    1134.59896
## 20    1134.59896
## 21      72.18379
## 22    -670.15854
## 23   -202.61394
## 24    1134.59896
## 25    1134.59896
## 26   -108.90481
## 27      72.18379
## 28    1134.59896
## 29    1044.98173
## 30  -1003.83845
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## 32  -1003.83845
## 33    -670.15854
## 34    -670.15854
## 35    1044.98173
## 36  -1003.83845
## 37    -670.15854
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## 911      72.18379
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## 945      72.18379
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```

```
## 965    1044.98173
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## 993      72.18379
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## 995    -670.15854
## 996    1044.98173
## 997    -670.15854
## 998   -1003.83845
## 999    1044.98173
## 1000   1134.59896
```

```
print("...")
```

```
## [1] "..."
```

Find linear OLS estimates if Q is used as the design matrix using the `lm` method. Is the OLS solution the same as the OLS solution for X?

```
#T0-D0
model_vanilla = lm(y_diamond ~ 0 + X_diamonds)
b = coef(model_vanilla)

model_ortho = lm(y_diamond ~ 0 + Q)
b_q = coef(model_ortho)
print(b)
```

```
## X_diamonds(Intercept)      X_diamondscolor.L      X_diamondscolor.Q
```

```
##          2378.76684          -640.05819          -259.98348
##    X_diamondscolor.C    X_diamondscolor^4    X_diamondscolor^5
##          153.23342          223.99725          -16.12721
##    X_diamondscolor^6
##          36.89513
```

```
print(b_q)
```

```
##          Q1          Q2          Q3          Q4          Q5          Q6          Q7
## 77671.2630 -5715.4205 -3299.1850  887.8146 2685.8521 -310.5586 452.0611
```

Use the predict function and ensure that the predicted values are the same for both linear models: the one created with X as its design matrix and the one created with Q as its design matrix.

```
#TO-DO
y_hat_vanilla = predict(model_vanilla, data.frame(X_diamonds))
y_hat_ortho = predict(model_ortho, data.frame(Q))

table(sum(abs(y_hat_vanilla - y_hat_ortho)))
```

```
##
## 2.55363374890294e-09
##          1
```

Clear the workspace and load the boston housing data and extract X and y. The dimensions are $n = 506$ and $p = 13$. Create a matrix that is $(p + 1) \times (p + 1)$ full of NA's. Label the columns the same columns as X. Do not label the rows. For the first row, find the OLS estimate of the y regressed on the first column only and put that in the first entry. For the second row, find the OLS estimates of the y regressed on the first and second columns of X only and put them in the first and second entries. For the third row, find the OLS estimates of the y regressed on the first, second and third columns of X only and put them in the first, second and third entries, etc. For the last row, fill it with the full OLS estimates.

```
#TO-DO
rm(list = ls())

# Load the MASS package
library(MASS)
# Load the Boston housing data
data(Boston)

# EXTRACT X AND y
X = Boston[, !(names(Boston) %in% 'medv')]
y = Boston$medv

# Extract Dimensions
n = nrow(X)
p = ncol(X)

# Create NA matrix of p+1 dim
na_matrix = matrix(NA, nrow=p+1, ncol=p+1)
```

```

# rename matrix colnames
col_names = colnames(X)
colnames(na_matrix) = c("Intercept", col_names)

# Sequentially regress y on an increasing number of predictors
for (i in 1:p) {
  # Regress y on the first i predictors
  X_sub = X[, 1:i, drop = FALSE] # TAKE ONE COLUMN AT A TIME
  model = lm(y ~ ., data = as.data.frame(X_sub)) # FIT A MODEL USING THE COLUMNS REMOVED

  # i + 1 => Accounts for intercept
  # Fill out per column
  na_matrix[i + 1, 1:(i + 1)] = coef(model)
}

# Regress y on all predictors for the last row
model_full = lm(y ~ ., data = as.data.frame(X))
na_matrix[p + 1, ] = coef(model_full)

head(na_matrix)

```

```

##      Intercept      crim      zn      indus      chas      nox rm age dis
## [1,]         NA         NA         NA         NA         NA      NA NA  NA  NA
## [2,]  24.03311 -0.4151903         NA         NA         NA      NA NA  NA  NA
## [3,]  22.48563 -0.3520783  0.11610909         NA         NA      NA NA  NA  NA
## [4,]  27.39465 -0.2486283  0.05850082 -0.4155778         NA      NA NA  NA  NA
## [5,]  27.11280 -0.2287981  0.05928665 -0.4403251  6.894059      NA NA  NA  NA
## [6,]  29.48994 -0.2185190  0.05511047 -0.3834805  7.026223 -5.424659  NA  NA  NA
##      rad tax ptratio black lstat
## [1,]  NA  NA      NA      NA      NA
## [2,]  NA  NA      NA      NA      NA
## [3,]  NA  NA      NA      NA      NA
## [4,]  NA  NA      NA      NA      NA
## [5,]  NA  NA      NA      NA      NA
## [6,]  NA  NA      NA      NA      NA

```

Why are the estimates changing from row to row as you add in more predictors?

The line fitted has to adjust, with more features to consider the fit will be different leading to different coefficients.

Create a vector of length $p+1$ and compute the R^2 values for each of the above models.

```

#TO-DO
r_squared = numeric(p + 1)

# Loop through each subset of predictors and compute R^2
for (i in 1:p) {
  X_sub <- X[, 1:i, drop = FALSE]

```

```

model <- lm(y ~ ., data = as.data.frame(X_sub)) # Fit the model

# Calculate R^2 and store it
r_squared[i + 1] <- summary(model)$r.squared
}

```

```

model_full <- lm(y ~ ., data = Boston) # Fit the full model
r_squared[1] <- summary(model_full)$r.squared # Store

```

```

## Warning in summary.lm(model_full): essentially perfect fit: summary may be
## unreliable

```

```
print(r_squared)
```

```

## [1] 1.0000000 0.1507805 0.2339884 0.2937136 0.3295277 0.3313127 0.5873770
## [8] 0.5894902 0.6311488 0.6319479 0.6396628 0.6703141 0.6842043 0.7406427

```

Is R^2 monotonically increasing? Why?

It is increasing because the more features in our model seems to help the model understand the relationship by accounting for the variance. The more features the better we can fit to our data

Create a 2x2 matrix with the first column 1's and the next column iid normals. Find the absolute value of the angle (in degrees, not radians) between the two columns in absolute difference from 90 degrees.

```

n = 100

X = matrix(rnorm(2 * n), ncol = 2)
acos(t(X[,1]) %*% X[,2] / sqrt(sum(X[, 1]^2) * sum(X[, 2]^2))) * 180 / pi

## [1]
## [1,] 82.10683

```

Repeat this exercise $N_{\text{sim}} = 1e5$ times and report the average absolute angle.

```

Nsim = 1e5
n = 100
angles = numeric(Nsim)

for (i in 1:Nsim) {
  X = matrix(rnorm(2 * n), ncol = 2) # Create random matrix of size (2*n)(2)
  angle = acos(t(X[,1]) %*% X[,2] / sqrt(sum(X[, 1]^2) * sum(X[, 2]^2))) * 180 / pi # This line calculates the angle
  angles[i] = abs(angle - 90)
}

average_angle = mean(angles)
print(average_angle)

```

```
## [1] 4.6029
```

Create a $n \times 2$ matrix with the first column 1's and the next column iid normals. Find the absolute value of the angle (in degrees, not radians) between the two columns. For $n = 10, 50, 100, 200, 500, 1000$, report the average absolute angle over $N_{\text{sim}} = 1e5$ simulations.

```
#TO-DO
n_values = c(10, 50, 100, 200, 500, 1000)
average_angles = numeric(length(n_values))

for (i in seq_along(n_values)) {
  n = n_values[i]
  angles = numeric(Nsim)

  for (j in 1:Nsim) {
    X = cbind(1, rnorm(n)) # Create the n x 2 matrix with 1's in the first column and iid normals in the second
    angle = acos(t(X[,1]) %*% X[,2] / sqrt(sum(X[, 1]^2) * sum(X[, 2]^2))) * 180 / pi
    angles[j] <- abs(angle) # Absolute value of the angle
  }

  average_angles[i] <- mean(angles)
}

# Print the results
result = data.frame(n_values, average_angles)
print(result)
```

```
##   n_values average_angles
## 1      10      89.92826
## 2      50      89.98565
## 3     100      89.95665
## 4     200      89.99340
## 5     500      89.98811
## 6    1000      89.99306
```

What is this absolute angle difference from 90 degrees converging to? Why does this make sense?

the convergence of the average absolute angle to 0 as n increases makes sense because it reflects the fact that the angle between the two columns of the matrix tends to be close to 90 degrees when n is large.