

MA334 Assignment Report

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1. Data Exploration

(a) Summary of the Data Set

```
library(knitr)

# Load the data
data <- read.csv("MA334-AU-7_2501629.csv")
data_pre <- head(data)

kable(data_pre, caption = "First 6 Observations")
```

Table 1: First 6 Observations

| carat | cut | color | clarity | depth | table | price | x | y | z |
|-------|-----------|-------|---------|-------|-------|-------|------|------|------|
| 0.53 | Premium | D | SI1 | 61.4 | 58 | 1566 | 5.19 | 5.23 | 3.20 |
| 0.85 | Ideal | G | SI1 | 62.2 | 56 | 3403 | 6.06 | 6.09 | 3.78 |
| 0.56 | Very Good | E | VS1 | 62.7 | 60 | 1931 | 5.25 | 5.28 | 3.30 |
| 0.70 | Very Good | I | VS1 | 63.3 | 55 | 1995 | 5.64 | 5.60 | 3.56 |
| 1.01 | Premium | I | SI2 | 62.6 | 58 | 3818 | 6.43 | 6.38 | 4.01 |
| 1.02 | Very Good | F | VS2 | 63.3 | 58 | 6007 | 6.36 | 6.38 | 4.03 |

```
# Number of observations and variables
no_of_observations <- nrow(data)
no_of_variables <- ncol(data)

# Creating a table to show the size of the dataset
dimensions_table <- data.frame(
  Item = c("Total Observations", "Total Variables"),
  Value = c(no_of_observations, no_of_variables)
)

kable(dimensions_table, caption = "Dataset Dimensions")
```

Table 2: Dataset Dimensions

| Item | Value |
|--------------------|-------|
| Total Observations | 1000 |
| Total Variables | 10 |

(Dataset Dimensions)

Table 1 shows that the dataset contains **1,000 observations** and **10 variables**.

```
# Variable names and types
variable_types <- data.frame(
  Variable = names(data),
  Type = sapply(data, class),
  row.names = NULL
)

kable(variable_types, caption = "Variable Names and Data Types")
```

Table 3: Variable Names and Data Types

| Variable | Type |
|----------|-----------|
| carat | numeric |
| cut | character |
| color | character |
| clarity | character |
| depth | numeric |
| table | numeric |
| price | integer |
| x | numeric |
| y | numeric |
| z | numeric |

(Variable Types)

Table 2 shows all the variables in the dataset and their data types. Some variables contain numbers (such as carat, depth, price, and size measurements), while others contain categories (cut, color, and clarity).

```
# Identifying qualitative variables
categorical_vars <- names(data)[sapply(data, function(x)
  is.character(x) | is.factor(x))]

# Converting qualitative variables to factors
data[categorical_vars] <- lapply(data[categorical_vars], as.factor)
```

```

# Create table for categories
categories_table <- do.call(rbind, lapply(categorical_vars, function(v) {
  data.frame(
    Variable = v,
    Categories = paste(levels(data[[v]]), collapse = ", "),
    stringsAsFactors = FALSE
  )
}))

kable(categories_table, caption = "Categories for Qualitative Variables")

```

Table 4: Categories for Qualitative Variables

| Variable | Categories |
|----------|--|
| cut | Fair, Good, Ideal, Premium, Very Good |
| color | D, E, F, G, H, I, J |
| clarity | I1, IF, SI1, SI2, VS1, VS2, VVS1, VVS2 |

(Categories)

Table 3 shows the categories for each qualitative variable. The variable **cut** has 5 categories, **color** has 7 categories, and **clarity** has 8 categories. These categories represent different levels of diamond quality.

(b) Location and Spread of Numeric Variables

```

# Select numeric variables only
numeric_data <- data[sapply(data, is.numeric)]

# Create a table of summary statistics
summary_table <- data.frame(
  Variable = names(numeric_data),
  Mean = sapply(numeric_data, mean),
  Median = sapply(numeric_data, median),
  Standard_Deviation = sapply(numeric_data, sd),
  Range = sapply(numeric_data, function(x) max(x) - min(x)),
  row.names = NULL
)

kable(summary_table, caption = "Summary Statistics for Numeric Variables")

```

Table 5: Summary Statistics for Numeric Variables

| Variable | Mean | Median | Standard_Deviation | Range |
|----------|----------|--------|--------------------|-------|
| carat | 0.78173 | 0.70 | 0.4603453 | 2.88 |
| depth | 61.75350 | 61.90 | 1.3726660 | 11.80 |

| Variable | Mean | Median | Standard_Deviation | Range |
|----------|------------|---------|--------------------|----------|
| table | 57.44450 | 57.00 | 2.1992604 | 24.00 |
| price | 3786.08000 | 2278.50 | 3833.2003865 | 18403.00 |
| x | 5.69129 | 5.64 | 1.1116064 | 9.15 |
| y | 5.69468 | 5.64 | 1.1025724 | 9.02 |
| z | 3.51608 | 3.46 | 0.6894836 | 5.98 |

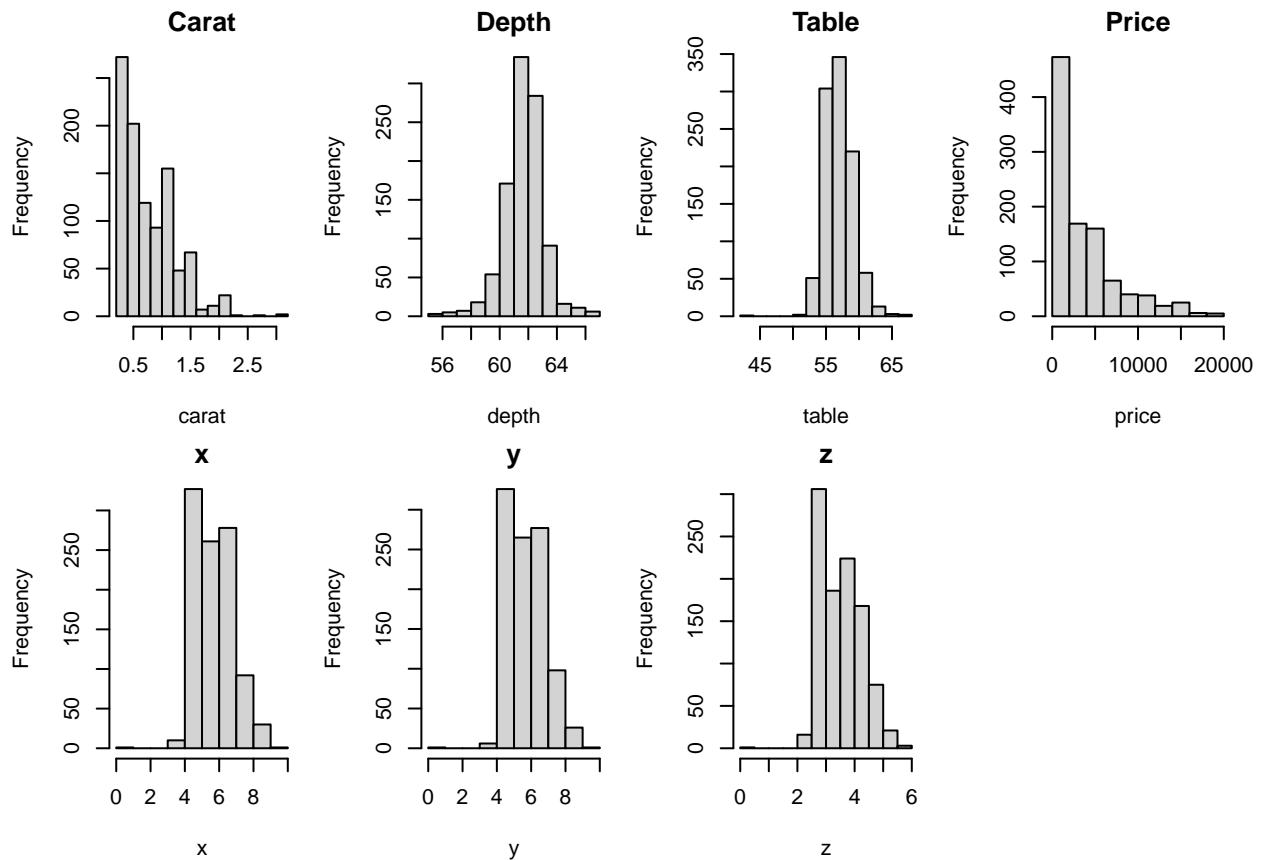
This table shows typical values (mean and median) and how spread out the values are (standard deviation and range) for each numeric variable.

(c) Visualisations of Variable Distributions

```
# ----- Numeric variables: Histograms (including z) -----
par(mfrow = c(2, 4), mar = c(4, 4, 2, 1))

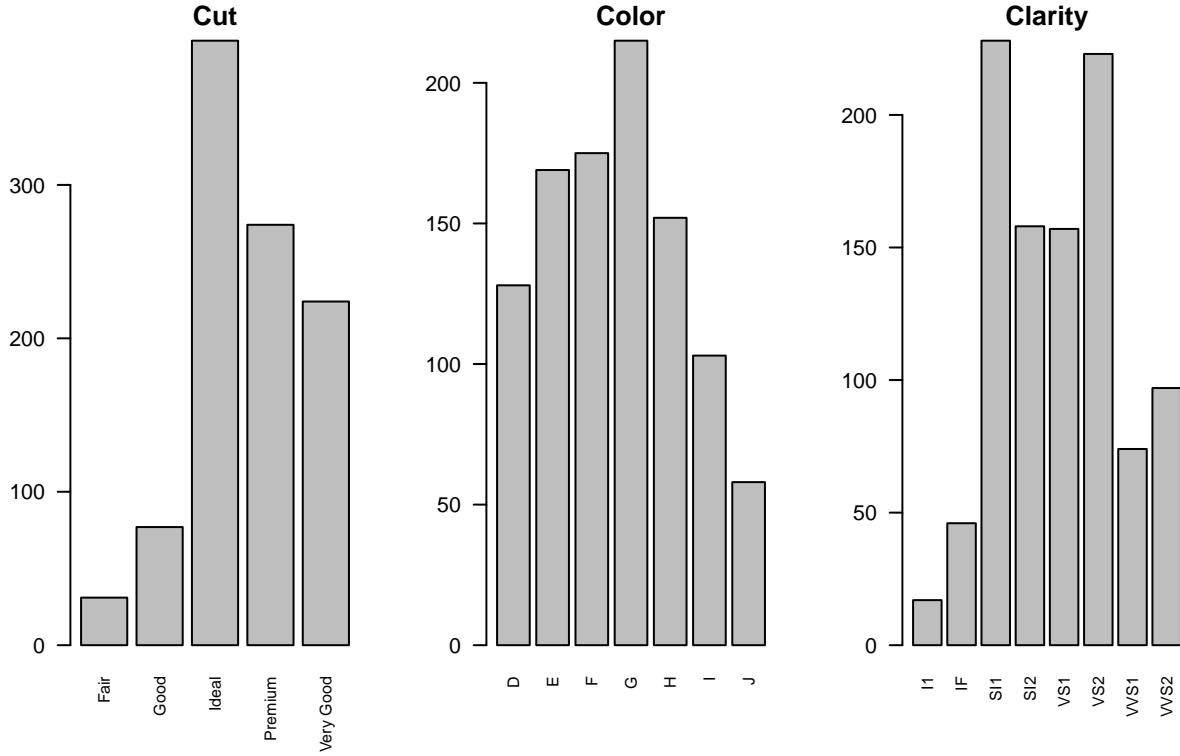
hist(data$carat, main = "Carat", xlab = "carat")
hist(data$depth, main = "Depth", xlab = "depth")
hist(data$table, main = "Table", xlab = "table")
hist(data$price, main = "Price", xlab = "price")
hist(data$x, main = "x", xlab = "x")
hist(data$y, main = "y", xlab = "y")
hist(data$z, main = "z", xlab = "z")

# Leave one panel empty (2x4 layout has 8 slots)
plot.new()
```



```
# ----- Categorical variables: Bar plots (labels visible) -----
par(mfrow = c(1, 3), mar = c(8, 4, 2, 1))

barplot(table(data$cut), main = "Cut", las = 2, cex.names = 0.8)
barplot(table(data$color), main = "Color", las = 2, cex.names = 0.8)
barplot(table(data$clarity), main = "Clarity", las = 2, cex.names = 0.8)
```



```

par(mfrow = c(1, 1))

### (d) Produce and interpret a correlation matrix between the numeric variables

# Select numeric variables only
numeric_data <- data[sapply(data, is.numeric)]

# Create correlation matrix
cor_matrix <- cor(numeric_data)

# Display correlation matrix
knitr::kable(round(cor_matrix, 2),
             caption = "Correlation Matrix of Numeric Variables")

```

Table 6: Correlation Matrix of Numeric Variables

| | carat | depth | table | price | x | y | z |
|-------|-------|-------|-------|-------|-------|-------|------|
| carat | 1.00 | 0.05 | 0.21 | 0.91 | 0.96 | 0.96 | 0.96 |
| depth | 0.05 | 1.00 | -0.31 | 0.00 | -0.03 | -0.03 | 0.09 |
| table | 0.21 | -0.31 | 1.00 | 0.15 | 0.23 | 0.22 | 0.18 |
| price | 0.91 | 0.00 | 0.15 | 1.00 | 0.85 | 0.85 | 0.84 |
| x | 0.96 | -0.03 | 0.23 | 0.85 | 1.00 | 1.00 | 0.99 |
| y | 0.96 | -0.03 | 0.22 | 0.85 | 1.00 | 1.00 | 0.99 |

| | carat | depth | table | price | x | y | z |
|---|-------|-------|-------|-------|------|------|------|
| z | 0.96 | 0.09 | 0.18 | 0.84 | 0.99 | 0.99 | 1.00 |

```
# Question 2 (Final)
# =====

# (a) Probabilities for one randomly chosen diamond
p_price_gt_10000 <- mean(data$price > 10000)      # i)
p_ideal <- mean(data$cut == "Ideal")                 # ii)

p_price_gt_10000

## [1] 0.093

p_ideal

## [1] 0.394

# (b) From a sample of 20 diamonds, P(more than 12 are Ideal)
# X ~ Binomial(n = 20, p = p_ideal)
prob_more_than_12_ideal <- 1 - pbinom(12, size = 20, prob = p_ideal)

prob_more_than_12_ideal

## [1] 0.01834124

# (c) 90% and 95% confidence intervals for mean carat weight (t-interval)
n <- length(data$carat)
xbar <- mean(data$carat)
s <- sd(data$carat)

ci_90 <- xbar + c(-1, 1) * qt(0.95, df = n - 1) * s / sqrt(n)
ci_95 <- xbar + c(-1, 1) * qt(0.975, df = n - 1) * s / sqrt(n)

ci_90

## [1] 0.757763 0.805697

ci_95

## [1] 0.7531634 0.8102966

# Which interval is wider?
width_90 <- ci_90[2] - ci_90[1]
width_95 <- ci_95[2] - ci_95[1]

width_90

## [1] 0.04793403
```

```

width_95

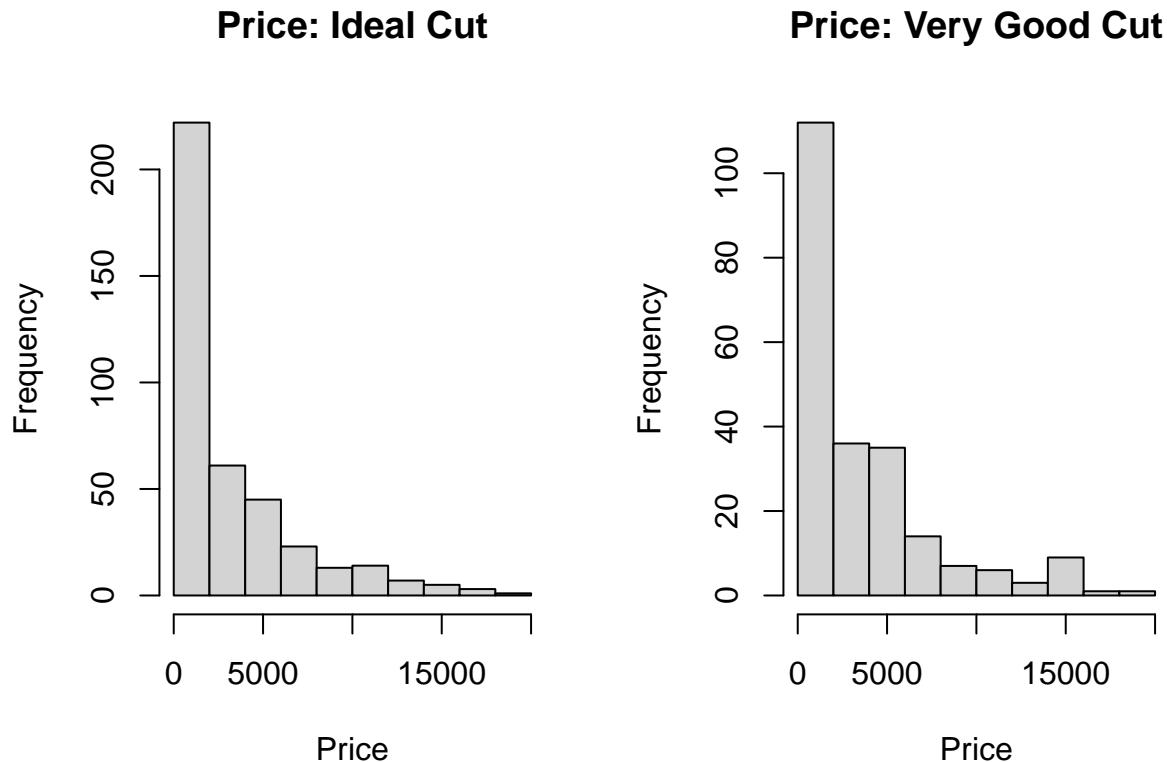
## [1] 0.05713317

# =====
# Question 3: Hypothesis Tests (a-e) in ONE chunk
# =====

# (a) i-ii: Distributions of price for Ideal and Very Good
ideal_price <- data$price[data$cut == "Ideal"]
vg_price <- data$price[data$cut == "Very Good"]

par(mfrow = c(1, 2))
hist(ideal_price, main = "Price: Ideal Cut", xlab = "Price")
hist(vg_price, main = "Price: Very Good Cut", xlab = "Price")

```



```

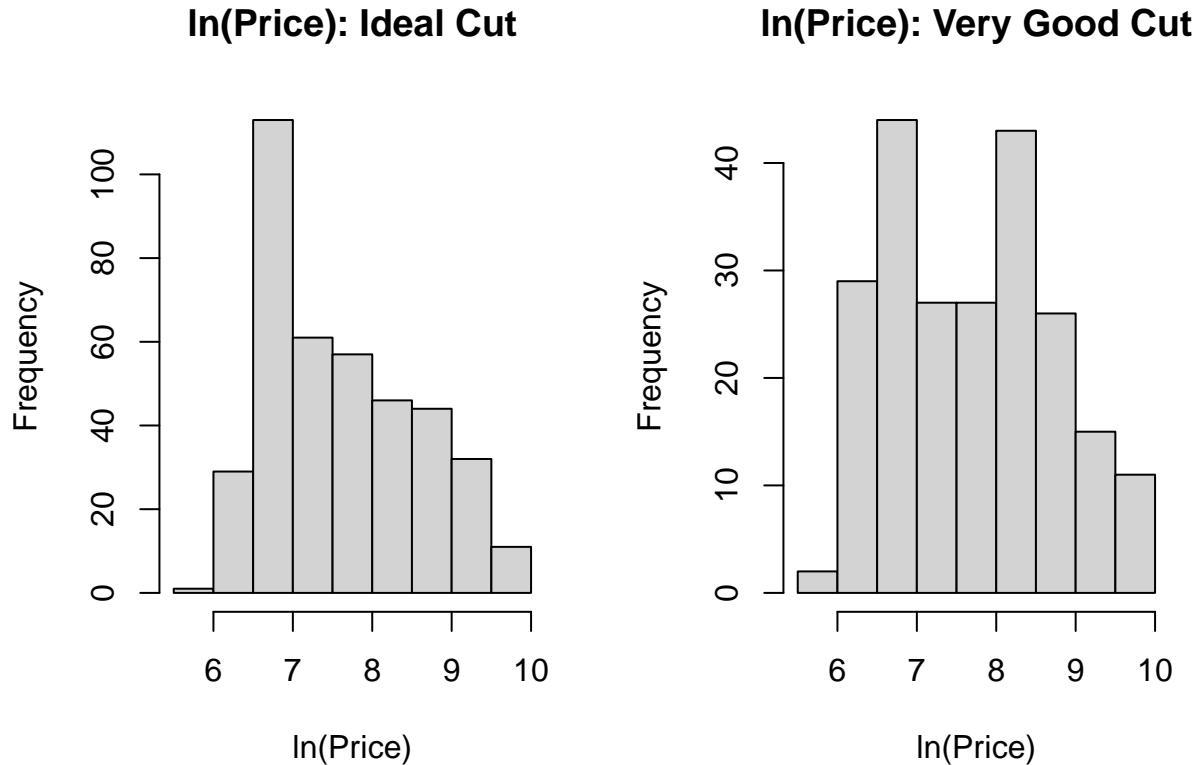
par(mfrow = c(1, 1))

# (a) iv: Create ln(price) and repeat plots
data$ln_price <- log(data$price)
ideal_ln <- data$ln_price[data$cut == "Ideal"]
vg_ln <- data$ln_price[data$cut == "Very Good"]

par(mfrow = c(1, 2))

```

```
hist(ideal_ln, main = "ln(Price): Ideal Cut", xlab = "ln(Price)")
hist(vg_ln, main = "ln(Price): Very Good Cut", xlab = "ln(Price)")
```



```
par(mfrow = c(1, 1))

# (b) t-test on price (5% level)
# H0: mean price (Ideal) = mean price (Very Good)
# H1: mean price (Ideal) != mean price (Very Good)
t_price <- t.test(ideal_price, vg_price, alternative = "two.sided", conf.level = 0.95)
t_price

##
##  Welch Two Sample t-test
##
## data: ideal_price and vg_price
## t = -1.2234, df = 431.73, p-value = 0.2219
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -1015.3727 236.2943
## sample estimates:
## mean of x mean of y
## 3275.563 3665.103
```

```

# (c) t-test on ln(price) (5% level)
# H0: mean ln(price) (Ideal) = mean ln(price) (Very Good)
# H1: mean ln(price) (Ideal) != mean ln(price) (Very Good)
t_ln <- t.test(ideal_ln, vg_ln, alternative = "two.sided", conf.level = 0.95)
t_ln

```

```

##
## Welch Two Sample t-test
##
## data: ideal_ln and vg_ln
## t = -0.92072, df = 432.88, p-value = 0.3577
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.24054911 0.08707367
## sample estimates:
## mean of x mean of y
## 7.620497 7.697235

```

```

# (d) Compare conclusions using p-values
t_price$p.value

```

```

## [1] 0.2218553

```

```

t_ln$p.value

```

```

## [1] 0.3577081

```

```

# (e) Chi-square independence test: price group (>5000) vs clarity
data$price_group <- ifelse(data$price > 5000, "High", "Low")
cont_table <- table(data$price_group, data$clarity)
cont_table

```

```

##
##          I1   IF SI1 SI2 VS1 VS2 VVS1 VVS2
## High     6    8  64  37  41  76    7   13
## Low     11   38 164 121 116 147   67   84

```

```

chisq.test(cont_table)

```

```

##
## Pearson's Chi-squared test
##
## data: cont_table
## X-squared = 29.959, df = 7, p-value = 9.661e-05

```

```

# =====
# Question 4: Linear Regression
# =====

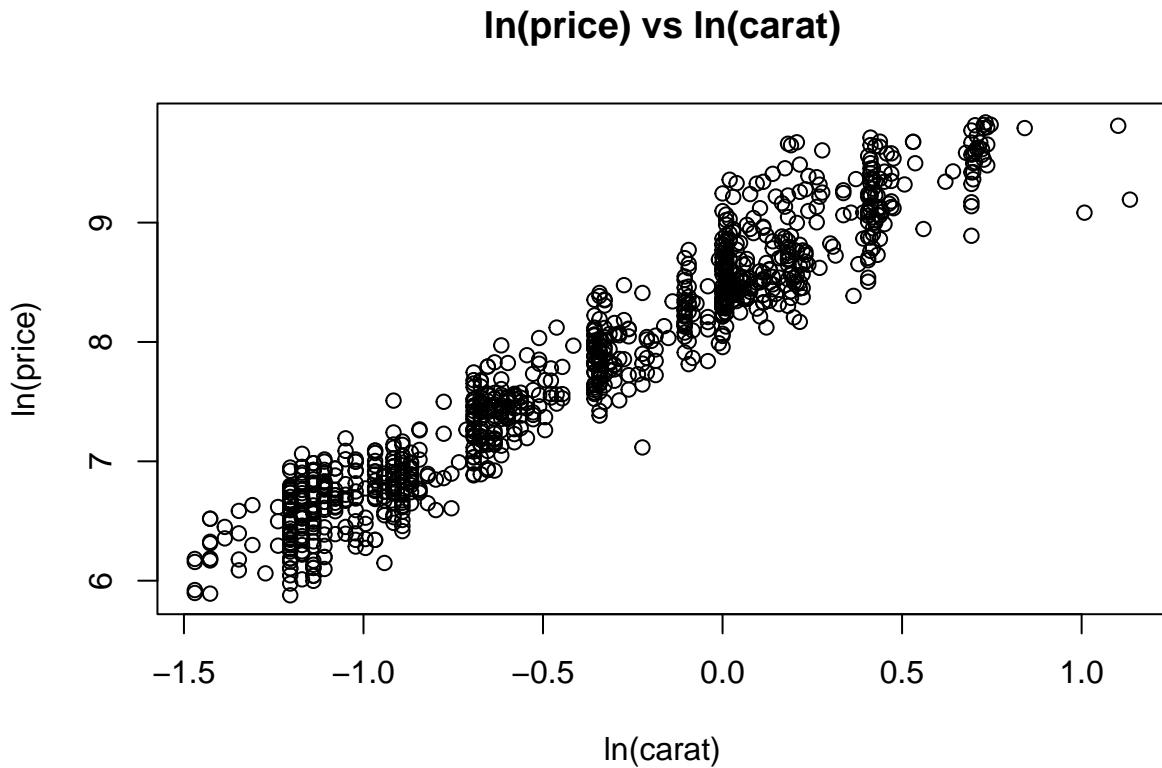
```

```

# Create log variables
data$ln_price <- log(data$price)
data$ln_carat <- log(data$carat)

# (a) Scatter plot of ln(price) vs ln(carat)
plot(data$ln_carat, data$ln_price,
      xlab = "ln(carat)", ylab = "ln(price)",
      main = "ln(price) vs ln(carat)")

```



```

# (b) Simple regression: ln(price) ~ ln(carat)
m1 <- lm(ln_price ~ ln_carat, data = data)
s1 <- summary(m1)

# (b)(i) slope
slope_m1 <- coef(m1)["ln_carat"]
slope_m1

## ln_carat
## 1.656087

# (b)(ii) p-value for ln(carat)
pval_m1 <- s1$coefficients["ln_carat", "Pr(>|t|)"]
pval_m1

## [1] 0

```

```

# (b)(iii) R-squared
r2_m1 <- s1$r.squared
r2_m1

## [1] 0.9279249

# (c) Multiple regression: ln(price) ~ ln(carat) + cut
m2 <- lm(ln_price ~ ln_carat + cut, data = data)
s2 <- summary(m2)
s2

##
## Call:
## lm(formula = ln_price ~ ln_carat + cut, data = data)
##
## Residuals:
##       Min      1Q  Median      3Q     Max
## -0.86330 -0.15805 -0.01857  0.16553  0.89623
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 8.15080   0.04588 177.644 < 2e-16 ***
## ln_carat    1.67895   0.01448 115.937 < 2e-16 ***
## cutGood     0.16828   0.05480  3.071  0.00219 **
## cutIdeal    0.36339   0.04851  7.491 1.51e-13 ***
## cutPremium   0.30738   0.04869  6.313 4.12e-10 ***
## cutVery Good 0.28626   0.04956  5.776 1.02e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.2553 on 994 degrees of freedom
## Multiple R-squared:  0.9336, Adjusted R-squared:  0.9333
## F-statistic:  2797 on 5 and 994 DF,  p-value: < 2.2e-16

# (d) Significant variables at 1% level
sig_1pct <- rownames(s2$coefficients)[s2$coefficients[, "Pr(>|t|)"] < 0.01]
sig_1pct

## [1] "(Intercept)"    "ln_carat"        "cutGood"         "cutIdeal"        "cutPremium"
## [6] "cutVery Good"

# (e) Hypothesis test for ln(carat) in model (c)
# H0: beta_ln_carat = 0
# H1: beta_ln_carat != 0

# (f) Result of the test (t-statistic and p-value for ln(carat) in model (c))
s2$coefficients["ln_carat", ]

##           Estimate Std. Error      t value      Pr(>|t|) 
## 1 1.67895120 0.01448158 115.93701166 0.00000000
```