

final_project

```
library(tidyverse)
```

Warning: package 'ggplot2' was built under R version 4.3.2

Warning: package 'tidyr' was built under R version 4.3.2

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr      1.1.4      v readr      2.1.4
v forcats    1.0.0      v stringr    1.5.1
v ggplot2    3.5.0      v tibble     3.2.1
v lubridate  1.9.3      v tidyr      1.3.1
v purrr      1.0.2
```

```
-- Conflicts ----- tidyverse_conflicts() --
```

```
x dplyr::filter() masks stats::filter()
```

```
x dplyr::lag()     masks stats::lag()
```

```
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become
```

```
simCorn <- function(overallEffect=0, fertilizerEffect=c(0,0,0), rowEffect=c(0,0,0), colEff
                      seed=NULL, dist = rnorm, ...) {
```

```
  if ((!is.numeric(seed)) && (!is.null(seed)))
    stop("You did not enter a valid seed")
```

```
  n <- length(fertilizerEffect)^2
```

```
  set.seed(seed)
  error <- dist(n, ...)
```

```

Fertilizer = factor(c("A", "B", "C", "C", "A", "B", "B", "C", "A"))
Row = factor(c(1, 1, 1, 2, 2, 2, 3, 3, 3))
Column = factor(c(1, 2, 3, 1, 2, 3, 1, 2, 3))
Yield = overallEffect + fertilizerEffect + rowEffect + colEffect + error
x <- data.frame(Fertilizer, Row, Column, Yield)

  return (x)
}

simCorn()

```

	Fertilizer	Row	Column	Yield
1	A	1	1	-0.44749897
2	B	1	2	-0.03641480
3	C	1	3	-0.51516386
4	C	2	1	0.02246551
5	A	2	2	-0.65584660
6	B	2	3	0.26256326
7	B	3	1	2.63643116
8	C	3	2	0.96921027
9	A	3	3	0.72645438

```
simCorn(overallEffect=10,seed=2123,dist=rgamma,shape=2)
```

	Fertilizer	Row	Column	Yield
1	A	1	1	14.83727
2	B	1	2	10.45424
3	C	1	3	13.13900
4	C	2	1	10.47095
5	A	2	2	10.90779
6	B	2	3	13.44940
7	B	3	1	10.77832
8	C	3	2	11.49251
9	A	3	3	10.62710

```

mu <- 7
alpha <- c(1,2,3)
beta <- c(2,2,1)

```

```

gamma <- c(3,3,2)
y <- simCorn(overallEffect=mu, fertilizerEffect=alpha, rowEffect=beta, colEffect=gamma,
             seed=29429, rnorm, mean=3, sd=2)
y

```

	Fertilizer	Row	Column	Yield
1	A	1	1	19.97551
2	B	1	2	16.08501
3	C	1	3	16.82301
4	C	2	1	13.00537
5	A	2	2	16.31535
6	B	2	3	13.89126
7	B	3	1	15.24435
8	C	3	2	16.98722
9	A	3	3	19.10382

```

pValue <- numeric(100)
p_vector_1 <- vector(mode = "numeric", length = 100)
set.seed(1331)
for (k in 1:100) {

  y <- simCorn(overallEffect=10, dist=rnorm)
  fitCorn <- lm(Yield ~ Fertilizer + Row + Column, data=y)
  pValue[k] <- anova(fitCorn)$"Pr(>F)"[1]
  p_vector_1[k] <- pValue[k]

}

p_vector_2 <- vector(mode = "numeric", length = 100)
set.seed(18694)

for (k in 1:100) {

  y <- simCorn(overallEffect=10, c(1,2,3), c(0,0,1), c(0,0,1), dist=rnorm)
  fitCorn <- lm(Yield ~ Fertilizer + Row + Column, data=y)
  pValue[k] <- anova(fitCorn)$"Pr(>F)"[1]
  p_vector_2[k] <- pValue[k]

}

```

```

}

p_vector_3 <- vector(mode = "numeric", length = 100)
set.seed(6516)

for (k in 1:100) {

  y <- simCorn(overallEffect=10, c(1,2,3), c(1,0,1), c(0,1,1), dist=rnorm)
  fitCorn <- lm(Yield ~ Fertilizer + Row + Column, data=y)
  pValue[k] <- anova(fitCorn)$"Pr(>F)"[1]
  p_vector_3[k] <- pValue[k]

}

p_vector_4 <- vector(mode = "numeric", length = 100)
set.seed(5)

for (k in 1:100) {

  y <- simCorn(overallEffect=10, c(1,2,3), c(1,0,1), c(0,1,1), dist=rnorm)
  fitCorn <- lm(Yield ~ Fertilizer + Row + Column, data=y)
  pValue[k] <- anova(fitCorn)$"Pr(>F)"[1]
  p_vector_4[k] <- pValue[k]

}

p_vector_5 <- vector(mode = "numeric", length = 100)
set.seed(574)
for (k in 1:100) {

  y <- simCorn(overallEffect=10, c(1,2,3), c(0,0,1), c(0,0,1), dist=rexp)
  fitCorn <- lm(Yield ~ Fertilizer + Row + Column, data=y)
  pValue[k] <- anova(fitCorn)$"Pr(>F)"[1]
  p_vector_5[k] <- pValue[k]

}

```

```

p_vector_6 <- vector(mode = "numeric", length = 100)
set.seed(9576)

for (k in 1:100) {

  y <- simCorn(overallEffect=10, c(1,2,3), c(1,0,1), c(0,1,1), dist=rexp)
  fitCorn <- lm(Yield ~ Fertilizer + Row + Column, data=y)
  pValue[k] <- anova(fitCorn)$"Pr(>F)"[1]
  p_vector_6[k] <- pValue[k]

}

p_vector_7 <- vector(mode = "numeric", length = 100)
set.seed(9743)

for (k in 1:100) {

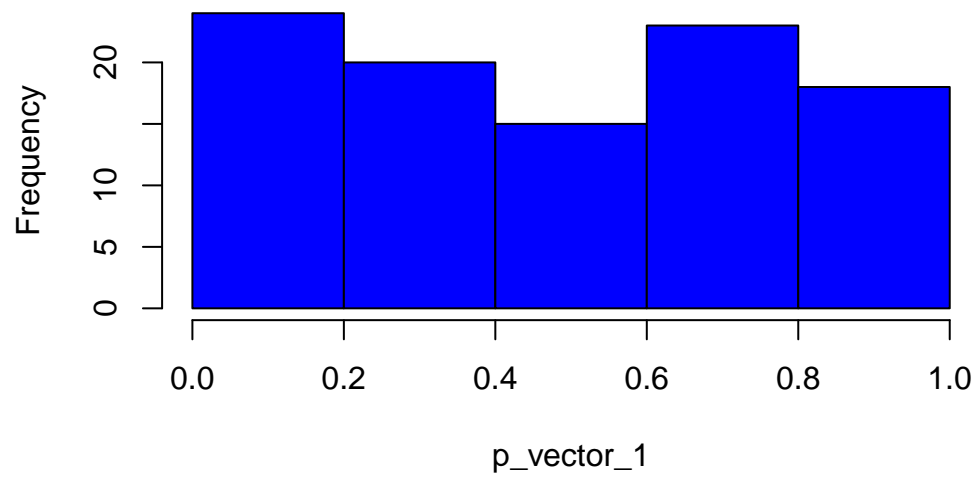
  y <- simCorn(overallEffect=10, c(1,2,3), c(0,1,0), c(0,1,0), dist=rexp)
  fitCorn <- lm(Yield ~ Fertilizer + Row + Column, data=y)
  pValue[k] <- anova(fitCorn)$"Pr(>F)"[1]
  p_vector_5[k] <- pValue[k]

}

hist(p_vector_1, breaks = c(0.0, 0.2, 0.4, 0.6, 0.8, 1.0), col = "blue", )

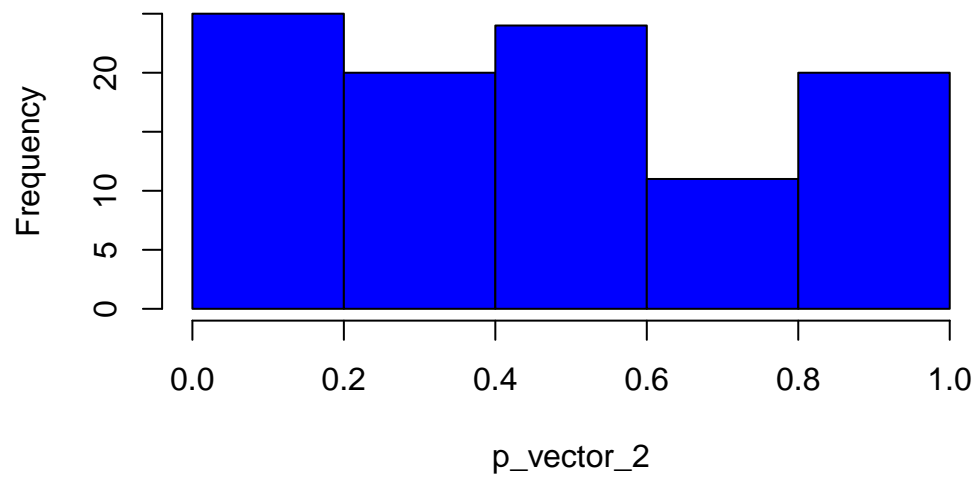
```

Histogram of p_vector_1



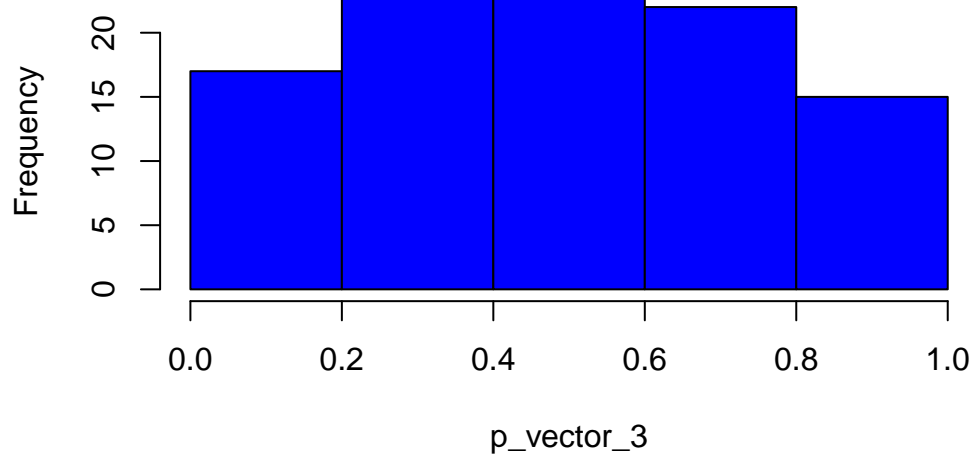
```
hist(p_vector_2, breaks = c(0.0, 0.2, 0.4, 0.6, 0.8, 1.0), col = "blue", )
```

Histogram of p_vector_2



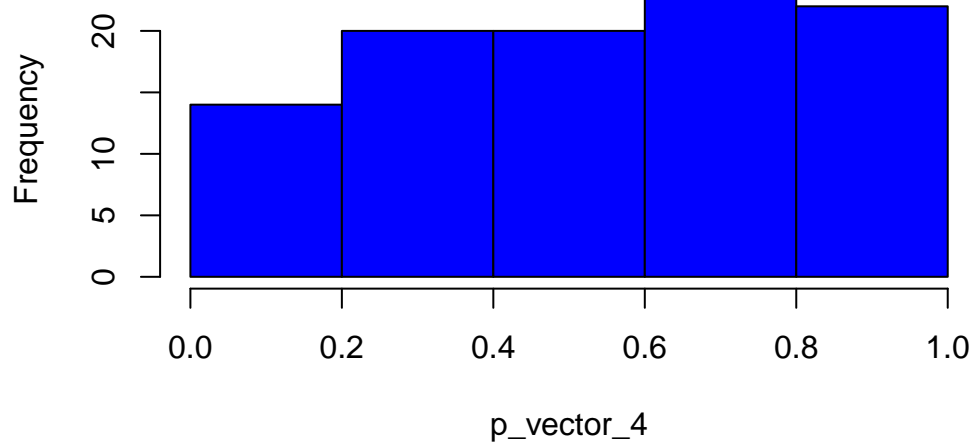
```
hist(p_vector_3, breaks = c(0.0, 0.2, 0.4, 0.6, 0.8, 1.0), col = "blue", )
```

Histogram of p_vector_3



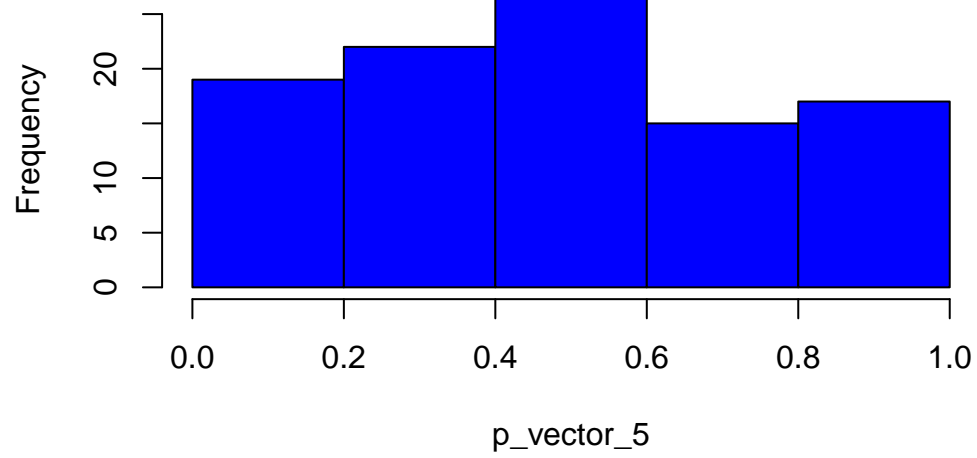
```
hist(p_vector_4, breaks = c(0.0, 0.2, 0.4, 0.6, 0.8, 1.0), col = "blue", )
```

Histogram of p_vector_4



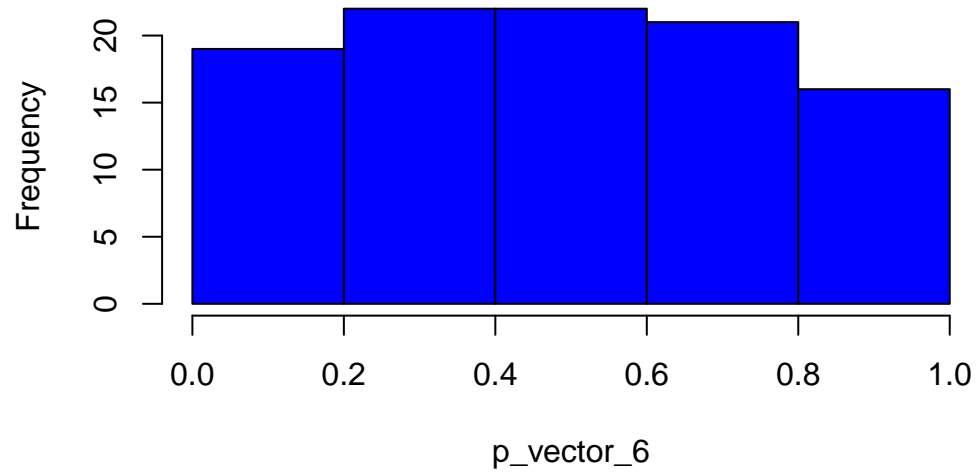
```
hist(p_vector_5, breaks = c(0.0, 0.2, 0.4, 0.6, 0.8, 1.0), col = "blue", )
```

Histogram of p_vector_5



```
hist(p_vector_6, breaks = c(0.0, 0.2, 0.4, 0.6, 0.8, 1.0), col = "blue", )
```

Histogram of p_vector_6



```
hist(p_vector_7, breaks = c(0.0, 0.2, 0.4, 0.6, 0.8, 1.0), col = "blue", )
```