

# DVM Reproducible Research

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## CO2 Uptake in Grass Plants

*This document provides a reproducible research of “Carbon Dioxide Uptake in Grass Plants” from a library dataset in “R” called “datasets”.*

*The research was compiled into a initial data frame called “CO2”*

*CO2 has 84 rows and 5 columns of data from an experiment on the cold tolerance of the grass species “Echinochloa crus-galli”.*

*CO2 was uptaken within six (6) plants from Quebec and six (6) plants from Mississippi.*

*It was measured at several levels of ambient CO2 concentration. Half the plants of each type were chilled overnight before the experiment was conducted.*

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## Summary of CO2

```
#install.packages("datasets")
suppressMessages(suppressWarnings(library(datasets)))
data(CO2)
summary(CO2)
```

```
##      Plant      Type      Treatment      conc
## Qn1      : 7  Quebec      :42  nonchilled:42  Min.      : 95
## Qn2      : 7  Mississippi:42  chilled   :42  1st Qu.: 175
## Qn3      : 7                                     Median : 350
## Qc1      : 7                                     Mean    : 435
## Qc3      : 7                                     3rd Qu.: 675
## Qc2      : 7                                     Max.    :1000
## (Other):42
##      uptake
## Min.      : 7.70
## 1st Qu.:17.90
## Median :28.30
## Mean     :27.21
## 3rd Qu.:37.12
## Max.     :45.50
##
```

## Summary Explained

- **Plant:** *an ordered factor with levels  $Qn1 < Qn2 < Qn3 < \dots < Mc1$  giving a unique identifier for each plant.*
- **Type:** *a factor with levels Quebec and Mississippi giving the origin of the plant*

- **Treatment:** a factor with levels nonchilled chilled
  - **Conc:** a numeric vector of ambient carbon dioxide concentrations (mL/L).
  - **Uptake:** a numeric vector of carbon dioxide uptake rates (umol/m<sup>2</sup> sec).
- 

## Relationship between “Type” and “Treatment”

### Table of CO2-Type

```
table(CO2$Type)
```

```
##
##      Quebec Mississippi
##      42             42
```

### Table of CO2-Treatment

```
table(CO2$Treatment)
```

```
##
## nonchilled    chilled
##      42       42
```

### Table of CO2-Type and Treatment

```
table(CO2$Type,CO2$Treatment)
```

```
##
##              nonchilled chilled
##   Quebec             21      21
##   Mississippi         21      21
```

---

## Cross Table of CO2 - Type and Treatment

### Relationship of Type and Treatment using the Cross Tabulations

```
#install.packages("gmodels")
suppressMessages(suppressWarnings(library("gmodels")))
#Contingency tables
joint.CO2<-CrossTable(CO2$Type,CO2$Treatment, prop.chisq = F)
```

```
##
##
##   Cell Contents
## |-----|
## |                      N |
## |          N / Row Total |
```

```

## |          N / Col Total |
## |          N / Table Total |
## |-----|
##
##
## Total Observations in Table:  84
##
##
##          | C02$Treatment
##    C02$Type | nonchilled |      chilled | Row Total |
## -----|-----|-----|-----|
##    Quebec |         21 |         21 |         42 |
##           |         0.500 |         0.500 |         0.500 |
##           |         0.500 |         0.500 |           |
##           |         0.250 |         0.250 |           |
## -----|-----|-----|-----|
## Mississippi |         21 |         21 |         42 |
##            |         0.500 |         0.500 |         0.500 |
##            |         0.500 |         0.500 |           |
##            |         0.250 |         0.250 |           |
## -----|-----|-----|-----|
## Column Total |         42 |         42 |         84 |
##            |         0.500 |         0.500 |           |
## -----|-----|-----|-----|
##
##

```

- **Note:** \*Based on PROC Frequency “PROC FREQ” in SAS or “CROSSTABS” in SPSS“\*

## Nested Contingency Tables

```
joint.C02
```

```

## $t
##           y
## x         nonchilled chilled
## Quebec           21      21
## Mississippi       21      21
##
## $prop.row
##           y
## x         nonchilled chilled
## Quebec           0.5      0.5
## Mississippi       0.5      0.5
##
## $prop.col
##           y
## x         nonchilled chilled
## Quebec           0.5      0.5
## Mississippi       0.5      0.5
##
## $prop.tbl
##           y
## x         nonchilled chilled

```

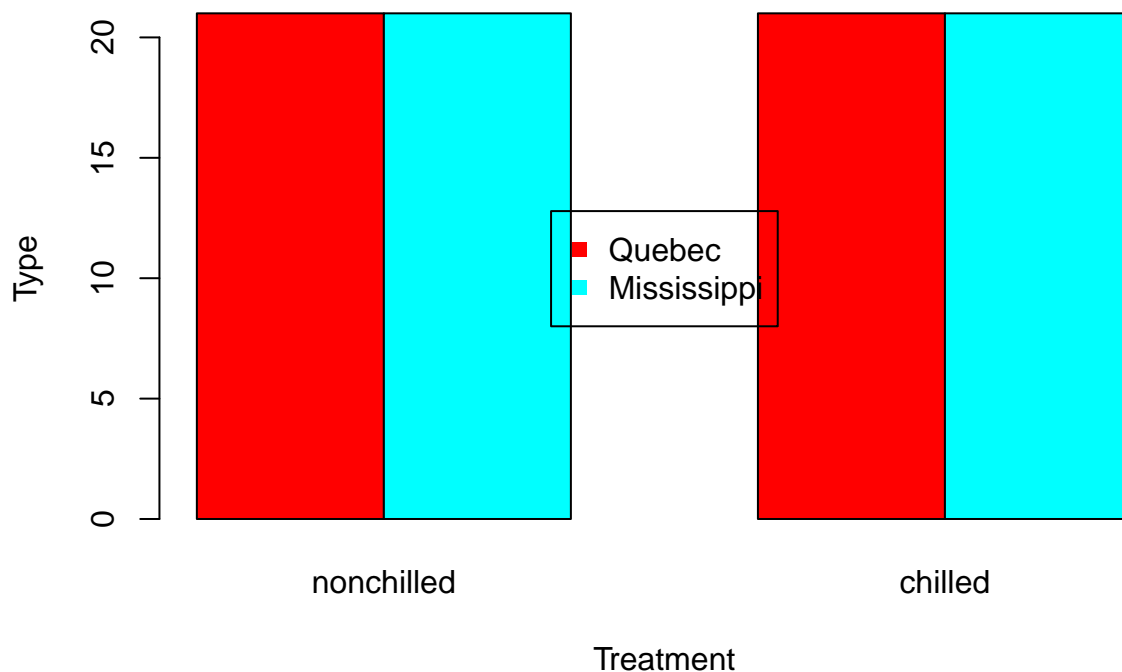
```
## Quebec      0.25    0.25
## Mississippi 0.25    0.25
```

- **Note:** \*Based on PROC Frequency “PROC FREQ” in SAS or “CROSSTABS” in SPSS“\*

---

## Barplot Relationship of “Type” and “Treatment”

```
par(mfrow=c(1,1))
joint.CO2_count<-joint.CO2$t
barplot(joint.CO2_count, beside = TRUE, col = rainbow(2),
        ylab = "Type", xlab = "Treatment")
legend("center", c("Quebec", "Mississippi"),
       pch = 15, col = rainbow(2))
```

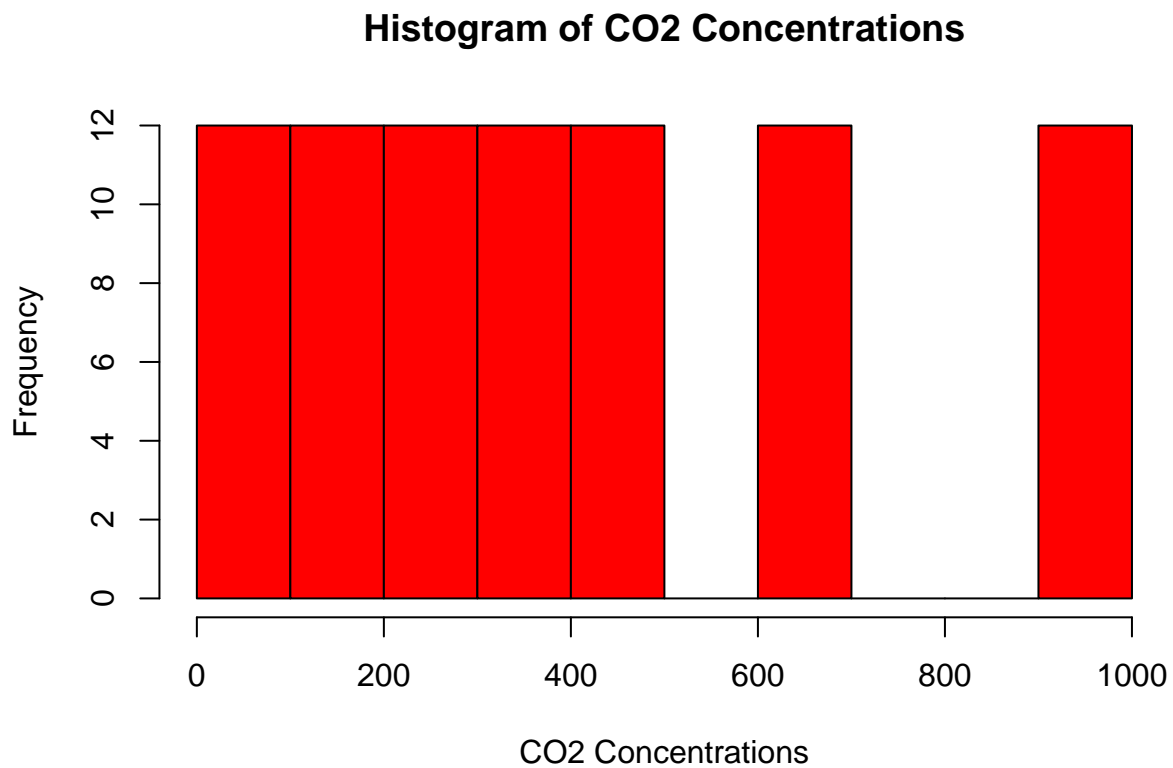



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## “Conc” and “UpTake” using Histograms

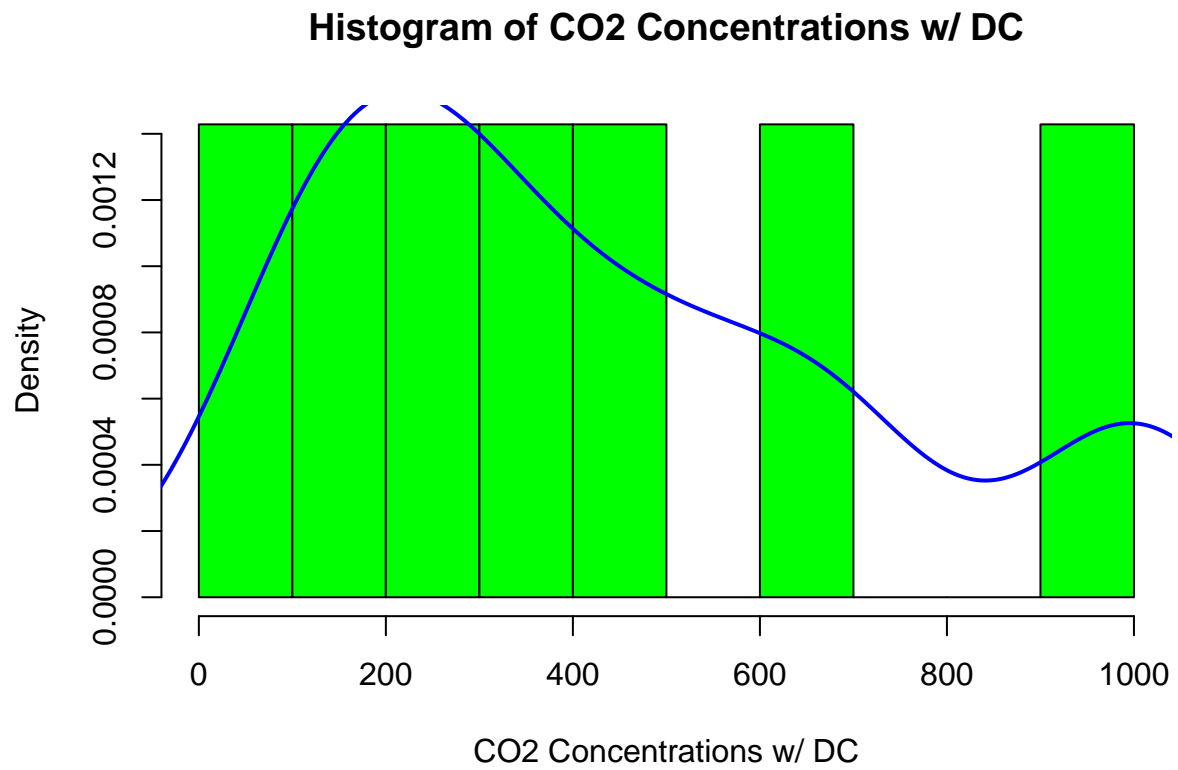
### Histogram of CO2 Concentrations

```
par(mfrow=c(1,1))
hist(CO2$conc, xlab="CO2 Concentrations",
     main="Histogram of CO2 Concentrations", col="red")
```



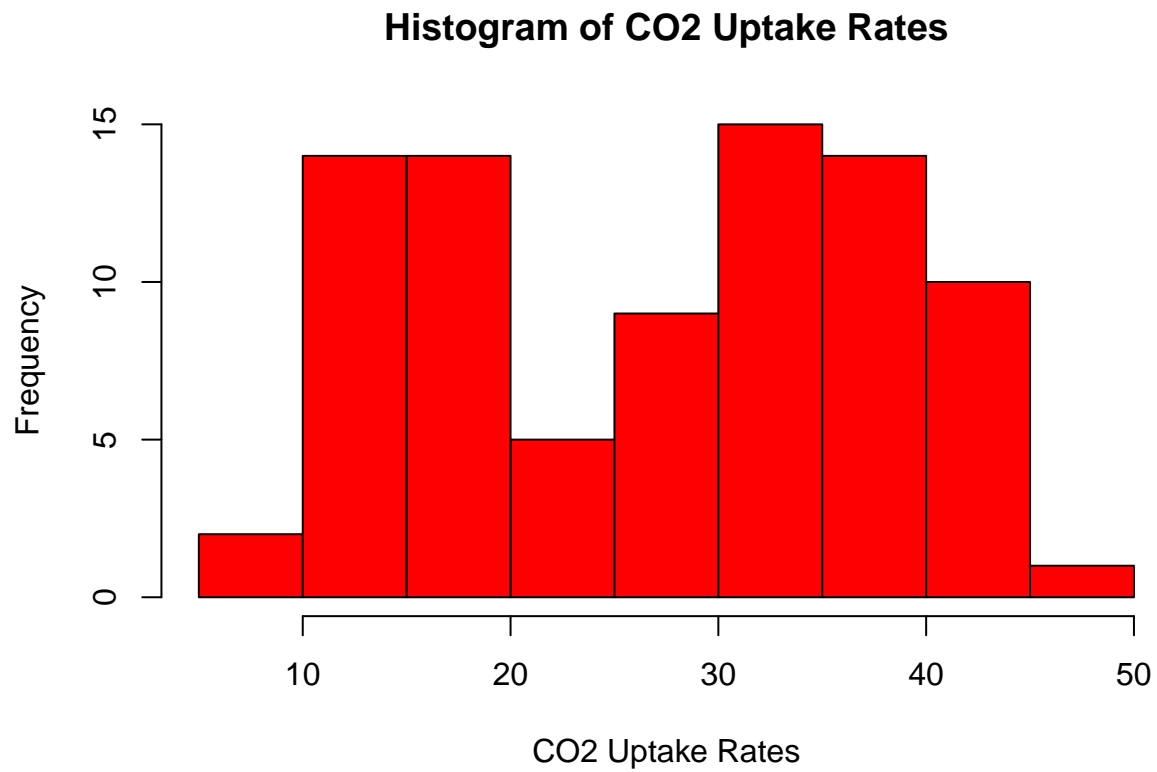
Histogram of CO2 Concentrations with it's Density Curve (DC)

```
par(mfrow=c(1,1))
x2 <- CO2$conc
hist(CO2$conc, xlab="CO2 Concentrations w/ DC",
     main="Histogram of CO2 Concentrations w/ DC",
     col="green", probability=TRUE)
lines(density(x2),col="blue", lwd=2)
```



Histogram of CO2 Uptake Rates

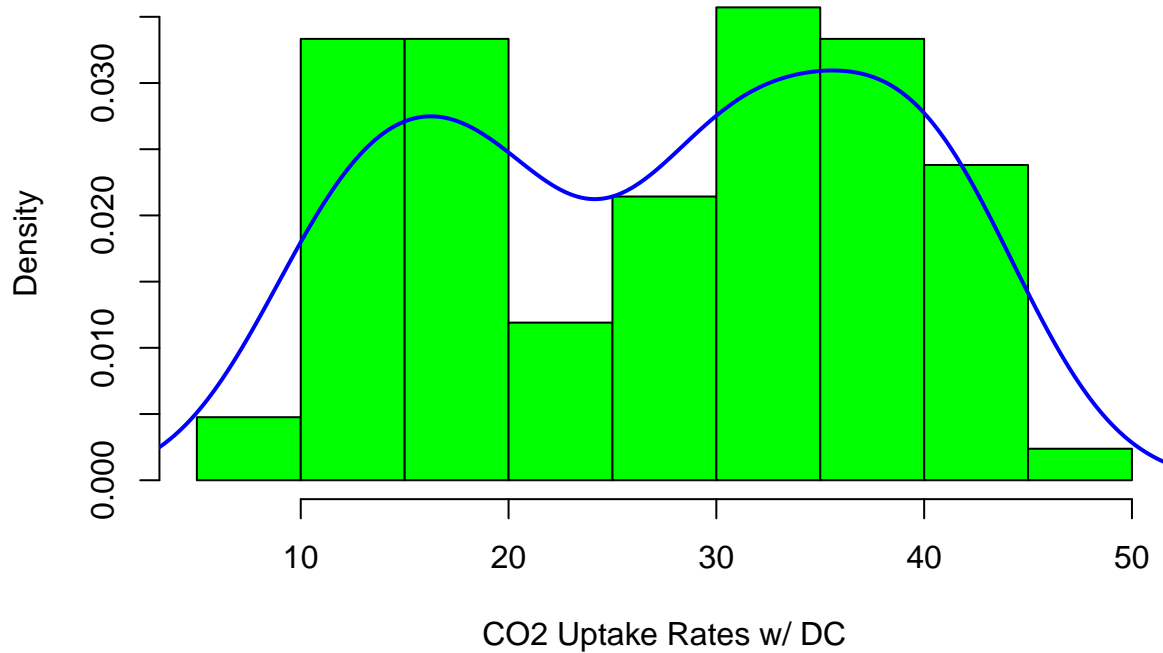
```
par(mfrow=c(1,1))  
hist(CO2$uptake, xlab="CO2 Uptake Rates",  
      main="Histogram of CO2 Uptake Rates", col="red")
```



Histogram of CO2 Uptake Rates with it's Density Curve (DC)

```
par(mfrow=c(1,1))
x3 <- CO2$uptake
hist(CO2$uptake, xlab="CO2 Uptake Rates w/ DC",
     main="Histogram of CO2 Uptake Rates w/ DC",
     col="green", probability=TRUE)
lines(density(x3),col="blue", lwd=2)
```

## Histogram of CO2 Uptake Rates w/ DC



## Linear Regression of “Conc” and “Uptake”

### Linear Regression Analysis of “Conc” and “Uptake”

```
fm <- lm(CO2$uptake ~ CO2$conc, data=CO2)
summary(fm)
```

```
##
## Call:
## lm(formula = CO2$uptake ~ CO2$conc, data = CO2)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -22.831  -7.729   1.483   7.748  16.394
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 19.500290   1.853080  10.523  < 2e-16 ***
## CO2$conc     0.017731   0.003529   5.024 2.91e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 9.514 on 82 degrees of freedom
```



```
## Multiple R-squared:  0.2354, Adjusted R-squared:  0.2261
## F-statistic: 25.25 on 1 and 82 DF,  p-value: 2.906e-06
```

### Linear Regression Model of “Conc” and “Uptake”

```
par(mfrow=c(1,1))
plot(CO2$conc, CO2$uptake,
     xlab="Ambient carbon dioxide concentrations (mL/L)",
     ylab="Carbon dioxide uptake rates (umol/m^2 sec)",
     main="Linear Regression Model of Conc and Uptake", col="red")
fm <- lm(CO2$uptake ~ CO2$conc, data=CO2)
abline(coef(fm), lty=4, col="blue")

#Extract coefficients of Conc and Uptake
fm_coef <- round(coef(fm), 3)

#Display the regression equation of the line
mtext(bquote(y == .(fm_coef[2])*x + .(fm_coef[1])),
      adj=1, padj=0)
```

