

ANOVA

Maxwel Coura Oliveira

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```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.2.1 --
## v ggplot2 3.2.1    v purrr  0.3.2
## v tibble  2.1.3    v dplyr  0.8.3
## v tidyr   1.0.0    v stringr 1.4.0
## v readr   1.3.1    v forcats 0.4.0

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

```
library(lme4)
```

```
## Loading required package: Matrix
##
## Attaching package: 'Matrix'
## The following objects are masked from 'package:tidyr':
##
##     expand, pack, unpack
```

```
library(nortest)
```

```
library(ggpubr)
```

```
## Loading required package: magrittr
##
## Attaching package: 'magrittr'
## The following object is masked from 'package:purrr':
##
##     set_names
## The following object is masked from 'package:tidyr':
##
##     extract
```

```
library(lmerTest)
```

```
##
## Attaching package: 'lmerTest'
## The following object is masked from 'package:lme4':
##
##     lmer
## The following object is masked from 'package:stats':
##
##     step
```

```

library(emmeans)

data <- read_csv("Yield.csv")

## Parsed with column specification:
## cols(
##   Plot = col_double(),
##   Rep = col_double(),
##   Location = col_character(),
##   Treatment = col_double(),
##   Cultivar = col_character(),
##   Seeding_Rate = col_double(),
##   Nitrogen_Rate = col_double(),
##   Grain_Yield = col_double(),
##   Fiber_Yield = col_double()
## )

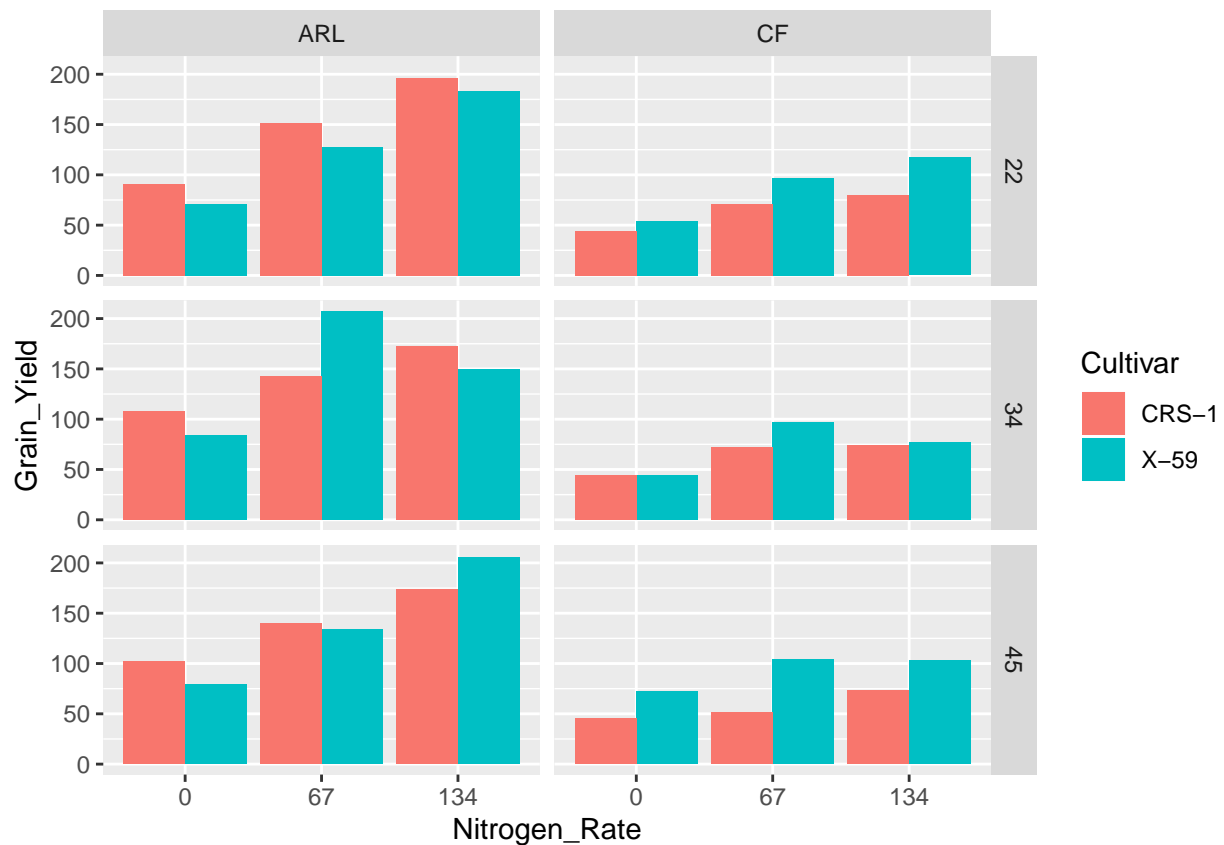
data$Seeding_Rate <- as.factor(data$Seeding_Rate) # put seeding rate as a factor
data$Nitrogen_Rate <- as.factor(data$Nitrogen_Rate)
glimpse(data)

## Observations: 144
## Variables: 9
## $ Plot      <dbl> 101, 304, 111, 316, 102, 403, 211, 310, 103, 407...
## $ Rep       <dbl> 1, 2, 3, 4, 1, 2, 3, 4, 1, 2, 3, 4, 1, 2, 3, 4, ...
## $ Location  <chr> "ARL", "ARL", "ARL", "ARL", "ARL", "ARL", "ARL", ...
## $ Treatment <dbl> 1, 1, 1, 1, 2, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, ...
## $ Cultivar  <chr> "X-59", "X-59", "X-59", "X-59", "X-59", "X-59", ...
## $ Seeding_Rate <fct> 22, 22, 22, 22, 22, 22, 22, 22, 22, 22, 22, 22, ...
## $ Nitrogen_Rate <fct> 0, 0, 0, 0, 67, 67, 67, 67, 134, 134, 134, 134, ...
## $ Grain_Yield <dbl> 29.09, 109.13, 38.24, 107.32, 180.88, 132.95, 78...
## $ Fiber_Yield <dbl> 9, 76, 27, 54, 132, 65, 58, 76, 189, 133, 122, 3...

ggplot(data, aes(x=Nitrogen_Rate, y=Grain_Yield, fill=Cultivar)) + geom_bar(stat="summary", position =

## No summary function supplied, defaulting to `mean_se()`
## No summary function supplied, defaulting to `mean_se()`
## No summary function supplied, defaulting to `mean_se()`
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```



```
Model <- lmer(Grain_Yield ~ Nitrogen_Rate * Seeding_Rate * Location * Cultivar + (1|Rep), data=data)
summary(Model)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## Grain_Yield ~ Nitrogen_Rate * Seeding_Rate * Location * Cultivar +
## (1 | Rep)
## Data: data
##
## REML criterion at convergence: 1125
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.93097 -0.52780 -0.07465  0.50753  2.17280
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Rep      (Intercept)          3.523   1.877
##  Residual                            1228.573  35.051
## Number of obs: 144, groups:  Rep, 4
##
## Fixed effects:
##
##              Estimate
## (Intercept)    91.308
## Nitrogen_Rate67  60.357
## Nitrogen_Rate134 104.932
```

## Seeding_Rate34	16.647	
## Seeding_Rate45	10.830	
## LocationCF	-47.543	
## CultivarX-59	-20.363	
## Nitrogen_Rate67:Seeding_Rate34	-25.950	
## Nitrogen_Rate134:Seeding_Rate34	-40.572	
## Nitrogen_Rate67:Seeding_Rate45	-23.080	
## Nitrogen_Rate134:Seeding_Rate45	-33.887	
## Nitrogen_Rate67:LocationCF	-32.945	
## Nitrogen_Rate134:LocationCF	-68.537	
## Seeding_Rate34:LocationCF	-16.382	
## Seeding_Rate45:LocationCF	-8.897	
## Nitrogen_Rate67:CultivarX-59	-3.687	
## Nitrogen_Rate134:CultivarX-59	7.243	
## Seeding_Rate34:CultivarX-59	-3.202	
## Seeding_Rate45:CultivarX-59	-2.370	
## LocationCF:CultivarX-59	30.413	
## Nitrogen_Rate67:Seeding_Rate34:LocationCF	27.015	
## Nitrogen_Rate134:Seeding_Rate34:LocationCF	33.935	
## Nitrogen_Rate67:Seeding_Rate45:LocationCF	1.325	
## Nitrogen_Rate134:Seeding_Rate45:LocationCF	24.650	
## Nitrogen_Rate67:Seeding_Rate34:CultivarX-59	92.480	
## Nitrogen_Rate134:Seeding_Rate34:CultivarX-59	-6.040	
## Nitrogen_Rate67:Seeding_Rate45:CultivarX-59	21.175	
## Nitrogen_Rate134:Seeding_Rate45:CultivarX-59	47.697	
## Nitrogen_Rate67:LocationCF:CultivarX-59	19.435	
## Nitrogen_Rate134:LocationCF:CultivarX-59	19.845	
## Seeding_Rate34:LocationCF:CultivarX-59	-7.025	
## Seeding_Rate45:LocationCF:CultivarX-59	18.572	
## Nitrogen_Rate67:Seeding_Rate34:LocationCF:CultivarX-59	-83.815	
## Nitrogen_Rate134:Seeding_Rate34:LocationCF:CultivarX-59	-17.770	
## Nitrogen_Rate67:Seeding_Rate45:LocationCF:CultivarX-59	-10.365	
## Nitrogen_Rate134:Seeding_Rate45:LocationCF:CultivarX-59	-71.235	
##	Std. Error	df
## (Intercept)	17.551	107.969
## Nitrogen_Rate67	24.785	105.000
## Nitrogen_Rate134	24.785	105.000
## Seeding_Rate34	24.785	105.000
## Seeding_Rate45	24.785	105.000
## LocationCF	24.785	105.000
## CultivarX-59	24.785	105.000
## Nitrogen_Rate67:Seeding_Rate34	35.051	105.000
## Nitrogen_Rate134:Seeding_Rate34	35.051	105.000
## Nitrogen_Rate67:Seeding_Rate45	35.051	105.000
## Nitrogen_Rate134:Seeding_Rate45	35.051	105.000
## Nitrogen_Rate67:LocationCF	35.051	105.000
## Nitrogen_Rate134:LocationCF	35.051	105.000
## Seeding_Rate34:LocationCF	35.051	105.000
## Seeding_Rate45:LocationCF	35.051	105.000
## Nitrogen_Rate67:CultivarX-59	35.051	105.000
## Nitrogen_Rate134:CultivarX-59	35.051	105.000
## Seeding_Rate34:CultivarX-59	35.051	105.000
## Seeding_Rate45:CultivarX-59	35.051	105.000
## LocationCF:CultivarX-59	35.051	105.000

## Nitrogen_Rate67:Seeding_Rate34:LocationCF	49.570	105.000
## Nitrogen_Rate134:Seeding_Rate34:LocationCF	49.570	105.000
## Nitrogen_Rate67:Seeding_Rate45:LocationCF	49.570	105.000
## Nitrogen_Rate134:Seeding_Rate45:LocationCF	49.570	105.000
## Nitrogen_Rate67:Seeding_Rate34:CultivarX-59	49.570	105.000
## Nitrogen_Rate134:Seeding_Rate34:CultivarX-59	49.570	105.000
## Nitrogen_Rate67:Seeding_Rate45:CultivarX-59	49.570	105.000
## Nitrogen_Rate134:Seeding_Rate45:CultivarX-59	49.570	105.000
## Nitrogen_Rate67:LocationCF:CultivarX-59	49.570	105.000
## Nitrogen_Rate134:LocationCF:CultivarX-59	49.570	105.000
## Seeding_Rate34:LocationCF:CultivarX-59	49.570	105.000
## Seeding_Rate45:LocationCF:CultivarX-59	49.570	105.000
## Nitrogen_Rate67:Seeding_Rate34:LocationCF:CultivarX-59	70.102	105.000
## Nitrogen_Rate134:Seeding_Rate34:LocationCF:CultivarX-59	70.102	105.000
## Nitrogen_Rate67:Seeding_Rate45:LocationCF:CultivarX-59	70.102	105.000
## Nitrogen_Rate134:Seeding_Rate45:LocationCF:CultivarX-59	70.102	105.000
##	t value	Pr(> t)
## (Intercept)	5.203	9.42e-07
## Nitrogen_Rate67	2.435	0.0166
## Nitrogen_Rate134	4.234	4.94e-05
## Seeding_Rate34	0.672	0.5033
## Seeding_Rate45	0.437	0.6630
## LocationCF	-1.918	0.0578
## CultivarX-59	-0.822	0.4132
## Nitrogen_Rate67:Seeding_Rate34	-0.740	0.4607
## Nitrogen_Rate134:Seeding_Rate34	-1.158	0.2497
## Nitrogen_Rate67:Seeding_Rate45	-0.658	0.5117
## Nitrogen_Rate134:Seeding_Rate45	-0.967	0.3359
## Nitrogen_Rate67:LocationCF	-0.940	0.3494
## Nitrogen_Rate134:LocationCF	-1.955	0.0532
## Seeding_Rate34:LocationCF	-0.467	0.6412
## Seeding_Rate45:LocationCF	-0.254	0.8001
## Nitrogen_Rate67:CultivarX-59	-0.105	0.9164
## Nitrogen_Rate134:CultivarX-59	0.207	0.8367
## Seeding_Rate34:CultivarX-59	-0.091	0.9274
## Seeding_Rate45:CultivarX-59	-0.068	0.9462
## LocationCF:CultivarX-59	0.868	0.3876
## Nitrogen_Rate67:Seeding_Rate34:LocationCF	0.545	0.5869
## Nitrogen_Rate134:Seeding_Rate34:LocationCF	0.685	0.4951
## Nitrogen_Rate67:Seeding_Rate45:LocationCF	0.027	0.9787
## Nitrogen_Rate134:Seeding_Rate45:LocationCF	0.497	0.6200
## Nitrogen_Rate67:Seeding_Rate34:CultivarX-59	1.866	0.0649
## Nitrogen_Rate134:Seeding_Rate34:CultivarX-59	-0.122	0.9033
## Nitrogen_Rate67:Seeding_Rate45:CultivarX-59	0.427	0.6701
## Nitrogen_Rate134:Seeding_Rate45:CultivarX-59	0.962	0.3381
## Nitrogen_Rate67:LocationCF:CultivarX-59	0.392	0.6958
## Nitrogen_Rate134:LocationCF:CultivarX-59	0.400	0.6897
## Seeding_Rate34:LocationCF:CultivarX-59	-0.142	0.8876
## Seeding_Rate45:LocationCF:CultivarX-59	0.375	0.7087
## Nitrogen_Rate67:Seeding_Rate34:LocationCF:CultivarX-59	-1.196	0.2345
## Nitrogen_Rate134:Seeding_Rate34:LocationCF:CultivarX-59	-0.253	0.8004
## Nitrogen_Rate67:Seeding_Rate45:LocationCF:CultivarX-59	-0.148	0.8827
## Nitrogen_Rate134:Seeding_Rate45:LocationCF:CultivarX-59	-1.016	0.3119
##		

```

## (Intercept) ***
## Nitrogen_Rate67 *
## Nitrogen_Rate134 ***
## Seeding_Rate34
## Seeding_Rate45
## LocationCF .
## CultivarX-59
## Nitrogen_Rate67:Seeding_Rate34
## Nitrogen_Rate134:Seeding_Rate34
## Nitrogen_Rate67:Seeding_Rate45
## Nitrogen_Rate134:Seeding_Rate45
## Nitrogen_Rate67:LocationCF
## Nitrogen_Rate134:LocationCF .
## Seeding_Rate34:LocationCF
## Seeding_Rate45:LocationCF
## Nitrogen_Rate67:CultivarX-59
## Nitrogen_Rate134:CultivarX-59
## Seeding_Rate34:CultivarX-59
## Seeding_Rate45:CultivarX-59
## LocationCF:CultivarX-59
## Nitrogen_Rate67:Seeding_Rate34:LocationCF
## Nitrogen_Rate134:Seeding_Rate34:LocationCF
## Nitrogen_Rate67:Seeding_Rate45:LocationCF
## Nitrogen_Rate134:Seeding_Rate45:LocationCF
## Nitrogen_Rate67:Seeding_Rate34:CultivarX-59 .
## Nitrogen_Rate134:Seeding_Rate34:CultivarX-59
## Nitrogen_Rate67:Seeding_Rate45:CultivarX-59
## Nitrogen_Rate134:Seeding_Rate45:CultivarX-59
## Nitrogen_Rate67:LocationCF:CultivarX-59
## Nitrogen_Rate134:LocationCF:CultivarX-59
## Seeding_Rate34:LocationCF:CultivarX-59
## Seeding_Rate45:LocationCF:CultivarX-59
## Nitrogen_Rate67:Seeding_Rate34:LocationCF:CultivarX-59
## Nitrogen_Rate134:Seeding_Rate34:LocationCF:CultivarX-59
## Nitrogen_Rate67:Seeding_Rate45:LocationCF:CultivarX-59
## Nitrogen_Rate134:Seeding_Rate45:LocationCF:CultivarX-59
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

##
## Correlation matrix not shown by default, as p = 36 > 12.
## Use print(x, correlation=TRUE) or
##      vcov(x)          if you need it

```

```
anova(Model)
```

```

## Type III Analysis of Variance Table with Satterthwaite's method
##
##              Sum Sq Mean Sq NumDF DenDF
## Nitrogen_Rate    104177     52089      2    105
## Seeding_Rate         27         13      2    105
## Location    159861    159861      1    105
## Cultivar         3403         3403      1    105
## Nitrogen_Rate:Seeding_Rate    11247         2812      4    105
## Nitrogen_Rate:Location    17495         8748      2    105
## Seeding_Rate:Location     1745         872      2    105

```

```
## Nitrogen_Rate:Cultivar          4813    2407     2    105
## Seeding_Rate:Cultivar           1660     830     2    105
## Location:Cultivar               6562    6562     1    105
## Nitrogen_Rate:Seeding_Rate:Location 2784     696     4    105
## Nitrogen_Rate:Seeding_Rate:Cultivar 5869    1467     4    105
## Nitrogen_Rate:Location:Cultivar    244     122     2    105
## Seeding_Rate:Location:Cultivar    2787    1393     2    105
## Nitrogen_Rate:Seeding_Rate:Location:Cultivar 4995    1249     4    105
##                               F value    Pr(>F)
## Nitrogen_Rate                42.3976  3.18e-14 ***
## Seeding_Rate                 0.0109   0.98920
## Location                    130.1196 < 2.2e-16 ***
## Cultivar                     2.7699   0.09904 .
## Nitrogen_Rate:Seeding_Rate    2.2885   0.06471 .
## Nitrogen_Rate:Location        7.1202   0.00126 **
## Seeding_Rate:Location         0.7101   0.49392
## Nitrogen_Rate:Cultivar        1.9588   0.14614
## Seeding_Rate:Cultivar         0.6754   0.51115
## Location:Cultivar            5.3408   0.02279 *
## Nitrogen_Rate:Seeding_Rate:Location 0.5665   0.68753
## Nitrogen_Rate:Seeding_Rate:Cultivar 1.1942   0.31774
## Nitrogen_Rate:Location:Cultivar 0.0993   0.90553
## Seeding_Rate:Location:Cultivar 1.1342   0.32560
## Nitrogen_Rate:Seeding_Rate:Location:Cultivar 1.0163   0.40247
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
emmip(Model, Nitrogen_Rate ~ Location | Location) # see the interactions
```

```
## NOTE: Results may be misleading due to involvement in interactions
```

