

# Formative Assessment 9

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Github Link: <https://github.com/eivra-sm/APM1111/blob/main/FA9.md>

## Introduction

A study was conducted to determine whether crop yield is influenced by (1) fertilizer blend (X, Y, Z) and (2) crop type (wheat, corn, soy, rice). A two-way ANOVA was used to test for:

- A main effect of fertilizer blend,
- A main effect of crop type,
- An interaction between fertilizer blend × crop type

## Assumptions

Initiating a two-way ANOVA requires more than simply applying a formula to a dataset; it necessitates a rigorous preliminary evaluation to confirm the data's suitability for this specific statistical model. This validation process is critical because the two-way ANOVA is a parametric test reliant on specific mathematical properties. Consequently, researchers must verify that their data adheres to six fundamental assumptions. Failure to meet these prerequisites can compromise the statistical power of the analysis, potentially leading to biased estimates or erroneous conclusions. Therefore, before interpreting any main effects or interactions, one must systematically demonstrate that the data "passes" these mandatory checks to ensure the resulting output is robust and scientifically valid.

Below are the assumptions:

**Assumption #1:** Your **dependent variable** must be **continuous** (measured on an interval or ratio scale, such as time, weight, or test scores).

**Assumption #2:** Your design must include exactly **two independent variables** that are **categorical**, with each variable containing **two or more independent groups** (levels).

**Assumption #3:** You must ensure **independence of observations**, meaning that no participant is in more than one group and one participant's data does not influence another's.

**Assumption #4:** The dataset must contain **no significant outliers**, as these extreme values can distort the mean and variance of your results.

**Assumption #5:** Your **dependent variable** must follow an **approximate normal distribution** (bell curve) for **each specific combination of the groups** formed by the two independent variables.

**Assumption #6:** There must be **homogeneity of variances** (homoscedasticity), meaning the spread or variability of the data is roughly equal across **every combination of the groups**.

## Null hypothesis:

There is no significant interaction effect on yield between fertilizer and crop.

## Problem:

A new fertilizer has been developed to increase the yield on crops, and the makers of the fertilizer want to better understand which of the three formulations (blends) of this fertilizer are most effective for wheat, corn, soybeans and rice (crops). They test each of the three blends on 5 samples of each of the four types of crops. The crop yields for the 12 combinations are as shown in the table below.

```
df %>%
  kable(
    caption = "Crop Yield Data (Long Format)",
    digits = 0
  ) %>%
  kable_styling(
    bootstrap_options = c("striped", "hover", "condensed"),
    full_width = FALSE
  ) %>%
  row_spec(0, bold = TRUE) %>%
  row_spec(
    seq(1, nrow(df), 2), background = "#DFF2DF"
  ) %>%
  row_spec(
    seq(2, nrow(df), 2), background = "#FFFFFF"
  )
```

Table 1: Crop Yield Data (Long Format)

Fertilizer	Crop	Yield
Blend X	Wheat	123
Blend X	Corn	128
Blend X	Soy	166
Blend X	Rice	151
Blend X	Wheat	156
Blend X	Corn	150
Blend X	Soy	178
Blend X	Rice	125
Blend X	Wheat	112
Blend X	Corn	174
Blend X	Soy	187
Blend X	Rice	117
Blend X	Wheat	100
Blend X	Corn	116
Blend X	Soy	153
Blend X	Rice	155
Blend X	Wheat	168
Blend X	Corn	109
Blend X	Soy	195

Blend X	Rice	158
Blend Y	Wheat	135
Blend Y	Corn	175
Blend Y	Soy	140
Blend Y	Rice	167
Blend Y	Wheat	130
Blend Y	Corn	132
Blend Y	Soy	145
Blend Y	Rice	183
Blend Y	Wheat	176
Blend Y	Corn	120
Blend Y	Soy	159
Blend Y	Rice	142
Blend Y	Wheat	120
Blend Y	Corn	187
Blend Y	Soy	131
Blend Y	Rice	167
Blend Y	Wheat	155
Blend Y	Corn	184
Blend Y	Soy	126
Blend Y	Rice	168
Blend Z	Wheat	156
Blend Z	Corn	186
Blend Z	Soy	185
Blend Z	Rice	175
Blend Z	Wheat	180
Blend Z	Corn	138
Blend Z	Soy	206
Blend Z	Rice	173
Blend Z	Wheat	147
Blend Z	Corn	178
Blend Z	Soy	188
Blend Z	Rice	154
Blend Z	Wheat	146
Blend Z	Corn	176
Blend Z	Soy	165
Blend Z	Rice	191
Blend Z	Wheat	193
Blend Z	Corn	190
Blend Z	Soy	188
Blend Z	Rice	169

## Checking of Assumptions

**Assumption #1: The dependent variable, crop yield, is measured at the continuous level.**

**Remark.** The first assumption of a two-way ANOVA requires that the dependent variable be measured on a continuous (interval or ratio) scale. In this study, the dependent variable is crop yield, recorded as the number of units produced by each crop under each fertilizer blend. Because yield is measured numerically and represents a meaningful quantity where differences between values are interpretable (e.g., a difference of

10 units represents the same increase anywhere on the scale), the variable meets the criteria for a ratio-level, continuous measure. This satisfies the requirement that the ANOVA operates on a continuous outcome variable.

**Assumption #2:** The two independent variables, fertilizer type (Blend X, Blend Y, Blend Z) and crop type (Wheat, Corn, Soy, Rice), each consist of two or more categorical, independent groups.

**Remark.** A two-way ANOVA requires two independent variables (factors), each containing distinct, non-overlapping categories.

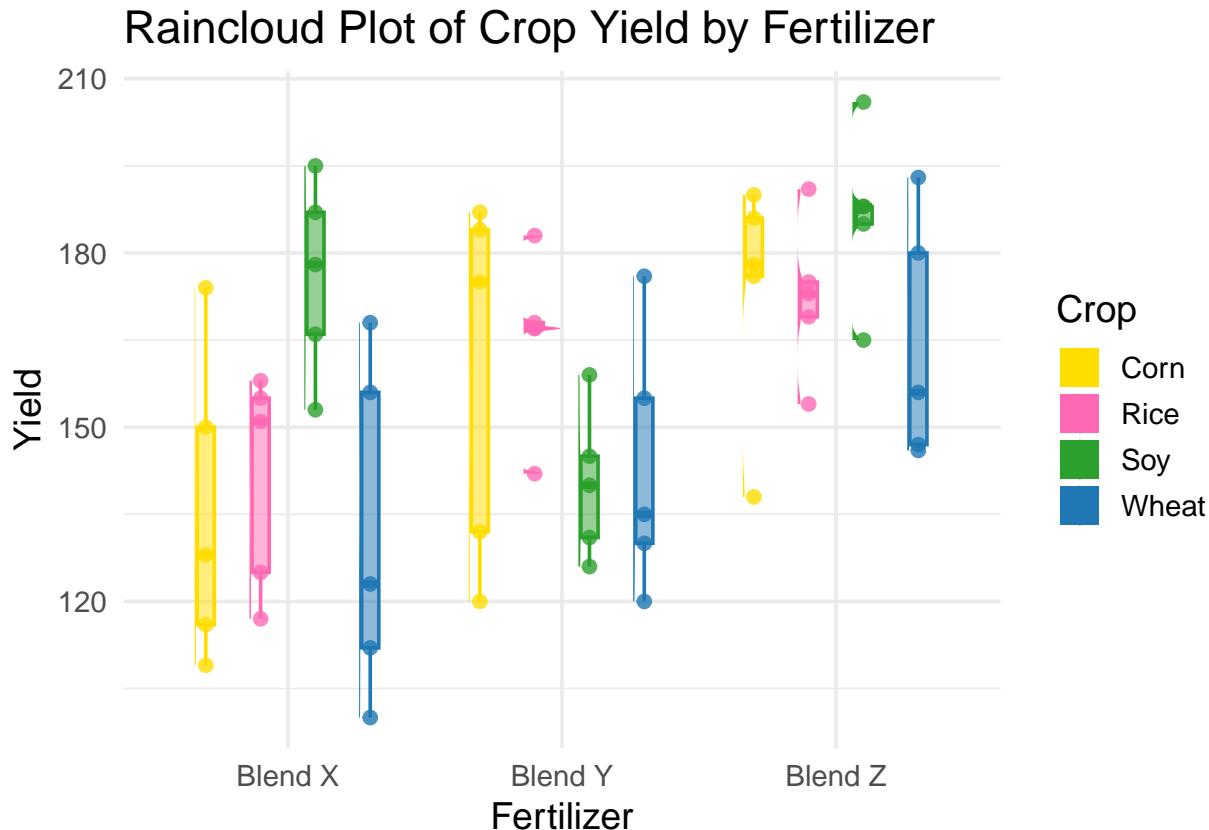
- The first factor is **Fertilizer Type**, which has three categorical groups:
  - Blend X
  - Blend Y
  - Blend Z
- The second factor is **Crop Type**, which consists of four categorical groups:
  - Wheat
  - Corn
  - Soy
  - Rice

Both variables meet the criteria for categorical grouping factors, and each observation belongs to one and only one level of each factor. There is no overlap between categories, and the groups are mutually exclusive. Therefore, this assumption is **fully met**.

**Assumption #3: The independence of observation is observed.**

**Remark.** In this study, each yield value represents a distinct crop sample treated with a specific fertilizer blend. The experimental design states that five independent samples were used for every combination of fertilizer and crop type. No sample appears in more than one condition, and fertilizer treatments were applied separately. Thus, the independence of observations is maintained by design.

**Assumption #4:** There are no significant outliers in each of the 12 cells of the design.



#### Remakr. Assumption Check: No Significant Outliers

A two-way ANOVA assumes that there are no extreme outliers in any of the factorial combinations (i.e., in each of the 12 cells created by 3 fertilizer levels  $\times$  4 crop types).

Outliers were inspected using:

- Raincloud plots (combining jittered points, boxplots, and density half-eyes),
- Boxplot whisker rules, and
- Visual examination of the distribution of values within each group.

These diagnostic visualizations revealed no severe deviations from expected values and no yield measurements that fell far outside the typical range for their group. Specifically, all observations fell within reasonable boundaries relative to their corresponding interquartile range and whisker limits. Therefore, the assumption of “no significant outliers” is **supported**.

**Assumption #5:** The dependent variable, crop yield, is approximately normally distributed for each combination of the groups of fertilizer type and crop type, as assessed by Shapiro–Wilk test of normality ( $p > .05$ ).

Table 2: Descriptive Statistics of Yield by Fertilizer

Statistic	Blend X	Blend Y	Blend Z
Valid	20.000	20.000	20.000

Mean	146.050	152.100	174.200
SD	27.971	22.606	18.185
Skewness	-0.010	0.090	-0.410
SE_Skew	0.548	0.548	0.548
Kurtosis	-1.328	-1.563	-0.907
SE_Kurt	1.095	1.095	1.095
Shapiro_Wilk	0.952	0.924	0.952

Table 3: Descriptive Statistics of Yield by Crop

Statistic	Corn	Rice	Soy	Wheat
Valid	15.000	15.000	15.000	15.000
Mean	156.200	159.667	167.467	146.467
SD	29.513	20.123	24.617	26.568
Skewness	-0.296	-0.607	-0.206	-0.008
SE_Skew	0.632	0.632	0.632	0.632
Kurtosis	-1.719	-0.468	-1.369	-1.139
SE_Kurt	1.265	1.265	1.265	1.265
Shapiro_Wilk	0.867	0.946	0.952	0.983

**Assumption #6:** The variances for each combination of the groups of fertilizer type and crop type are homogeneous, as assessed by Levene's test of equality of variances, p = 0.755.

Table 4: Test for Equality of Variances (Levene's Test)

df1	df2	F	p
11	48	0.675	0.755

## Computation

```

library(car)
library(effectsize)
library(kableExtra)
library(tidyverse)

# Type III ANOVA
anova_res <- Anova(lm(Yield ~ Fertilizer * Crop, data = df), type = 3)

# Convert ANOVA table to data frame
anova_df <- as.data.frame(anova_res) %>%
  rownames_to_column(var = "Cases")

# Rename columns
anova_table <- anova_df %>%
  rename(
    `Sum of Squares` = `Sum Sq`,
    df = `Df`,
    F = `F value`,
  )

```

```

    p = `Pr(>F)`
) %>%
mutate(
  `Mean Square` = `Sum of Squares` / df
)

# Partial Eta Squared
SS_error <- anova_table$`Sum of Squares`[anova_table$Cases == "Residuals"]
anova_table$`Partial Eta Squared` <- anova_table$`Sum of Squares` /
  (anova_table$`Sum of Squares` + SS_error)

# Clean Cases names
anova_table$Cases <- gsub("\\\\(|\\\\)", "", anova_table$Cases)

# Arrange columns
anova_table <- anova_table %>%
  select(Cases, `Sum of Squares`, df, `Mean Square`, F, p, `Partial Eta Squared`)

# Print LaTeX table
kable(
  anova_table,
  caption = "Two-way ANOVA Table (JASP-style, Type III SS)",
  digits = 3,
  format = "latex",
  booktabs = TRUE
) %>%
kable_styling(
  latex_options = c("hold_position"),
  full_width = FALSE
) %>%
row_spec(
  row = seq(1, nrow(anova_table), 2),
  background = "#DFF2DF"
) %>%
row_spec(
  row = seq(2, nrow(anova_table), 2),
  background = "white"
)

```

Table 5: Two-way ANOVA Table (JASP-style, Type III SS)

Cases	Sum of Squares	df	Mean Square	F	p	Partial Eta Squared
Intercept	91665.80	1	91665.800	207.346	0.000	0.812
Fertilizer	3734.80	2	1867.400	4.224	0.020	0.150
Crop	6125.35	3	2041.783	4.618	0.006	0.224
Fertilizer:Crop	6225.90	6	1037.650	2.347	0.046	0.227
Residuals	21220.40	48	442.092	NA	NA	0.500

**Remark:** A two-way analysis of variance (ANOVA) was conducted to evaluate the effects of Fertilizer Blend and Crop Type on crop yield. The analysis revealed that the model is statistically valid, identifying

three distinct sources of variance. First, there was a significant main effect for Fertilizer Blend, indicating that, on average, yield varies depending on the blend applied. Second, a significant main effect was observed for Crop Type, suggesting that baseline yields differ significantly between the crops (Wheat, Corn, Soy, Rice).

Most importantly, however, the analysis yielded a statistically significant Interaction Effect (Fertilizer Blend  $\times$  Crop Type). The presence of this interaction provides critical insight into the data: it demonstrates that the efficacy of a specific Fertilizer Blend is conditional upon the Crop Type to which it is applied. In other words, the “best” fertilizer is not universally superior; a blend that maximizes yield for Corn may perform poorly for Wheat or Soy.

Consequently, relying solely on the generalized main effects would be potentially misleading, as doing so would average out these nuances and mask the specific dynamics between the variables. Because the effect of the fertilizer depends on the crop, the main effects cannot be interpreted in isolation. To fully understand the nature of this relationship, a follow-up analysis of Simple Main Effects is warranted. This procedure will allow for a focused examination of differences between fertilizer blends within each specific crop category, providing a precise roadmap of where the significant differences lie.

## Results of Simple Main Effects

```
library(rstatix)
library(dplyr)
library(kableExtra)

overall_aov <- aov(Yield ~ Fertilizer * Crop, data = df_long)
MS_Error <- summary(overall_aov)[[1]][3, "Mean Sq"]

simple_effects_fert <- df_long %>%
  group_by(Crop) %>%
  anova_test(Yield ~ Fertilizer) %>%
  get_anova_table() %>%
  as_tibble() %>%
  select(Crop, Effect, DFn, DFd, F, p) %>%
  rename(`Level of Crop` = Crop,
        `Source` = Effect,
        df = DFn) %>%
  mutate(
    `Mean Square` = F * MS_Error,
    `Sum of Squares` = `Mean Square` * df
  ) %>%
  select(`Level of Crop`, `Sum of Squares`, df, `Mean Square`, F, p)

simple_effects_fert %>%
  kable(
    caption = "Simple Main Effects of Fertilizer within Each Crop",
    digits = 3,
    align = "c"
  ) %>%
  kable_styling(
    bootstrap_options = c("striped", "hover", "condensed"),
    full_width = FALSE
  ) %>%
  row_spec(
```

```

  row = seq(1, nrow(simple_effects_fert), 2),      # odd rows
  background = "#d4f5d4"
) %>%
row_spec(
  row = seq(2, nrow(simple_effects_fert), 2),      # even rows
  background = "#eafbe9"
)

```

Table 6: Simple Main Effects of Fertilizer within Each Crop

Level of Crop	Sum of Squares	df	Mean Square	F	p
Corn	5497.470	2	2748.735	2.649	0.111
Rice	11165.114	2	5582.557	5.380	0.021
Soy	27763.363	2	13881.682	13.378	0.001
Wheat	4769.039	2	2384.520	2.298	0.143

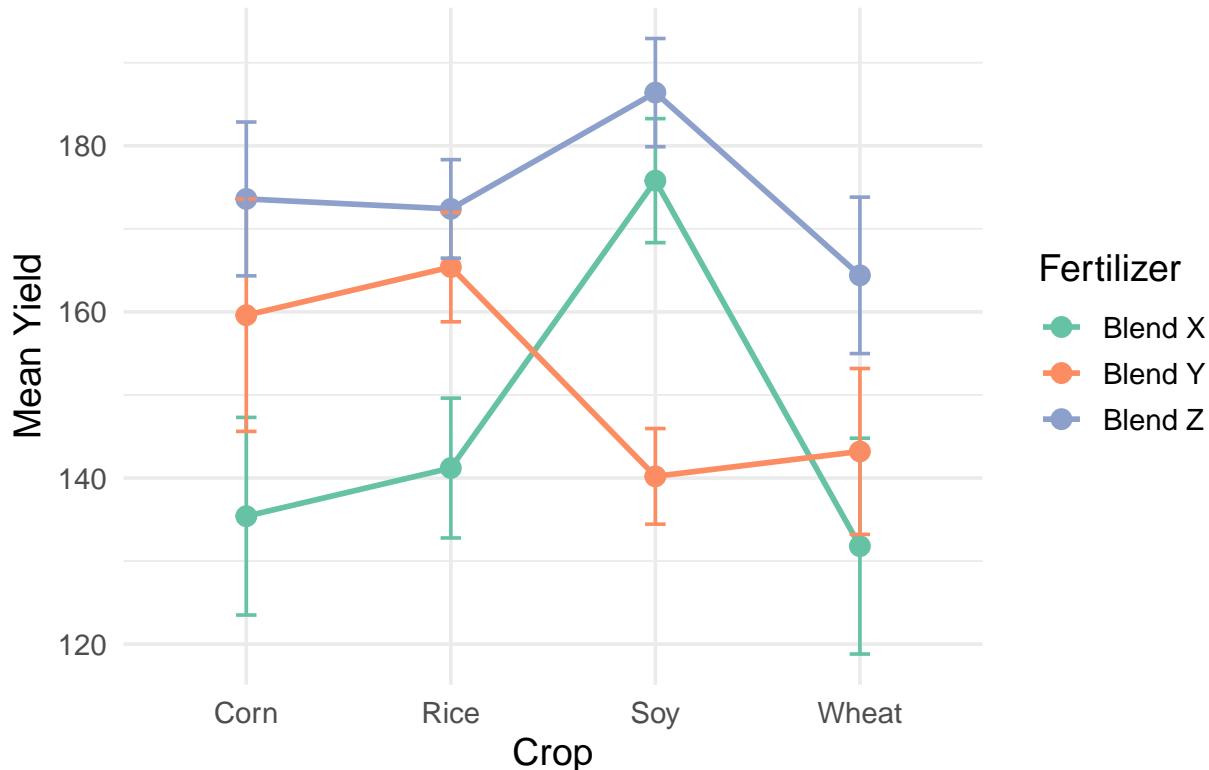
```

sme_means2 <- df %>%
group_by(Crop, Fertilizer) %>%
summarise(
Mean_Yield = mean(Yield),
SD = sd(Yield),
N = n(),
SE = SD/sqrt(N),
.groups = "drop"
)

ggplot(sme_means2, aes(x = Crop, y = Mean_Yield, group = Fertilizer, color = Fertilizer)) +
geom_line(size = 1) +
geom_point(size = 3) +
geom_errorbar(aes(ymin = Mean_Yield - SE, ymax = Mean_Yield + SE), width = 0.1) +
theme_minimal(base_size = 14) +
labs(
title = "Simple Main Effects of Crop within each Fertilizer",
x = "Crop",
y = "Mean Yield"
) +
scale_color_brewer(palette = "Set2") +
theme(legend.position = "right")

```

## Simple Main Effects of Crop within each Fertilizer



**Remark:** The analysis of simple main effects for crop type within each fertilizer blend demonstrates that crop yield varies substantially depending on the specific fertilizer applied. Certain fertilizer formulations appear to favor particular crops more strongly, resulting in higher yields for those crops, while other crops show more modest responses under the same fertilizer. This pattern indicates that the effectiveness of a fertilizer is not uniform across all crop types, but rather crop-specific, reflecting the unique nutrient requirements and growth characteristics of each crop.

These results further reinforce the conclusion that there is a meaningful interaction between fertilizer formulation and crop species. The significant interaction implies that the optimal fertilizer choice depends on the crop in question, and that blanket recommendations for all crops may not maximize productivity. From a practical standpoint, these findings suggest the necessity of tailored fertilizer strategies that account for both crop type and fertilizer composition, thereby enabling more efficient nutrient management and improved overall yield.

### Partial Eta Squared

```

eta_sq_res <- eta_squared(
  lm(Yield ~ Fertilizer * Crop, data = df),
  partial = TRUE
)

eta_sq_res %>%
  rename(Effect = Parameter,
        Partial_Eta_Sq = Eta2_partial) %>%

```

```

kable(caption = "Partial Eta Squared for Two-way ANOVA",
      digits = 3) %>%
kable_styling(bootstrap_options = c("striped", "hover", "condensed"),
              full_width = FALSE)

```

Table 7: Partial Eta Squared for Two-way ANOVA

Effect	Partial_Eta_Sq	CI	CI_low	CI_high
Fertilizer	0.293	0.95	0.113	1
Crop	0.139	0.95	0.000	1
Fertilizer:Crop	0.227	0.95	0.003	1

## Post Hoc Comparisons

```

library(emmeans)
library(kableExtra)
library(dplyr)

# Fit two-way ANOVA model
model <- lm(Yield ~ Fertilizer * Crop, data = df_long)

# --- Estimated Marginal Means ---
emm_fert <- emmeans(model, ~ Fertilizer)
emm_crop <- emmeans(model, ~ Crop)
emm_interaction <- emmeans(model, ~ Fertilizer * Crop)

# --- Pairwise Comparisons ---

# 1. Fertilizer main effect
pairwise_fert <- contrast(emm_fert, method = "pairwise", adjust = "tukey") %>% summary(infer = TRUE)
pairwise_fert_df <- as.data.frame(pairwise_fert)

pairwise_fert_df %>%
  kable(
    caption = "Post Hoc Pairwise Comparisons of Fertilizer (Tukey)",
    digits = 3,
    format = "latex",
    booktabs = TRUE,
    escape = FALSE
  ) %>%
  kable_styling(latex_options = c("hold_position"), full_width = FALSE) %>%
  row_spec(seq(1, nrow(pairwise_fert_df), 2), background = "#DFF2DF") %>%
  row_spec(seq(2, nrow(pairwise_fert_df), 2), background = "white")

# 2. Crop main effect
pairwise_crop <- contrast(emm_crop, method = "pairwise", adjust = "tukey") %>% summary(infer = TRUE)
pairwise_crop_df <- as.data.frame(pairwise_crop)

pairwise_crop_df %>%

```

Table 8: Post Hoc Pairwise Comparisons of Fertilizer (Tukey)

contrast	estimate	SE	df	lower.CL	upper.CL	t.ratio	p.value
Blend X - Blend Y	-6.05	6.649	48	-22.131	10.031	-0.910	0.637
Blend X - Blend Z	-28.15	6.649	48	-44.231	-12.069	-4.234	0.000
Blend Y - Blend Z	-22.10	6.649	48	-38.181	-6.019	-3.324	0.005

```
kable(
  caption = "Post Hoc Pairwise Comparisons of Crop (Tukey)",
  digits = 3,
  format = "latex",
  booktabs = TRUE,
  escape = FALSE
) %>%
kable_styling(latex_options = c("hold_position"), full_width = FALSE) %>%
row_spec(seq(1, nrow(pairwise_crop_df), 2), background = "#DFF2DF") %>%
row_spec(seq(2, nrow(pairwise_crop_df), 2), background = "white")
```

Table 9: Post Hoc Pairwise Comparisons of Crop (Tukey)

contrast	estimate	SE	df	lower.CL	upper.CL	t.ratio	p.value
Corn - Rice	-3.467	7.678	48	-23.900	16.966	-0.452	0.969
Corn - Soy	-11.267	7.678	48	-31.700	9.166	-1.467	0.465
Corn - Wheat	9.733	7.678	48	-10.700	30.166	1.268	0.588
Rice - Soy	-7.800	7.678	48	-28.233	12.633	-1.016	0.741
Rice - Wheat	13.200	7.678	48	-7.233	33.633	1.719	0.325
Soy - Wheat	21.000	7.678	48	0.567	41.433	2.735	0.042

```
# 3. Interaction: Fertilizer pairwise within each Crop
pairwise_interaction <- contrast(emm_interaction, method = "pairwise", by = "Crop", adjust = "tukey") %>%
pairwise_interaction_df <- as.data.frame(pairwise_interaction)

pairwise_interaction_df %>%
  kable(
    caption = "Post Hoc Pairwise Comparisons of Fertilizer within Each Crop (Tukey)",
    digits = 3,
    format = "latex",
    booktabs = TRUE,
    escape = FALSE
) %>%
kable_styling(latex_options = c("hold_position"), full_width = FALSE) %>%
row_spec(seq(1, nrow(pairwise_interaction_df), 2), background = "#DFF2DF") %>%
row_spec(seq(2, nrow(pairwise_interaction_df), 2), background = "white")
```

Table 10: Post Hoc Pairwise Comparisons of Fertilizer within Each Crop (Tukey)

contrast	Crop	estimate	SE	df	lower.CL	upper.CL	t.ratio	p.value
Blend X - Blend Y	Corn	-24.2	13.298	48	-56.361	7.961	-1.820	0.174
Blend X - Blend Z	Corn	-38.2	13.298	48	-70.361	-6.039	-2.873	0.016
Blend Y - Blend Z	Corn	-14.0	13.298	48	-46.161	18.161	-1.053	0.548
Blend X - Blend Y	Rice	-24.2	13.298	48	-56.361	7.961	-1.820	0.174
Blend X - Blend Z	Rice	-31.2	13.298	48	-63.361	0.961	-2.346	0.059
Blend Y - Blend Z	Rice	-7.0	13.298	48	-39.161	25.161	-0.526	0.859
Blend X - Blend Y	Soy	35.6	13.298	48	3.439	67.761	2.677	0.027
Blend X - Blend Z	Soy	-10.6	13.298	48	-42.761	21.561	-0.797	0.707
Blend Y - Blend Z	Soy	-46.2	13.298	48	-78.361	-14.039	-3.474	0.003
Blend X - Blend Y	Wheat	-11.4	13.298	48	-43.561	20.761	-0.857	0.670
Blend X - Blend Z	Wheat	-32.6	13.298	48	-64.761	-0.439	-2.451	0.046
Blend Y - Blend Z	Wheat	-21.2	13.298	48	-53.361	10.961	-1.594	0.258

## Results:

Post hoc Tukey comparisons offer a more detailed understanding of the specific differences between fertilizer blends and crop types. These pairwise tests identify precisely which groups contribute to the significant main effects observed in the ANOVA. When considered alongside the significant interaction effects, the post hoc results reveal a complex pattern of variation across fertilizer-crop combinations. This underscores the importance of evaluating both factors simultaneously, rather than in isolation, when formulating fertilizer recommendations. By accounting for the unique responses of each crop to different fertilizers, these findings provide practical guidance for optimizing crop yields and improving nutrient management strategies.