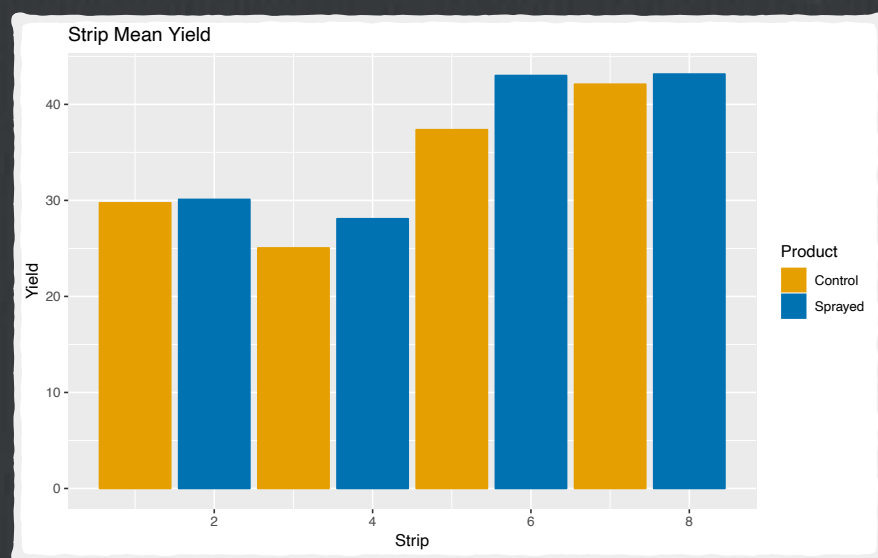


We fail to reject the null hypothesis



Even though :

- For each pair of strips, the treated strip had greater yield
- Average yield in the treated strips was greater (36.1 vs 33.6 bu/acre - ~ 7.5% increase)

```
> wilcox.test(Yield ~ Product, paired=TRUE,...)
Wilcoxon signed rank test
data: Yield by Product
V = 0, p-value = 0.125
```

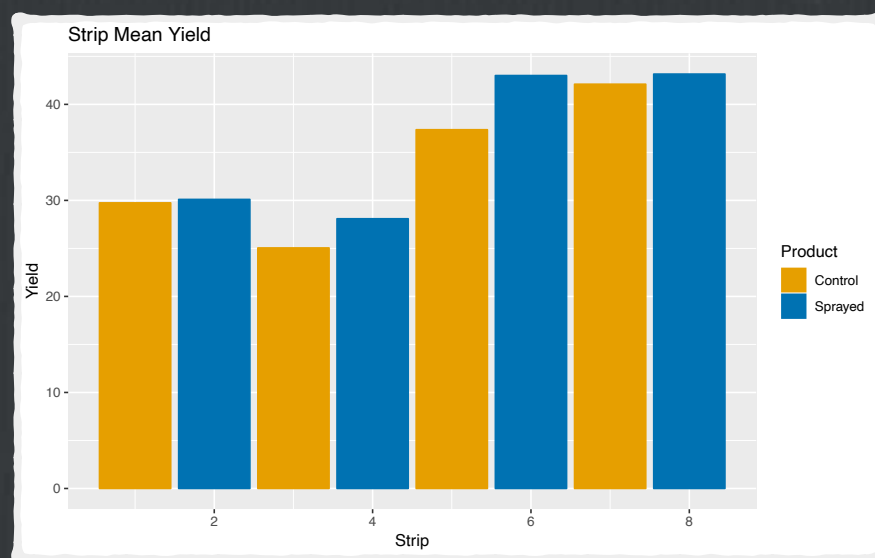
```
> t.test(Yield ~ Product, paired=TRUE, ...)
Paired t-test
data: Yield by Product
t = -2.1319, df = 3, p-value = 0.1228
sample estimates:
mean in group Control mean in group Sprayed
33.56637 36.09020
```

```
> friedman.test(Yield ~ Block | Product, ...)
Friedman rank sum test
data: Yield and Block and Product
Friedman chi-squared = 6, df = 3, p-value = 0.1116
```

```
> anova(Yield ~ Block + Product, ...)
Analysis of Variance Table

Response: Yield
Df Sum Sq Mean Sq F value Pr(>F)
Block 3 363.52 121.173 43.230 0.005732 **
Product 1 12.74 12.739 4.545 0.122791
Residuals 3 8.41 2.803
```

We fail to reject the null hypothesis



- This result is disappointing, but may be the best outcome for a designed based analysis of these data.

- ```
> wilcox.test(Yield ~ Product, paired=TRUE,...)
```

**Wilcoxon signed rank test**

data: Yield by Product  
**V = 0, p-value = 0.125**
- ```
> t.test(Yield ~ Product, paired=TRUE, ...)
```

Paired t-test

data: Yield by Product
t = -2.1319, df = 3, p-value = 0.1228

sample estimates:
mean in group Control mean in group Sprayed
33.56637 36.09020
- ```
> friedman.test(Yield ~ Block | Product, ...)
```

**Friedman rank sum test**

data: Yield and Block and Product  
Friedman chi-squared = 6, df = 3, **p-value = 0.1116**
- ```
> anova(Yield ~ Block + Product, ...)
```

Analysis of Variance Table

Response: Yield

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Block	3	363.52	121.173	43.230	0.005732 **
Product	1	12.74	12.739	4.545	0.122791
Residuals	3	8.41	2.803		