

ANOVA as Linear Models

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ANOVA

What is ANOVA, really?

The one-way ANOVA we've seen is just a linear model with one categorical predictor:

$$y = \beta_0 + \beta_1 x + \epsilon$$

which can be written as

$$y_{ij} = \mu + \alpha_i + \epsilon_{ij}$$

- ▶ μ is the baseline mean
- ▶ α_i is the effect of category i

Example

```
anova(aov(weight ~ feed, chickwts))
```

Is exactly the same as

```
anova(lm(weight ~ feed, chickwts))
```

```
## Analysis of Variance Table
```

```
##
```

```
## Response: weight
```

```
##           Df Sum Sq Mean Sq F value    Pr(>F)
```

```
## feed         5 231129   46226  15.365 5.936e-10 ***
```

```
## Residuals 65 195556     3009
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Two-Way ANOVA

We can extend the one-way ANOVA to other settings, such as the two-way ANOVA.

- ▶ This is a linear model with two categorical predictors, plus their interaction

$$y_{ij} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \epsilon_{ij}$$

- ▶ ANOVA adds predictors into the model using *sequential hypothesis testing*.

Two-Way ANOVA

```
anova(lm(len ~ supp*as.factor(dose), ToothGrowth))
```

```
## Analysis of Variance Table
```

```
##
```

```
## Response: len
```

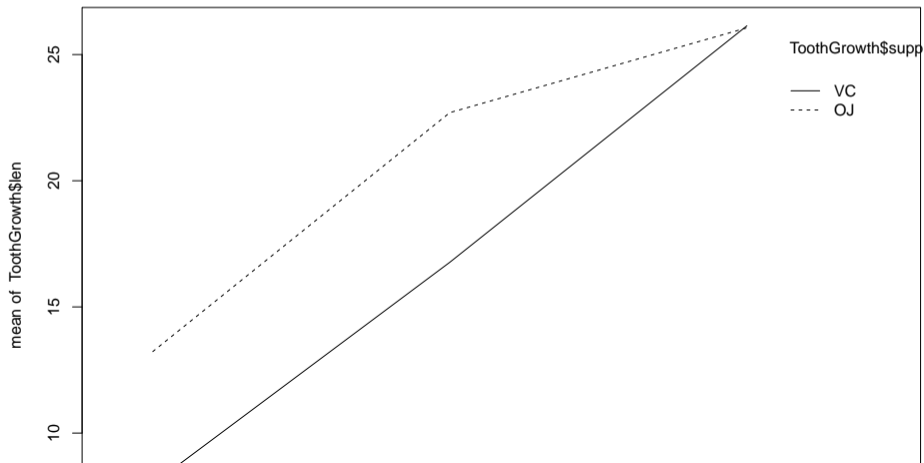
##		Df	Sum Sq	Mean Sq	F value	Pr(>F)	
##	supp	1	205.35	205.35	15.572	0.0002312	***
##	as.factor(dose)	2	2426.43	1213.22	92.000	< 2.2e-16	***
##	supp:as.factor(dose)	2	108.32	54.16	4.107	0.0218603	*
##	Residuals	54	712.11	13.19			

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Interaction Plot

```
interaction.plot(as.factor(ToothGrowth$dose), ToothGrowth$supp, ToothGrowth$len)
```



Sequential Hypothesis Testing

	Null Model	Alternative
1.	$y \sim 1$	$y \sim \text{supp}$
2.	$y \sim \text{supp}$	$y \sim \text{supp} + \text{dose}$
3.	$y \sim \text{supp} + \text{dose}$	$y \sim \text{supp} + \text{dose} + \text{interaction}$
