

# Workshop 7.4a: Single factor ANOVA

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# Section 1

## Revision

# Estimation

Y	X
3	0
2.5	1
6	2
5.5	3
9	4
8.6	5
12	6

$$3.0 = \beta_0 \times 1 + \beta_1 \times 0 + \varepsilon_1$$

$$2.5 = \beta_0 \times 1 + \beta_1 \times 1 + \varepsilon_1$$

$$6.0 = \beta_0 \times 1 + \beta_1 \times 2 + \varepsilon_2$$

$$5.5 = \beta_0 \times 1 + \beta_1 \times 3 + \varepsilon_3$$

# Estimation

$$\begin{aligned}3.0 &= \beta_0 \times 1 + \beta_1 \times 0 + \varepsilon_1 \\2.5 &= \beta_0 \times 1 + \beta_1 \times 1 + \varepsilon_1 \\6.0 &= \beta_0 \times 1 + \beta_1 \times 2 + \varepsilon_2 \\5.5 &= \beta_0 \times 1 + \beta_1 \times 3 + \varepsilon_3 \\9.0 &= \beta_0 \times 1 + \beta_1 \times 4 + \varepsilon_4 \\8.6 &= \beta_0 \times 1 + \beta_1 \times 5 + \varepsilon_5 \\12.0 &= \beta_0 \times 1 + \beta_1 \times 6 + \varepsilon_6\end{aligned}$$

$$\underbrace{\begin{pmatrix} 3.0 \\ 2.5 \\ 6.0 \\ 5.5 \\ 9.0 \\ 8.6 \\ 12.0 \end{pmatrix}}_{\text{Response values}} = \underbrace{\begin{pmatrix} 1 & 0 \\ 1 & 1 \\ 1 & 2 \\ 1 & 3 \\ 1 & 4 \\ 1 & 5 \\ 1 & 6 \end{pmatrix}}_{\text{Model matrix}} \times \underbrace{\begin{pmatrix} \beta_0 \\ \beta_1 \end{pmatrix}}_{\text{Parameter vector}} + \underbrace{\begin{pmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \varepsilon_3 \\ \varepsilon_4 \\ \varepsilon_5 \\ \varepsilon_6 \end{pmatrix}}_{\text{Residual vector}}$$

# Matrix algebra

$$\underbrace{\begin{pmatrix} 3.0 \\ 2.5 \\ 6.0 \\ 5.5 \\ 9.0 \\ 8.6 \\ 12.0 \end{pmatrix}}_{\text{Response values}} = \underbrace{\begin{pmatrix} 1 & 0 \\ 1 & 1 \\ 1 & 2 \\ 1 & 3 \\ 1 & 4 \\ 1 & 5 \\ 1 & 6 \end{pmatrix}}_{\text{Model matrix}} \times \underbrace{\begin{pmatrix} \beta_0 \\ \beta_1 \end{pmatrix}}_{\text{Parameter vector}} + \underbrace{\begin{pmatrix} \epsilon_1 \\ \epsilon_2 \\ \epsilon_3 \\ \epsilon_4 \\ \epsilon_5 \\ \epsilon_6 \end{pmatrix}}_{\text{Residual vector}}$$

$$Y = X\beta + \epsilon$$

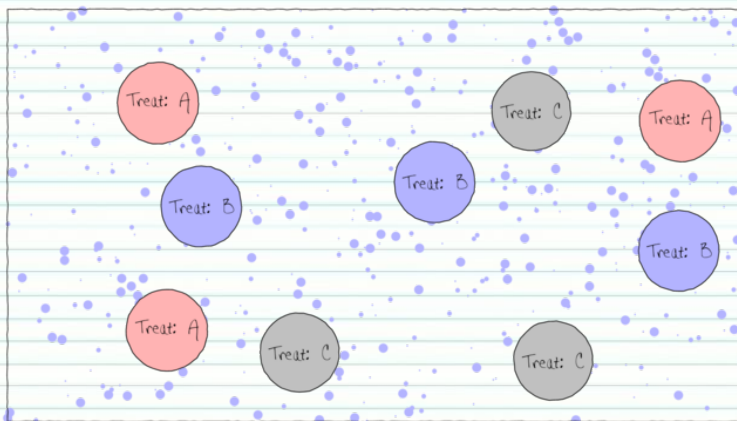
$$\hat{\beta} = (X'X)^{-1}X'Y$$

# Section 2

## Anova Parameterization

# Simple ANOVA

Three treatments (One factor - 3 levels), three replicates



# Simple ANOVA

Two treatments, three replicates





# Categorical predictor

Y	A	dummy1	dummy2	dummy3
2	G1	1	0	0
3	G1	1	0	0
4	G1	1	0	0
6	G2	0	1	0
7	G2	0	1	0
8	G2	0	1	0
10	G3	0	0	1
11	G3	0	0	1
12	G3	0	0	1

$$y_{ij} = \mu + \beta_1(\text{dummy}_1)_{ij} + \beta_2(\text{dummy}_2)_{ij} + \beta_3(\text{dummy}_3)_{ij} + \varepsilon_{ij}$$

# Overparameterized

$$y_{ij} = \mu + \beta_1(\text{dummy}_1)_{ij} + \beta_2(\text{dummy}_2)_{ij} + \beta_3(\text{dummy}_3)_{ij} + \varepsilon_{ij}$$

Y	A	Intercept	dummy1	dummy2	dummy3
2	G1	1	1	0	0
3	G1	1	1	0	0
4	G1	1	1	0	0
6	G2	1	0	1	0
7	G2	1	0	1	0
8	G2	1	0	1	0
10	G3	1	0	0	1
11	G3	1	0	0	1
12	G3	1	0	0	1

# Overparameterized

$$y_{ij} = \mu + \beta_1(\text{dummy}_1)_{ij} + \beta_2(\text{dummy}_2)_{ij} + \beta_3(\text{dummy}_3)_{ij} + \varepsilon_{ij}$$

- three treatment groups
- four parameters to estimate
- need to re-parameterize

# Categorical predictor

$$y_i = \mu + \beta_1(\text{dummy}_1)_i + \beta_2(\text{dummy}_2)_i + \beta_3(\text{dummy}_3)_i + \varepsilon_i$$

**MEANS** **PARAMETERIZATION**

$$y_i = \beta_1(\text{dummy}_1)_i + \beta_2(\text{dummy}_2)_i + \beta_3(\text{dummy}_3)_i + \varepsilon_{ij}$$

$$y_{ij} = \alpha_i + \varepsilon_{ij} \quad i = p$$

# Categorical predictor

## MEANS PARAMETERIZATION

$$y_i = \beta_1(\text{dummy}_1)_i + \beta_2(\text{dummy}_2)_i + \beta_3(\text{dummy}_3)_i + \varepsilon_i$$

Y	A	dummy1	dummy2	dummy3
2	G1	1	0	0
3	G1	1	0	0
4	G1	1	0	0
6	G2	0	1	0
7	G2	0	1	0
8	G2	0	1	0
10	G3	0	0	1
11	G3	0	0	1
12	G3	0	0	1

# Categorical predictor DD

## MEANS PARAMETERIZATION

$$y_i = \alpha_1 D_{1i} + \alpha_2 D_{2i} + \alpha_3 D_{3i} + \varepsilon_i$$

$$y_i = \alpha_p + \varepsilon_i,$$

	Y	A
1	2.00	G1
2	3.00	G1
3	4.00	G1
4	6.00	G2
5	7.00	G2
6	8.00	G2
7	10.00	G3
8	11.00	G3
9	12.00	G3

where  $p$  = number of levels of the factor and  $D$  = dummy variables

$$\begin{pmatrix} 2 \\ 3 \\ 4 \\ 6 \\ 7 \\ 8 \\ 10 \\ 11 \\ 12 \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 1 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \\ 0 & 0 & 1 \\ 0 & 0 & 1 \end{pmatrix} \times \begin{pmatrix} \alpha_1 \\ \alpha_2 \\ \alpha_3 \end{pmatrix} + \begin{pmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \varepsilon_3 \\ \varepsilon_4 \\ \varepsilon_5 \\ \varepsilon_6 \\ \varepsilon_7 \\ \varepsilon_8 \\ \varepsilon_9 \end{pmatrix}$$

# Categorical predictor

## MEANS PARAMETERIZATION

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Parameter	Estimates	Null Hypothesis
$\alpha_1^*$	mean of group 1	$H_0: \alpha_1 = \alpha_1 = 0$
$\alpha_2^*$	mean of group 2	$H_0: \alpha_2 = \alpha_2 = 0$
$\alpha_3^*$	mean of group 3	$H_0: \alpha_3 = \alpha_3 = 0$

---

```
> summary(lm(Y~-1+A))$coef
```

	Estimate	Std. Error	t value	Pr(> t )
AG1	3	0.5773503	5.196152	2.022368e-03
AG2	7	0.5773503	12.124356	1.913030e-05
AG3	11	0.5773503	19.052559	1.351732e-06

# Categorical predictor

$$y_i = \mu + \beta_1(\text{dummy}_1)_i + \beta_2(\text{dummy}_2)_i + \beta_3(\text{dummy}_3)_i + \varepsilon_i$$

**EFFECTS**      **PARAMETERIZATION**

$$y_{ij} = \mu + \beta_2(\text{dummy}_2)_{ij} + \beta_3(\text{dummy}_3)_{ij} + \varepsilon_{ij}$$

$$y_{ij} = \mu + \alpha_i + \varepsilon_{ij} \quad i = p - 1$$



# Categorical predictor

## EFFECTS    PARAMETERIZATION

$$y_i = \alpha + \beta_2(\text{dummy}_2)_i + \beta_3(\text{dummy}_3)_i + \varepsilon_i$$

Y	A	alpha	dummy2	dummy3
2	G1	1	0	0
3	G1	1	0	0
4	G1	1	0	0
6	G2	1	1	0
7	G2	1	1	0
8	G2	1	1	0
10	G3	1	0	1
11	G3	1	0	1
12	G3	1	0	1

# Categorical predictor

EFFECTS

PARAMETERIZATION

	Y	A
1	2.00	G1
2	3.00	G1
3	4.00	G1
4	6.00	G2
5	7.00	G2
6	8.00	G2
7	10.00	G3
8	11.00	G3
9	12.00	G3

$$y_i = \alpha + \beta_2 D_{2i} + \beta_3 D_{3i} + \varepsilon_i$$

$$y_i = \alpha_p + \varepsilon_i,$$

where  $p$  = number of levels of the factor minus 1 and

$D$  = dummy variables

$$\begin{pmatrix} 2 \\ 3 \\ 4 \\ 6 \\ 7 \\ 7 \\ 8 \\ 10 \\ 11 \\ 12 \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 1 & 0 \\ 1 & 1 & 0 \\ 1 & 1 & 0 \\ 1 & 1 & 0 \\ 1 & 0 & 1 \\ 1 & 0 & 1 \\ 1 & 0 & 1 \end{pmatrix} \times \begin{pmatrix} \mu \\ \alpha_2 \\ \alpha_3 \end{pmatrix} + \begin{pmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \varepsilon_3 \\ \varepsilon_4 \\ \varepsilon_5 \\ \varepsilon_6 \\ \varepsilon_7 \\ \varepsilon_8 \\ \varepsilon_9 \end{pmatrix}$$

# Categorical predictor

## TREATMENT CONTRASTS

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Parameter	Estimates	Null Hypothesis
Intercept	mean of control group	$H_0: \mu = \mu_1 = 0$
$\alpha_2^*$	mean of group 2 minus mean of control group	$H_0: \alpha_2 = \alpha_2 = 0$
$\alpha_3^*$	mean of group 3 minus mean of control group	$H_0: \alpha_3 = \alpha_3 = 0$

---

```
> contrasts(A) <-contr.treatment  
> contrasts(A)
```

# Categorical predictor

## TREATMENT CONTRASTS

---

Parameter	Estimates	Null Hypothesis
Intercept	mean of control group	$H_0: \mu = \mu_1 = 0$
$\alpha_2^*$	mean of group 2 minus mean of control group	$H_0: \alpha_2 = \alpha_2 = 0$
$\alpha_3^*$	mean of group 3 minus mean of control group	$H_0: \alpha_3 = \alpha_3 = 0$

---

```
> summary(lm(Y~A))$coef
```

# Categorical predictor

## USER DEFINED CONTRASTS

User defined contrasts

Grp2 vs Grp3

Grp1 vs (Grp2 & Grp3)

	Grp1	Grp2	Grp3
$\alpha_2^*$	0	1	-1
$\alpha_3^*$	1	-0.5	-0.5

```
> contrasts(A) <- cbind(c(0,1,-1),c(1, -0.5, -0.5))  
> contrasts(A)
```

```
      [,1] [,2]  
G1      0  1.0
```

# Categorical predictor

## USER DEFINED CONTRASTS

- $p - 1$  comparisons (contrasts)
- all contrasts must be orthogonal

# Categorical predictor

## ORTHOGONALITY

Four groups (A, B, C, D)

$p - 1 = 3$  comparisons

1. A vs B ::  $A > B$

2. B vs C ::  $B > C$

3. A vs C ::

# Categorical predictor

## USER DEFINED CONTRASTS

```
> contrasts(A) <- cbind(c(0,1,-1),c(1, -0.5, -0.5))  
> contrasts(A)
```

	[,1]	[,2]
G1	0	1.0
G2	1	-0.5
G3	-1	-0.5

$$\begin{aligned}0 \times 1.0 &= 0 \\1 \times -0.5 &= -0.5 \\-1 \times -0.5 &= 0.5 \\ \text{sum} &= 0\end{aligned}$$



# Categorical predictor

## USER DEFINED CONTRASTS

```
> contrasts(A) <- cbind(c(0,1,-1),c(1, -0.5, -0.5))  
> contrasts(A)
```

```
  [,1] [,2]  
G1    0  1.0  
G2    1 -0.5  
G3   -1 -0.5
```

```
> crossprod(contrasts(A))
```

```
  [,1] [,2]  
[1,]   2  0.0  
[2,]   0  1.5
```

```
> summary(lm(Y~A))$coef
```

# Categorical predictor

## USER DEFINED CONTRASTS

```
> contrasts(A) <- cbind(c(1, -0.5, -0.5),c(1,-1,0))  
> contrasts(A)
```

```
      [,1] [,2]  
G1  1.0    1  
G2 -0.5   -1  
G3 -0.5    0
```

```
> crossprod(contrasts(A))
```

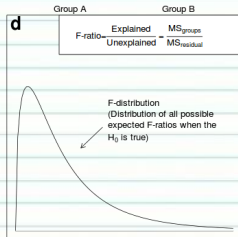
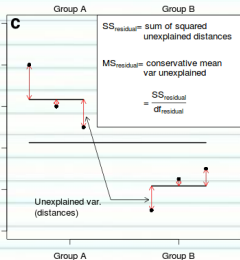
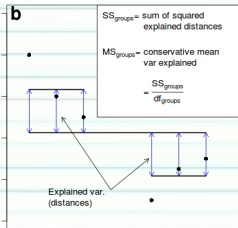
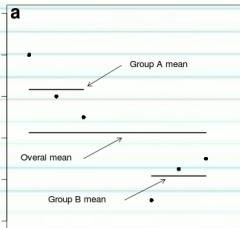
```
      [,1] [,2]  
[1,]  1.5  1.5  
[2,]  1.5  2.0
```

# Section 3

## Partitioning of variance (ANOVA)

# ANOVA

## PARTITIONING VARIANCE



# ANOVA

## PARTITIONING VARIANCE

```
> anova(lm(Y~A))
```

Analysis of Variance Table

Response: Y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
A	2	96	48	48	0.0002035 ***
Residuals	6	6	1		

---

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

# Categorical predictor

## POST-HOC COMPARISONS

No. of Groups	No. of comparisons	Familywise Type I error probability
3	3	0.14
5	10	0.40
10	45	0.90

# Categorical predictor

## POST-HOC COMPARISONS

Bonferoni

```
> summary(lm(Y~A))$coef
```

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	7	0.3333333	21.000000	7.595904e-07
A1	-8	0.9428090	-8.485281	1.465426e-04
A2	4	0.8164966	4.898979	2.713682e-03

```
> 0.05/3
```

```
[1] 0.01666667
```

# Categorical predictor

## POST-HOC COMPARISONS

Tukey's test

```
> library(multcomp)
> data.lm<-lm(Y~A)
> summary(glht(data.lm, linfct=mcp(A="Tukey")))
```

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

Fit: lm(formula = Y ~ A)

Linear Hypotheses:

	Estimate	Std. Error	t value	Pr(> t )	
G2 - G1 == 0	4.0000	0.8165	4.899	0.00653	**
G3 - G1 == 0	8.0000	0.8165	9.798	< 0.001	***
G3 - G2 == 0	4.0000	0.8165	4.899	0.00679	**



# Assumptions

- Normality
- Homogeneity of variance
- Independence
  
- As for regression

# Section 4

## Worked Examples

# Worked Examples

```
> day <- read.csv('../data/day.csv', strip.white=T)
> head(day)
```

	TREAT	BARNACLE
1	ALG1	27
2	ALG1	19
3	ALG1	18
4	ALG1	23
5	ALG1	25
6	ALG2	24

# Worked Examples

Question: what effects do different substrate types have on barnacle recruitment

Linear model:

$$\text{Barnacle}_i = \mu + \alpha_j + \varepsilon_i \quad \varepsilon \sim \mathcal{N}(0, \sigma^2)$$

# Worked Examples

```
> partridge <- read.csv('../data/partridge.csv', strip.white=T)
> head(partridge)
```

	GROUP	LONGEVITY
1	PREG8	35
2	PREG8	37
3	PREG8	49
4	PREG8	46
5	PREG8	63
6	PREG8	39

```
> str(partridge)
```

```
'data.frame': 125 obs. of 2 variables:
```

```
$ GROUP : Factor w/ 5 levels "NONE0","PREG1",...: 3 3 3 3 3 3 3 3 3 3 ...
```

```
$ LONGEVITY: int 35 37 49 46 63 39 46 56 63 65 ...
```

# Worked Examples

Question: what effects does mating have on the longevity of male fruitflies

Linear model:

$$\text{Longevity}_i = \mu + \alpha_j + \varepsilon_i \quad \varepsilon \sim \mathcal{N}(0, \sigma^2)$$