Fruit Bursts

I need 4 volunteers...

How many choices does the **4th** person have in their choice of Fruit Burst?



Fruit Bursts

I need 4 volunteers...

How many choices does the 4th person have?... Only (4 - 1) = 3 of you have the freedom to choose your Fruit Burst!

Constraint: there are only 4 Fruit Bursts and each person "has" to eat a unique one

Degrees of freedom

What is a degree of freedom?

The degree of freedom is

- the number of variables that are free to vary without breaking a constraint,
- the number of independent observations or measurements that can be made in order to calculate some statistic.

Example

Given data sample $x = \{1, 5, 6\}$

How many elements of x are **free to vary** (independently)?

Example (1-SAMPLE T TEST)

Given data sample $x = \{1, 5, 6\}$

Constraint: we would like to estimate the mean of the population.

Due to maths... mean(x) * (number of observations in x) = sum of observations

So sum of observations (must) equal 4 * 3 = 12

How many elements of x are **free to vary** (independently)?

```
> t.test(x = c(1, 5, 6), mu = 0)
```

One Sample t-test

data: c(1, 5, 6) t = 2.6186, df = 2, p-value = 0.1201

alternative hypothesis: true mean is not equal to 0 95 percent confidence interval:

-2.572411 10.572411

sample estimates:

mean of x

Example (2 SAMPLE T TEST)

Given data sample $x = \{1, 5, 6\}$ and $k = \{7, 9, 11\}$

Constraint: we would like to estimate the differences between means.

Due to maths... mean(.) * (number of observations in .) = sum of observations

So, sum of x (must) equal 4 * 3 = 12 & sum of k (must) equal 9 * 3 = 27

How many elements of x and k combined are **free to vary** (independently)?

```
> t.test(x = c(1, 5, 6), y = c(7, 9, 11), var.equal = TRUE)
```

Two Sample t-test

data: c(1, 5, 6) and c(7, 9, 11)

t = -2.6112, df = 4, p-value = 0.05935

alternative hypothesis: true difference in means is not equal to 0

95 percent confidence interval:

-10.3164876 0.3164876

sample estimates:

mean of x mean of y

Example (PAIRED T TEST)

Given data sample $x = \{1, 5, 6, 7, 9, 11\}$ measurements taken from three different individuals before and after some event

Constraint: we would like to estimate the differences between/change in means.

Due to maths... mean(.) * (number of observations in .) = sum of observations

How many elements of x are **free to vary** (**independently**)?

```
> t.test(x = c(1, 5, 6), y = c(7, 9, 11), var.equal = TRUE, paired = TRUE)
```

Paired t-test

alternative hypothesis: true mean difference is not equal to 0

data: c(1, 5, 6) and c(7, 9, 11) t = -8.6603, df = 2, p-value = 0.01307

95 percent confidence interval:

- 5

-7.484138 -2.515862 sample estimates: mean difference

Example (ANOVA)

Given data sample $x = \{1, 5, 6\}, k = \{7, 9, 11\}$ and $h = \{1, 10, 22\}$

Constraint: we would like to estimate the differences between means.

Due to maths... mean(.) * (number of observations in .) = sum of observations

So, sum of x (must) equal 4 * 3 = 12 & sum of k (must) equal 9 * 3 = 27 & sum of h (must) equal 11 * 3 = 33

How many elements of x, k, and h combined are **free to vary** (independently)?

```
> data <- data.frame(y = c(1, 5, 6, 7, 9, 11, 1, 10, 22), group = rep(c("x", "k", "h"), each = 3))
> aov(y ~ group, data = data) |> summary()
           Df Sum Sq Mean Sq F value Pr(>F)
            2 78 39.00 0.959 0.435
group
Residuals 6 244 40.67
> lm(y ~ group, data = data) |> summary()
Call:
lm(formula = v \sim group, data = data)
Residuals:
  Min
          10 Median
                       30
                             Max
   -10 -2
                              11
```

```
(Intercept) 11.000 3.682 2.988 0.0244 * groupk -2.000 5.207 -0.384 0.7141 groupx -7.000 5.207 -1.344 0.2274
```

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

Estimate Std. Error t value Pr(>|t|)

Residual standard error: 6.377 on 6 degrees of freedom Multiple R-squared: 0.2422, Adjusted R-squared: -0.01035

Coefficients:

F-statistic: 0.959 on 2 and 6 DF, p-value: 0.4351

| Example (REGRESSION) | |
|--|--|
| <pre>lm(logAUC ~ Disease*Organ, data = data)</pre> | |

| Healthy |
|---------|
| Healthy |
| Diabeti |

Disease

Organ

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

Animal

1

2

3

4

5

6

7

8

10

11

12

Sample

1

2

1

2

1

2

1

2

1

2

1

2

logAUC

9.40

8.83

10.33

10.49

9.74

10.98

7.92

9.37

8.69

11.31

7.01

9.29

```
Rows: 12 Columns: 5
0s— Column specification
Delimiter: ","
chr (2): Disease, Organ
dbl (3): Animal, Sample, logAUC
i Use `spec()` to retrieve the full column specification for this data.
i Specify the column types or set `show col types = FALSE` to quiet this message.
> aov(logAUC ~ Disease*Organ, data = data) |> summary()
             Df Sum Sq Mean Sq F value Pr(>F)
Disease
             1 3.183 3.183 3.630 0.0932 .
Organ
             1 4.296 4.296 4.900 0.0578 .
Disease:Organ 1 2.539 2.539 2.896 0.1272
Residuals 8 7.014 0.877
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> lm(logAUC ~ Disease*Organ, data = data) |> summarv()
Call:
lm(formula = logAUC ~ Disease * Organ, data = data)
Residuals:
    Min
              10 Median
                               30
                                       Max
-1.27000 -0.64000 -0.01833 0.58417 1.32000
Coefficients:
                           Estimate Std. Error t value Pr(>|t|)
                            7.8733
(Intercept)
                                       0.5406 14.565 4.84e-07 ***
DiseaseHealthy
                            1.9500
                                       0.7645 2.551 0.0341 *
OrganouterLV
                            2.1167
                                      0.7645 2.769 0.0243 *
DiseaseHealthy:OrganouterLV -1.8400
                                      1.0812 -1.702 0.1272
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.9363 on 8 degrees of freedom
Multiple R-squared: 0.5882, Adjusted R-squared: 0.4338
```

F-statistic: 3.809 on 3 and 8 DF, p-value: 0.05788

> data <- readr::read csv("https://raw.githubusercontent.com/STATS-UOA/databunker/master/data/factorial expt.csv")</pre>

| Example (REGRESSION) | |
|---|--|
| <pre>lm(logAUC ~ Disease*Organ + Animal, data = data)</pre> | |

| LIXAL | ibie (wrowrooiou | |
|----------------------------|----------------------------------|--|
| <pre>lm(logAUC data)</pre> | ~ Disease*Organ + Animal, data = | |
| | | |
| | | |
| | | |



Disease

Organ

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

innerLV

outerLV

Animal

1

1

2

2

3

3

4

4

5

5

6

6

logAUC

9.40

8.83

10.33

10.49

9.74

10.98

7.92

9.37

8.69

11.31

7.01

9.29

Sample

1

2

1

2

1

2

1

2

1

2

1

2

```
> data <- readr::read csv("https://raw.githubusercontent.com/STATS-UOA/databunker/master/data/split plot.csv")</pre>
Rows: 12 Columns: 5
 0s— Column specification
Delimiter: ","
chr (2): Disease, Organ
dbl (3): Animal, Sample, logAUC
i Use `spec()` to retrieve the full column specification for this data.
i Specify the column types or set `show col types = FALSE` to quiet this message.
> aov(logAUC ~ Disease*Organ + Animal, data = data) |> summary()
             Df Sum Sq Mean Sq F value Pr(>F)
Disease
              1 3.183 3.183 3.309 0.1117
              1 4.296 4.296 4.467 0.0724 .
Organ
Animal
              1 0.281 0.281 0.292 0.6054
Disease:Organ 1 2.539 2.539 2.640 0.1482
Residuals
            7 6.732 0.962
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> lm(logAUC ~ Disease*Organ + Animal, data = data) |> summary()
Call:
lm(formula = logAUC ~ Disease * Organ + Animal, data = data)
Residuals:
    Min
              10 Median
                                30
                                        Max
-1.08250 -0.54625 -0.00083 0.55312 1.32000
Coefficients:
                           Estimate Std. Error t value Pr(>|t|)
```

1.8238 3.803 0.00669 **

1.3127 1.914 0.09718 .

0.8007 2.643 0.03326 *

1.1324 -1.625 0.14822

0.541 0.60544

0.3467

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

Residual standard error: 0.9807 on 7 degrees of freedom

Multiple R-squared: 0.6047, Adjusted R-squared: 0.3788

F-statistic: 2.677 on 4 and 7 DF, p-value: 0.121

6.9358

2.1167

0.1875

2.5125

(Intercept)

OrganouterLV

Animal

DiseaseHealthy

DiseaseHealthy:OrganouterLV -1.8400