



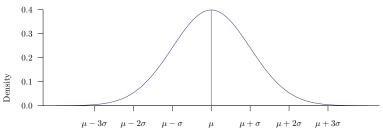
Analysis of Variance (ANOVA)

Cancer Research UK – 19^{th} of July 2017 D.-L. Couturier / M. Dunning / R. Nicholls

Quick review: Normal distribution

$$\begin{split} X \sim N(\mu, \sigma^2), \qquad f_X(x) &= \frac{1}{\sqrt{2\pi\sigma^2}} \; e^{-\frac{(x-\mu)^2}{2\sigma^2}} \\ \mathrm{E}[X] &= \mu, \qquad \mathrm{Var}[X] = \sigma^2, \\ Z &= \frac{X-\mu}{\sigma} \sim N(0,1), \qquad f_Z(z) = \frac{1}{\sqrt{2\pi}} \; e^{-\frac{x^2}{2}}. \end{split}$$

Probability density function of a normal distribution:



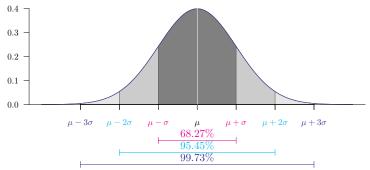
Quick review: Normal distribution

$$X \sim N(\mu, \sigma^2), \qquad f_X(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

$$\mathsf{E}[X] = \mu, \qquad \mathsf{Var}[X] = \sigma^2,$$

$$Z = \frac{X - \mu}{\sigma} \sim N(0, 1), \qquad f_Z(z) = \frac{1}{\sqrt{2\pi}} e^{-\frac{x^2}{2}}.$$

Probability density function of a normal distribution:

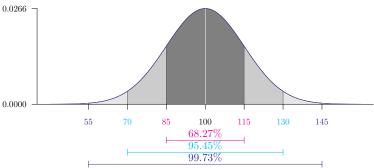




Quick review: Normal distribution

$$\begin{split} X \sim N(\mu, \sigma^2), \qquad f_X(x) &= \frac{1}{\sqrt{2\pi\sigma^2}} \; e^{-\frac{(x-\mu)^2}{2\sigma^2}} \\ \mathrm{E}[X] &= \mu, \qquad \mathrm{Var}[X] = \sigma^2, \\ Z &= \frac{X-\mu}{\sigma} \sim N(0,1), \qquad f_Z(z) = \frac{1}{\sqrt{2\pi}} \; e^{-\frac{x^2}{2}}. \end{split}$$

Suitable modelling for a lot of phenomena: IQ $\sim N(100,15^2)$.





Quick review: Tests

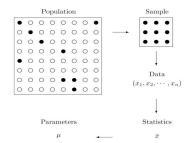
▶ Define hypotheses

$$\triangleright \ \mathbf{H0:} \ \mu = \mu_0 \quad \triangleright \ \mathbf{H1:} \ \begin{matrix} \mu < \mu_0 \\ \mu \neq \mu_0 \\ \mu > \mu_0 \end{matrix}$$

- ightharpoonup Chose α
- ightharpoonup Collect data and estimate T_{obs}
- ▶ Define p-value, knowing the distribution of T under **H0**

$$p-\text{value} = \begin{cases} P(T < T_{obs}) & \textbf{H1: } \mu < \mu_0, \\ P(|T| > T_{obs}) & \textbf{H1: } \mu \neq \mu_0, \\ P(T > T_{obs}) & \textbf{H1: } \mu > \mu_0. \end{cases}$$

- ▶ Compare p-value to α
 - ightharpoonup if p- value $< lpha \longrightarrow {\sf accept}\ {\sf H1}$ at the lpha level
 - if p value $> \alpha \longrightarrow$ do not reject H0 at the α level





 $f_T(t)$

Diet B

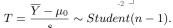
Assumed model

$$Y_i = \mu + \epsilon_i,$$
 where $i = 1, ..., n$ and $\epsilon_i \sim N(0, \sigma^2).$

Hypotheses ▶ **H0:** $\mu = 0$,

One Sample t-test

- ▶ **H1:** μ < 0.
- ► Test statistic's distribution under H0

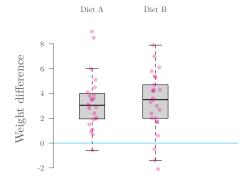


6 -

2 -

```
data: dietB
t = 6.6301, df = 24, p-value = 3.697e-07
alternative hypothesis: true mean is greater than 0
95 percent confidence interval:
 2.424694
               Inf
sample estimates:
mean of x
    3.268
```





Two-sample location tests: t-tests and Mann-Whitney-Wilcoxon's test

Two independent sample Student's t-test

▶ Assumed model

$$\begin{split} Y_{i(g)} &= \mu_g + \epsilon_{i(g)}, \\ &= \mu + \delta_g + \epsilon_{i(g)}, \\ \text{where } g &= A, B, \ i = 1, ..., n_g, \\ \epsilon_{i(g)} &\sim N(0, \sigma^2) \text{ and } \sum n_g \delta_g = 0. \end{split}$$

Hypotheses

 \triangleright **H0**: $\mu_A = \mu_B$,

 \triangleright **H1**: $\mu_A \neq \mu_B$.

3.268

► Test statistic's distribution under H0

$$T = \frac{(\overline{Y}_A - \overline{Y}_B) - (\mu_A - \mu_B)}{s_p \sqrt{n_A^{-1} + n_B^{-1}}} \sim Student(n_A + n_B - 2).$$

Weight difference

Diet. A

Diet B

Two Sample t-test

3.300

```
data: dietA and dietB
t = 0.0475, df = 47, p-value = 0.9623
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-1.323275   1.387275
sample estimates:
mean of x mean of y
```



Two independent sample Welch's t-test

Assumed model

$$\begin{split} Y_{i(g)} &= \mu_g + \epsilon_{i(g)}, \\ &= \mu + \delta_g + \epsilon_{i(g)}, \\ \text{where } g &= A, B, \ i = 1, ..., n_g, \\ \epsilon_{i(g)} &\sim N(0, \sigma_g^2) \text{ and } \sum n_g \delta_g = 0. \end{split}$$

Hypotheses

mean of x mean of y 3.300

▶ **H0**: $\mu_A = \mu_B$,

 \triangleright **H1:** $\mu_A \neq \mu_B$.

3.268

Test statistic's distribution under H0

Diet. A

Diet B

```
Welch Two Sample t-test
data: dietA and dietB
t = 0.047594, df = 46.865, p-value = 0.9622
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-1.320692 1.384692
sample estimates:
```

 $T = \frac{(\overline{Y}_A - \overline{Y}_B) - (\mu_A - \mu_B)}{\sqrt{s_X^2/n_X + s_Y^2/n_Y}} \sim Student(\mathrm{df}).$



Two independent sample Mann-Whitney-Wilcoxon test

Assumed model

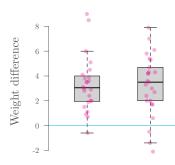
$$\begin{split} Y_{i(g)} &= \theta_g + \epsilon_{i(g)}, \\ &= \theta + \delta_g + \epsilon_{i(g)}, \\ \text{where } g &= A, B, \ i = 1, ..., n_g, \\ \epsilon_{i(g)} &\sim iid(0, \sigma^2) \ \text{and} \ \sum n_g \delta_g = 0. \end{split}$$

Hypotheses

 \triangleright **H0**: $\theta_A = \theta_B$,

 \triangleright **H1:** $\theta_A \neq \theta_B$.

► Test statistic's distribution under **H0**



Diet. A

Diet B

$$z = \frac{\sum_{i=1}^{n_B} R_{i(g)} - [n_B(n_A + n_B + 1)/2]}{\sqrt{n_A n_B(n_A + n_B + 1)/12}},$$

where

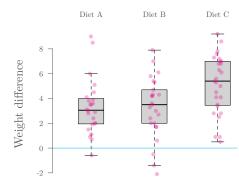
 $ightharpoonup R_{i(g)}$ denotes the global rank of the ith observation of group g.

Wilcoxon rank sum test with continuity correction

data: dietA and dietB
W = 277, p-value = 0.6526

W = 277, p-value = 0.6526 alternative hypothesis: true location shift is not equal to 0





Two or more sample location tests: one-way ANOVA & multiple comparisons

More than two sample case: Fisher's one-way ANOVA

Assumed model

$$\begin{split} Y_{i(g)} &= \mu_g + \epsilon_{i(g)}, \\ &= \mu + \delta_g + \epsilon_{i(g)}, \\ \text{where } g &= 1, ..., G, \ i = 1, ..., n_g, \\ \epsilon_{i(g)} &\sim N(0, \sigma^2) \text{ and } \sum n_g \delta_g = 0. \end{split}$$

► Hypotheses

$$\triangleright$$
 H0: $\mu_1 = \mu_2 = ... = \mu_G$,

ightharpoonup H1: $\mu_k
eq \mu_l$ for at least one pair (k,l).

► Test statistic's distribution under H0

$$F = \frac{Ns_{\overline{Y}}^2}{s_n^2} \sim Fisher(G - 1, N - G),$$

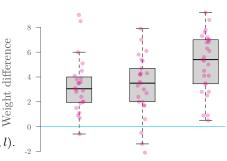
diet.type

Residuals

where

$$ightharpoonup s_p^2 = \frac{1}{N-G} \sum_{g=1}^G (n_g - 1) s_g^2,$$

$$N = \sum n_g, \ \overline{\overline{Y}} = \frac{1}{N} \sum_{g=1}^G n_g \overline{Y}_g.$$



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

30.264

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

Diet B

Diet C

Diet A



5.383 0.0066 **

More than two sample case: Welch's one-way ANOVA

Assumed model

$$\begin{split} Y_{i(g)} &= \mu_g + \epsilon_{i(g)}, \\ &= \mu + \delta_g + \epsilon_{i(g)}, \\ \text{where } g &= 1, ..., G, \ i = 1, ..., n_g, \\ \epsilon_{i(g)} &\sim N(0, \sigma_g^2) \text{ and } \sum n_g \delta_g = 0. \end{split}$$

Hypotheses

$$ho$$
 H0: $\mu_1 = \mu_2 = ... = \mu_G$,

 \triangleright **H1:** $\mu_k \neq \mu_l$ for at least one pair (k, l).

Test statistic's distribution under H0

$$F^* = \frac{s_{\overline{Y}}^{*2}}{1 + \frac{2(G-2)}{2}} \sim Fisher(G-1, \Delta),$$

Veight difference

where

$$\mathbf{w}_g = \frac{n_g}{s_g^2}, \ \overline{\overline{Y}}^{\star} = \sum_{g=1}^G \frac{w_g \overline{Y}_g}{\sum_i w_g}.$$

Diet A

Diet B

Diet C

data: weight.diff and diet.type F = 5.2693, num df = 2.00, denom df = 48.48, p-value = 0.008497



More than two sample case: Kruskal-Wallis test

Assumed model

$$\begin{split} Y_{i(g)} &= \theta_g + \epsilon_{i(g)}, \\ &= \theta + \delta_g + \epsilon_{i(g)}, \\ \text{where } g &= 1, ..., G, \ i = 1, ..., n_g, \\ \epsilon_{i(g)} &\sim iid(0, \sigma^2) \ \text{and} \ \sum n_g \delta_g = 0. \end{split}$$

Hypotheses

$$\triangleright$$
 H0: $heta_1= heta_2=...= heta_G$,

 \triangleright **H1:** $\theta_k \neq \theta_l$ for at least one pair (k, l).

Test statistic's distribution under **H0**

Diet. A

Diet B

Diet C

$$H = \frac{\frac{12}{N(N+1)} \sum_{g=1}^{G} \frac{\overline{R}_g}{n_g} - 3(N-1)}{1 - \frac{\sum_{v=1}^{V} t_v^3 - v}{N^3 - N}} \sim \chi(G-1),$$

where

- $\overline{R}_g = \frac{1}{n_g} \sum_{i=1}^{n_g} R_{i(g)}$ and $R_{i(g)}$ denotes the global rank of the *i*th observation of group g,
- $lackbox{$V$}$ is the number of different values/levels in f y and t_v denotes the number of times a given value/level occurred in f y.

Kruskal-Wallis rank sum test

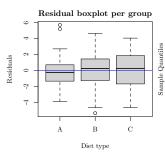


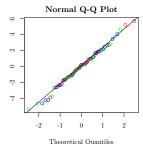
Model check: Residual analysis

$$Y_{i(g)} = \theta_g + \epsilon_{i(g)}$$
$$\hat{\epsilon}_{i(g)} = Y_{i(g)} - \hat{\theta}_g,$$

where

- $ightharpoonup \widehat{\epsilon}_{i(q)} \sim N(0, \widehat{\sigma}^2)$ for Fisher's ANOVA
- $lackbox{}\widehat{\epsilon}_{i(g)}\sim N(0,\widehat{\sigma}_g^2)$ for Welch's ANOVA
- $\widehat{\epsilon}_{i(g)} \sim iid(0,\widehat{\sigma}^2)$ for Kruskal-Wallis' ANOVA





Shapiro-Wilk normality test

data: diet\$resid.mean
W = 0.99175, p-value = 0.9088

Bartlett test of homogeneity of variances

data: diet\$resid.mean by as.numeric(diet\$diet.type)
Bartlett's K-squared = 0.21811, df = 2, p-value = 0.8967



Finding different pairs: Multiple comparisons

► All-pairwise comparison problem:

Interested in finding which pair(s) are different by testing

$$ho$$
 H0₁: $\mu_1 = \mu_2$, ho H0₂: $\mu_1 = \mu_3$, ... ho H0_K: $\mu_{G-1} = \mu_G$, leading to a total of $K = G(G-1)/2$ pairwise comparisons.

▶ Family-wise type I error for K tests, α_K

For each test, the probability of rejecting H0 when H0 is true equals α . For K independent tests, the probability of rejecting H0 at least 1 time when H0 is true, α_K , is given by

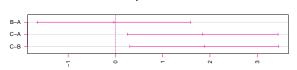
$$\alpha_K = 1 - (1 - \alpha)^K$$
. $\Rightarrow \alpha_1 = 0.05,$
 $\Rightarrow \alpha_2 = 0.0975,$
 $\Rightarrow \alpha_{10} = 0.4013.$

► Multiplicity correction

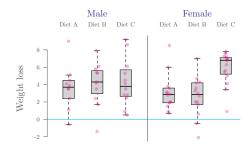
Principle: change the level of each test so that $\alpha_K = 0.05$, for example:

- **b** Bonferroni's correction (indep. tests): $\alpha = \alpha_K/K$,
- Dunn-Sidak's correction (indep. tests): $\alpha = 1 (1 \alpha_K)^{1/K}$,
- Tukey's correction (dependent tests).

95% family-wise confidence level







Two or more sample location tests: two-way ANOVA

More than one factor: Fisher's two-way ANOVA

Assumed model

$$Y_{i(g)} = \mu_{gk} + \epsilon_{i(gk)},$$

= $\mu + \delta_g + \delta_k + \delta_{gk} + \epsilon_{i(gk)},$

- ightharpoonup g = 1, ..., G, k = 1, ..., K,
- $\begin{array}{ll}
 \bullet & i = 1, ..., n_g, \\
 \bullet & \epsilon_{i(gk)} \sim N(0, \sigma^2)
 \end{array}$
- $\sum n_q \delta_q = \sum n_k \delta_k = \sum n_{qk} \delta_{qk} = 0.$

Hypotheses

$$ho$$
 H0₁: $\delta_g = 0 \ \forall \ g$, ho H1₁: H0₁ is false.

 \triangleright H0₂: $\delta_k = 0 \ \forall \ k$, \triangleright H1₂: H0₂ is false.

Male

Diet B

Diet C

Diet A



Female

Diet B

Diet C

More than one factor: Fisher's two-way ANOVA

Assumed model

$$Y_{i(g)} = \mu_{gk} + \epsilon_{i(gk)},$$

= $\mu + \delta_g + \delta_k + \delta_{gk} + \epsilon_{i(gk)},$

- ightharpoonup g = 1, ..., G, k = 1, ..., K,
- $i = 1, ..., n_g,$ $\epsilon_{i(gk)} \sim N(0, \sigma^2)$
- $\sum n_g \delta_g = \sum n_k \delta_k = \sum n_{gk} \delta_{gk} = 0.$

Hypotheses

 \triangleright H0₁: $\delta_g = 0 \ \forall \ g$, \triangleright H1₁: H0₁ is false.

 \triangleright H0₂: $\delta_k = 0 \ \forall \ k$, \triangleright H1₂: H0₂ is false.

Veight loss

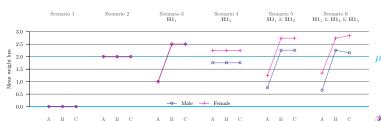
 \triangleright H0₃: $\delta_{gk} = 0 \ \forall \ g, k$, \triangleright H1₃: H0₃ is false.

Male

Diet B

Diet C

Diet A





Female

Diet B

Diet C

More than one factor: Fisher's two-way ANOVA

Assumed model

- $i = 1, ..., n_q$
- $ightharpoonup \epsilon_{i(qk)} \sim N(0, \sigma^2)$

Hypotheses

```
\triangleright H0<sub>1</sub>: \delta_q = 0 \ \forall \ q ,
                                                                       \triangleright H0<sub>2</sub>: \delta_k = 0 \ \forall \ k,
▶ H1₁: H0₁ is false.
                                                                       \triangleright H1<sub>2</sub>: H0<sub>2</sub> is false.
```

 \triangleright H0₃: $\delta_{qk} = 0 \ \forall \ q, k$, ▶ H13: H03 is false.

Male

Diet B

Diet C

Diet A

```
Df Sum Sq Mean Sq F value Pr(>F)
                     60.5
                           30.264
diet.type
                                     5.629 0.00541 **
gender
                      0.2
                           0.169 0.031 0.85991
diet.type:gender
                     33.9
                           16.952
                                     3.153 0.04884 *
                    376.3
                           5.376
Residuals
                 70
```

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



Female

Diet B

Summary

