



Analysis of Variance (ANOVA)

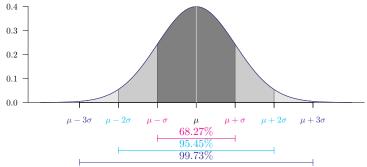
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Quick review: Normal distribution

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

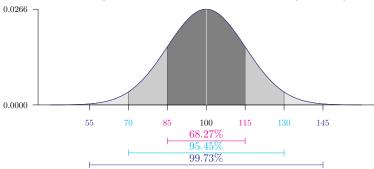
Probability density function of a normal distribution:



Quick review: Normal distribution

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

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

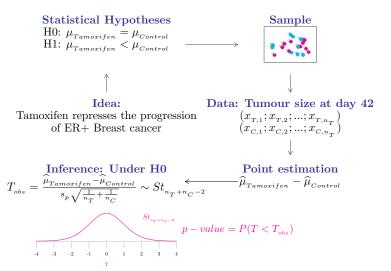


Grand Picture of Statistics

$$\begin{array}{c} \text{Statistical Hypotheses} \\ \text{H0: } \mu_{Tamoxifen} = \mu_{Control} \\ \text{H1: } \mu_{Tamoxifen} < \mu_{Control} \\ \end{array} \\ \rightarrow \begin{array}{c} \text{Idea:} \\ \text{Tamoxifen represses the progression} \\ \text{of ER+ Breast cancer} \end{array} \\ \begin{array}{c} \text{Data: Tumour size at day 42} \\ (x_{T,1}; x_{T,2}; ...; x_{T,n_T}) \\ (x_{C,1}; x_{C,2}; ...; x_{C,n_T}) \\ \end{array} \\ \downarrow \\ T_{obs} = \frac{\widehat{\mu}_{Tamoxifen} - \widehat{\mu}_{Control}}{\sum_{s_p \sqrt{\frac{1}{n_r} + \frac{1}{n_C}}}} \sim St_{n_T + n_C - 2} \\ \end{array} \\ \begin{array}{c} \text{Point estimation} \\ \widehat{\mu}_{Tamoxifen} - \widehat{\mu}_{Control} \\ \end{array}$$



Grand Picture of Statistics



One-sample Student's t-test

▶ Assumed model

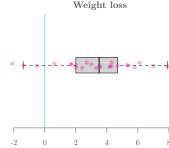
$$Y_i = \mu + \epsilon_i,$$
 where $i = 1, ..., n$ and $\epsilon_i \sim N(0, \sigma^2).$

Hypotheses

▶ **H0**: $\mu = 0$,

ightharpoonup H1: $\mu > 0$.

► Test statistic's distribution under H0

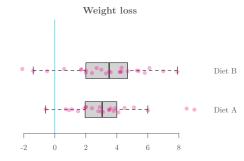


Diet B



One Sample t-test





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

 $\triangleright \mathbf{H0}: \ \mu_A = \mu_B,$

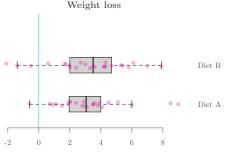
 \triangleright **H1**: $\mu_A \neq \mu_B$.

► Test statistic's distribution under H0

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

Two Sample t-test

```
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
3.300 3.268
```





Two independent sample Welch's t-test

Assumed model

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

Hypotheses

 $\triangleright \mathbf{H0}: \ \mu_A = \mu_B,$

3.268

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

► Test statistic's distribution under H0

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

Weight loss

Welch Two Sample t-test

3.300

```
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 \quad 1.384692 sample estimates: mean of x mean of y
```



Diet B

Two independent sample Mann-Whitney-Wilcoxon test

Assumed model

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

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

Hypotheses

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

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



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

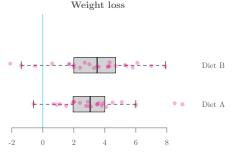
where

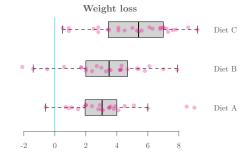
 $ightharpoonup R_{i(g)}$ denotes the global rank of the ith observation of group g.

Wilcoxon rank sum test with continuity correction

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

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

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

 $= \mu + \delta_g + \epsilon_{i(g)},$

where g=1,...,G, $i=1,...,n_g$, $\epsilon_{i(g)}\sim N(0,\sigma^2)$ and $\sum n_g\delta_g=0$.

Hypotheses

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

▶ **H1:** $\mu_k \neq \mu_l$ for at least one pair (k, l). ▶ Test statistic's distribution under **H0**

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

where

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

Weight loss

Diet C

Diet B

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: $\mu_1 = \mu_2 = \dots = \mu_G$, ho **H1**: $\mu_k \neq \mu_l$ for at least one pair (k, t^2) .

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

$$F^{\star} = \frac{s_{\overline{Y}}^{\star^2}}{1 + \frac{2(G-2)}{3\Delta}} \sim Fisher(G-1, \Delta),$$

where

$$\Delta = \left[\frac{3}{G^2 - 1} \sum_{n=1}^{G} \frac{1}{n_g} \left(1 - \frac{w_g}{\sum_{w_g}} \right) \right]^{-1},$$

One-way analysis of means (not assuming equal variances)

Weight loss

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



Diet C

Diet B

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: $\theta_1 = \theta_2 = ... = \theta_G$.

 \triangleright **H1**: $\theta_k \neq \theta_l$ for at least one pair $(k, l)^2$.

Test statistic's distribution under H0

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

Weight loss

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

Kruskal-Wallis rank sum test

data: weight.loss by diet.type Kruskal-Wallis chi-squared = 9.4159, df = 2, p-value = 0.009023



Diet C

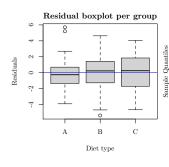
Diet B

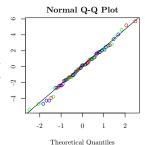
Model check: Residual analysis

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

where

- $lackbox{}\widehat{\epsilon}_{i(g)}\sim N(0,\widehat{\sigma}^2)$ for Fisher's ANOVA
- $m{\epsilon}_{i(g)} \sim N(0,\widehat{\sigma}_g^2)$ 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.

ightharpoonup 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

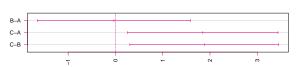
$$\alpha_K = 1 - (1 - \alpha)^K$$
. $\Rightarrow \alpha_1 = 0.05,$
 $\Rightarrow \alpha_2 = 0.0975,$
 $\Rightarrow \alpha_{10} = 0.4013.$

► Multiplicity correction

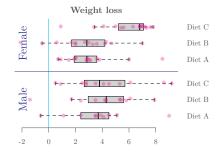
Principle: change the level of each test so that $\alpha_K = 0.05$, for example:

- 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







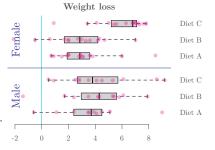
Two or more sample location tests: two-way ANOVA

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

Assumed model

$$\begin{split} Y_{i(g)} &= \mu_{gk} + \epsilon_{i(gk)}, \\ &= \mu + \delta_g + \delta_k + \delta_{gk} + \epsilon_{i(gk)}, \end{split}$$

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



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.

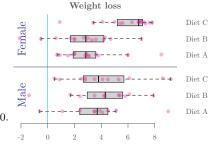
$$ightharpoonup H0_3$$
: $δ_{gk} = 0 ∀ g, k$, $ightharpoonup H1_3$: $H0_3$ is false.

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

Assumed model

$$\begin{split} Y_{i(g)} &= \mu_{gk} + \epsilon_{i(gk)}, \\ &= \mu + \delta_g + \delta_k + \delta_{gk} + \epsilon_{i(gk)}, \end{split}$$

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

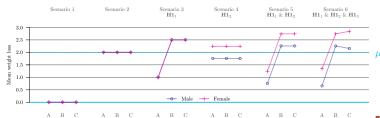


Hypotheses

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

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

 $\begin{array}{l} \rhd \ \mathbf{H0}_3 \text{:} \ \delta_{g\,k} = 0 \ \forall \ g,k \ \text{,} \\ \rhd \ \mathbf{H1}_3 \text{:} \ \mathbf{H0}_3 \ \text{is false}. \end{array}$



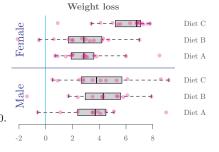


More than one factor: Fisher's two-way ANOVA

Assumed model

$$\begin{split} Y_{i(g)} &= \mu_{gk} + \epsilon_{i(gk)}, \\ &= \mu + \delta_g + \delta_k + \delta_{gk} + \epsilon_{i(gk)}, \end{split}$$

- p = 1, ..., G, k = 1, ..., K,
- $i = 1, ..., n_q$
- $ightharpoonup \epsilon_{i(gk)} \sim N(0, \sigma^2)$



Hypotheses

```
\triangleright H0<sub>1</sub>: \delta_g = 0 \ \forall \ g, \triangleright H1<sub>1</sub>: H0<sub>1</sub> is false.
```

```
\triangleright H0<sub>2</sub>: \delta_k = 0 \ \forall \ k, \triangleright H1<sub>2</sub>: H0<sub>2</sub> is false.
```

 $ightharpoonup H0_3$: $δ_{gk} = 0 ∀ g, k$, $ightharpoonup H1_3$: $H0_3$ is false.

```
Df Sum Sq Mean Sq F value Pr(>F)
diet.type
                      60.5
                            30.264
                                     5.629 0.00541
                      0.2
gender
                           0.169
                                    0.031 0.85991
diet.type:gender
                     33.9
                           16.952
                                    3.153 0.04884 *
Residuals
                70
                    376.3
                           5.376
               0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' ' 1
Signif. codes:
```



Summary

