## 2-sample t-test

 $\bar{y} = \frac{1}{n} \sum_{i=1}^{n} y_i$  Estimates the population mean  $\mu$ .  $S^2 = \frac{1}{n-1} \sum_{i=1}^{n} (y_i - \bar{y})^2$  Estimates the variance  $\sigma^2$ . The Hypothesis test is:  $H_0$ :  $\mu_1 = \mu_2$ ;  $H_a$ :  $\mu_1 \neq \mu_2$ , or < (left-tailed) or > (right-tailed).

Known Variances use the z test;  $Z = (\bar{y}_1 - \bar{y}_2) / \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}$  (two population means), or  $Z = (\bar{y} - \mu_0) / \frac{\sigma}{\sqrt{n}}$  (one).  $100(1 - \alpha\%)$  C.I. for  $\mu_1 - \mu_2$ :  $(\bar{y}_1 - \bar{y}_2) \pm Z_{\alpha/2} \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}$ . C.R. where  $H_a$ :  $\neq \implies Z_{\alpha/2}$ .  $\iff Z_{\alpha} > \implies Z_{\alpha}$ 

Unknown Variances  $(\sigma_1^2 = \sigma_2^2)$   $t = (\bar{y}_1 - \bar{y}_2)/S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}, d.f = v = n_1 + n_2 - 2.$   $S_p^2 = \frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2}$   $(S_p = \sqrt{S_p^2})$   $100(1 - \alpha)\%$  C.I. for  $\mu_1 - \mu_2$ :  $(\bar{y}_1 - \bar{y}_2) \pm t_{\alpha/2,v} S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$  (two means), or  $\bar{y} \pm t_{\alpha,n-1} \frac{\sigma}{\sqrt{n}}$  (one mean).

Unknown Variances  $(\sigma_1^2 \neq \sigma_2^2)$   $t = (\bar{y}_1 - \bar{y}_2) / \sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}, d.f = v = \frac{((S_1^2/n_1) + (S_2^2/n_2))^2}{\frac{(S_1^2/n_1)^2}{n_1 - 1} + \frac{(S_2^2/n_2)^2}{n_2 - 1}}.$  (round down v to the nearest int.) C.R. where  $H_a$ :  $\neq \implies t_{\alpha/2,v}$ .  $H_a$ :  $< \implies -t_{\alpha,v}$ .  $H_a$ :  $> \implies t_{\alpha,v}$ 

Tests on Variances The Hypothesis test is:  $H_0$ :  $\sigma_1^2 = \sigma_2^2$ ;  $H_a$ :  $\sigma_1^2 \neq \sigma_2^2$ , or < (left-tailed) or > (right-tailed). two-tailed( $\neq$ ):  $F = \frac{S_1^2}{S_2^2} > F_{\alpha/2,n_1-1,n_2-1}$ , left-tail(<):  $F = \frac{S_2^2}{S_1^2} > F_{\alpha,n_2-1,n_1-1}$ , right-tail (>):  $F = \frac{S_1^2}{S_2^2} > F_{\alpha,n_1-1,n_2-1}$ .

#### Paired t-test (2 measurements from each exp unit)

Advantage: less variability than a design with 2 separate samples, test is more sensitive, ie more powerful. Hypothesis:  $H_0$ :  $\mu_{\delta} = 0$ .  $H_a$ :  $\mu_{\delta} \neq 0$ . Where  $\mu_{\delta} = \mu_2 - \mu_1$  (before - after). Test statistic:  $t = \frac{\bar{d}}{S_d/\sqrt{n}}$  Where  $\bar{d} = \frac{1}{n} \sum_{j=1}^n d_j$  is the sample mean. And  $S_d^2 = \frac{1}{n-1} \sum_{j=1}^n (d_j - \bar{d})^2$ . (note: use  $S_d = \sqrt{S_d^2}$ ) Reject  $H_0$  if  $|t| > t_{\alpha/2, n-1}$ .

### **ANOVA**

The overall mean  $\mu$  is  $\frac{1}{a}\sum_{i=1}^{a}\mu_{i}$ . Note that  $\sum_{i=1}^{a}\tau_{i}=0$ . LS estimates:  $\hat{\mu}=\bar{y}_{..}, \hat{\mu}_{i}=\bar{y}_{i.}, \hat{\tau}=\bar{y}_{i}.-\bar{y}_{..}$ , to estimate common population variances  $\sigma^{2}$ , use  $S_{p}^{2}=\frac{\sum_{i=1}^{a}(n_{i}-1)s_{i}^{2}}{\sum_{i=1}^{a}(n_{i}-1)}$ 

# Anova table (one-way layout)

Source	df	~~	MS	F	Note: $N$ is the total number of observations,
Treatments	a-1	SSTr	$MSTr = \frac{SSTr}{a-1}$	$F = \frac{MSTr}{MSE}$	Pooled sample variance $S_p^2 = \frac{SSE}{N-g} = MSE$ .
Error	N-a	SSE	$MSE = \frac{\overset{a-1}{SSE}}{N-a}$		Reject $H_0$ if $F > F_{\alpha,a-1,N-a}$ , or p-value $< \alpha$ .
Total	N-1	SST			

Hypotheses:  $H_0$ :  $\mu_1 = \mu_2 = \cdots = \mu_a$ .  $H_a$ : Not all  $\mu_i$  are equal. Or  $H_0$ :  $\tau_1 = \tau_2 = \cdots = \tau_a = 0$ .  $H_a$ : Not all  $\tau_i = 0$ .

$$100(1-\alpha)\%$$
 C.I. for  $\mu_i$ :  $\bar{y}_{i\cdot} \pm t_{\alpha/2,N-a} \sqrt{\frac{MSE}{n}}$ 

To check model assumptions use a normal probability plot and normality test (Anderson-Darling or some other), and a plot of residuals vs. predicted value (residual plot). The residuals should look like they are from a normal distribution if the model assumptions were met. If the model assumptions were met, the residual plot should look like a random scatter. if the plot shows a 'funnel' shape, this is an indication of non-constant variance (heteroscedasticity). The remedy is typically a transformation of the response.

# Normal probability plot

To estimate the standard deviation  $\sigma$ , note that the slope  $\approx \frac{1}{\sigma}$ , only with z-score. Or you can estimate the distance from 16% to 50% as one standard deviation. To estimate the mean  $\mu$  look at the 50<sup>th</sup> percentile. Hypotheses:  $H_0$ : The data are drawn from a normal distribution. Large p-value means normal.

Standard to Percentile:
Standard Deviations
-2 -1 0 1 2 

Percentile
2.27% 15.88% 50% 84.23% 97.72%

Note that  $\Phi$ (standard deviation) = percentile, and  $\Phi^{-1}$ (percentile) = standard deviation.

#### Contrasts

 $\Gamma = \sum_{i=1}^{a} c_i \mu_i$ , where  $c_i$ 's are constants such that  $\sum_{i=1}^{a} c_i = 0$ . To estimate  $\Gamma$  use  $\hat{\Gamma} = \sum_{i=1}^{a} c_i \bar{y}_i$ . Hypotheses:  $H_0$ :  $\sum_{i=1}^{a} c_i \mu_i = 0$ .  $H_a$ :  $\sum_{i=1}^{a} c_i \mu_i \neq 0$ . (or < or >).

Test statistic:  $t = (\sum_{i=1}^{a} c_i \bar{y}_{i\cdot})/(\sqrt{\frac{MSE}{n}} \sum_{i=1}^{a} c_i^2)$ . Reject  $H_0$  if t in C.R. of  $t_{\alpha,N-a}$  (or  $\alpha/2$  for 2-tail) or if p-value  $< \alpha$ . Note:  $F = (\sum_{i=1}^{a} c_i \bar{y}_{i\cdot})^2/(\frac{MSE}{n} \sum_{i=1}^{a} c_i^2)$  can be used to test 2-tailed. Reject  $H_0$  if  $F > F_{\alpha,1,N-a}$  or if p-value  $< \alpha$ .

#### Pairwise comparisons

Note: LSD is most powerful, but worst at guarding against Type I errors. Tukey's is most balanced for equal sample sizes. And Bonferroni's is least likely to make a Type I error.

**Fishers LSD** For every pair of treatments test the hypothesis at  $\alpha$ .  $H_0$ :  $\mu_i = \mu_j$ .  $H_a$ :  $\mu_i \neq \mu_j$ .

Test statistic  $t = |\bar{y}_{i\cdot} - \bar{y}_{j\cdot}| / \sqrt{MSE\left(\frac{1}{n_i} + \frac{1}{n_j}\right)}$ . Reject  $H_0$  is  $|t| > t_{\alpha/2, N-a}$ .

Therefore, The two treatment means are significantly different if  $|\bar{y}_{i\cdot} - \bar{y}_{j\cdot}| > t_{\alpha/2,N-a} \sqrt{MSE\left(\frac{1}{n_i} + \frac{1}{n_j}\right)}$ .

**Bonferroni's Method** Denote the FWE by  $\alpha$ . Identical to Fisher's LSD except that

$$|\bar{y}_{i\cdot} - \bar{y}_{j\cdot}| > t_{\alpha/2k,N-a} \sqrt{MSE\left(\frac{1}{n_i} + \frac{1}{n_j}\right)}$$
. Where  $k = \binom{a}{2}$ .

**Tukey's Method** Means are significantly different at  $FWE\alpha$  when:  $|\bar{y}_{i\cdot} - \bar{y}_{j\cdot}| > q_{\alpha,a,N-a} \sqrt{\frac{MSE}{2} \left(\frac{1}{n_i} + \frac{1}{n_j}\right)}$ . 100(1 –  $\alpha$ )% Simultaneous C.I's for all possible treatment differences  $\mu_i - \mu_j$  using Tukeys Method:

$$\bar{y}_{i.} - \bar{y}_{j.} \pm q_{\alpha,a,N-a} \sqrt{\frac{MSE}{2} \left(\frac{1}{n_i} + \frac{1}{n_j}\right)}$$

Graphical results Note that a line under the treatments indicates that they are not significantly different.

(5)

 $(2) \qquad \qquad (1)$ 

 $(4) \qquad \qquad (3)$ 

Treatments should be ordered from the smallest sample mean to the largest sample mean. This graph indicates that treatment 2 and 1 are not significantly different, and 4 and 3, and 5 are not significantly different.

# Test of equality of variances

Hypothesis:  $H_0$ :  $\sigma_1^2 = \sigma_2^2 = \cdots = \sigma_a^2$ .  $H_a$ : at least one  $\sigma_i^2$  is different.

Bartlett's test (use if data is normal) assumes normality for each treatment. Very sensitive to non-normality.

**Modified Levene's test** (use if data  $\approx$  normal) robust for non-normality. For data from a normal dist, not as powerful as Bartlett's.

## Conclusions

**Rejection Methods** Reject  $H_0$  if p-value  $\leq \alpha$ . Reject  $H_0$  if the test statistic falls within the critical region.

 $100(1-\alpha)\%$  C.I. "We are  $100(1-\alpha)\%$  confident that the true {test} is between {upper bound} and {lower bound}."

**Reject**  $H_0$  "There is enough statistical evidence to support (the statement of  $H_a$ )."

Fail to reject  $H_0$  "There is not enough statistical evidence to support (the statement of  $H_a$ )." or "The evidence from the data is consistent with (the statement of  $H_0$ )"