Master Program in *Data Science and Business Informatics*  **Statistics for Data Science** Lesson 32 - Multiple-sample tests of the mean and applications to classifier comparison

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### The multiple comparisons problem

• Single test  $H_0: \mu = 0$ , with significance level  $\alpha = 0.05$ 

[false positive rate]

- test is called *significant* when we reject  $H_0$
- $\alpha$  is Type I error, probability of rejecting  $H_0$  when it is true
- Multiple tests, say m = 20

▶ E.g.,  $H_0^i : \mu_i = 0$  for i = 1, ..., m where  $\mu_i$  is the expectation of a subpopulation

- What is the probability of rejecting at least one  $H_0^i$  when all of them are true?
  - ► For independent tests:  $P(\cup_{i=1}^{m} \{p_i \leq \alpha\}) = 1 P(\cap_{i=1}^{m} \{p_i > \alpha\}) = 1 (1 \alpha)^m$ and then  $1 - (0.95)^{20} \approx 0.64$
  - For dependent tests:  $P(\bigcup_{i=1}^{m} \{p_i \leq \alpha\}) \leq \sum_i P(\{p_i \leq \alpha\}) = m \cdot \alpha$ , and then  $\leq 20 \cdot 0.05 = 1$

#### Family-wise error rate (FWER)

The FWER is the probability of making at least one Type I error in a family of m tests. If the tests are independent:

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\alpha_{FWER} = 1 - (1 - \alpha)^m
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If the test are dependent:  $\alpha_{FWER} \leq m \cdot \alpha$ 

### Multiple comparisons: corrections

Question: what should be  $\alpha$  such that  $\alpha_{FWER} \leq b$ ?

- Bonferroni correction (most conservative one):
  - scale significance level  $\alpha = b/m$  [invert  $b = m \cdot \alpha$ ]
  - thus  $\alpha_{FWER} \leq m \cdot \alpha = b$

Notice:  $p \leq \alpha$  is equivalent to scale p-values and test  $p \cdot m \leq b$ 

- *Šidák correction* (exact for independent tests):
  - ▶ scale significance level  $\alpha = 1 (1 b)^{1/m}$  [invert  $b = 1 (1 \alpha)^m$ ]
  - thus  $\alpha_{FWER} = 1 (1 \alpha')^m = b$

Notice:  $p \leq \alpha$  is equivalent to scale p-values and test  $1 - (1 - p)^m \leq b$ 

### False Discovery Rate and *q*-values

		True state of nature	
		$H_0$ is true	$H_1$ is true
Our decision on the basis of the data	Reject $H_0$	False Positive	True Positive
	Not reject $H_0$	True Negative	False Negative

See R script

- False Positive Rate: FPR = FP/(FP + TN)
  - Corrections control for FPR since  $FWER = P(FP > 0|H_0^i \ i = 1, ..., m)$
- Drawback: acting on  $\alpha$  increases FNR = FN/(FN + TP)
- False Discovery Rate: FDR = FP/(FP + TP)
  - FDR = 0.05 means 5% of rejected  $H_0$ 's are actually true
- *q*-value is  $P(H_0|T \ge t)$ 
  - *FDR* can be controlled by requiring  $q \leq$  threshold

[Korthauer et al, 2019]

[vs.  $p = P(T \ge t | H_0)$ ]

### Omnibus tests and post-hoc tests

- $H_0: \theta_1 = \theta_2 = \ldots = \theta_k \ [= 0]$
- $H_1: \theta_i \neq \theta_j$  for some  $i \neq j$
- Omnibus tests detect any of several possible differences
  - Advantage: no need to pre-specify which treatments are to be compared ... ... and then no need to adjust for making multiple comparisons
- If  $H_0$  is rejected (test significant), a *post-hoc test* to find which  $\theta_i \neq \theta_j$ 
  - Everything to everything post-hoc compare all pairs
  - One to everything post-hoc compare a new population to all the others
- We distinguish a few cases:
  - Multiple linear regression (normal errors + homogeneity of variances, i.e., U<sub>i</sub> ~ N(0, σ<sup>2</sup>)):
    Γ-test + t-test
  - Equality of means (normal distributions + homogeneity of variances):
    - $\ \ \square \ \ ANOVA + Tukey/Dunnett$
  - Equality of means (general distributions):
    - $\Box$  Friedman + Nemenyi

## F-test for multiple linear regression

• 
$$\boldsymbol{Y} = \boldsymbol{X} \cdot \boldsymbol{\beta} + \boldsymbol{U}$$
, where  $\boldsymbol{Y} = (Y_1, \dots, Y_n)$ ,  $\boldsymbol{U} = (U_1, \dots, U_n)$ , and  $\boldsymbol{X} = (\boldsymbol{x}_1, \dots, \boldsymbol{x}_n)$ 

- $\boldsymbol{\beta}^{T} = (\alpha, \beta_1, \dots, \beta_k) \text{ and } \boldsymbol{x}_i = (1, x_i^1, \dots, x_i^k)$
- Unexplained (residual) error  $SSE = S(\beta) = \sum_{i=1}^{n} (y_i \mathbf{x}_i \cdot \beta)^2$
- Null model (or intercept-only model):  $\textbf{Y} = \textbf{1} \cdot \alpha + \textbf{U}$ 
  - ► Total error  $SST = S(\alpha) = \sum_{i=1}^{n} (y_i \bar{y}_n)^2$  [residuals of the null model]
- Explained error  $SSR = SST SSE = \sum_{i=1}^{n} (\bar{y}_n \boldsymbol{x}_i \cdot \boldsymbol{\beta})^2$
- Coefficient of determination  $R^2 = SSR/SST = 1 SSE/SST$  [See Lesson 20]
  - Is the model useful? Fraction of explained error
- Is the model statistically significant? [vs a specific  $\beta_i$  significant? See Lesson 29]
- $H_0: \beta_1 = \ldots = \beta_k = 0$   $H_1: \beta_i \neq 0$  for some  $i = 1, \ldots, k$
- Test statistic:  $F = \frac{SSR}{SSE} \frac{n-k-1}{k} \sim F(k, n-k-1)$

See R script

# Equality of means: ANOVA

- $H_0: \mu_1 = \mu_2 = \ldots = \mu_k$
- $H_1: \mu_1 \neq \mu_2$  for some  $i \neq j$
- datasets  $y_1^j, \ldots, y_{n_j}^j$  for  $j = 1, \ldots, k$ 
  - Assumption: normality (Shapiro-Wilk test) + homogeneity of variances (Bartlett test)
  - ▶ responses of k-1 treatments and 1 control group
  - accuracies of k classifiers over  $n_j = n$  datasets [repeated measures/two way ANOVA]
- Linear regression model over dummy encoded *j*:

$$Y = \alpha + \beta_1 x_1 + \ldots + \beta_{k-1} x_{k-1}$$

- $\alpha = \mu_k$  is the mean of the reference group (j = k)
- $\blacktriangleright \ \beta_j = \mu_j \mu_k$
- ▶ in R:  $lm(Y \sim Group)$  where Group contains the labels of j = 1, ..., k
- F-test (over linear regression):  $H_0: \beta_1 = \ldots = \beta_k = 0$ , i.e.,  $\mu_j = \mu_k$  for  $j = 1, \ldots, k$
- Tukey HSD (Honest Significant Differences) is an all-pairs post-hoc test
- Dunnet test is a one-to-everything test

### See R script

[generalization of two sample t-test]

[one wav ANOVA]

### Non-parametric test of equality of means: Friedman

• 
$$H_0: \mu_1 = \mu_2 = \ldots = \mu_k$$

- $H_1: \mu_1 \neq \mu_2$  for some  $i \neq j$
- datasets  $x_1^j, \ldots, x_n^j$  for  $j = 1, \ldots, k$ 
  - ► accuracies of k classifiers over n datasets
- Let r<sub>i</sub><sup>j</sup> be the rank of x<sub>i</sub><sup>j</sup> in x<sub>i</sub><sup>1</sup>,...,x<sub>i</sub><sup>k</sup>
  e.g., j<sup>th</sup> classifier w.r.t. i<sup>th</sup> dataset
- Average rank of classifier:  $R_i = \frac{1}{n} \sum_{i=1}^{n} r_i^j$
- Under  $H_0$ , we have  $R_1 = \ldots = R_k$  and, for *n* and *k* large:

$$\chi_F^2 = \frac{12n}{k(k+1)} \left( \sum_{j=1}^k R_j^2 - \frac{k(k+1)^2}{4} \right) \sim \chi^2(k)$$

- Nemenyi test is an all-pairs post-hoc test
- Bonferroni correction is a one-to-everything test
- For unpaired observations, use Kruskal-Wallis test instead of Friedman test

#### See R script

[paired observations/repeated measures]

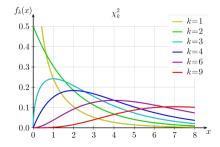
### Chi-square distribution

### Chi-square distribution

The Chi-square distribution with k degrees of freedom  $\chi^2(k)$  has density:

$$f(x) = \frac{1}{2^{k/2} \Gamma(k/2)} x^{k/2-1} e^{-x/2}$$

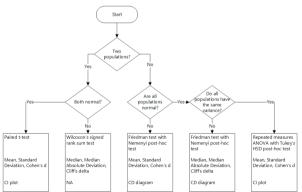
Let 
$$X_1, \ldots, X_k \sim \mathcal{N}(0, 1)$$
. Then  $Y = \sum_{i=1}^k X_i^2 \sim \chi^2(k)$ 



# Comparing classifiers: Summary



The SCMAMP package in R

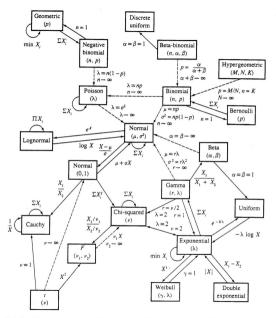


#### The AutoRank package in Python

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### Common distributions

- Probability distributions at Wikipedia
- Probability distributions in R
- C. Forbes, M. Evans, N. Hastings, B. Peacock (2010) Statistical Distributions, 4th Edition Wiley



Relationships among common distributions. Solid lines represent transformations and special cases, dashed lines represent limits. Adapted from Leemis (1986). 11 /

- On confidence intervals and statistical tests (with R code)
- Myles Hollander, Douglas A. Wolfe, and Eric Chicken (2014) Nonparametric Statistical Methods. 3rd edition, John Wiley & Sons, Inc.
  - On False Discovery Rate
- Keegan Korthauer, Patrick K. Kimes, Claire Duvallet, Alejandro Reyes, Ayshwarya Subramanian, Mingxiang Teng, Chinmay Shukla, Eric J. Alm, and Stephanie C. Hicks (2019)
   A practical guide to methods controlling false discoveries in computational biology. Genome Biology 20, article 118