



Method validation & correlation

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With material from

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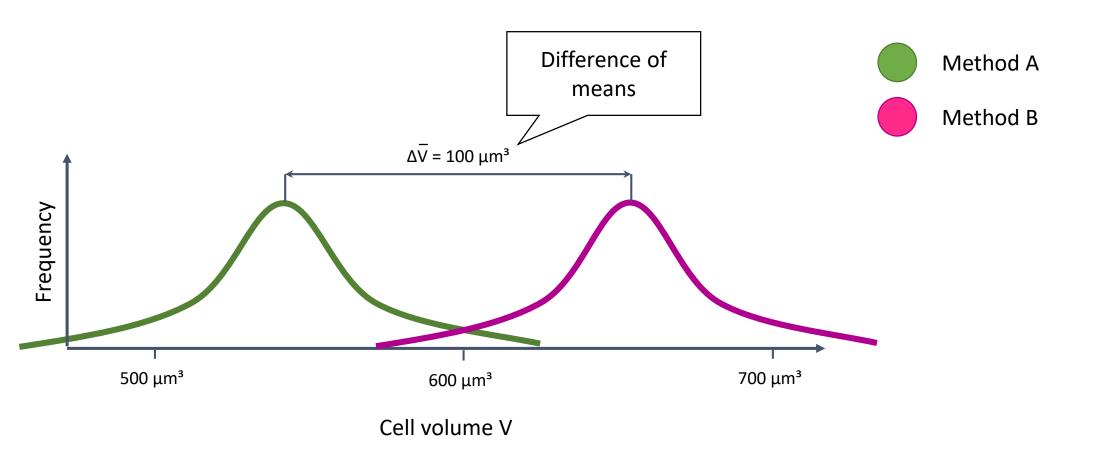
Douglas G. Altman



Comparison of means



• Comparing mean measurements appears reasonable on the first view.

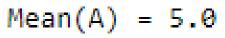


Comparison of means



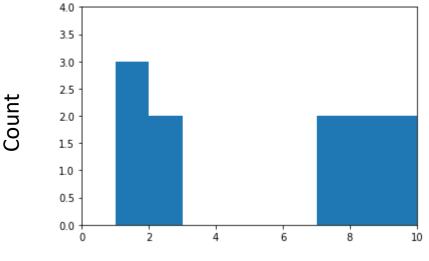
Are two methods doing the same if their mean measurement is similar?

- А В
- 1 4
- 9 5
- 7 5
- 1 7
- 2 4
- 8 5
- 9 4
- _ .
- 1 6
- 7 5
- 8 4



$$Mean(B) = 5.0$$

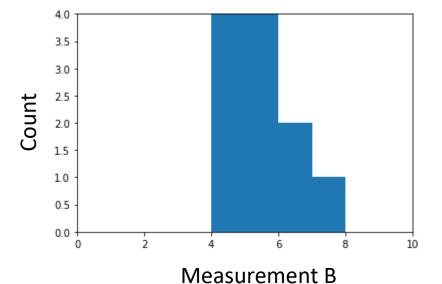
• Draw histograms! How can two methods do the same if histograms from their measurements are different?





Similar means is a necessary condition, but it is NOT sufficient!

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Comparison of means



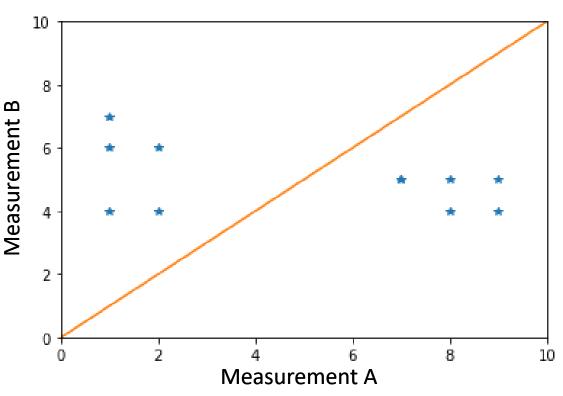
Are two methods doing the same if their mean measurement is similar?

$$Mean(A) = 5.0$$

$$Mean(B) = 5.0$$

Plot the measurements against each other. What does it mean if they lie on a straight line? What if not?

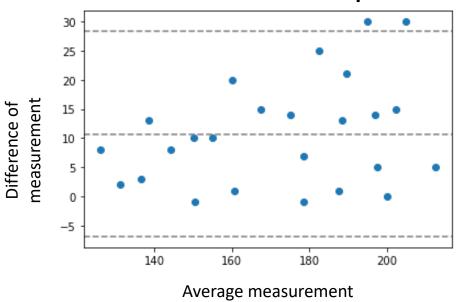
- 7 5
- 1 7
- 2 4
- 8 5
- 9 4
- 2 6
- 1 6
- 7 5
- 8 4



The "criterion of agreement was that the two methods gave the same mean measurement; 'the same' appears to stand for 'not significantly different'. Clearly, this approach tells us very little about the accuracy of the methods." 1

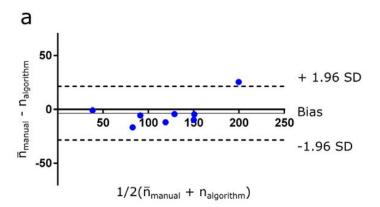


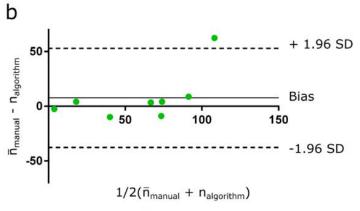
Bland-Altman plot

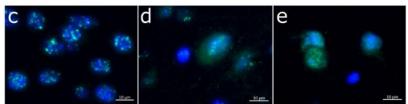


Practical application:

- Script counts cell damage (gH2AX expression) in nucleus
- Damage and nuclei are counted independently and require settings some parameters
- Bland-Altman analysis used to validate parameters





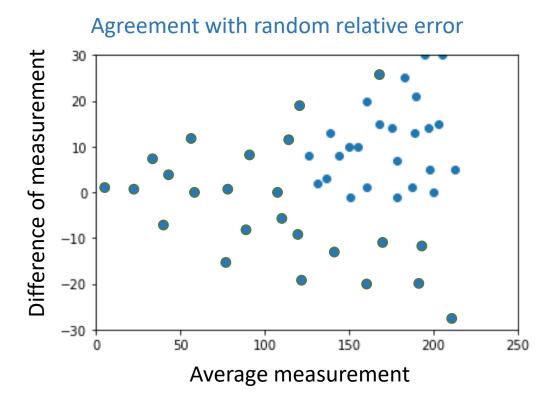


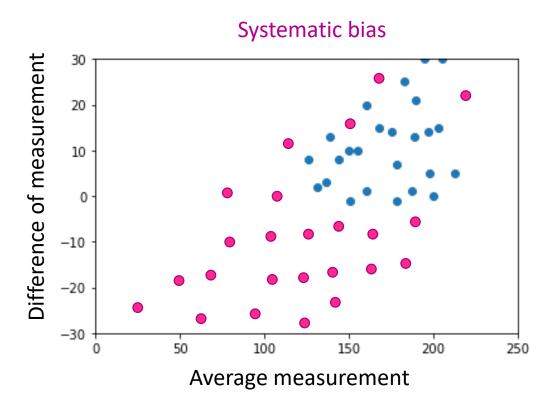
Suckert, Beyreuther, Müller et al. Rad. Onc. (2020)

In practice:



Bland-Altman plots allow us to differentiate various kinds of bias.



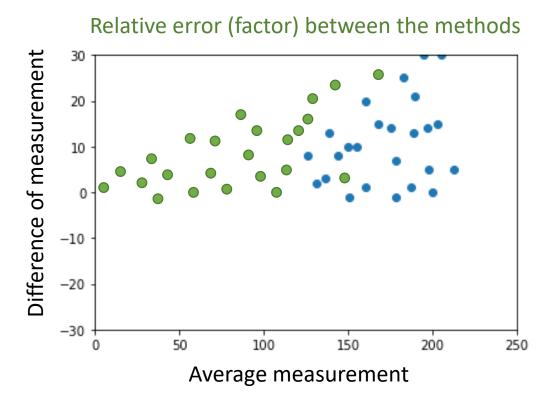


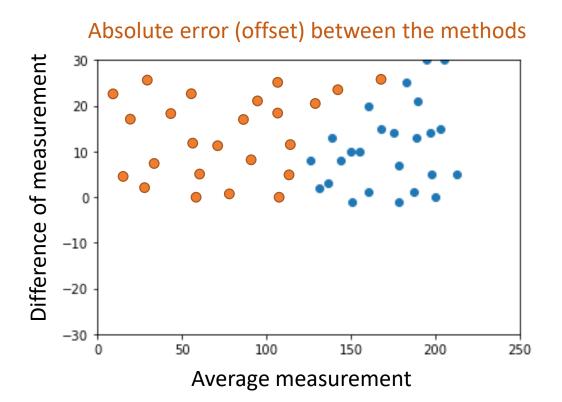
• Both distributions could have the same mean difference and confidence interval.

In practice:



Bland-Altman plots allow us to differentiate various kinds of bias.





• Both effects can be corrected by calibration.

The confidence interval





• "The British Standards Institution (1979) define a coefficient of repeatability as 'the value below which the difference between two single test results ... may be expected to lie with a specified probability; in the absence of other indications, the probability is 95 per cent'."1

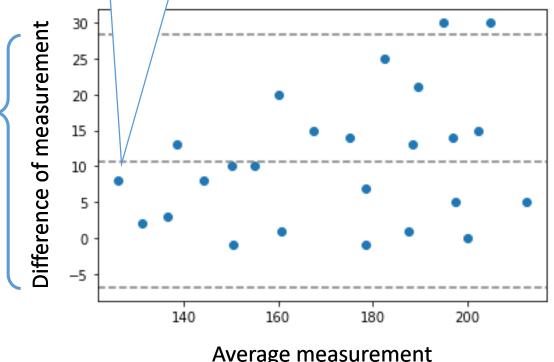
> The mean difference between the methods

Bland-Altman plot

The confidence interval *CI* of agreement

$$CI = (\mu - 2\sigma, \mu + 2\sigma)$$

Mean difference Standard deviation of differences



Average measurement

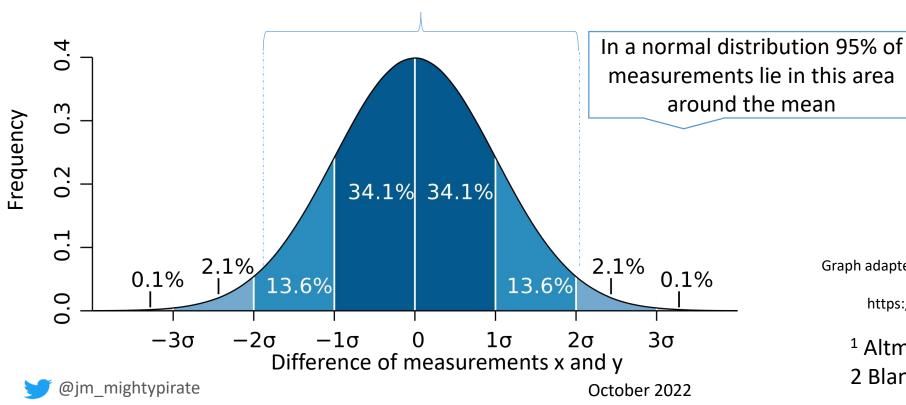
[@]jm_mightypirate

The confidence interval & the coefficient of repeatability.





- "The British Standards Institution (1979) define a coefficient of repeatability as 'the value below which the difference between two single test results ... may be expected to lie with a specified probability; in the absence of other indications, the probability is 95 per cent'."
- If the two measurements come from the same method which just repeated twice, we can assume that the mêan difference is zero. The coefficient of repeatability *CR* can then be estimated: It's the standard deviation of differences.²



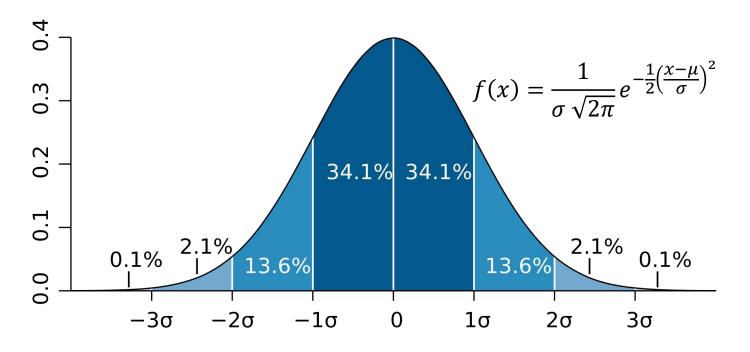
 $CR(X,Y) = \sqrt{\sum_{x \in X, y \in Y} \frac{(x-y)^2}{n}}$

Graph adapted from: M. W. Toews - Own work, based (in concept) on figure by Jeremy Kemp, on 2005-02-09, CC BY 2.5, https://commons.wikimedia.org/w/index.php?curid=1903871

¹ Altman & Bland, The Statistician 32, 1983 2 Bland & Altman, Lancet, 1986

Parametric vs. non-parametric





Normal distribution:

- Can be completely described by mean μ and standard deviation σ
- Allows comparing distributions (e.g., with two-sided/paired t-test)

Ranked distribution:

- Replace each value with its "rank"
- Rank = index of value in sorted list
- Robust to outliers
- Independent on underlying distribution

Value	Rank
10	1
15	2
3	0
97	3

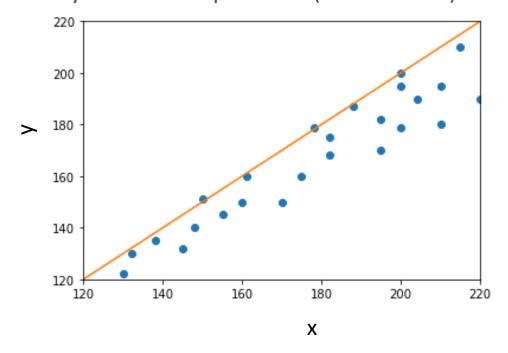
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Pearson Correlation



- Are two methods doing the same if they correlate?
 - Correlation: Any kind of relationship.
 - Measurable; e.g. using Pearson's Correlation Coefficient r enumerated linear correation.

Comparison of two methods of measuring systolic blood pressure (Data from 1)



Expectation E

Mean average µ

$$r(X,Y) = \frac{E[(X - \mu_X)(Y - \mu_y)]}{\sigma_X \sigma_Y}$$

Disclosure: Mean and standard deviation must be obtained from the whole population or from a sample set which is sufficiently large.

Standard deviation o → Unit independence

In practice *E* is the weighted sum:

$$r(X,Y) = \frac{\sum_{x \in X, y \in Y} \frac{(x - \mu_X)(y - \mu_Y)}{n}}{\sigma_X \sigma_Y}$$

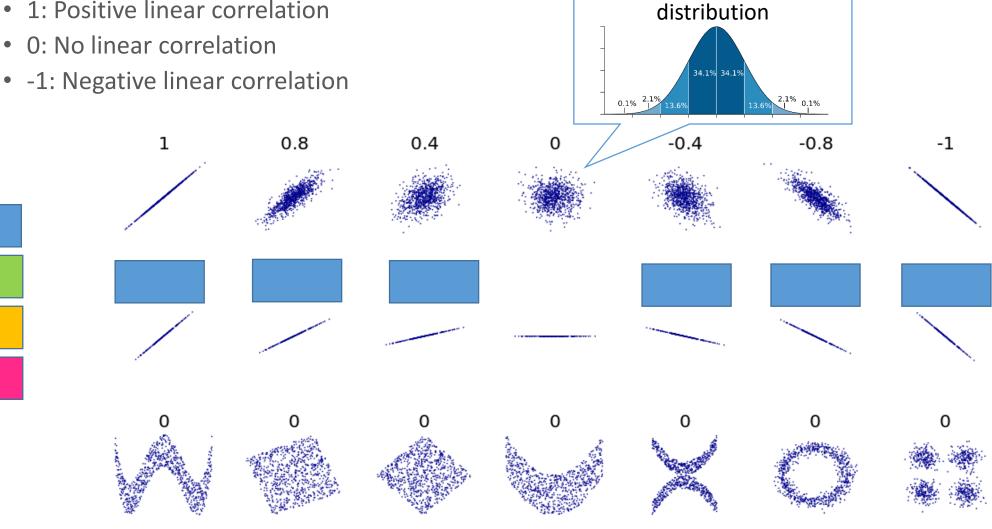
Number of measurements *n*



¹ Altman & Bland, The Statistician 32, 1983

- Pearson's r lies between -1 and 1
 - 1: Positive linear correlation

 - -1: Negative linear correlation



2-dimensional normal

0.4

Correlation: Spearman's r



Value x	Rank x'
10	1
15	2
3	0
97	3
	•••

- Spearman's *r* lies between -1 and 1
 - 1: Positive monotonous correlation
 - 0: No monotonous correlation
 - -1: Negative monotonous correlation

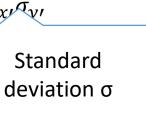
Expectation E

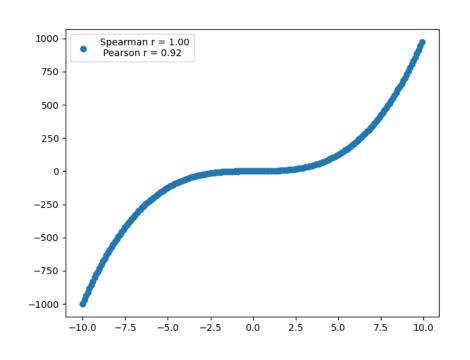
Mean average μ

$$r_{Spearman}(X,Y) = \frac{E[(X' - \mu_{x'})(Y' - \mu_{y'})]}{\sigma_{x'}\sigma_{y'}}$$

Spearman's *r* is equivalent to using Pearson's r on ranked data:

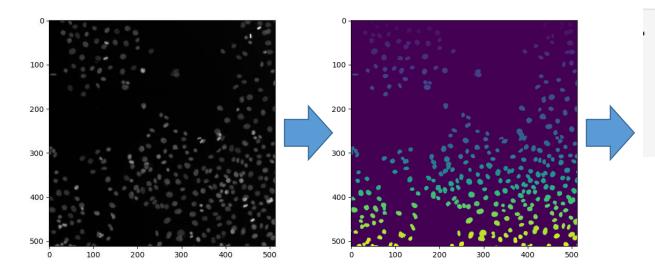
- μ_x : Mean of Samples in X
- μ_x : Mean of ranks of samples in X





Applications:

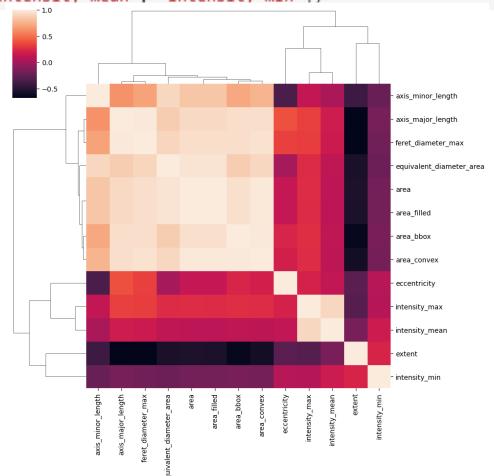






Feature selection: Measuring many features usually brings along some redundancies

- → Use the correlation coefficient to remove or group such features
 - → Create meta-feature (linear combination, mean, etc.) from correlating features (scaling!)
 - → Pick one
- → Downstream analysis works better with fewer, relevant features







Hypothesis testing

Johannes Müller

With material from Anna Poetsch

We need to talk



result = stats.pearsonr(x,y) result PearsonRResult(statistic=-0.886888157935661, pvalue=2.595689084498263e-14)

P-values: Probability that the **null hypothesis** H₀ is true, but rejected by chance

General: It is (usually) much easier to falsify a statement than proving it true

Example 1: $x^n + y^n = z^n$ for $n \ge 3$ and $x, y, z \in Z$

→ this took 358 years to prove – If we could have found just a single combination of x,y & z, we would immediately be done

Example 2: Albert Hammond (1972): It never rains in southern california

→ Very hard to prove – very easy to disprove

 H_0 hypothesis: A treatment agent is ineffective/There is no difference between two groups/Cell fate is not correlated to feature_x

https://en.wikipedia.org/wiki/Fermat%27s Last Theorem

In the context of correlation





https://docs.scipy.org/doc/scipy/reference/generated/scipy.stats.

```
result = stats.pearsonr(x,y)
result
```

PearsonRResult(statistic=-0.8868881579356613, pvalue=2.595689084498263e-14)

H₀ **hypothesis:** correlation coefficient r=0

In scipy: alternative : {'two-sided', 'less', 'greater'}, optional

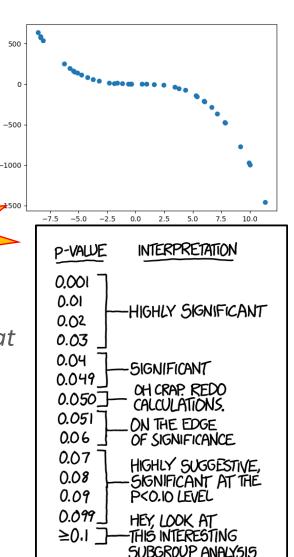
Defines the alternative hypothesis. Default is 'two-sided'. The following options are available:

- 'two-sided': the correlation is nonzero H_0 ?
- 'less': the correlation is negative (less than zero) H_0 ?
- 'greater': the correlation is positive (greater than zero) H₀?

P-value: Probability that correlation coefficient r≠0 although r=0 "We just happened to draw an unfortunate selection of points from our data that looked like correlation – the odds of this happening was p"

How small should the p-value be to confidently reject H_0 ? \rightarrow alpha-value

- → Don't set a threshold just report
- \rightarrow Some pleasant number (0.05, 0.001, etc.
- \rightarrow A common value in the field (0.05, 5 σ , etc.)

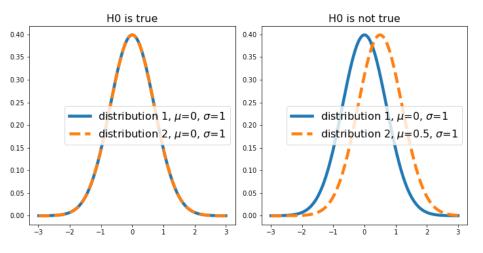


False

positive

Other tests

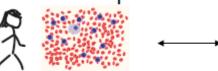




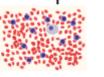
Comparing two (normal) distributions

- → Unpaired t-test (H0: The means are different)
- \rightarrow Paired t-test (H0: $X_{after} X_{before} = 0$)

Before sport



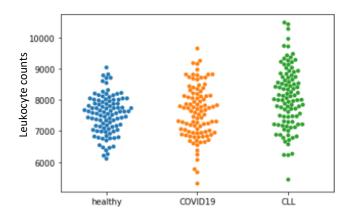
After sport



Alternative: Wilcoxon-Mann-Whitney-Test if assumptions are violated

Many observations don't follow normal distributions

- → (Cell) count data: Poisson distribution
- → Binary outcomes (e.g., coin flip): Binomial distribution
- → Each provides appropriate tests



Comparing multiple groups:

- → ANOVA (analysis of variances), H0: No differences between distributions
- → Requires "post-hoc" tests to find out which groups are different

Data skewed by outliers:

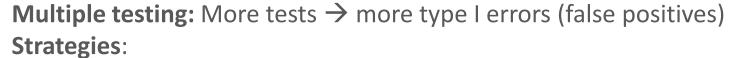
→ Consider comparing ranks rather than raw data

Pitfalls

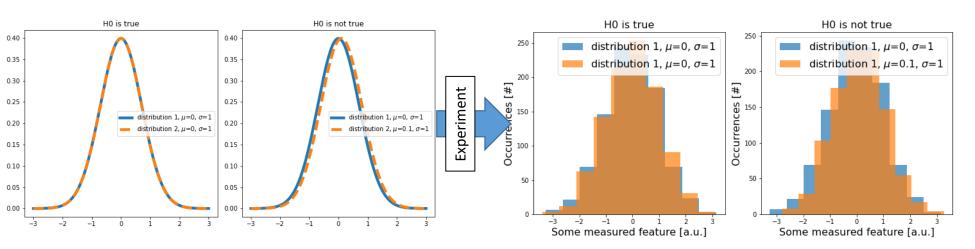




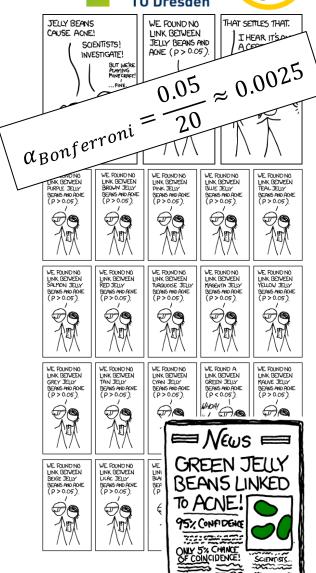




- 1. Control family-wise error rate FWER = $P(n_{false\ positives} \ge 1) = 1 (1 \alpha)^N$ Bonferroni correction: $\alpha_{adj} = \frac{\alpha}{N}$
 - → Prevents false positives (type I error)
 - → Introduces false negatives (type II error)
- 2. Benjamini-Hochberg adjustment: Control false discovery rate $FDR = \frac{FP}{FP+TP}$
 - \rightarrow Find largest k so that $p_k \leq \frac{k}{m} \alpha$ (p_k: p-value of rank k)
- 3. Tukey range test: Typically done after ANOVA, controls type I errors

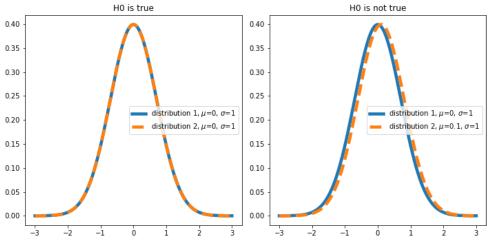


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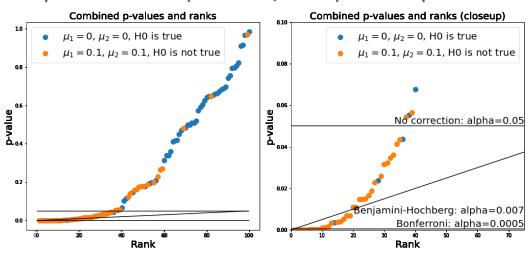


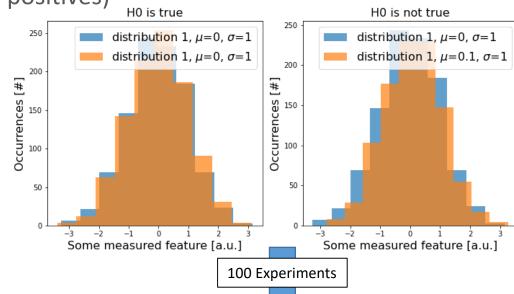


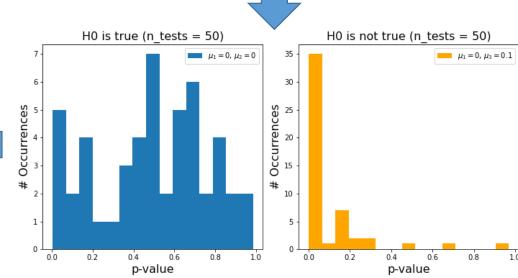


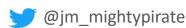
Multiple testing correction:

Separate cases (H0 true/false) in this plot:









Rank p-values

Experiment

Pitfalls



Multiple testing: More tests \rightarrow more type I errors (false positives)

Distribution type: T-test assumes normal distribution of data

- → Some data may follow different distributions (Poisson, binomial, etc.)
- → The equivalent for a t-test exists for all other distributions, too!
- → Less strict test types exist ask your statistician!

Sample size: Do not perform statistical test with small (n < 10) sample sizes.

→ If you work in this region (experiments expensive, animals, etc): Consult your local statistician!

Sample independence:

→ T-tests are only valid if samples are independent: "Two events are independent [...] if [...] the occurrence of one does not affect the probability of occurrence of the other"

Examples:

Histological slices from same animal: Not independent Same blood test derived from two patients: Independent





A small p-value indicates....

A big difference between datasets

Small probability of false positives

Small standard deviations of the compared groups



