

Inference

how surprising is your statistic? (thresholding)





Outline

- Null-hypothesis and Null-distribution
- Multiple comparisons and Family-wise error
- Different ways of being surprised
 - Voxel-wise inference (Maximum z)
 - Cluster-wise inference (Maximum size)
- Parametric vs non-parametric tests
- Enhanced clusters
- FDR False Discovery Rate



Outline

- Null-hypothesis and Null-distribution
- Multiple comparisons and Family-wise error
- Different ways of being surprised
 - Voxel-wise inference (Maximum z)
 - Cluster-wise inference (Maximum size)
- Parametric vs non-parametric tests
- Enhanced clusters
- FDR False Discovery Rate



The task of classical inference

• Given some data we want to know if (e.g.) a mean is different from zero or if two means are different





I.A null-hypothesis

Typically the opposite of what we actually "hope", e.g.

There is **no** effect of treatment: $\mu = 0$

There is **no** difference between groups: $\mu_1 = \mu_2$







- I.A null-hypothesis
- 2. A test-statistic

Assesses "trustworthiness"





I.A null-hypothesis

t =

2. A test-statistic

Assesses "trustworthiness"

 $\frac{x_1 - \overline{x_2}}{\overline{x_1}}$

A *t*-statistic reflects precisely this

Large difference: Trustworthy

Many measurements: Trustworthy Small variability: Trustworthy



- I.A null-hypothesis
- 2. A test-statistic

Or expressed in GLM lingo





- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution

Let us assume there is no difference, i.e. the null-hypothesis is true.





- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution

We might then get these data





- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution

or we could have gotten these





- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution



maybe these



-5

Tools of classical inference

I.A null-hypothesis 2. A test-statistic or perhaps these 3. A null-distribution $\begin{array}{|c|c|c|} & \beta_1 \\ & \beta_2 \end{array} + \mathbf{e} \end{array}$ t = 2.19 σ^2

5

0

= 1.22

onstant



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution



etc



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution





And if we do this til the cows come home



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution



So, why is this helpful?



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution



Well, it for example tells us that in ~1% of the cases t > 3.00, even when the null-hypothesis is true.



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution



Or that in ~5% of the cases t > 1.99. When the nullhypothesis is true.



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution



And best of all:This distribution is known *i.e.* one can calculate it. Much as one can calculate sine or cosine



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution



And best of all: This distribution is known *i.e.* one can calculate it. Much as one can calculate sine or cosine

Provided that $\mathbf{e} \sim N(0,\sigma^2)$

I.A null-hypothesis

$$H_0: \overline{x}_1 = \overline{x}_2$$
, $H_1: \overline{x}_1 > \overline{x}_2$

- 2. A test-statistic
- 3. A null-distribution

So, with these tools let us do an experiment

 $H_0: \overline{x}_1 = \overline{x}_2$, $H_1: \overline{x}_1 > \overline{x}_2$

 $t_8 = 2.64$

- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution

So, with these tools let us do an experiment



- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution

$$H_0: \overline{x}_1 = \overline{x}_2$$
, $H_1: \overline{x}_1 > \overline{x}_2$
 $t_8 = 2.64$

So, with these tools let us do an experiment



If the null-hypothesis is true, we would expect to have a ~1.46% chance of finding a t-value this large or larger

- I.A null-hypothesis
- 2. A test-statistic
- 3. A null-distribution

$$H_0: \overline{x}_1 = \overline{x}_2 , H_1: \overline{x}_1 > \overline{x}_2$$
$$t_8 = 2.64$$
$$t_8 = 2.64*$$

So, with these tools let us do an experiment



There is ~1.46% risk that we reject the nullhypothesis (i.e. claim we found something) when the null is actually true. We can live with that (well, I can).

FSIL

False positives/negatives

- I am sure you have all heard about "false positives" and "false negatives".
- But what does that actually mean?

- I am sure you have all heard about "false positives" and "false negatives".
- But what does that actually mean?
- We want to perform an experiment and as part of that we define a null-hypothesis, e.g. $H_0: \mu = 0$
- Now what can happen?

- I am sure you have all heard about "false positives" and "false negatives".
- But what does that actually mean?
- We want to perform an experiment and as part of that we define a null-hypothesis, e.g. $H_0: \mu = 0$
- Now what can happen?

 H_0 is true B True state of affairs H_0 is false B

- I am sure you have all heard about "false positives" and "false negatives".
- But what does that actually mean?
- We want to perform an experiment and as part of that we define a null-hypothesis, e.g. $H_0: \mu = 0$
- Now what can happen?

 H_0 is true B True state of affairs H_0 is false B

We don't reject H_0 We reject H_0 Our decision



 H_0 is true B True state of affairs H_0 is false B

We don't reject H_0 Gur Gur





 H_0 is true B True state of affairs H_0 is false B

We don't reject H_0 Gur Gur





 H_0 is true H_0 is false H_0 is false

We don't reject H_0 We reject H_0 Our decision





 H_0 is true H_0 is false H_0 is false

We don't reject H_0 Gur Gur





Outline

- Null-hypothesis and Null-distribution
- Multiple comparisons and Family-wise error
- Different ways of being surprised
 - Voxel-wise inference (Maximum z)
 - Cluster-wise inference (Maximum size)
- Parametric vs non-parametric tests
- Enhanced clusters
- FDR False Discovery Rate



Multiple Comparisons

 In neuroimaging we typically perform <u>many</u> tests as part of a study





What happens when we apply this to imaging data?



z-map where each voxel ~N. Null-hypothesis true everywhere, i.e. NO ACTIVATIONS



Ζ

z-map thresholded at I.64



16 clusters288 voxels~5.5% of the voxels

That's a LOT of false positives



Italians doing maths: The Bonferroni correction

Bonferroni says threshold at α divided by # of tests



5255 voxels $0.05/5255 \approx 10^{-5}$



z-map thresholded at 5.65

No false positives. Hurrah for Italy!



But ... doesn't 5.65 sound very high?



So what do we want then?



Family-wise error

Let's say we perform a series of identical studies



Each z-map is the end result of a study

Let us further say that the null-hypothesis is true We want to threshold the data so that only once in 20 studies do we find a voxel above this threshold



But how do we find such a threshold?