

The problem of reproducibility for Imaging genetics

Jean-Baptiste Poline

Helen Wills Neuroscience Institute, UC Berkeley

Reproducibility - preliminary remarks

- Reminding ourselves : Reproducibility is the backbone of scientific activity
- Reproducibility versus replicability
- Is there a problem ?
- Not everybody is convinced that there is a problem
- Do we have hard evidence ?

Reproducibility - evidence of the problem

- In epidemiology
 - Ioannidis 2005
- In social sciences and in psychology
 - Reproducibility Project: Psychology (<https://osf.io/ezcuj/wiki/home/>)
 - Simmons, J. P., L. D. Nelson, and U. Simonsohn. “False-Positive Psychology: Undisclosed Flexibility in Data Collection and Analysis Allows Presenting Anything as Significant.” 2011. Shows an easy 60% FPR instead of 5%.
- In cognitive neuroscience
 - Barch, Deanna M., and Tal Yarkoni. “... Special Issue on Reliability and Replication in Cognitive and Affective Neuroscience Research.” 2013.

Reproducibility - evidence of the problem

- In brain imaging
 - Functional and Structural Neuroimaging: Reproducibility Issues in Multicentre MRI Studies, Jorge Jovicich, Univ. of Trento
- In genetics
 - Cf Jason talk (many old references and warning here, eg: “Drinking from the fire hose ...” by Hunter and Kraft 2007)
- And not in imaging genetics ?

Why do we have a problem?

- Things are getting complex
- Publication pressure is high
- Mistakes are done
- Power issues

Why do we have a problem?

Things are getting complex

- Data complexity (eg: chip idiosyncrasis, format, preprocessings, etc)
- Data need to be linked appropriately
- Data size: number of variables - files you cannot check visually
- Methods: we have to trust external software
- Methods: complexity higher

Why do we have a problem?

Publication pressure is high

- There's no way there isn't a paper out of this data set.
- You will not get your Phd if you don't publish this study
- You won't get tenure
- You won't get funding and your peers admiration and consideration
- Conclusion: the pressure is *very* high

Why do we have a problem?

Mistakes are done

The “Mistakes” argument : an unpopular topic.

- Ioannidis 2005
- Anatomy of an Error
- The Left/Right issue
- The ADHD 1000 connectome

The power issue

- Studies of low power have low probability of detecting an effect (duh!)
- Studies of low power have low positive predictive value :
 $PPV = P(H1 \text{ True} | \text{detection})$
 - If we have 4 chances over 5 that H_0 is true (odd ratio = 1/4), and 1/5 that H_1 true, with 30% power we have $PPV = 50\%$.

odd ratio=0.25	power=0.10,	alpha=0.05	PPV=0.33
odd ratio=0.25	power=0.30,	alpha=0.05	PPV=0.60
odd ratio=0.25	power=0.50,	alpha=0.05	PPV=0.71
odd ratio=0.25	power=0.70,	alpha=0.05	PPV=0.78
odd ratio=0.25	power=0.90,	alpha=0.05	PPV=0.82

The power issue

What is the estimated power in common meta analyses?

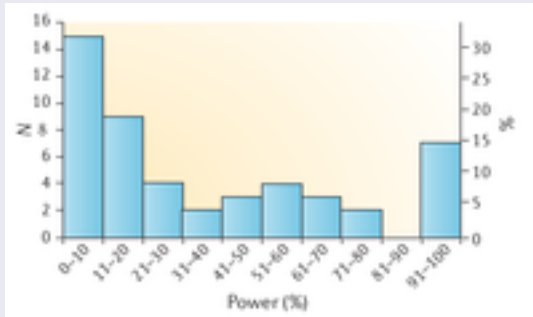


Figure: Button et al. NRN, 2013

The power issue

Studies of low power inflate the detected effect (1)

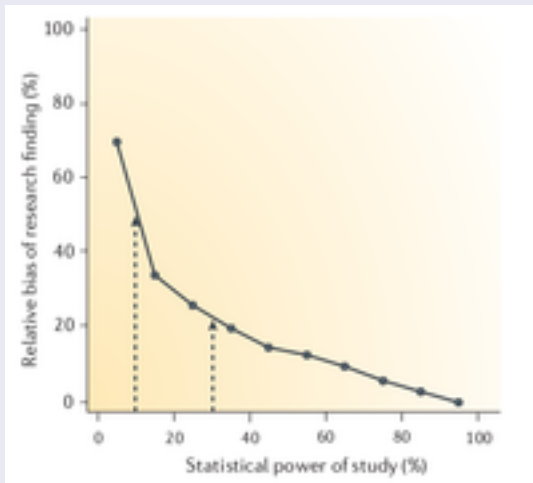


Figure: Button et al. NRN, 2013

The power issue

Studies of low power inflate the detected effect (2)

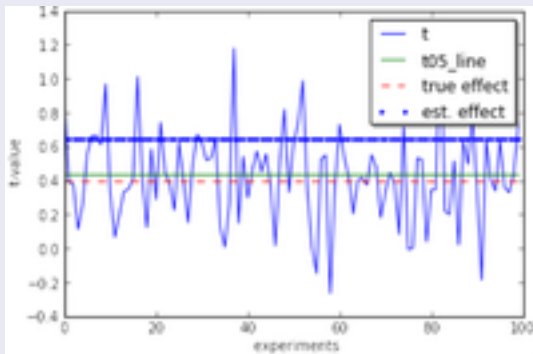


Figure: A quick simulation

What is specific to Imaging Genetics

- Combination of imaging and of genetics issues: “AND” (if independent: would probability of getting it right would multiply: $.7 * .7 = .5$)
- The combination of having to get very large number of subjects for GWAS and not being able to get them in imaging
- The multiple comparison issues
- The “trendiness” of the field
- The rapidity of the changes
- The capacity to “rationalize findings” (eg: noise in brain images is always interpretable)

What are the solutions: technical

- Pre-register hypotheses
- Statistics:
 - Always try to get a sense of the power
 - Take the right statistical tool
 - Meta analysis if you can
 - Replication always
 - Effect size variation estimation (bootstrapping)

What are the solutions: learning

- Learn the right tools:
 - How can I check my code ? How can I go back to a certain state ? (learn git/mercurial, learn git Annex or others)
 - How can others check my analyses? Learn the emerging social open science frameworks
- Learn “one layer below” (A. Martelli)

Train the new generation

- Statistics: in depth
- Computing: in depth
- A more collaborative and a more open science model
- Work such that the next post-doc will need weeks to start progress - not months
- Work such that others in the community can reproduce **and** build upon

What are the solutions: social

- Put some pressure on editors to
 - Accept replication studies
 - Accept preregistration
 - Increase the verifiability of analyses (code and data available)
- Share data / share intermediate results
- Increase the capacity of the community to verify
- Increase capacity to do meta/mega analyses
- Because we should be more interested in replication and new findings than our own publication record
- Put pressure to change their evaluation criteria - Decrease publication pressure

Conclusion : learn from Donoho et al.

An article about computational science in a scientific publication is not the scholarship itself, it is merely advertising of the scholarship. The actual scholarship is the complete software development environment and the complete set of instructions which generated the figures.

—D. Donoho

Figure: Donoho on publication