

Assessing publication bias in coordinate-based meta-analysis techniques

Freya Acar¹, Ruth Seurinck¹, Simone Kühn², Beatrijs Moerkerke¹

¹ Ghent University, Department of Data Analysis, ² Max Planck Institute for Human Development, Berlin

Ghent University



Meta-analysis of functional Magnetic Resonance Imaging (fMRI)

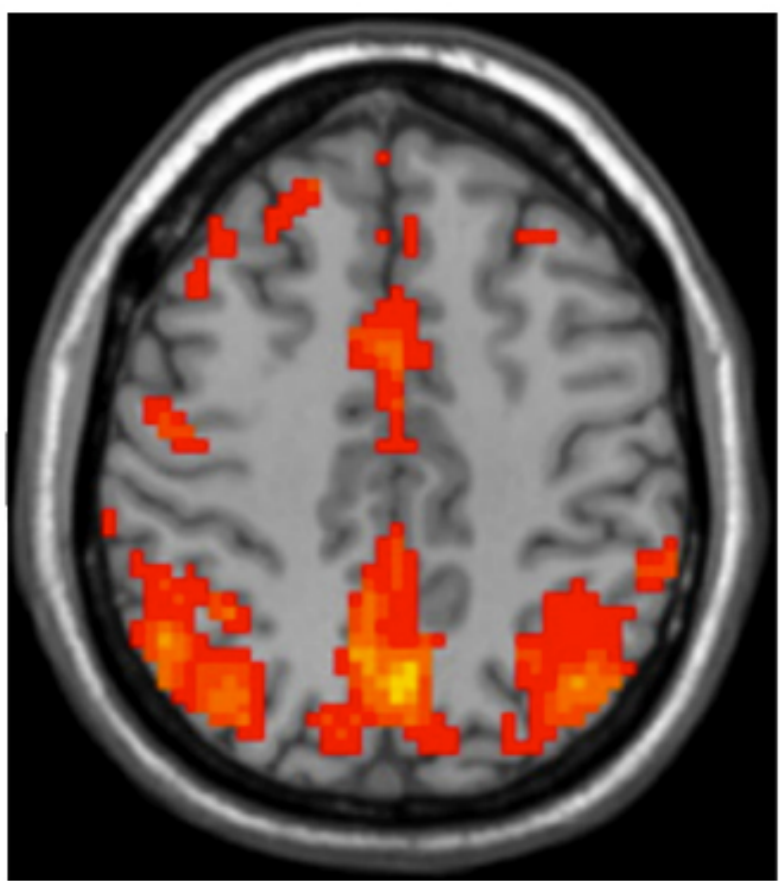
Coordinate-based meta-analysis is a popular method for fMRI, these toolboxes have been developed:

- Activation Likelihood Estimation (ALE) [2,3]
- Multi-level Kernel Density Analysis (MKDA) [4]
- Seed-based d Mapping [5] (uses peak height as effect size when available)

Meta-analyses require publication bias diagnostics

- Publication bias: studies that fail to show significance in a certain region fail to get published
- This study introduces publication bias measures for coordinate-based meta-analysis methods that do not rely on effect sizes (e.g. ALE)

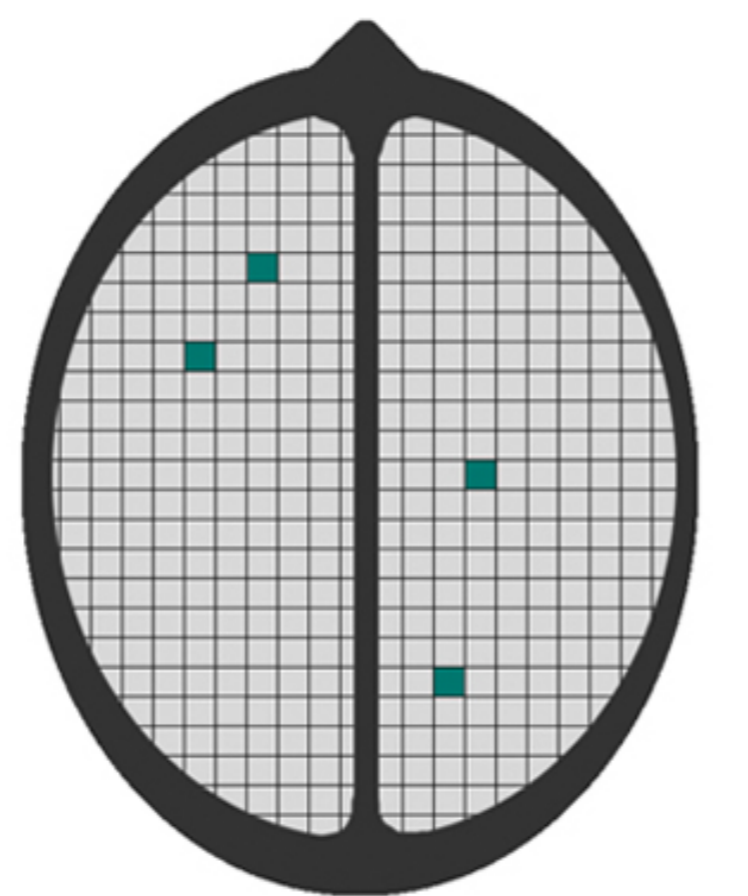
WHOLE BRAIN MAP



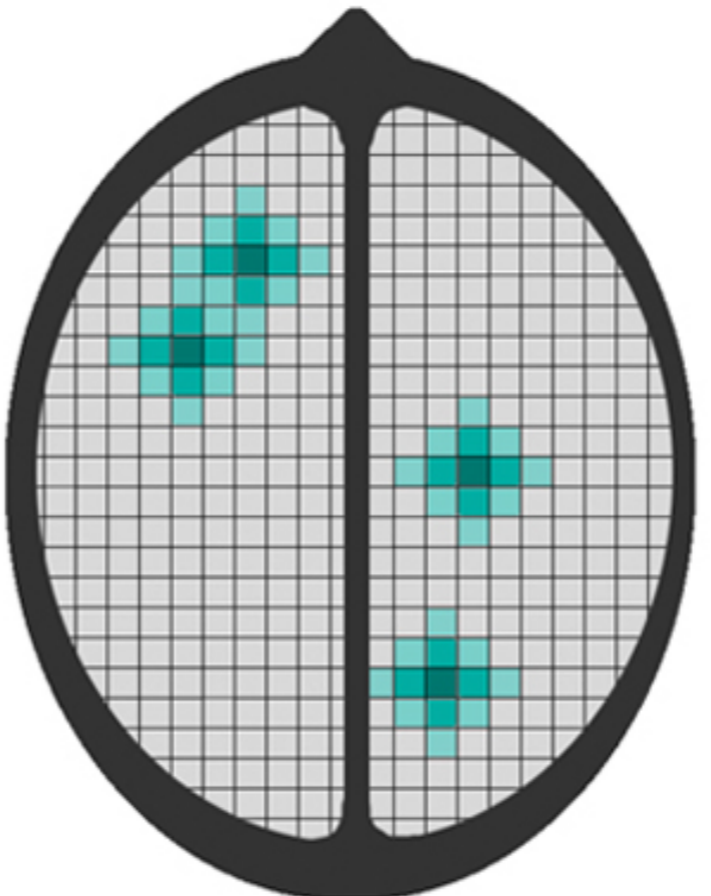
RESULTS

t(1,41)	Talairach coordinates (reported in LPI)		
	X	Y	Z
4.65	48.9	-2.8	31.8
4.75	-45.1	-11.0	32.0
4.63	-47.0	-13.0	34.8
4.51	48.7	-6.9	35.0
4.31	-27.9	-7.1	-9
3.62	-7.2	-5	49.2
4.26	12.0	-13.9	38.6
5.11	-35.5	-2.0	-8.5
5.35	50.0	2.7	-22.9

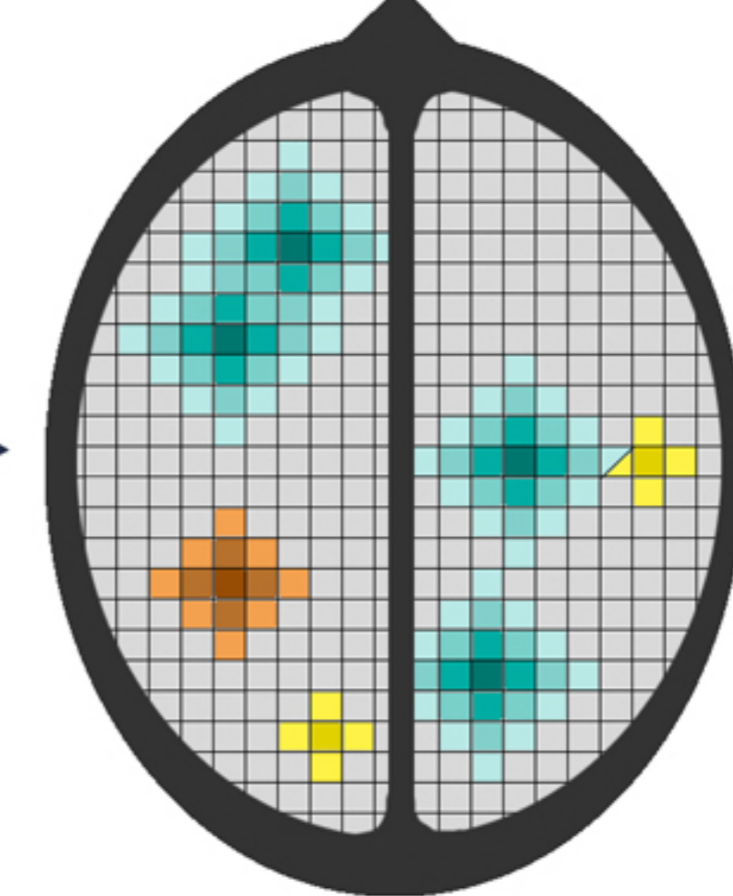
PEAK COORDINATES



GAUSSIAN KERNEL



MULTIPLE STUDIES



UNION OF ACTIVATIONS

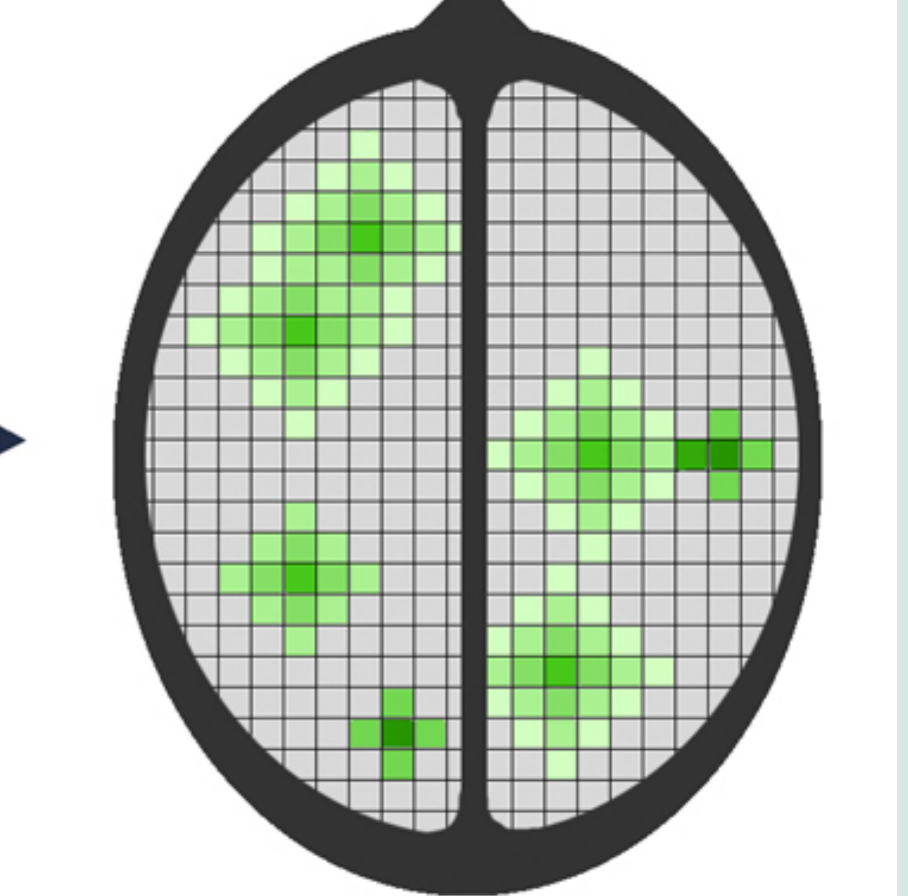


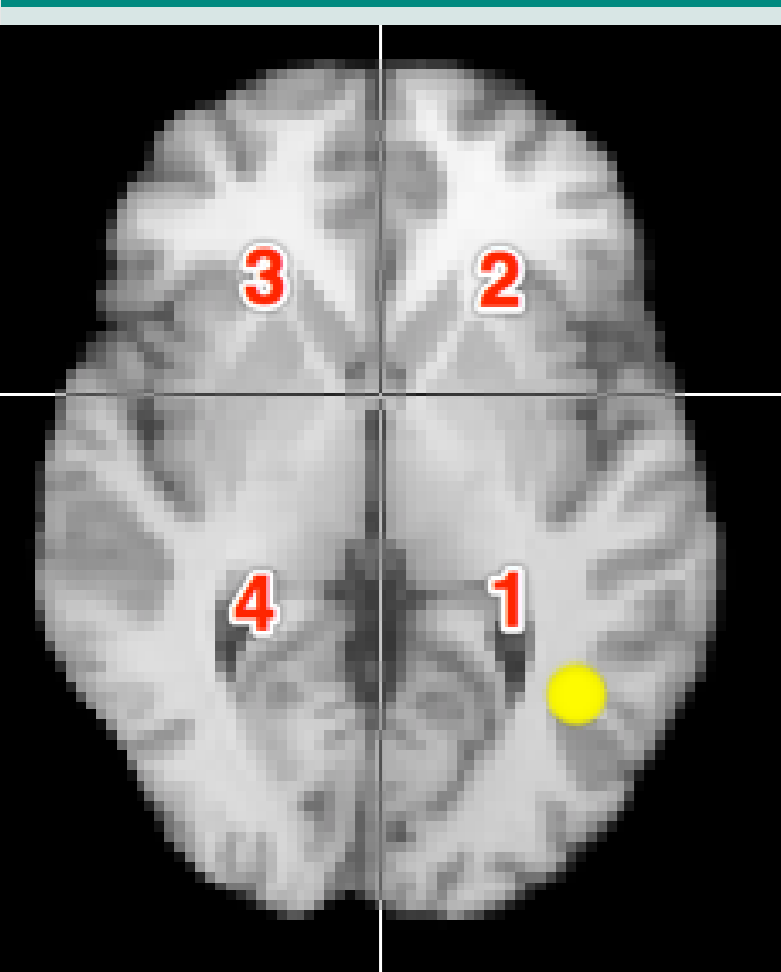
Figure 1: Overview of ALE meta-analysis.

Introduction of 2 methods for the assessment of publication bias:

① Fail-Safe N [6] (test for % contributing studies)

How many null studies can be added to a meta-analysis showing a significant effect in a region before the result is no longer statistically significant?

Simulation study



- 3 'real' studies with 1 peak in target region distance on average 3mm from location true activation
- Null studies each 1 peak in quadrant 2,3 or 4
- Effect of sample size: small (n~10), medium (n~20) or large (n~30), se=1
- Effect of thresholding: 7 thresholding methods

Figure 2: The brain is divided in 4 quadrants for simulations, true activation at the location of the yellow dot.

Results and discussion

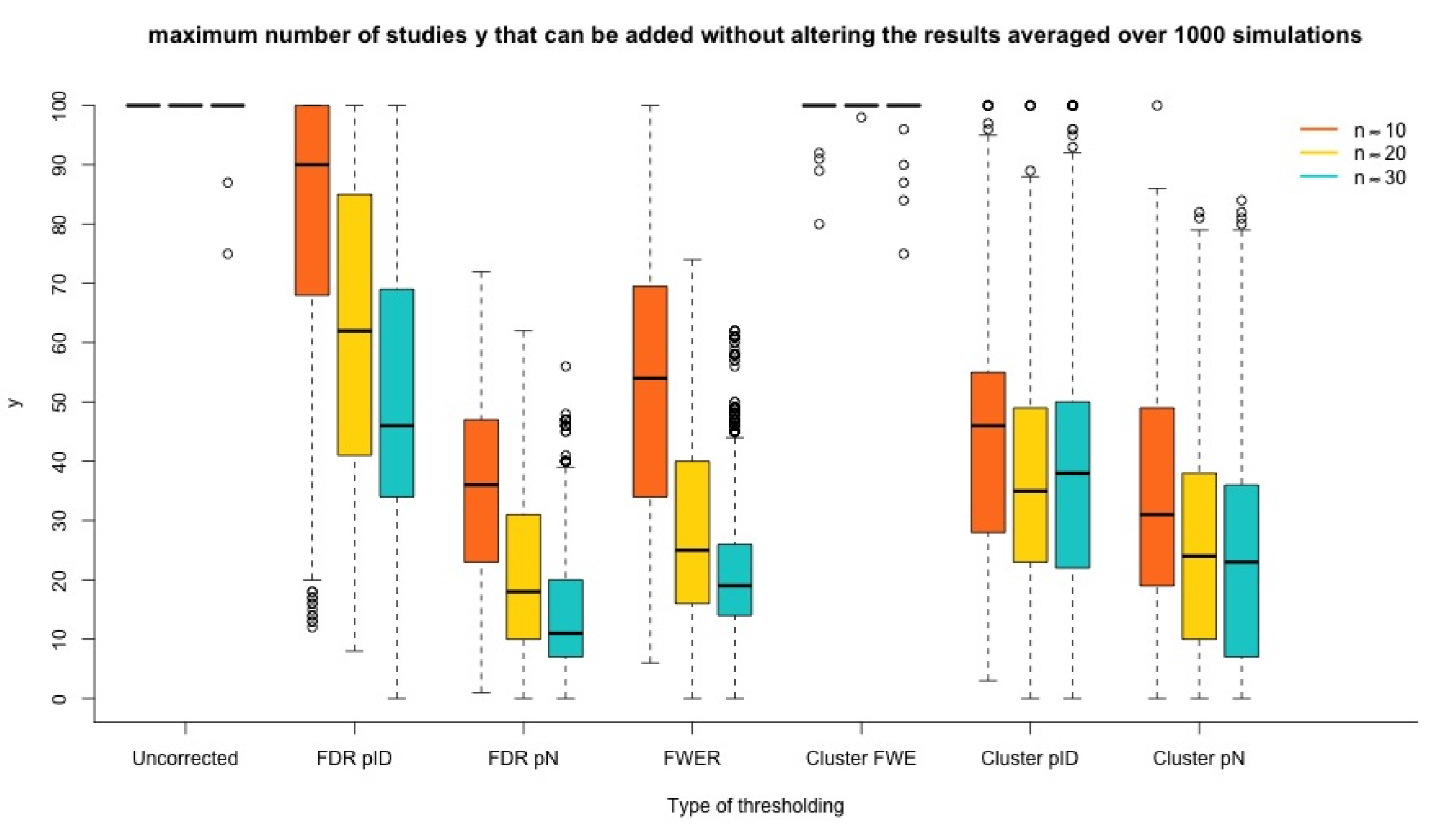


Figure 3: Amount of null studies that can be added to a meta-analysis of 3 studies before the target cluster is no longer statistically significant, by thresholding method and average sample size.

Robustness versus leniency:

- Big influence of both thresholding method and sample size
- A lot of variability within conditions
- What is an acceptable number of null studies that can be added without altering the results?
 - Too low? Points at non-robust results. (In spirit of classic Fail-Safe N [5])
 - Too high? One or a small number of studies drives the entire analysis
 - Results for sample size: contra-intuitive for robustness but intuitive for leniency
- Uncorrected thresholding shows large influence of small number of studies [9]

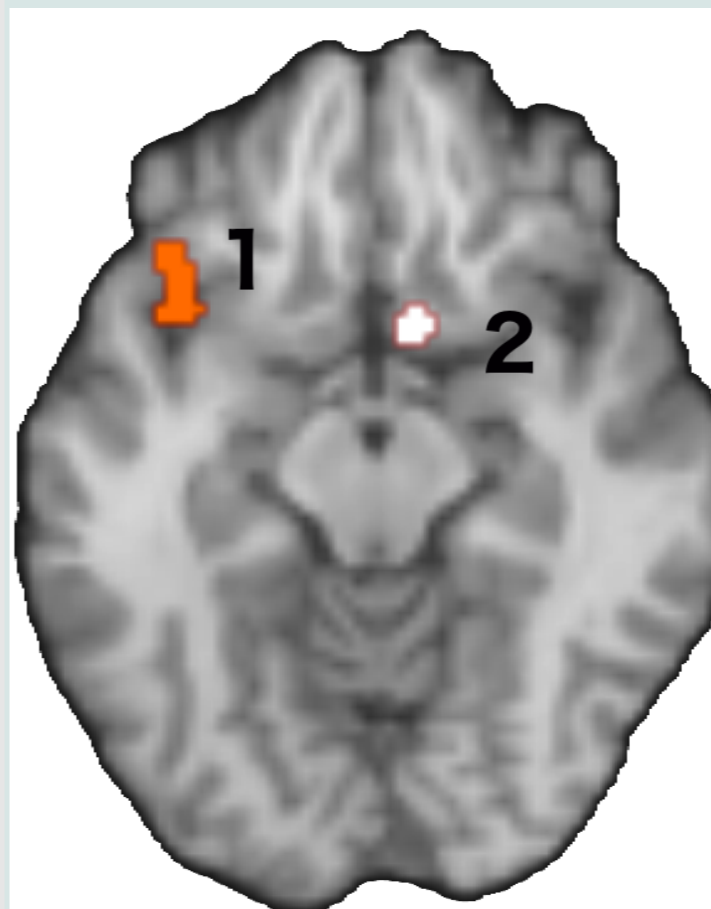
② Regression test for sample size of contributing studies [8]

Verifies whether the resulting clusters of a meta-analysis are caused by activation foci stemming from small studies (small sample bias).

How does it work?

- Sleuth database was searched for experiments with paradigm class 'taste'.
- Contrast taste > no taste was selected (87 studies, 529 foci).
- Voxel FWE < 0.05 thresholding resulted in 4 statistically significant clusters.
- Two of these clusters are plotted below and checked for a small sample bias.

This meta-analysis is conducted solely for demonstration, references on request.



Study	n	Contribution to cluster?	
		1	2
1	12	Yes	No
2	8	Yes	No
3	11	No	Yes
4	6	Yes	Yes
5	15	No	Yes
6	14	No	Yes
...

- Given a significant ALE cluster, included studies get a value of 0 (did not contribute to cluster) or 1 (contributed to cluster)
- activation (x-axis) is plotted against sample size (y-axis)
- slope gives an indication about publication bias

Figure 4: Two significant clusters and a selection of studies, their sample sizes and whether they have foci that contributed to the clusters.

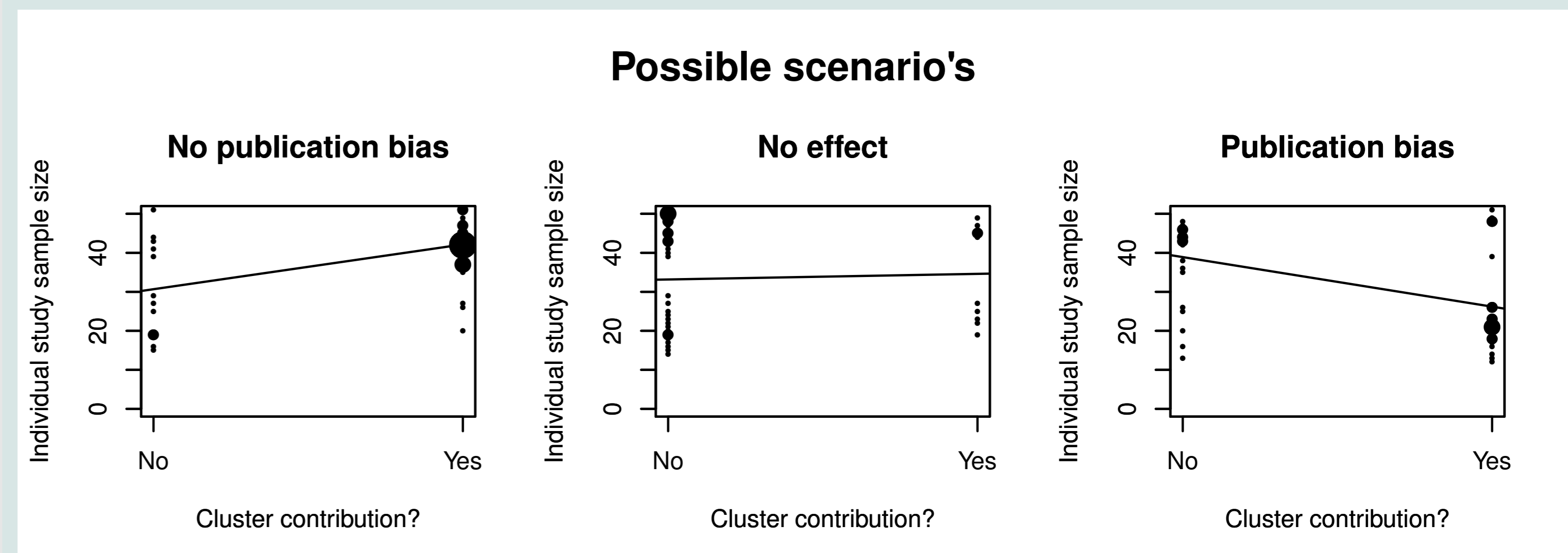


Figure 5: Possible patterns based on presence or absence of publication bias.

Results and discussion

- Results indicate a small sample bias for the first cluster, not for the second.
- Effect of thresholding at study level?

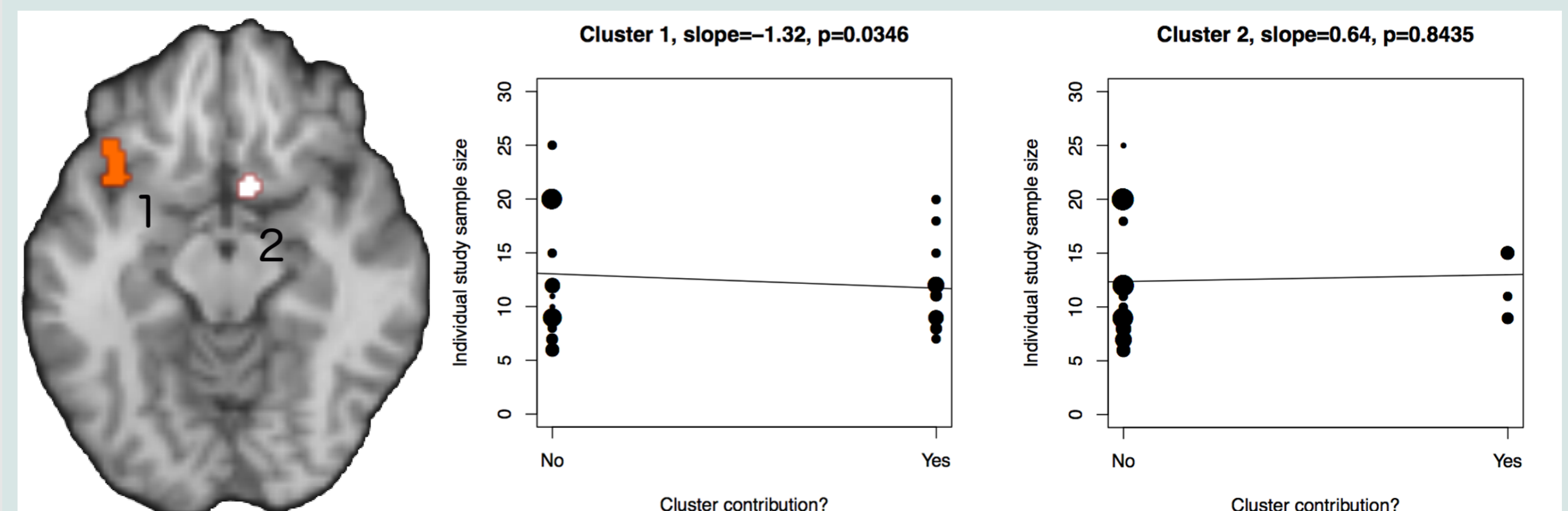


Figure 6: Regression test results for each of the two depicted clusters.

¹ Han et al. (2015) *Progress in Neuro-Psychopharm. & Biol. Psychiatry*, 59.
² Eickhoff et al., (2009, 2012). *Human Brain Mapping*, 30; *Neuroimage*, 59.
³ Turkeltaub et al., (2012). *Human Brain Mapping*, 33.
⁴ Wager et al., (2009). *Neuroimage*, 45.
⁵ Radua et al., (2009, 2010, 2012, 2014). *Br J Psychiatry*, 195; *Arch Gen Psychiatry*, 67; *Eur Psychiatry*, 59; *Front Psychiatry*, 59.
⁶ Rosenthal, (1979). *Human Brain Mapping*, 33.
⁷ Light & Pillimer, (1984).
⁸ Egger & Smith, (1997). *BMJ*, 315.
⁹ Eickhoff et al., (2016). *Neuroimage*, 173.

