

# Is Z enough? Impact of Meta-Analysis using only Z/T images in lieu of estimates and standard errors

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**Abstract.** The abstract should summarize the contents of the paper using at least 70 and at most 150 words. It will be set in 9-point font size and be inset 1.0 cm from the right and left margins. There will be two blank lines before and after the Abstract. ...

**Keywords:** TODO

## 1 Introduction

## 2 Methods

### 2.1 Theory

Given a set of  $k$  studies, we denote for each study  $i$ : its contrast estimate by  $Y_i$ , its contrast variance estimate by  $V_{Y_i}$ , its standardized statistical map by  $Z_i$  and its sample size by  $n_i$ .

*Combining contrast estimates and their standard error* The gold standard approach to combine contrast estimates and their standard errors is to input them into a GLM [3], creating effectively the third-level of a hierarchical model (level 1: subject; level 2: study; level 3: meta-analysis). The general formulation is:

$$Y = X\beta + \epsilon \quad (1)$$

where  $\beta$  is the meta-analytic parameter to be estimated,  $Y = [Y_1 \dots Y_k]^T$  is the vector of contrast estimates and  $\epsilon \sim \mathcal{N}(0, W)$  is the residual term. Eq. (1) can be solved by weighted least square giving:

$$\hat{\beta} = (X^T W X)^{-1} X^T W Y \quad (2)$$

$$\text{Var}(\hat{\beta}) = (X^T W X)^{-1} \quad (3)$$

In a random-effects model, we have  $W = \text{diag}(\sigma_1^2 + \tau^2 \dots \sigma_k^2 + \tau^2)$  where  $\tau^2$  denotes the between-studies variance. Approximating  $\sigma_i^2$  by  $V_{Y_i}$  and given  $\hat{\tau}^2$  an

	Statistic	Disribution under $H_0$
FFX GLM	$\frac{1}{\sqrt{\sum_{i=1}^k 1/V_{Y_i}}} \sum_{i=1}^k \frac{Y_i}{V_{Y_i}}$	$\mathcal{T}_{(\sum_{i=1}^k n_i - 1) - 1}$
MFX GLM	$\frac{1}{\sqrt{\sum_{i=1}^k 1/(V_{Y_i} + \hat{\tau}^2)}} \sum_{i=1}^k \frac{Y_i}{V_{Y_i} + \hat{\tau}^2}$	$\mathcal{T}_{k-1}$
RFX GLM	$\frac{1}{\hat{\sigma}_C^2/\sqrt{k}} \sum_{i=1}^k \frac{Y_i}{k}$	$\mathcal{T}_{k-1}$
Contrast Permutation	$\frac{1}{\hat{\sigma}_C^2/\sqrt{k}} \sum_{i=1}^k \frac{Y_i}{k}$	Determined through permutations with sign switching.
Fisher's	$-2 \sum_{i=1}^k \ln(\Phi(-Z_i))$	$\chi_{(2k)}^2$
Stouffer's	$\frac{\sum_{i=1}^k Z_i}{\sqrt{k}}$	$\mathcal{N}(0, 1)$
Optimally weighted-Z	$\frac{\sum_{i=1}^k \sqrt{n_i} Z_i}{\sqrt{\sum_{i=1}^k n_i}}$	$\mathcal{N}(0, 1)$
Stouffer's MFX	$\frac{\sum_{i=1}^k Z_i}{\sqrt{k\hat{\sigma}}}$	$\mathcal{T}_{k-1}$
Z Permutation	$\frac{\sum_{i=1}^k Z_i}{\sqrt{k}}$	Determined through permutations with sign switching.

**Table 1.** Statistics for one-sample meta-analysis tests and distributions under the null hypothesis.

estimate of  $\tau^2$  we obtain the statistics detailed in table 1 for a one sample test. This reference approach will be referred to as **Mixed-effects (MFX) GLM**.

In a fixed-effects model (i.e. assuming no between-study variances), we have  $W = \text{diag}(\sigma_1^2 \dots \sigma_k^2)$  where  $\sigma_i^2$  denotes the contrast variance for study  $i$ . This approach will be referred to as **Fixed-effects (FFX) GLM**.

*Combining contrast estimates* In the absence of standard error, the contrast estimates  $Y_i$  can be combined by assuming that the within-study variance  $\sigma_i^2$  is roughly constant ( $\sigma_i^2 \simeq \sigma^2 \forall 1 \leq i \leq k$ ) or negligible by comparison to the between-study variance ( $\sigma_i^2 \ll \tau^2 \forall 1 \leq i \leq k$ ). Then  $W = \text{diag}(\sigma_C^2 \dots \sigma_C^2)$  where  $\sigma_C^2$  is the combined within and between-subject variance such as  $\sigma_C^2 \simeq \tau^2$  or  $\sigma_C^2 \simeq \tau^2 + \sigma^2$ . Under these assumptions, eq. (1) can be solved by ordinary least square giving:

$$\hat{\beta} = (X^T X)^{-1} X^T Y \quad (4)$$

$$\text{Var}(\hat{\beta}) = (X^T W X)^{-1} \quad (5)$$

Given  $\hat{\sigma}_C^2$ , an estimate of  $\sigma_C^2$  we obtain the statistics presented in table 1 for one sample tests. This approach will be referred to **Random-effects (RFX) GLM** in the following.

As an alternative to parametric approaches, non-parametric statistics [5, 6] can be computed by comparing the RFX GLM T-statistic to the distribution obtained by permuting the sign of each sample included in the analysis. This approach will be referred to as **Contrast permutation**.

*Combining standardised statistics* In the presence of standardised statistical estimates, **Fisher's** meta-analysis provide a statistic to combine the associated p-values [4]. **Stouffer's** approach combines directly the standardised statistic [7]. In [9] following [2], the author proposed a weighted method that weights each study's  $Z_i$  by the square root of its sample size [3,7]. This approach will be referred to as **Optimally weighted-Z**. All these meta-analytic statistics assumes fixed-effects (no between-study variance) and are suited only for one-sample tests. The corresponding statistics are presented in table 1.

As suggested in [1], to get a kind of MFX with Stouffer's approach, the standardised statistical estimates  $Z_i$  can be combined in an OLS analysis. The corresponding estimate, referred as **Stouffer's MFX** is also provided in 1

As an alternative to parametric approaches, a non-parametric distribution [5, 6] can be estimated by permutation on the  $Z_i$ 's. This approach will be referred to as **Z permutation**.

## 2.2 Experiments

**Simulations** To verify the validity of each estimator under the null hypothesis we estimated the false positive rate at  $p < 0.05$  uncorrected. For each meta-analysis, we simulated a contrast estimate and a variance estimates such as:

$$Y_i \sim \mathcal{N}(0, \frac{\sigma_i^2}{n_i} + \tau^2) \quad (6)$$

$$V_{Y_i} \sim \frac{\sigma_i^2}{n_i - 1} \chi_{(n_i - 1)}^2 \quad (7)$$

where  $\sigma_i^2 \in [1/2, 1, 2, 4]$  is the within-study variance,  $\tau^2 \in [0, 1]$  is the between-study variance (fixed-effects if  $\tau^2$  is 0, random-effects otherwise). We simulated different number of studies per meta-analysis:  $k \in [5, 10, 25, 50]$  and the number of subjects per studies  $n_i$  was selected such as we would have varying number of subjects per studies in given meta-analysis across the common range of subjects involved in neuroimaging studies. In each simulated meta-analysis we simulated one study with exactly 20, 25, 10 and 50 subjects. For the remaining studies the number of subjects were drawn from uniform distributions a quarter from  $\mathcal{U}(11, 20)$ , a quarter from  $\mathcal{U}(26, 50)$  and the remaining from  $\mathcal{U}(21, 25)$ . A total of 32 parameter sets ( $4 \sigma_i^2 \times 2 \tau^2 \times 4 k$ ) was therefore tested, 71 repeats with 5041 samples per repeats were simulated.

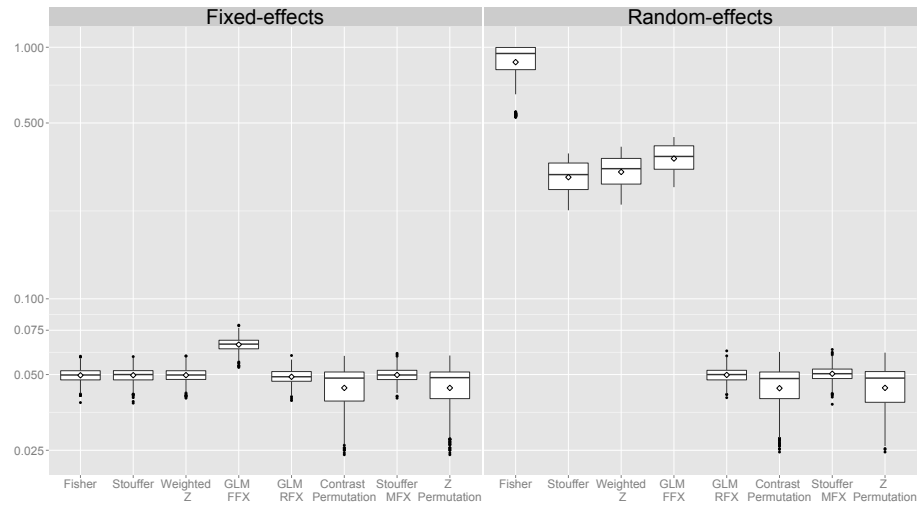
**Real data** We first compared the Z-scores obtained by the three approaches using a Bland-Altman plot. Then, as results are usually presented as a thresholded map, we computed the dice similarity score between thresholded maps obtained with Stouffer’s and weighted-Z FFX with FLAME FFX for three (uncorrected) thresholds:  $p \leq 0.001$ ,  $0.01$  and  $0.05$ . Finally, as results are best reported using a multiple comparison correction, we defined ground truth activations as the FLAME FFX analysis FDR-corrected at a threshold of  $p \leq 0.05$  and plotted Receiver-Operating-Characteristics (ROC) curves of Stouffer’s and weighted-Z FFX.

All plots were generated using ggplot [8].

### 3 Results

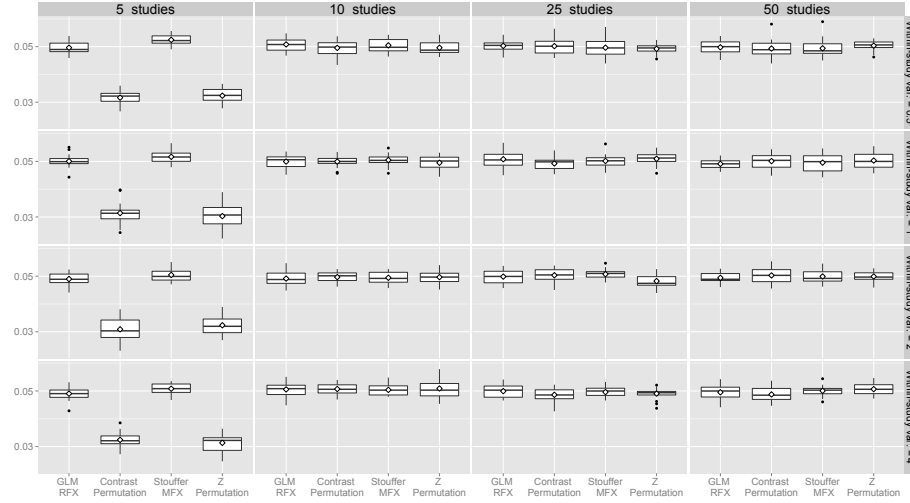
#### 3.1 Simulations

Fig. 1 displays the false positive rate at  $p < 0.05$  obtained for the eight estimators over all set of parameters in the absence and presence of random-effects. From this graph, it is clear that the fixed-effects meta-analytic summary statistics, i.e. Fisher’s, Stouffer’s and weighted-z estimates are overly liberal in the presence of random-effects. As expected the original Fisher’s approach is the most invalid. Surprisingly, FFX GLM is also invalid under fixed-effects, maybe suggesting inaccurate degrees of freedoms (here set to  $(\sum_{i=1}^k n_i - 1) - 1$ ). Stouffer’s MFX, GLM RFX and permutations of  $Y_i$ ’s or  $Z_i$ ’s provide valid estimates. The permutation estimates present the largest sampling variance.



**Fig. 1.** False positive rates of the meta-analytic estimators under the null hypothesis for  $p < 0.05$ .

The impact of the number of studies involved in the meta-analysis and of the size of the within-study variance are investigated in fig. 2. The permutation estimates appears conservative (FPR  $\simeq 0.03$ ) when 5 studies are involved. All approaches perform equally as soon as 10 or more studies are included in the meta-analysis.



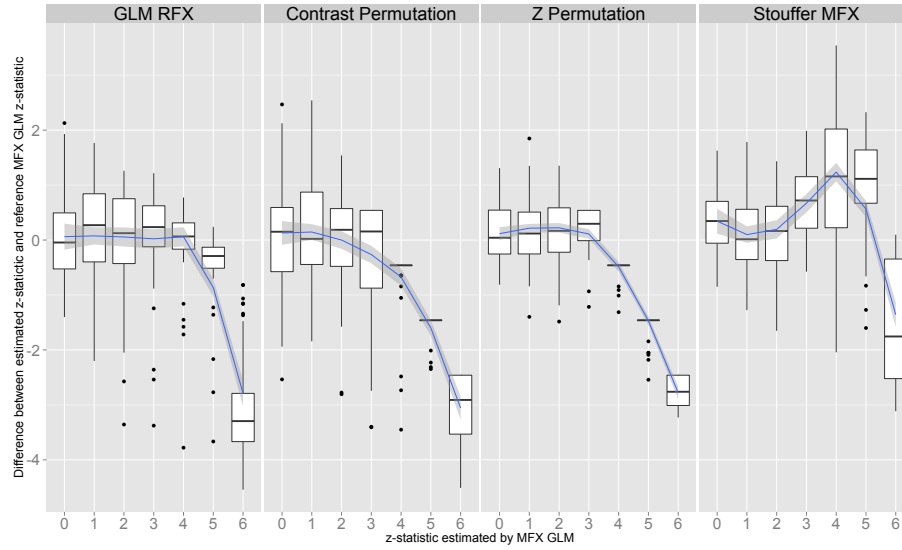
**Fig. 2.** False positive rates of the valid random-effects meta-analytic estimators under the null hypothesis for  $p < 0.05$  as a function of the number of studies and the within-study variance.

### 3.2 Real data

Fig. 3 plots the difference between the z-score estimated by each meta-analytic approach and the reference z-score computed with MFX GLM. GLM RFX and contrast permutations provide z-scores estimate that are equal or smaller than the reference. Z permutation provides slightly larger z-scores between 1 and 3 (reference p-values between 0.16 and 0.0013) but is mostly in agreement with the reference z-scores. On the other hand, Stouffer's MFX is more liberal than the reference for z-score ranging from 3 to 5 (reference p-values between 0.0013 and  $2.9e-07$ ) and more stringent for z-scores smaller than 5.

Dice among valids

1. StouffersMFX: 0.9454
2. PermutZ: 0.9450
3. GLMRFX: 0.8994
4. PermutCon: 0.8991



**Fig. 3.** Difference between the z-score estimated from each meta-analytic approach and the reference z-score from MFX GLM as a function of reference z-score

1. WeightedZ: 0.9244
2. Stouffers: 0.9184
3. GLMFFX: 0.8972
4. fishers: 0.8382

AUC between 0 and 0.1 among valids

1. StouffersMFX: 0.8924
  2. PermutZ: 0.8919
  3. GLMRFX: 0.7809
  4. PermutCon: 0.7815
- 
1. WeightedZ: 0.8293
  2. Stouffers: 0.8619
  3. fishers: 0.6329
  4. GLMFFX: 0.6111

## 4 Conclusion

We have found appreciable differences between the Z-score only approaches as compared to a gold-standard approach. Overall the weighted-Z method provided results that were closer to the ground truth than Stouffer's approach. We hypothesize that Stouffer's methods may be attributing greater weights to less-representative subsets of the data. All three procedures are valid, but the

gold-standard should be giving the most faithful representation of the population effect. This advocates over the development of tools supporting the sharing E+SE's.

## 5 Acknowledgements

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## References

1. Meta-analysis of neuroimaging data: a comparison of image-based and coordinate-based pooling of studies. *NeuroImage*, 45(3):810–23, 2009.
2. Peter Cummings. On the combination of independent tests. *Magyar Tud. Akad. Mat. Kutato Int. Kozl.*, 3:171–197, 1958.
3. Peter Cummings. Meta-analysis based on standardized effects is unreliable. *Archives of pediatrics & adolescent medicine*, 158(6):595–7, 2004.
4. R.A. Fisher. *Statistical Methods for Research Workers*. Oliver and Boyd, Edinburgh, 1932.
5. A. P. Holmes, R. C. Blair, G. J.D. Watson, and I. Ford. Nonparametric Analysis of Statistic Images from Functional Mapping Experiments. *Journal of Cerebral Blood Flow and Metabolism*, 16:7–22, 1996.
6. Thomas E. Nichols and Andrew P Holmes. Nonparametric permutation tests for functional neuroimaging: a primer with examples. *Human brain mapping*, 15(1):1–25, January 2002.
7. S. Stouffer, L. DeVinney, and E. Suchmen. *The American Soldier: Adjustment During Army Life*, volume 1. Princeton University Press, Princeton, NJ, 1949.
8. H. Wickham. *ggplot2: elegant graphics for data analysis*. Springer New York, 2009.
9. D V Zaykin. Optimally weighted Z-test is a powerful method for combining probabilities in meta-analysis. *Journal of evolutionary biology*, 24(8):1836–41, 2011.