



Evidence-based Second-Generation p-values on Functional Magnetic Resonance Imaging Data

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Introduction to fMRI data

ENAR
2 of 17

Introduction to fMRI data

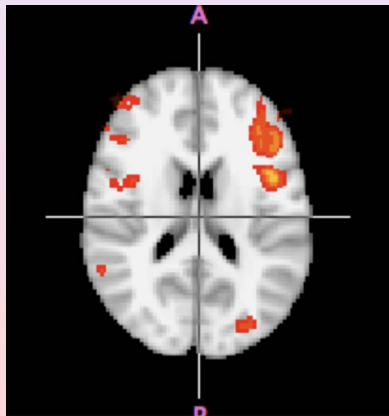
Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
ready to use
in practice?

1. Task-induced fMRI
 - Participants are engaged with tasks
 - Look for localized brain activation patterns
2. Statistical inference relies on p-values





P-values as an inference tool - good or bad?

Advantages:

- Simple computation
- Widely used in all fields

Disadvantages:

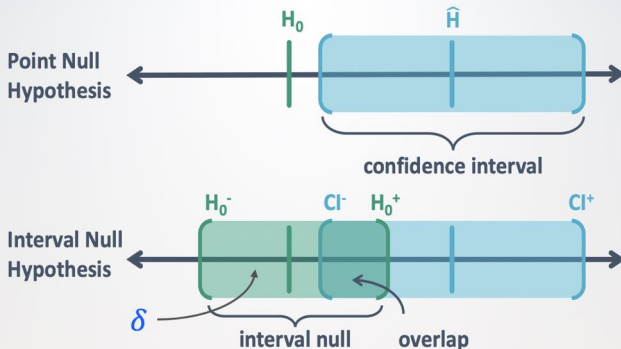
- Interpretation issues¹
- Statistical v.s. Clinical significance²

¹Hubbard et al. (2003)

²Mark et al. (2016); Ranganathan et al. (2015)



SGPV brings in interval null testing



Point null hypothesis H_0 and interval null hypothesis $[H_0^-, H_0^+]$

Data-supported hypothesis \hat{H} and confidence interval $[CI^-, CI^+]$



Simple interpretation as fraction of overlap

ENAR
5 of 17

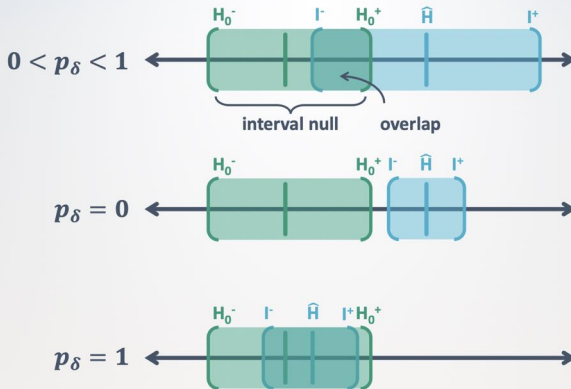
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Inference Tools

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Works with confidence, credible, and support intervals



The computation of SGPV is concise

Second-generation p-value (SGPV)

$$p_{\delta} = \frac{|I \cap H_0|}{|I|} \times \max\left\{\frac{|I|}{2|H_0|}, 1\right\}$$

Proportion of data-supported
hypotheses that are also
null hypotheses

**Small-sample
correction factor**

shrinks proportion
to $\frac{1}{2}$ when $|I|$ wide

when $|I| > 2|H_0|$



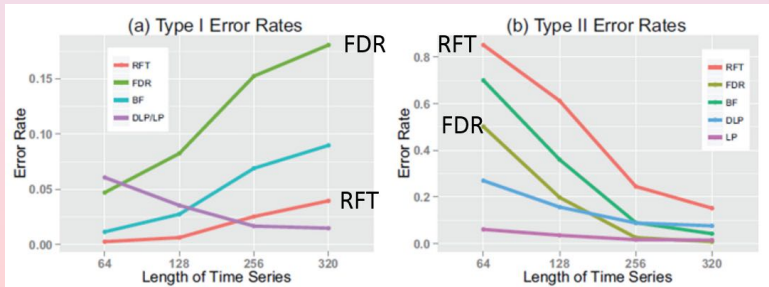
Balancing power and Type I error rate is challenging in fMRI analysis

Nature of the data

- Conduct analysis on large number of voxels
- Data are noisy

Multiple comparison adjustments

- Control FWER (Random Field Theory (RFT))
- Control False Discovery Rate (FDR)



¹Single subject simulation results from Kang et al. (2015)



Settings to simulate task-induced fMRI data

ENAR

8 of 17

Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
ready to use
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- Spatio-temporally correlated data
- 32×32 , 2-D images with two active regions
- At each voxel and a time, with P stimuli,
$$Y_v(t) = \sum_{p=1}^P X_p(t) \beta_v^p + \epsilon_v(t)$$
- Clinically null region
 - No neurons in Cerebrospinal fluid (CSF)
→ any signal is **noise**
 - 10×10 region
 - $\beta^1 = \beta^2 = 0$
- Vary time lengths and sample sizes
- Gaussian kernel with FWHM = 8 mm



Visualize the true image

ENAR
9 of 17

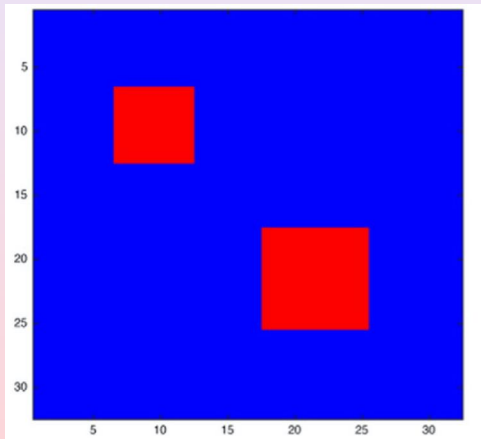
Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

**Statistical
inference in
fMRI data
analysis**

Is SGPV
ready to use
in practice?





Estimation and Inference of simulation

ENAR
10 of 17

Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
ready to use
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1. Fit the linear model at each voxel to estimate $\beta^2 - \beta^1$
2. SGPV clinical null region:
 - $0 \pm \text{IQR}(\text{all } \hat{\beta}^2 - \hat{\beta}^1 \text{ in CSF}) / 6$
3. Methods compared:
 - SGPV: Compute SGPV with 95% CI
 - FDR : Compute p-values; control at FDR = 0.05
 - RFT : Compute z-scores; control error probability at 0.05
4. Fair comparison: Dichotimized SGPV (D-SecondP)
 - Voxels with SGPV = 0 are deemed as **significant**
 - Voxels with SGPV = 1 are deemed as **insignificant**
 - Voxels with $0 < \text{SGPV} < 1$ (inconclusive region) are deemed as **insignificant**



SGPV obtains power and keep Type I error rate steady

ENAR
11 of 17

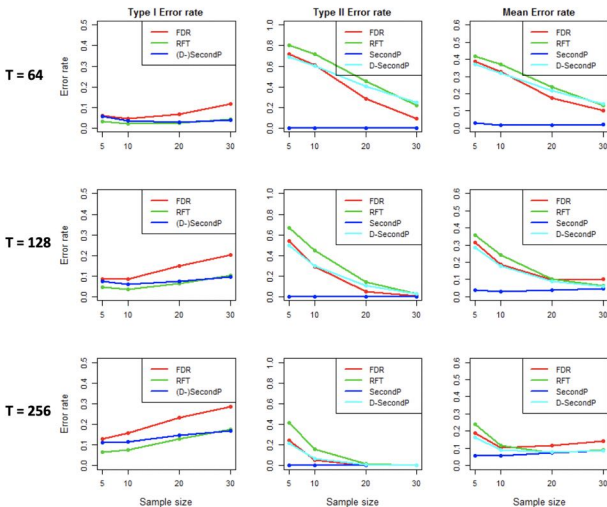
Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
ready to use
in practice?





Participants and task in real fMRI data

ENAR
12 of 17

Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
ready to use
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Data:

- 29 women with major depression disorders, aged between 45 and 75
- Spatial attention task that measures attention bias

Analysis:

1. Linear model at each voxel to estimate $\beta^2 - \beta^1$
2. Compute SGPV with clinical null region
 - $0 \pm \text{IQR}/6$ using estimates from voxels in CSF
3. For voxels outside of CSF
 - Compute SGPV with 95% CI
 - Control z-scores with RFT at 0.05
 - Control p-values with FDR at 0.05
4. Data Decimation



SGPV method is the most robust method

ENAR
13 of 17

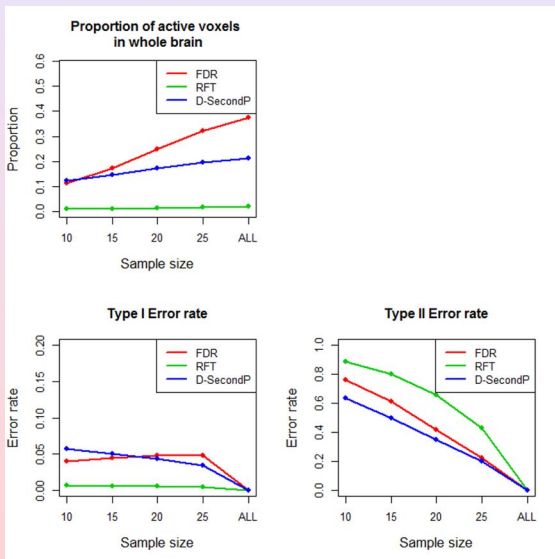
Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
ready to use
in practice?





Visualization of the results

ENAR
14 of 17

Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

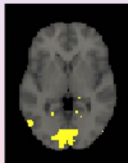
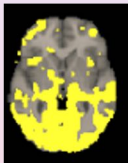
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FDR

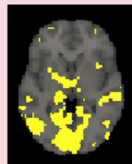
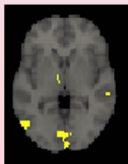
RFT

D-SecondP

N = 29



N = 15





SGPV offers good properties in fMRI analysis

ENAR
15 of 17

Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
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1. Convenient and simple interpretation
2. Incorporate clinical information (in fMRI: data collected from CSF)
3. Provide transparency, rigor and reproducibility of scientific results
4. Reduce false positives in fMRI data analysis



Tools have been developed; more are coming

ENAR
16 of 17

Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

Is SGPV
ready to use
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1. Method:

- “*An Introduction to Second-Generation p-Values*”, Blume et al. (2019)
- “*Second-generation p-values: Improved rigor, reproducibility, transparency in statistical analyses*”, Blume et al. (2018)

2. R package:

- github.com/weltybiostat/sgpv

3. R-shiny app for fMRI analysis (work in progress)

- Offer clear explanation of SGPV
- Provide interactive visualization
- Download results with format compatible with other imaging software



Acknowledgments

ENAR
17 of 17

Introduction
to fMRI data

Inference
Tools

Second-
generation
p-values
(SGPV)

Statistical
inference in
fMRI data
analysis

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ready to use
in practice?

Advisor: Prof. Hakmook Kang, Department of Biostatistics, Vanderbilt University

SGPV method:

- Prof. Jeffrey Blume, Department of Biostatistics, Vanderbilt University
- Valerie Welty, Department of Biostatistics, Vanderbilt University

Collaborators:

- Dr. Warren Taylor, Department of Psychiatry, Vanderbilt University
- Brian Boyd, Department of Psychiatry, Vanderbilt University