#### Advanced Methods in Applied Statistics

#### Christian Starup & Loui Wentzel

Niels Bohr Institute

March 8, 2018

◆□▶ ◆□▶ ◆臣▶ ◆臣▶ 臣 のへで

### Journal Article

# Combining dependent *P*-values with an empirical adaptation of Brown's method

## William Poole, David L. Gibbs, Ilya Shmulevich, Brady Bernard $^{\rm t}$ and Theo A. Knijnenburg\*, $^{\rm t}$

Institute for Systems Biology, Seattle, WA 98109-5263, USA

\*To whom correspondence should be addressed.

<sup>†</sup>The authors wish it to be known that, in their opinion, the last two authors should be regarded as joint Last Authors.

#### Abstract

Motivation: Combining *P*-values from multiple statistical tests is a common exercise in bioinformatics. However, this procedure is non-trivial for dependent *P*-values. Here, we discuss an empirical adaptation of Brown's method (an extension of Fisher's method) for combining dependent *P*-values which is appropriate for the large and correlated datasets found in high-throughput biology.

Results: We show that the Empirical Brown's method (EBM) outperforms Fisher's method as well as alternative approaches for combining dependent *P*-values using both noisy simulated data and gene expression data from The Cancer Genome Atlas.

Availability and Implementation: The Empirical Brown's method is available in Python, R, and MATLAB and can be obtained from https://github.com/llyaLab/CombiningDependentPvalues UsingEBM. The R code is also available as a Bioconductor package from https://www.bioconduc tor.org/packages/devel/bioc/html/EmpiricalBrownsMethod.html.

Contact: Theo.Knijnenburg@systemsbiology.org

Supplementary information: Supplementary data are available at *Bioinformatics* online.

#### Problem

Given a dataset where one needs to calculate several or many p-values. Should one account for a possible correlation between data variables?



◆□▶ ◆□▶ ◆三▶ ◆三▶ 三三 - のへで

#### No Correlation solution

If the P-values are not correlated, then according to  $H_0$  the distribution of each P-value should be uniform, and the product of P-values should then be drawn from the distribution of N products of uniform numbers:

$$P = \int_0^{\prod} \frac{(-1)^{N-1}}{(N-1)!} \cdot \ln(u)^{N-1} du$$
 (1)

This is equivalent to a  $\chi^2$ -test with 2k degrees of freedom called Fishers Method:

$$\Psi = \sum_{i=1}^{N} -2\log(P_i)$$
(2)
$$P = \phi_{i}(W) = \int_{-\infty}^{\infty} e^{2\pi i W} dW$$
(2)

$$P = \phi_{2k}(\Psi) = \int_{\Psi}^{\infty} \chi_{2k}^2(x) dx \tag{3}$$

▲□▶ ▲□▶ ▲□▶ ▲□▶ ▲□ ● ● ●

#### Correlation solution

However, if the data is correlated, we can't assume a uniform distribution of P-values.

Brown therefore expanded Fisher's method to include a re-scaling factor, c, such that  $\Psi \sim c \chi^2_{2f}$ .

$$f = \frac{E[\Psi]^2}{var[\Psi]} \quad c = \frac{Var[\Psi]}{2E[\Psi]} = \frac{k}{f} \quad Var[\Psi] = 4k + 2\sum_{i < j} cov(W_i, W_j)$$

With  $W_i = -2 \log(P_i)$ ,  $E[\Psi] = 2k$  (assuming a  $\chi^2$  distribution), k is the Fisher's DoF and f the re-scaled Brown's DoF. The combined P-value is then:

$$P_{combined} = 1 - \Phi_{2f}(\Psi/c)$$

with  $\Psi = \sum W_i$ ,  $\Phi_{2k}$  being the cumulative distribution function of  $\chi^2_{2f}$ .

#### Correlation solution continued

The articles contribution to Browns' method is to calculate the covariance matrix by an empirical approximation, thereby the Empirical Brown's method (EBM):

$$egin{aligned} \mathsf{cov}(W_i, W_j) &pprox \mathsf{cov}(w_i, w_j) \ w_i &= -2\log(1 - \mathcal{F}(\overrightarrow{x_i})) \end{aligned}$$

Kost's method uses another approach to calculate the covariance:

$$cov(W_i, W_j) \approx 3.263 \rho_{ij} + 0.710 \rho_{ij}^2 + 0.027 \rho_{ij}^3$$

The EBM is a non-parametric approach, where  $F(\vec{x_i})$  is the right-sided empirical cumulative distribution function.

#### Simulating data

Parameters were  $\mu_i = 0$ , a = 0.8, n = 4.  $b_j$  was randomly sampled from [-0.5; 0.5]. Each sample had 200 entries.

$$M = \begin{bmatrix} 1 & b_2 & \dots & b_j & \dots & b_n \\ b_2 & 1 & \dots & a & \dots & a \\ \vdots & \vdots & \ddots & \vdots & \ddots & \vdots \\ b_j & a & \dots & 1 & \dots & a \\ \vdots & \vdots & \ddots & \vdots & \ddots & \vdots \\ b_n & a & \dots & a & \dots & 1 \end{bmatrix}$$
(4)

From any sample  $\vec{y}$  drawn from this distribution, *n*-dimensional uniform noise from [-1; 1] was added:

$$\vec{x} = \vec{y} + \xi \vec{U} \tag{5}$$

They draw numbers from one axis on the multivariate normal distribution (axis 1 with correlations  $b_j$  to the others) and test the correlation to the other axes using Pearsons correlation test.



Fig. 2. Pvalues from simulated data using Fisher's method and EBM. (a) Line plot of histogram counts of Pvalues from Fisher's method applied on simulated null data with varying degrees of covariance as represented by a. The histogram was created by binning the Pvalues in 20 bins of size 0.05 from 0 to 1. (b) Similar to (a) built of Pvalues derived with the Empirical Brown's method

A D > A B > A B > A B >

ж

#### Ground Truth P-values

To test the different tests against correlated data, it should yield the same results as if the data was uncorrelated.

- Shuffle  $\vec{y}_1$
- Calculate Ψ<sup>\*</sup> as earlier
- Repeat *M* times

The ground truth P-value is then

$$P_{ground} = \frac{\sum_{m=1}^{M} I(\Psi_m^* \ge \Psi)}{M}$$
(6)

▲□▶ ▲□▶ ▲□▶ ▲□▶ ▲□ ● ● ●

Notice this gives a resolution in the ground truth P-value by 1/M.

#### Performance results as a function of Signal to Noise ratio

