

Optimal Sample Size for Multiple Testing: the Case of Gene Expression Microarrays

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Outline

Sample size choice for massive multiple comparisons.

I. Decision problems:

1. Terminal decision

2. Sample size

3. Simulation: preposterior MCMC

II. Prob Model:

4. A hierarchical Gamma/Gamma model

5. A mixture model extension

6. Results

Summary

- Discussion indep of prob model
- Two dec problems: sample size & multiple comparison
- One loss function for both.
- Terminal decision under reasonable loss functions:
Reject if marg posterior prob > threshold t.

Summary (ctd.)

- Sample size: based on same loss function
- Supplemented by power calculation and sensitivity analysis
- Evaluation by (easy) preposterior M.C.
- Prob models: (mixture of) gamma/gamma

0. Microarrays

Statistically, **massive multiple testing**: comparison of genes under two (or more) conditions to detect differentially expressed genes

	Gene A	Gene B	Gene C	...
Patient I	1	0	0	...
Patient II	1	0	1	...
Patient III	0	0	0	...
...

Huge number of genes and moderate to large number of cases
Costly preliminary step to even more costly assays (e.g., RT-PCR)

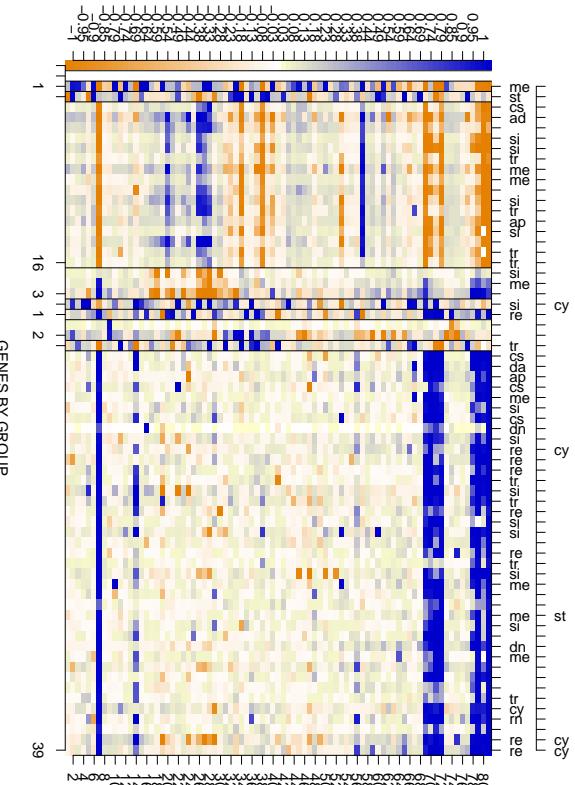
[Parmigiani et al., 2002 JRSSB]

PractBayes 5, Milton Keynes, 2003

Müller, Parmigiani, Robert & Rousseau 7

PractBayes 5, Milton Keynes, 2003

Müller, Parmigiani, Robert & Rousseau 8



I. Decision Problems

Sequential decision problem:

First sample size, then terminal decision

1. Sample size choice:

- *Before* experiment, marginalize over **putative** data
- Decide # arrays J

2. Multiple comparisons:

- *After* experiment, conditional on **observed** data
- Decide about n genes: differential vs. non-differential

1. Terminal Decision

Decision and Hypotheses: n comparisons, $i = 1, \dots, n$:

$$\text{truth} \quad z_i \in \{0, 1\}$$

$$\text{decision} \quad d_i \in \{0, 1\}$$

n **huge** plus hierarchical structure:

isolated hypothesis testing one gene at a time very inefficient

Natural Bayesian framework to pull strength from all observations with help from Decision Theory (loss function)

A central notion: FDR & FNR

[Benjamini & Hochberg, 95, JRSSB]

False Discovery Rate: generalizes **type I error** for multiple comparisons.

Let $D = \sum d_i = \# \text{ rejections}$.

$$\text{FDR}(d, z) = \frac{\sum d_i(1 - z_i)}{D} = \frac{FD}{D}$$

False negative rate: akin to **type II error**

$$\text{FNR}(d, z) = \frac{\sum (1 - d_i)z_i}{n - D} = \frac{FN}{n - D}$$

Posterior mean FDR & FNR:

[Genovese & Wasserman, Valencia 7 & JRSSb 02]

Posterior expected FDR

Let $v_i = \Pr(z_i = 1 | y)$:

$$\overline{\text{FDR}} = \mathbb{E}(\text{FDR} | y) = \mathbb{E}\left(\frac{\sum d_i(1 - z_i)}{D} | y\right) = \frac{\sum d_i(1 - v_i)}{D} = \frac{\overline{\text{FD}}}{D}$$

Simplification : only need marginal post probs v_i .

Same for FNR

$$\overline{\text{FNR}} = \mathbb{E}(\text{FNR} | y) = \frac{\sum(1 - d_i)v_i}{n - D} = \frac{\overline{\text{FN}}}{D}$$

Optimal terminal decision

Can show: optimal terminal decision ($\arg \min$ posterior expected loss)

$$d_i = \mathbb{I}_{(v_i > t)}$$

under **all** four loss functions.

Threshold t_L^* : Loss functions differ only in $t = t_L^*$:

$$t_N^*(y_J) = c/(1 + c), \quad [\text{constant}]$$

$$t_R^*(y_J) = \dots \quad [\text{implicit}]$$

$$t_{2R}^*(y_J) = \min\{t : \overline{\text{FDR}}(t, y_J) \leq \alpha\},$$

$$t_{2N}^*(y_J) = \min\{t : \overline{\text{FD}}(t, y_J) \leq \alpha\},$$

(Expected) Loss functions

Combine FD(R) and FN(R)

1. $L_N(d, y) = c \overline{\text{FD}} + \overline{\text{FN}} \quad (c > 0)$
2. $L_R(d, y) = c \overline{\text{FDR}} + \overline{\text{FNR}} \quad (c > 0)$
3. $L_{2N}(d, y) = (\overline{\text{FD}}, \overline{\text{FN}})$
4. $L_{2R}(d, y) = (\overline{\text{FDR}}, \overline{\text{FNR}})$

Multicriteria decision problems:

$$L_{2N}: \min_d \overline{\text{FN}} \quad \text{subject to} \quad \overline{\text{FD}} \leq \alpha.$$

$$L_{2R}: \min_d \overline{\text{FNR}} \quad \text{subject to} \quad \overline{\text{FDR}} \leq n \alpha.$$

[Back to basics?!]

For

$$\begin{aligned} L_R &= c \frac{\sum d_i(1 - v_i)}{D} + \frac{\sum(1 - d_i)v_i}{n - D} \\ &= C_1(D) - C_2(D) \sum d_i v_i + C_3(D) \sum v_i \frac{\sum(1 - d_i)v_i}{n - D} \\ &\geq C_1(D) + C_2(D) \sum_{i=n-D+1}^n v_{(i)} + C_3(D) \sum_{i=1}^{n-D} v_{(i)} \\ &\geq C_1(D^*) + C_2(D^*) \sum_{i=n-D^*+1}^n v_{(i)} + C_3(D^*) \sum_{i=1}^{n-D^*} v_{(i)} \end{aligned}$$

we derive

$$t^* = v_{(n-D^*)}$$

Properties

Decision: $d_i = \mathbb{I}_{(v_i > t)}$

L_N : t is constant and FDR $\rightarrow 0$

L_{2R} : $\overline{\text{FDR}}$ is constant (??)

L_{2N} : $\overline{\text{FD}}$ is constant (??)

L_R : ???

Rates of Convergence (in J)

$$P(z=1|y, \eta) = \frac{1}{1 + e^{-J\Delta} \sqrt{J}}$$

but ... Big... bummer:

$$\overline{\text{FN}}(y_J, t^*) = O_P(\sqrt{\log J/J})$$

under L_{2R}, L_{2N}, L_N

and

$$\overline{\text{FN}}(y_J, t^*) = O_P(n \sqrt{\log J/J})$$

under L_R

Much too slow/flat!!

2. Sample Size Determination

Preposterior mean loss \rightarrow sample size J .

$$L_R^m(J) = \mathbb{E}_{y_J} [\min_d \{L_R(d, y_J)\}],$$

$$L_N^m(J) = \mathbb{E}_{y_J} [\min_d \{L_N(d, y_J)\}],$$

and

$$L_{2R}^m(J) = \mathbb{E}_{y_J} \left[\min_d \{ \overline{\text{FN}}(d, y_J) \mid \overline{\text{FDR}}(d, y_J) \leq \alpha \} \right]$$

$$L_{2N}^m(J) = \mathbb{E}_{y_J} \left[\min_d \{ \overline{\text{FN}}(d, y_J) \mid \overline{\text{FD}}(d, y_J) \leq \alpha \} \right]$$

Expectation: $\mathbb{E}[\cdot]$ w.r.t. y_J

(model parameters already integrated in L)

Nested optimization: \min w.r.t. d

Power/Sensitivity

Power: prob accept as function of true effect & J

$$\beta(\rho) = \Pr\{v_i(y) > t(y) \mid \rho\} = \int \mathbb{I}_{v_i(y) > t(y)} dp(y|\rho)$$

where ρ level of differential expression

($\rho = 0$ meaning no difference, i.e. $z = 0$)

Incorporation in the design and possible calibration of loss

3. Simulation

Compute L and L^m by preposterior simulation

Simulation: Loop over repeated simulations

1. Prior: $(\omega^o, z^o) \sim p(\omega, z)$.
 2. Data: $y_{J_1} \sim p(y_{J_1} | \omega^o, z^o)$.
 3. Grid: $J = J_0, \dots, J_1$, let $y_J \subset y_{J_1}$
[requires only one simulation run]
- Posterior MCMC: Compute $v_i = \Pr(z_i = 1 | y_J)$.
[easy MCMC, starting with true ω^o]
 - Thresholds: Compute the cutoffs t_L^*
 - Save $\bar{FN}(t^*, y_J)$ and $\bar{FNR}(t^*, y_J)$.
 - Save (J, ρ_i^o, d_i) .

Simulation (ctd.)

Curve Fitting of Monte Carlo Experiments:

1. Expected loss $L^m(J), \bar{FN}_m$ and \bar{FNR}_m :

Curve through (J, \bar{FNR}) and (J, \bar{FN}) using $\sqrt{\log J/J}$

2. Power: Fit curve through (J, ρ_i^o, d_i) .

Optimal sample size:

Use $\hat{L}^m(J)$ and power curves for an informed choice.

II. Probability Model

[Newton et al. (01, J Comp Bio), Newton & Kendziorski (03)]

4. Gamma/Gamma hierarchical model

Observed gene expression:

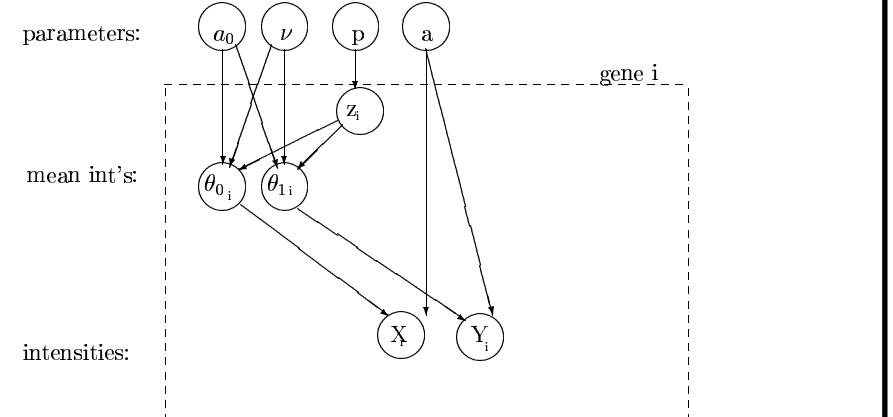
$$X_{ij} \sim Ga(a, \theta_{0i}) \text{ and } Y_{ij} \sim Ga(a, \theta_{1i}).$$

Mean expressions:

$$\begin{aligned} \theta_{0i} &\sim Ga(a_0, \nu) \\ \theta_{1i} &= \begin{cases} \theta_{0i} & \text{if } z_i = 0 \\ \sim Ga(a_0, \nu) & \text{if } z_i = 1 \end{cases} \end{aligned}$$

Differential expression:

$$\Pr(\theta_{0i} = \theta_{1i}) = \Pr(z_i = 0) = p.$$



Closed Form Expressions

We find

$$p(X_i, Y_i | z_i = 0, \eta) = \left\{ \frac{\Gamma(2Ja + a_0)}{\Gamma(a)^{2J} \Gamma(a_0)} \right\} \frac{(\nu)^{a_0} (\prod_j X_{ij} \prod_j Y_{ij})^{a-1}}{[(\sum_j X_i + \sum_j Y_i + \nu)]^{2a+a_0}}$$

and

$$p(X_i, Y_i | z_i = 1, \eta) = \left\{ \frac{\Gamma(aJ + a_0)}{\Gamma(a)^J \Gamma(a_0)} \right\}^2 \frac{(\nu\nu)^{a_0} (\prod_j X_{ij} \prod_j Y_{ij})^{a-1}}{[(\sum_j X_{ij} + \nu)(\sum_j Y_{ij} + \nu)]^{a+a_0}},$$

5. Mixture of Ga/Ga hierarchical model

- *Mixture of Gammas:*

$$\begin{aligned} X_{ij} &\sim \int \text{Ga}(a, \theta_{0i} r_{ij}) dp(r_{ij}|w) \\ &= \sum_{k=1}^K w_k \text{Ga}(a, \theta_{0i} r_{ij}^k) \end{aligned}$$

and

$$Y_{ij} \sim \int \text{Ga}(a, \theta_{0i} s_{ij}) dp(r_{ij}|w)$$

Addresses issues of noise in data collection and experimental conditions, based on a pilot run on control tissue

- Plus slide specific mixture:

$$(X_{ij} | r_{ij}, g_j) \sim \text{Ga}(a, \theta_{0i} g_j r_{ij})$$

and

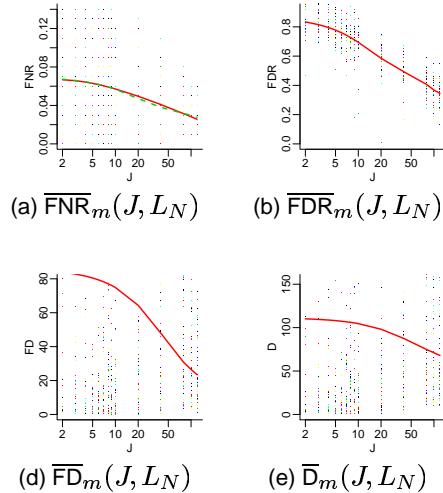
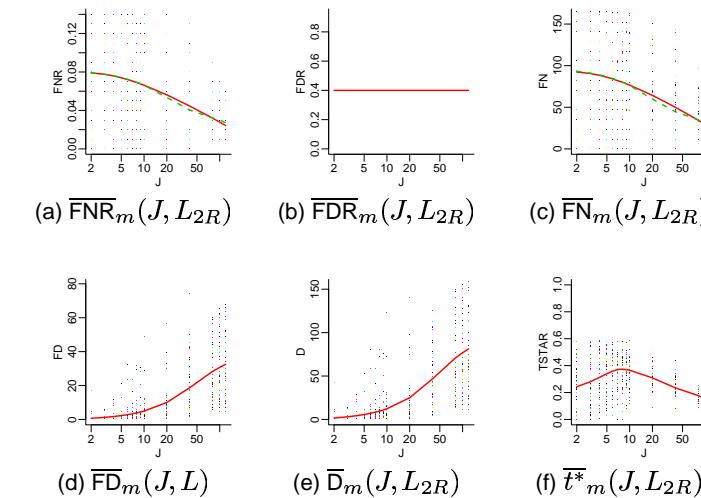
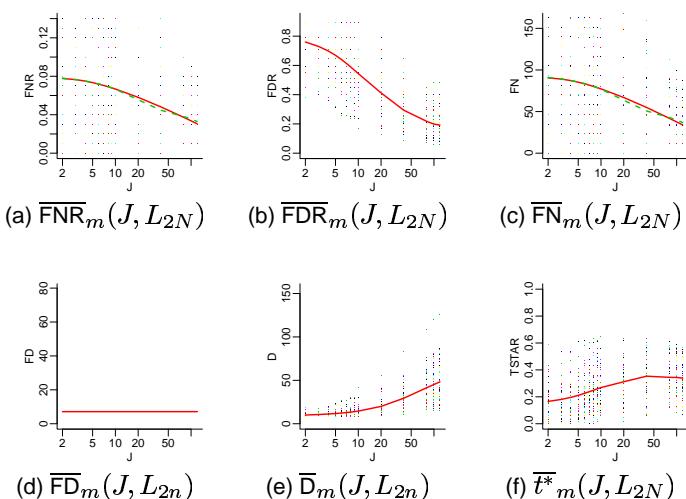
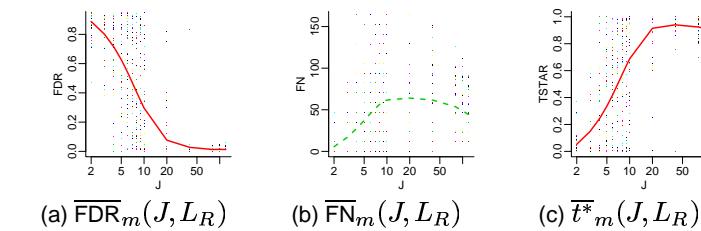
$$(Y_{ij} | s_{ij}, g_j) \sim \text{Ga}(a, \theta_{1i} g_j s_{ij}),$$

for outlier slides, again based on pilot data.

- MCMC: remains (almost) the same (add'l RJMCMC bits for # of components on both mixtures, found to be $K = 3$ and $L = 2$ on the dataset used in Newton et al., 2001)

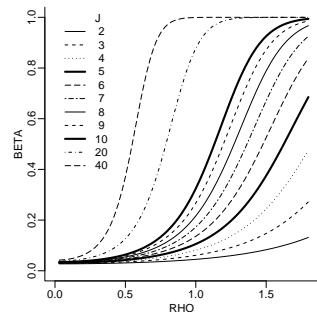
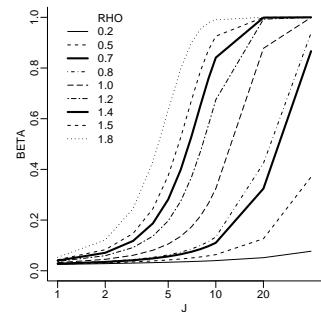
6. Results

Simulation from the estimated mixtures ($K = 3, L = 2$) rather than from the prior (use of pilot dataset)

Results L_N : t^*_N fixed by design**Results L_{2R} : \overline{FDR}_m fixed by design $\alpha = .4$** **Results L_{2N} : \overline{FD} fixed by design $\alpha_N = 0.1n\bar{p}\alpha/(1 - \alpha)$** **Results L_R : arrrrgh!!!**

Awkward jump in \overline{FDR}_m (and $\overline{t^*}_m$).

Power $\beta = \mathbb{E}_{y_J} \{\Pr(d = 1 | \rho, y_J)\}$ against $\rho = \log(\theta_{0i}/\theta_{1i})$ and J

(a) β against ρ (by J)(b) β against J (by ρ)

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- Two dec problems: sample size & multiple comparison
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