



Weierstrass Institute for
Applied Analysis and Stochastics



Simultaneous Statistical Inference

Thorsten Dickhaus

WIAS Berlin, Research Group “Stochastic Algorithms and Nonparametric Statistics”

- 1 Multiple comparisons (multiple tests)**
- 2 Simultaneous confidence regions**
- 3 Classification and Model selection**
- 4 Simultaneous inference group at WIAS**

We assume a statistical model (statistical experiment)

$$(\Omega, \mathcal{F}, (\mathbb{P}_\vartheta)_{\vartheta \in \Theta})$$

More concrete scenario (chosen for exemplary purposes):

Balanced ANOVA1 model:

$$X = (X_{ij})_{i=1, \dots, k, j=1, \dots, n}, X_{ij} = \mathcal{N}(\mu_i, \sigma^2),$$

X_{ij} stochastically independent random variables on \mathbb{R} ,

$\mu_i \in \mathbb{R} \forall 1 \leq i \leq k, \sigma^2 > 0$ (known or unknown) variance

$k \geq 3, n \geq 2, \nu = k(n - 1)$ (degrees of freedom)

$$\Omega = \mathbb{R}^{k \cdot n}, \mathcal{F} = \mathbb{B}^{k \cdot n}$$

$$\vartheta = (\mu_1, \dots, \mu_k, \sigma^2) \in \mathbb{R}^k \times (0, \infty) = \Theta$$

1 Multiple comparisons (multiple tests)

2 Simultaneous confidence regions

3 Classification and Model selection

4 Simultaneous inference group at WIAS

$$\mathcal{H}_m = (H_i)_{i=1,\dots,m}$$

Family of null hypotheses with $\emptyset \neq H_i \subset \Theta$
and alternatives $K_i = \Theta \setminus H_i$

$$(\Omega, \mathcal{F}, (\mathbb{P}_{\vartheta})_{\vartheta \in \Theta}, \mathcal{H}_m)$$

multiple test problem

$$\varphi = (\varphi_i : i = 1, \dots, m)$$

multiple test for \mathcal{H}_m

		Test decision		
		0	1	
Hypotheses	true	U_m	V_m	m_0
	false	T_m	S_m	m_1
		W_m	R_m	m

Wanted features of φ :

No (or only minor) contradictions among the individual test decisions

Control of the probability of erroneous decisions

Concepts of Contradiction: Let $I = \{1, \dots, m\}$.

- The multiple test φ is **coherent** if $\forall i, j \in I$ with $H_i \subseteq H_j$:
 $\{\varphi_j = 1\} \subseteq \{\varphi_i = 1\}$.
- The multiple test φ is **consonant** if $\forall i \in I$ with $\exists j \in I : H_i \subset H_j$:
 $\{\varphi_i = 1\} \subseteq \bigcup_{j: H_j \supset H_i} \{\varphi_j = 1\}$.

(Type I) error measures / concepts:

$$\text{FWER}_m(\varphi) = \mathbb{P}_{\vartheta}(V_m > 0) \stackrel{!}{\leq} \alpha \quad \forall \vartheta \in \Theta$$

(Strong) control of the **Family-Wise Error Rate (FWER)**

$$\text{FDR}_m(\varphi) = \mathbb{E}_{\vartheta} \left[\frac{V_m}{R_m \vee 1} \right] \stackrel{!}{\leq} \alpha \quad \forall \vartheta \in \Theta$$

Control of the **False Discovery Rate (FDR)**

FWER-controlling multiple testing principles:

- Historical single-step tests (Bonferroni, Šidák, Tukey, Scheffé, Dunnett)
- Closed test principle (Marcus, Peritz, Gabriel (1976))
- Intersection-union principle (generalized closed testing)
- Partitioning principle (Finner and Straßburger 2001, Hsu 1996)
- Projection methods under asymptotic normality (Bretz, Hothorn and Westfall)
- Multivariate methods based on copula modeling (Dickhaus et al. 2013/2014)
- Resampling-based multiple testing (Westfall & Young 1993, Dudoit & van der Laan 2008)

FDR-controlling multiple tests:

- Linear step-up (under positive dependency, Benjamini and Hochberg 1995)
- Adaptive LSU (BH 2000, Storey 2002, 2003, 2004, Benjamini, Krieger, Yekutieli 2006, Sarkar 2008, Blanchard and Roquain (2008, 2009), ...)
- Asymptotically optimal rejection curve
(under independence; Finner, Dickhaus, Roters 2009)
- Resampling-based multiple testing (Dudoit & van der Laan 2008)

Due to rapid technical developments in many scientific fields, the number m of hypotheses to be tested simultaneously can nowadays become **almost arbitrary large**:

- Genetics, microarrays: $m \sim 30,000$ genes / hypotheses
- Genetics, SNPs: $m \sim 500,000$ SNPs / hypotheses
- Proteomics: $m \sim 5000$ protein spots per gel sheet
- Cosmology: Signal detection, $m \sim 10^6$ pixels / hypotheses
- Neurology: Identification of active brain loci, $m \sim 10^3$ voxels
- Biometry: pairwise comparisons of many means, tests for correlations

Analyses have (in a first step) typically **explorative character**.

⇒ Control of the FWER much too conservative goal!

Θ	Parameter space
H_1, \dots, H_m	Null hypotheses
$\varphi = (\varphi_1, \dots, \varphi_m)$	Multiple test procedure
$V_m = \{i : \varphi_i = 1 \text{ and } H_i \text{ true }\} $	Number of falsely rejected, true nulls
$R_m = \{i : \varphi_i = 1\} $	Total number of rejections

$$\text{FDR}_\vartheta(\varphi) = \mathbb{E}_\vartheta \left[\frac{V_m}{R_m \vee 1} \right]$$

False Discovery Rate (FDR) given $\vartheta \in \Theta$

Definition: Let $\alpha \in (0, 1)$ fixed.

The multiple test φ controls the FDR at level α if

$$\text{FDR}(\varphi) = \sup_{\vartheta \in \Theta} \text{FDR}_\vartheta(\varphi) \leq \alpha.$$

p_1, \dots, p_m (marginal) p -values for hypotheses H_1, \dots, H_m

H_i true for $i \in I_{m,0}$, H_i false for $i \in I_{m,1}$

$$I_{m,0} + I_{m,1} = \mathbb{N}_m = \{1, \dots, m\}, m_0 = |I_{m,0}|$$

$p_i \sim \text{UNI}[0, 1], i \in I_{m,0}$ stochastically independent (I1)

$(p_i : i \in I_{m,0}), (p_i : i \in I_{m,1})$ stoch. independent vectors (I2)

$p_{1:m} \leq \dots \leq p_{m:m}$ ordered p -values

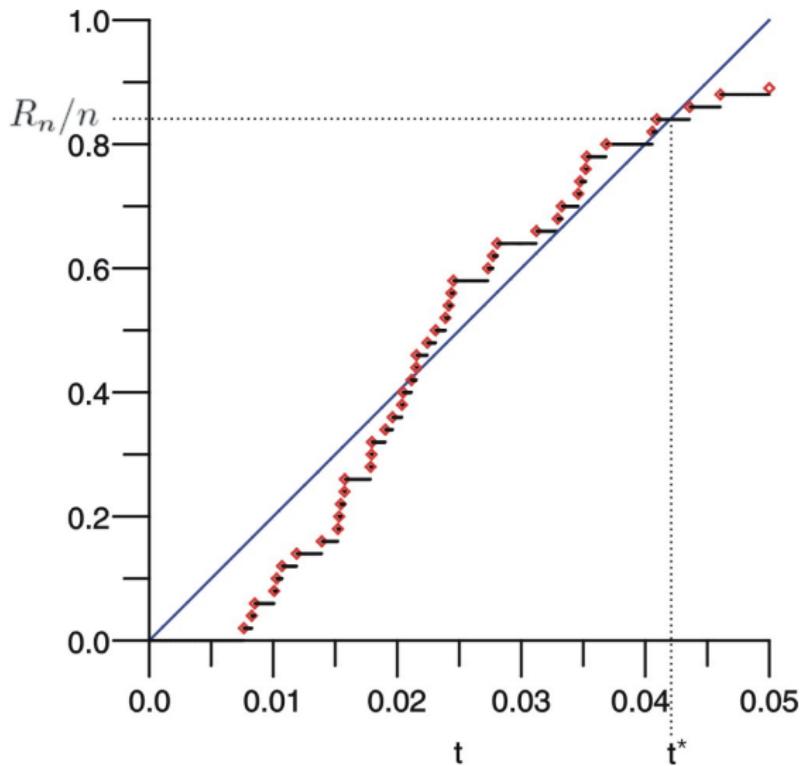
Linear step-up procedure φ^{LSU} with Simes' crit. values $\alpha_{i:m} = i\alpha/m$:

Reject all H_i with $p_i \leq \alpha_{\overline{m}:m}$, where $\overline{m} = \max\{j : p_{j:m} \leq j\alpha/m\}$.

Then it holds:

$$\forall \vartheta \in \Theta : \text{FDR}_\vartheta(\varphi^{\text{LSU}}) = \mathbb{E}_\vartheta \left[\frac{V_m}{R_m \vee 1} \right] = \frac{m_0}{m} \alpha.$$

Google Scholar, 22.04.2014, 11am: Paper cited 21,614 times!



Benjamini, Y. & Yekutieli, D. (2001) / Sarkar, S. K. (2002):

Proofs for FDR control in presence of special dependency structures:

$$\forall \vartheta \in \Theta : \text{FDR}_\vartheta(\varphi) \leq \frac{m_0}{m} \alpha.$$

for stepwise test procedures employing Simes' critical values.

Model assumptions: **MTP₂ oder PRDS**

Examples:

Multivariate normal distributions with non-negative correlations,
certain multivariate (absolute) *t*-distributions

(Finner, Dickhaus, Roters (2007), The Annals of Statistics **35**, 1432-1455)

Since under positive dependency the FDR of the LSU- procedure is bounded by

$$\frac{m_0}{m} \alpha$$

for any $m > 1$ and given $\alpha \in (0, 1)$, φ^{LSU} does **not exhaust** the FDR level α in case of $m_0 < m$.

It may be asked:

**Is it possible to derive a better rejection curve
circumventing the factor m_0/m appearing in
the FDR of φ^{LSU} ?**

Assumptions:

Independent p -values p_1, \dots, p_m ;

$m_0 = m_0(m)$ null hypotheses true with

$$\lim_{m \rightarrow \infty} \frac{m_0(m)}{m} = \zeta \in (0, 1].$$

m_0 of the p -values UNI[0, 1]-distributed (corresp. hypotheses true)

$m_1 = m - m_0$ of the p -values δ_0 -distributed (corresp. hypotheses false)

Then the ecdf of the p -values F_m (say) converges (Glivenko-Cantelli) for $m \rightarrow \infty$ to

$$G_\zeta(x) = (1 - \zeta) + \zeta x \text{ for all } x \in [0, 1].$$

Assume we reject all H_i with $p_i \leq x$ for some $x \in (0, 1)$.

Then the FDR (depending on ζ and x) under $\text{DU}(\zeta)$ is asymptotically given by

$$\text{FDR}_\zeta(x) = \frac{\zeta x}{(1 - \zeta) + \zeta x}.$$

Aim: Find an optimal threshold x_ζ (say), such that

$$\text{FDR}_\zeta(x_\zeta) \equiv \alpha \text{ for all } \zeta \in (\alpha, 1).$$

We obtain:

$$\text{FDR}_\zeta(x_\zeta) = \alpha \iff x_\zeta = \frac{\alpha(1 - \zeta)}{\zeta(1 - \alpha)}.$$

Ansatz: Rejection curve f_α and G_ζ shall cross each other in x_ζ , i.e.,

$$f_\alpha(x_\zeta) = G_\zeta(x_\zeta).$$

Plugging in x_ζ derived above yields

$$f_\alpha \left(\frac{\alpha(1-\zeta)}{\zeta(1-\alpha)} \right) = \frac{1-\zeta}{1-\alpha}.$$

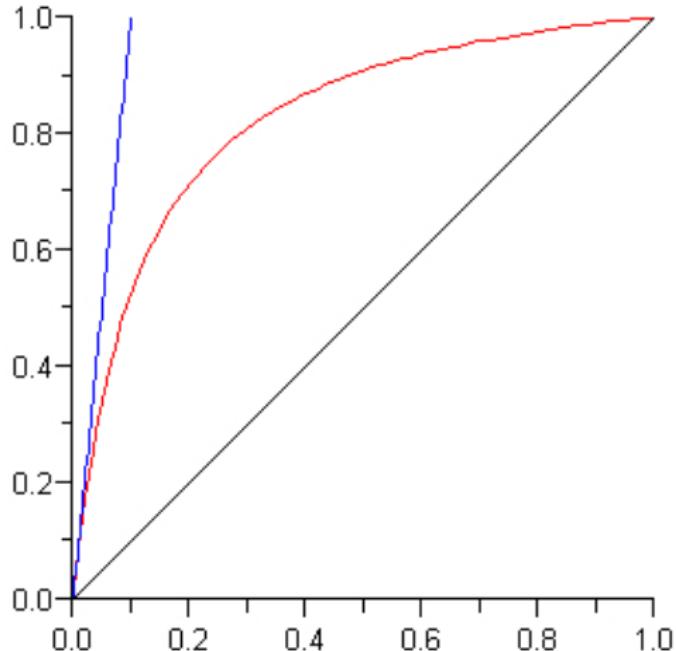
Substituting $t = \frac{\alpha(1-\zeta)}{\zeta(1-\alpha)}$ $\iff \zeta = \frac{\alpha}{(1-\alpha)t + \alpha}$,

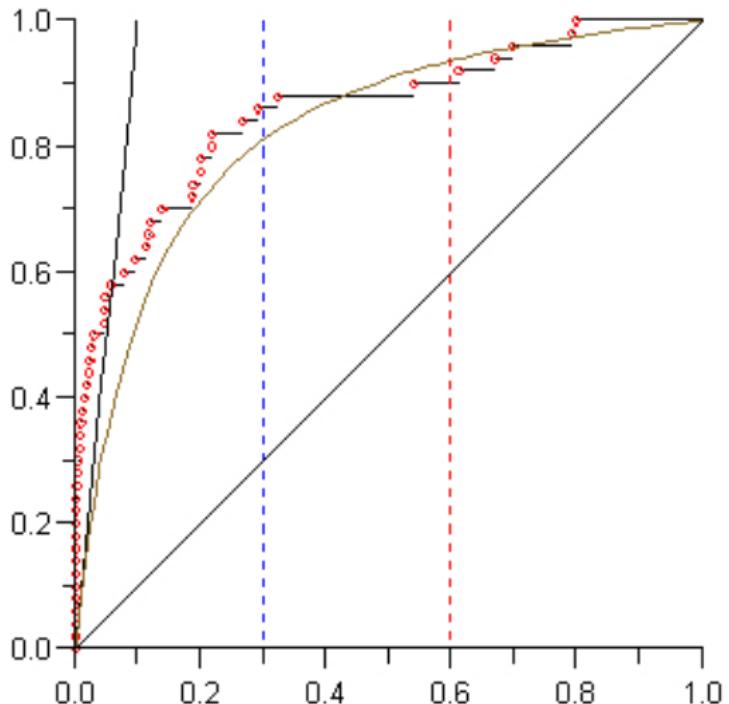
we get that

$$f_\alpha(t) = \frac{t}{(1-\alpha)t + \alpha}, \quad t \in [0, 1],$$

is the curve solving the problem!

AORC: Finner, Dickhaus, Roters (2009), Annals of Statistics 37, 596-618





1 Multiple comparisons (multiple tests)

2 Simultaneous confidence regions

3 Classification and Model selection

4 Simultaneous inference group at WIAS

Consider the ANOVA1 model from above.

Define $\theta_{ij} = \mu_i - \mu_j, 1 \leq i < j \leq k$ (pairwise mean differences).

A simultaneous confidence region $C_{ij}(X)$ is a subset of \mathbb{R}
with the property ($\alpha \in (0, 1)$ fixed) that

$$\forall \mu \in \mathbb{R}^k : \forall \sigma^2 > 0 : \mathbb{P}_{\mu, \sigma^2}(C_{ij}(X) \ni \theta_{ij} \forall 1 \leq i < j \leq k) \geq 1 - \alpha. \quad (1)$$

Duality of multiple tests and simultaneous confidence regions:

Tukey's test \iff Tukey confidence intervals,

Scheffé's simultaneous coverage thm. \iff Scheffé test, etc.

Remark. Equation (1) is an analog of the FWER.

Let $\bar{Y}_i = \bar{X}_i - \mu_i \Rightarrow \bar{X}_i - \bar{X}_j - \theta_{ij} = \bar{Y}_i - \bar{Y}_j, i = 1, \dots, k$. Setting

$$T_{ij}(X) = \sqrt{n} \frac{|\bar{Y}_i - \bar{Y}_j|}{S}, 1 \leq i < j \leq k,$$

the statistic $\max_{1 \leq i < j \leq k} T_{ij}(X)$ has a known distribution, namely, the

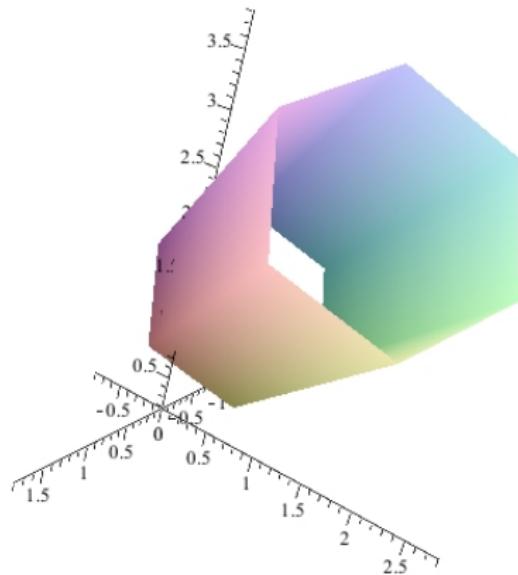
Studentized range distribution $q_{k,k(n-1)}$ with k groups and $k(n-1)$ degrees of freedom.

The simultaneous multiple test based on the $T_{ij}(X)$ and the quantiles of $q_{k,k(n-1)}$ is due to Tukey (1953).

Noticing that

$$\sqrt{n} \frac{|\bar{y}_i - \bar{y}_j|}{s} \leq q_{k,k(n-1);\alpha} \iff \theta_{ij} \in \left[(\bar{x}_i - \bar{x}_j) \pm \frac{s}{\sqrt{n}} q_{k,k(n-1);\alpha} \right],$$

we obtain **simultaneous confidence intervals** for the pairwise mean differences and, consequently, a **simultaneous confidence region** for the mean vector in \mathbb{R}^k !



(Maple code taken from Jason C. Hsu, 1996)

1 Multiple comparisons (multiple tests)

2 Simultaneous confidence regions

3 Classification and Model selection

4 Simultaneous inference group at WIAS

- Abramovich et al. (2006), Bogdan et al. (2011), Neuvial and Roquain (2012):
FDR-thresholding (with φ^{LSU}) asymptotically achieves Bayes-optimal classification risk under sparsity
- Storey (2003), Dickhaus et al. (2013):
Weighted average of pFDR and pFNR is a proxy for cost-weighted Bayes classification risk in non-sparse models
- Donoho and Jin (2008):
Higher criticism thresholding leads to optimal feature selection for classification in rare/weak models

- Bauer et al. (1988), Bunea et al. (2006), Furmańczyk (2013):
Model selection consistency of multiple test procedures
- Zuber and Strimmer (2011):
Optimization of information criteria AIC, BIC, RIC, ... is equivalent to single-step multiple testing
- Abramovich et al. (2006), Benjamini and Gavrilov (2009):
Model selection performance of stepwise rejective multiple tests
- Berk et al. (2013):
Simultaneous inference implies valid post-selection inference
- Qiu and Hwang (2007), Benjamini and Yekutieli (2005), ...
Valid inference for selected parameters based on FWER / FDR control

-
- 1 Multiple comparisons (multiple tests)**
 - 2 Simultaneous confidence regions**
 - 3 Classification and Model selection**
 - 4 Simultaneous inference group at WIAS**

- **Jens Stange**, Nina Loginova:

Multiple testing for behavioural genetics (DFG grant DI 1723/3-1)

- **Konstantin Schildknecht**, Tina Baumann:

Multiple testing for epigenetic methylation data
(BMBF project EPILYZE with Epiontis GmbH)

- Simultaneous statistical inference in multivariate time series models

(Project A3 in IRTG 1792 "High Dimensional Non Stationary Time Series")

Still at Humboldt-University Berlin:

- **Taras Bodnar**, Yuriy Kopanskyy:

Multiple testing under unspecified dependency structure
(DFG Research Unit FOR 1735)

