

A Maximum Wilcoxon-Mann-Whitney Test In High Dimensions

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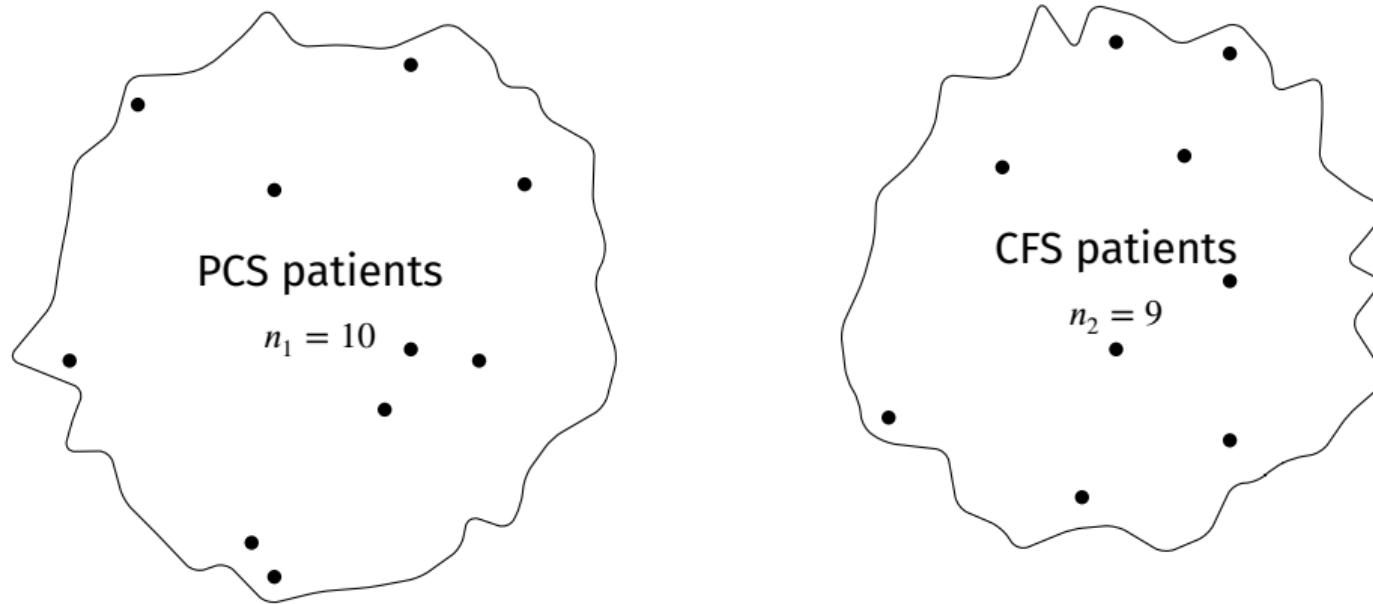
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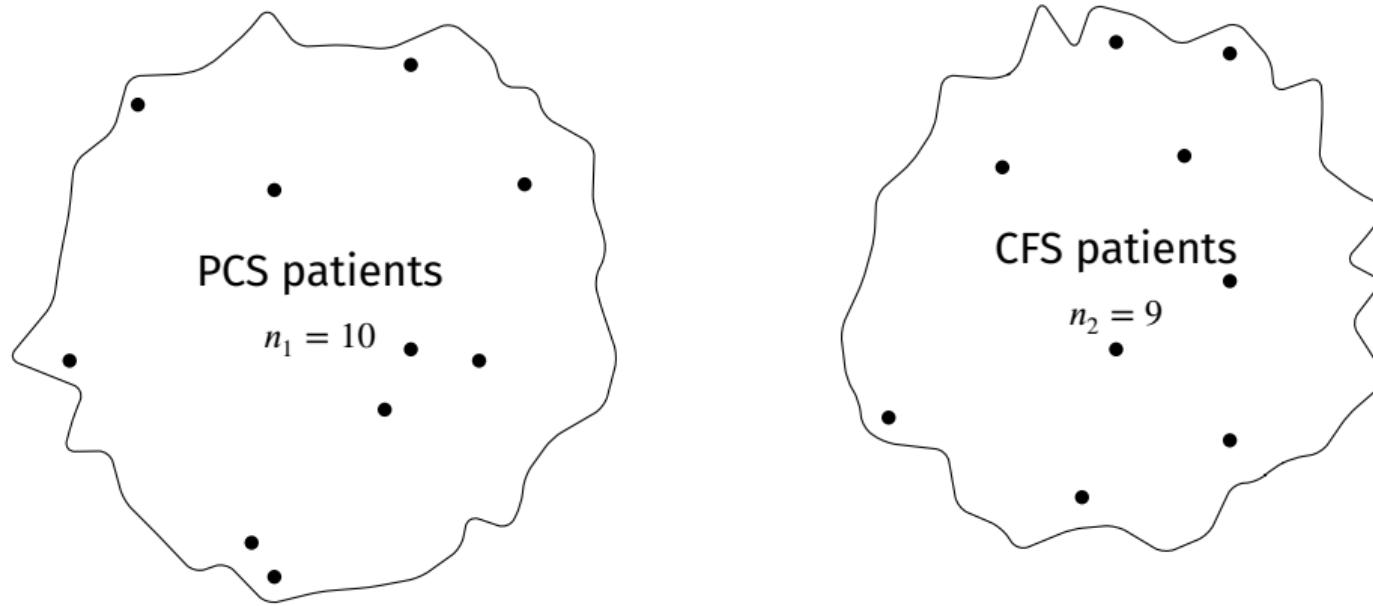
Example:

Post Covid Syndrome (PCS) and Chronique Fatigue Syndrome (CFS)

SETTING



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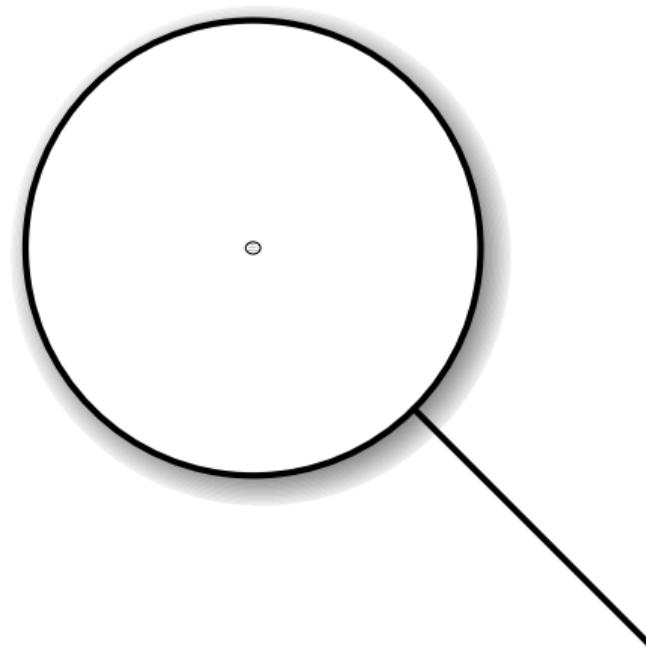


→ Goal: compare both groups

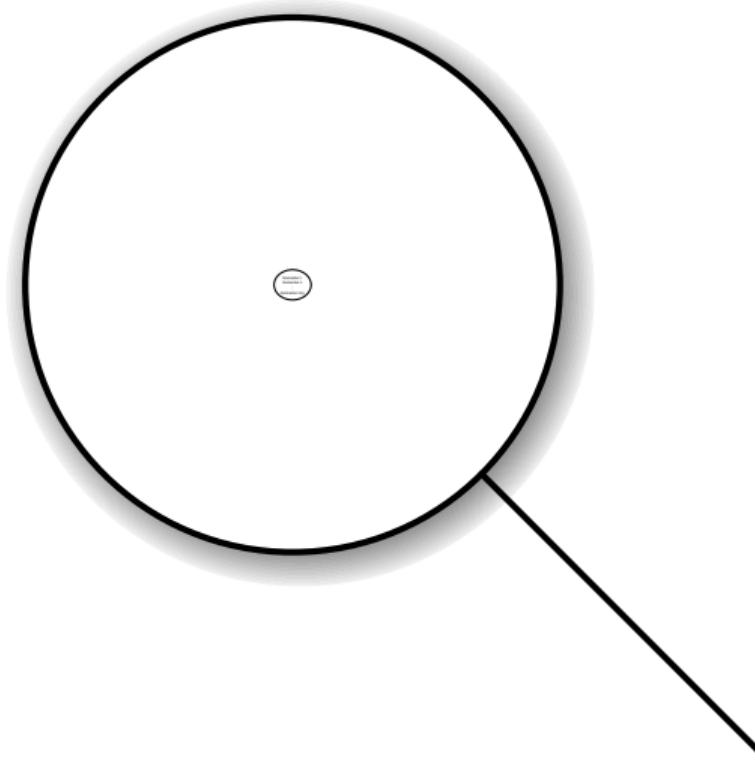
A CLOSER LOOK



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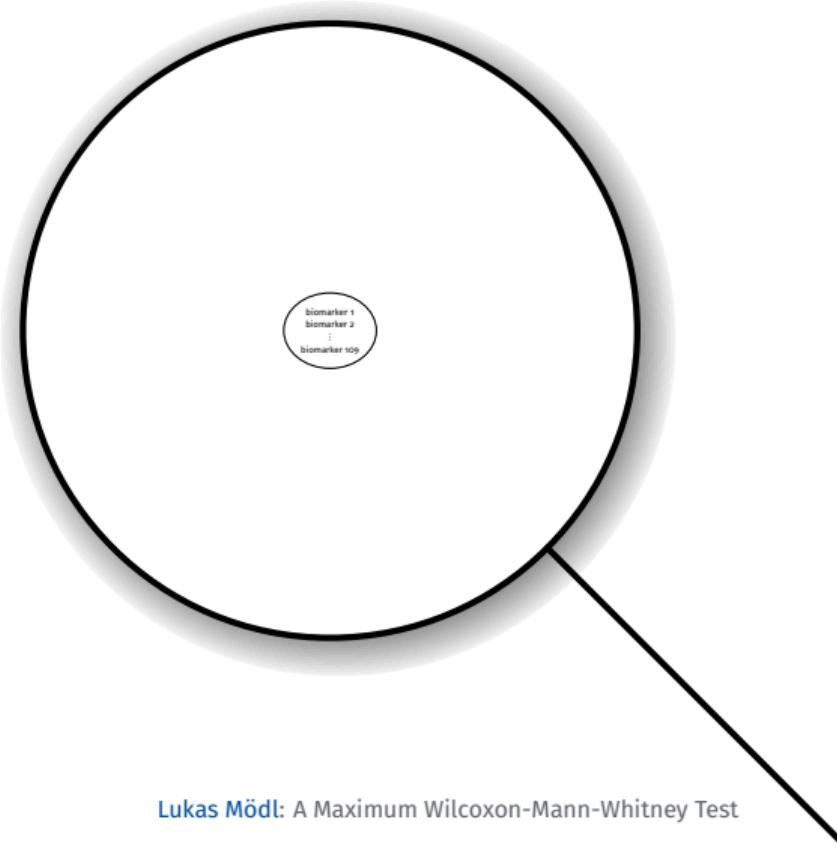


A CLOSER LOOK

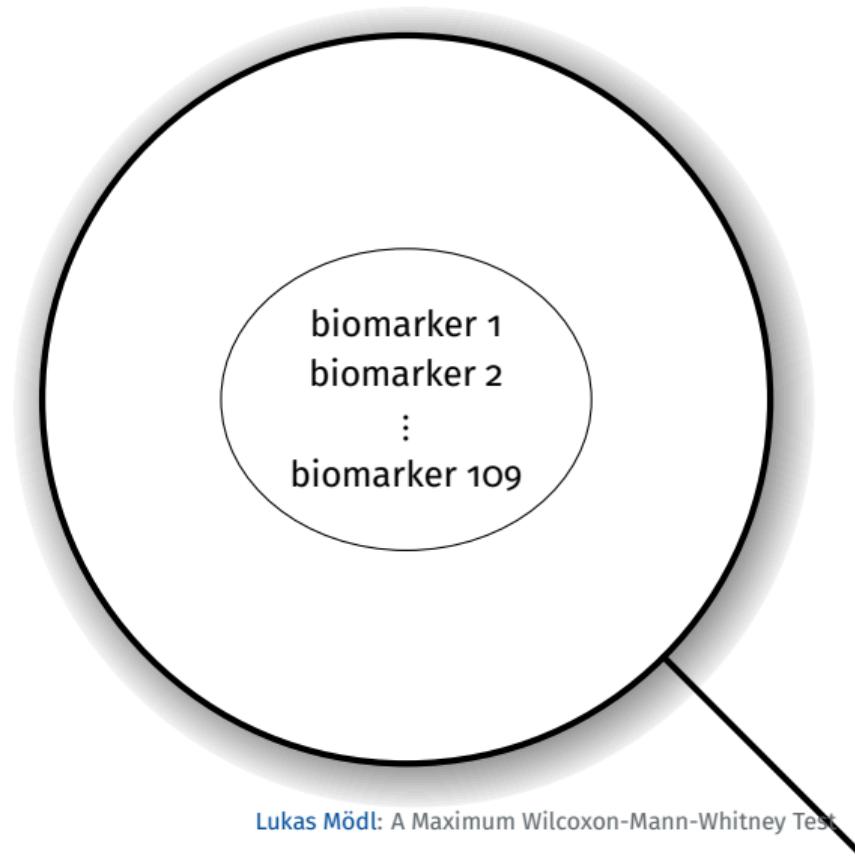


Lukas Mödl: A Maximum Wilcoxon-Mann-Whitney Test

A CLOSER LOOK



A CLOSER LOOK



COMPARING THE GROUPS

Test the global hypothesis

distribution of biomarker 1 in PCS patients = distribution of biomarker 1 in CFS patients

distribution of biomarker 2 in PCS patients = distribution of biomarker 2 in CFS patients

⋮

distribution of biomarker 109 in PCS patients = distribution of biomarker 109 in CFS patients

at a type 1 error rate α

How to test this hypothesis?

COMMON APPROACHES

- Bonferroni-type test
 - Run 109 separate Wilcoxon-Mann-Whitney tests with adjusted type 1 error rate α
 - Inference is based on the univariate standard normal distribution

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- ANOVA-type test
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 - Inference is based on the centered 109-variate normal distribution with the (scaled) empirical correlation matrix of ranks as variance-covariance matrix

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- MCTP-type test
 - Run 109 separate Wilcoxon-Mann-Whitney tests and take the ℓ^∞ -norm
 - Inference is based on the centered 109-variate normal distribution with the empirical correlation matrix of ranks as variance-covariance matrix

TYPE 1 ERROR RATE CONTROL

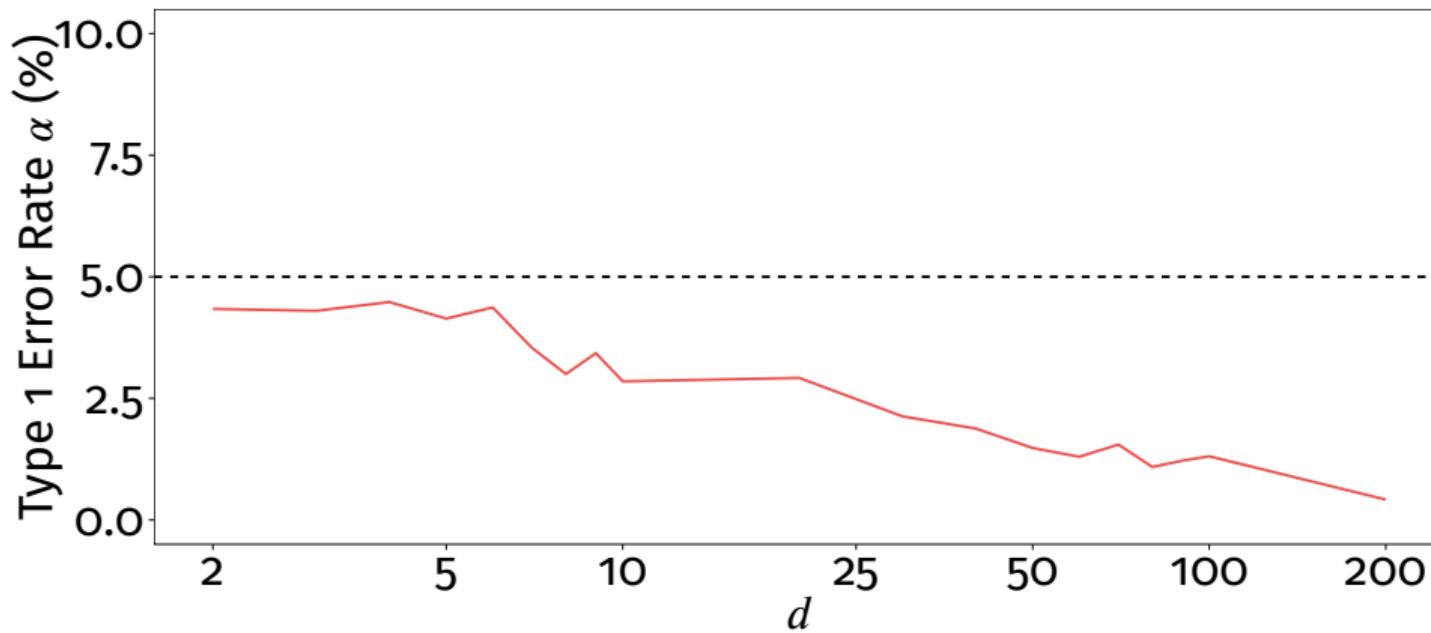


Fig. 1: Simulated data from multivariate normal distribution with $n_1 = 10$, $n_2 = 9$; 10,000 runs for each d

Can we do something about it?

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~~> Wild Bootstrap

STEP 1

Data matrix:

X_{111}	X_{121}	...	X_{1d1}
X_{112}	X_{122}	...	X_{1d2}
\vdots	\vdots	\ddots	\vdots
X_{11n_1}	X_{12n_1}	...	X_{1dn_1}
X_{211}	X_{221}	...	X_{2d1}
X_{212}	X_{222}	...	X_{2d2}
\vdots	\vdots	\ddots	\vdots
X_{21n_2}	X_{22n_2}	...	X_{2dn_2}

STEP 1

Assign ranks:

R_{111}	R_{121}	...	R_{1d1}
R_{112}	R_{122}	...	R_{1d2}
:	:	:	:
R_{11n_1}	R_{12n_1}	...	R_{1dn_1}
R_{211}	R_{221}	...	R_{2d1}
R_{212}	R_{222}	...	R_{2d2}
:	:	:	:
R_{21n_2}	R_{22n_2}	...	R_{2dn_2}

STEP 2

Center ranks by rank mean $c := \frac{n_1+n_2+1}{2}$:

$(R_{111} - c)$	$(R_{121} - c)$	\dots	$(R_{1d1} - c)$
$(R_{112} - c)$	$(R_{122} - c)$	\dots	$(R_{1d2} - c)$
\vdots	\vdots	\ddots	\vdots
$(R_{11n_1} - c)$	$(R_{12n_1} - c)$	\dots	$(R_{1dn_1} - c)$
$(R_{211} - c)$	$(R_{221} - c)$	\dots	$(R_{2d1} - c)$
$(R_{212} - c)$	$(R_{222} - c)$	\dots	$(R_{2d2} - c)$
\vdots	\vdots	\ddots	\vdots
$(R_{21n_2} - c)$	$(R_{22n_2} - c)$	\dots	$(R_{2dn_2} - c)$

STEP 3

Perturbate centered ranks of each row by a random sign W_{ik} (+ or -):

$W_{11}(R_{111} - c)$	$W_{11}(R_{121} - c)$...	$W_{11}(R_{1d1} - c)$
$W_{12}(R_{112} - c)$	$W_{12}(R_{122} - c)$...	$W_{12}(R_{1d2} - c)$
⋮	⋮	⋮	⋮
$W_{1n_1}(R_{11n_1} - c)$	$W_{1n_1}(R_{12n_1} - c)$...	$W_{1n_1}(R_{1dn_1} - c)$
$W_{21}(R_{211} - c)$	$W_{21}(R_{221} - c)$...	$W_{21}(R_{2d1} - c)$
$W_{22}(R_{212} - c)$	$W_{22}(R_{222} - c)$...	$W_{22}(R_{2d2} - c)$
⋮	⋮	⋮	⋮
$W_{2n_2}(R_{21n_2} - c)$	$W_{2n_2}(R_{22n_2} - c)$...	$W_{2n_2}(R_{2dn_2} - c)$

STEP 4

Compute column means for both groups:

$W_{11}(R_{111} - c)$	$W_{11}(R_{121} - c)$...	$W_{11}(R_{1d1} - c)$
$W_{12}(R_{112} - c)$	$W_{12}(R_{122} - c)$...	$W_{12}(R_{1d2} - c)$
⋮	⋮	⋮	⋮
$W_{1n_1}(R_{11n_1} - c)$	$W_{1n_1}(R_{12n_1} - c)$...	$W_{1n_1}(R_{1dn_1} - c)$
$W_{21}(R_{211} - c)$	$W_{21}(R_{221} - c)$...	$W_{21}(R_{2d1} - c)$
$W_{22}(R_{212} - c)$	$W_{22}(R_{222} - c)$...	$W_{22}(R_{2d2} - c)$
⋮	⋮	⋮	⋮
$W_{2n_2}(R_{21n_2} - c)$	$W_{2n_2}(R_{22n_2} - c)$...	$W_{2n_2}(R_{2dn_2} - c)$

STEP 4

Compute column means for both groups:

$$W_{11}(R_{111} - c)$$

$$W_{11}R_{112} - c)$$

$$\underline{M}_{11}$$

$$W_{1n_1}(R_{11n_1} - c)$$


$$W_{21}(R_{211} - c)$$

$$W_{21}R_{212} - c)$$

$$\underline{M}_{21}$$

$$W_{2n_2}(R_{21n_2} - c)$$


$$W_{11}(R_{121} - c)$$

$$W_{11}R_{122} - c)$$

$$\underline{M}_{12}$$

$$W_{1n_1}(R_{12n_1} - c)$$


$$W_{21}(R_{221} - c)$$

$$W_{21}R_{222} - c)$$

$$\underline{M}_{22}$$

$$W_{2n_2}(R_{22n_2} - c)$$


$$\dots$$

$$\dots$$

$$\ddots$$

$$\dots$$

$$\dots$$

$$\ddots$$

$$\dots$$

$$W_{1d}(R_{1d1} - c)$$

$$W_{1d}R_{1d2} - c)$$

$$\underline{M}_{1d}$$

$$W_{1n_1}(R_{1dn_1} - c)$$


$$W_{21}(R_{2d1} - c)$$

$$W_{21}R_{2d2} - c)$$

$$\underline{M}_{2d}$$

$$W_{2n_2}(R_{2dn_2} - c)$$


STEP 5

Compute standard deviations column-wise:

$W_{11}(R_{111} - c)$	$W_{11}(R_{121} - c)$...	$W_{11}(R_{1d1} - c)$
$W_{12}(R_{112} - c)$	$W_{12}(R_{122} - c)$...	$W_{12}(R_{1d2} - c)$
⋮	⋮	⋮	⋮
$W_{1n_1}(R_{11n_1} - c)$	$W_{1n_1}(R_{12n_1} - c)$...	$W_{1n_1}(R_{1dn_1} - c)$
$W_{21}(R_{211} - c)$	$W_{21}(R_{221} - c)$...	$W_{21}(R_{2d1} - c)$
$W_{22}(R_{212} - c)$	$W_{22}(R_{222} - c)$...	$W_{22}(R_{2d2} - c)$
⋮	⋮	⋮	⋮
$W_{2n_2}(R_{21n_2} - c)$	$W_{2n_2}(R_{22n_2} - c)$...	$W_{2n_2}(R_{2dn_2} - c)$

STEP 5

Compute standard deviations column-wise:

$$W_{11}(R_{111} - c)$$

$$W_{11}(R_{112} - c)$$

 \vdots

$$W_{11} S_1(R_{11n_1} - c)$$

$$W_{12} S_1(R_{211} - c)$$

$$W_{12} S_1(R_{212} - c)$$

 \vdots

$$W_{2n_2} S_1(R_{21n_2} - c)$$

$$W_{11}(R_{121} - c)$$

$$W_{11}(R_{122} - c)$$

 \vdots

$$W_{11} S_2(R_{12n_1} - c)$$

$$W_{12} S_2(R_{221} - c)$$

$$W_{12} S_2(R_{222} - c)$$

 \vdots

$$W_{2n_2} S_2(R_{22n_2} - c)$$

 \dots
 \dots
 \ddots
 \dots
 \dots
 \ddots
 \dots

$$W_{11}(R_{1d1} - c)$$

$$W_{11}(R_{1d2} - c)$$

 \vdots

$$W_{11} S_d(R_{1dn_1} - c)$$

$$W_{12} S_d(R_{2d1} - c)$$

$$W_{12} S_d(R_{2d2} - c)$$

 \vdots

$$W_{2n_2} S_d(R_{2dn_2} - c)$$

STEP 6

Repeat steps 3 to 5 B times and obtain the empirical p-value by the proportion of

$$T_0 < T_0^*,$$

where

- $T_0 := \max \{ |T_1|, |T_2|, \dots, |T_d| \}$ with T_j being the usual Wilcoxon-Mann-Whitney test statistic for testing the local hypothesis

distribution of outcome j in group 1 = distribution of outcome j in group 2

- $T_0^* := \max \{ |T_1^*|, |T_2^*|, \dots, |T_d^*| \}$ with

$$T_j^* := \sqrt{\frac{n_1 n_2}{n_1 + n_2}} \frac{M_{1j} - M_{2j}}{S_j}$$

TYPE 1 ERROR RATE CONTROL REVISITED

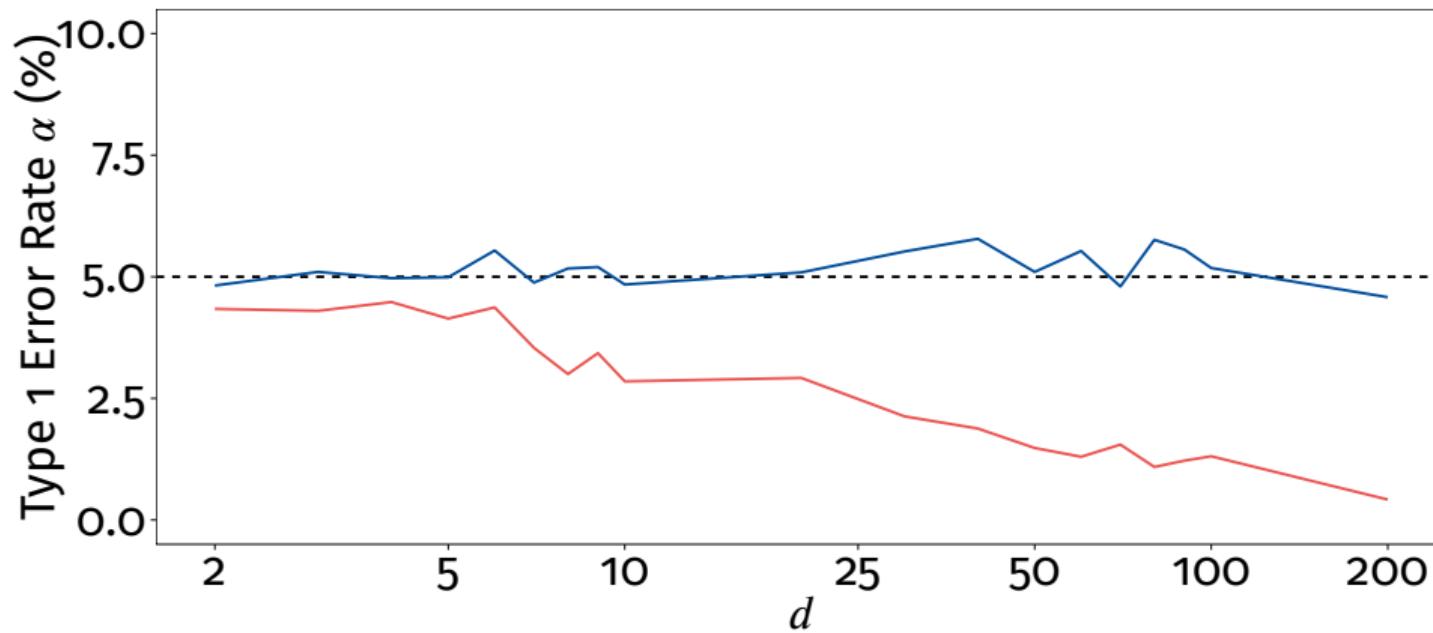


Fig. 2: Same setting as for Fig. 1; $b = 10,000$ were used for the wild bootstrap approach

Conclusion

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Summary:

- The wild bootstrap approach controls type 1 error rate α quite well
→ even in high dimensions

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- The wild bootstrap approach controls type 1 error rate α quite well
→ even in high dimensions

Outlook:

- Extend to multiple groups
- Consider other tests (e.g., Brunner-Munzel test)
- Consider other norms
- ...

Discussion

References

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-  Brunner E, Bathke A, Konietzschke F. *Rank and Pseudo-Rank Procedures for Independent Observations in Factorial Designs: Using R and SAS*. Springer International Publishing. 2019.
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Appendix

APPENDIX A

Theorem 1 – Consistency of Wild Bootstrap Approach

Let Q_n denote the joint distribution of (T_1, T_2, \dots, T_d) , Q_n^* the joint distribution of $(T_1^*, T_2^*, \dots, T_d^*)$ given the data, and suppose that $\frac{n_1+n_2}{n_i} \rightarrow \lambda_i$ with $0 < \lambda_i < 1$ for $i = 1, 2$ as $n_1 \wedge n_2 \rightarrow \infty$. Then

$$\varrho(Q_n, Q_n^*) \rightarrow 0$$

in probability as $n_1 \wedge n_2 \rightarrow \infty$, where ϱ denotes the Lévy-Prokhorov metric on the space of probability measures defined on the usual Borel σ -algebra over \mathbb{R}^d .