

A Primer on Permutation Tests (not only) for MVPA

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(with slides by Carsten Allefeld)

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introduction

why use a permutation test?

- ▶ sometimes precise distributions are **not known**, especially in MVPA
- ▶ a permutation test makes **weaker assumptions** about distributions than parametric tests
- ▶ permutation tests provide **exact inference**
- ▶ permutation testing applies **in the same way** to univariate and multivariate analysis

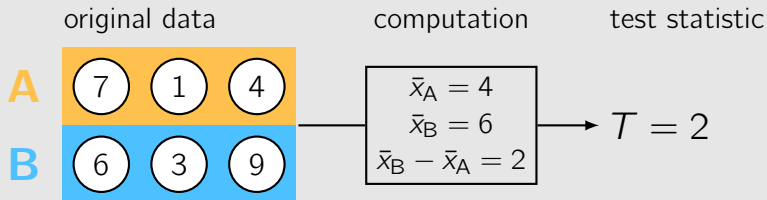
this talk

- ▶ no recipes, but the **basic principles** underlying permutation tests, especially exchangeability

A simple example

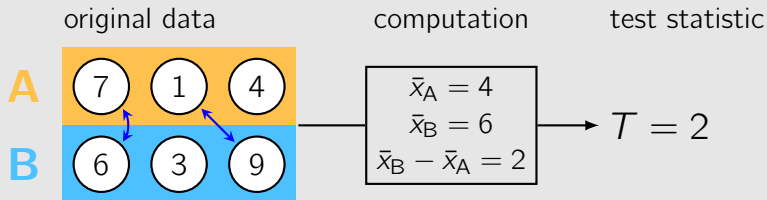
a univariate example: two-sample test

is there a mean difference between groups A and B?



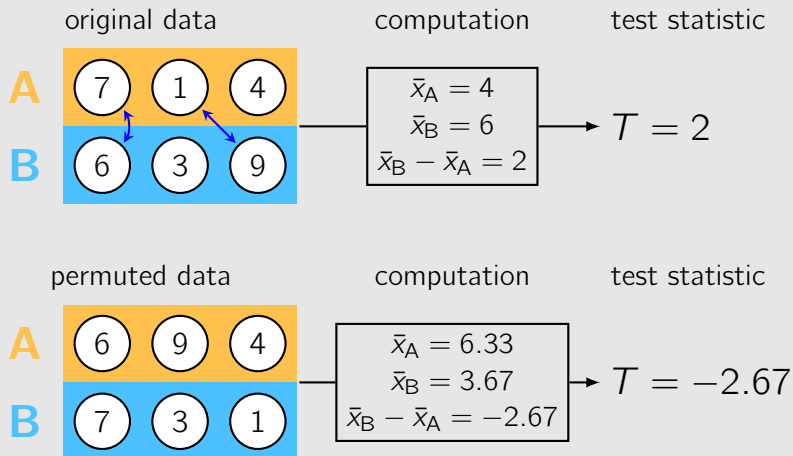
two-sample test

if not, values can be **exchanged** between conditions



two-sample test

the same test statistic is computed from **permuted data**



two-sample test

compute the test statistic for **all permutations**

perm	data		$T = \bar{x}_B - \bar{x}_A$
1	7 1 4	6 3 9	2.00
2	7 1 6	4 3 9	0.67
3	7 1 3	4 6 9	2.67
4	7 1 9	4 6 3	-1.33
5	7 4 6	1 3 9	-1.33
6	7 4 3	1 6 9	0.67
7	7 4 9	1 6 3	-3.33
8	7 6 3	1 4 9	-0.67
9	7 6 9	1 4 3	-4.67
10	7 3 9	1 4 6	-2.67
11	1 4 6	7 3 9	2.67
12	1 4 3	7 6 9	4.67
13	1 4 9	7 6 3	0.67
14	1 6 3	7 4 9	3.33
15	1 6 9	7 4 3	-0.67
16	1 3 9	7 4 6	1.33
17	4 6 3	7 1 9	1.33
18	4 6 9	7 1 3	-2.67
19	4 3 9	7 1 6	-0.67
20	6 3 9	7 1 4	-2.00

two-sample test

determine **ranks** by sorting in descending order

perm	data	$T = \bar{x}_B - \bar{x}_A$	sorted	rank	
1	7 1 4	6 3 9	2.00	4.67	1
2	7 1 6	4 3 9	0.67	3.33	2
3	7 1 3	4 6 9	2.67	2.67	3
4	7 1 9	4 6 3	-1.33	2.67	4
5	7 4 6	1 3 9	-1.33	2.00	5
6	7 4 3	1 6 9	0.67	1.33	6
7	7 4 9	1 6 3	-3.33	1.33	7
8	7 6 3	1 4 9	-0.67	0.67	8
9	7 6 9	1 4 3	-4.67	0.67	9
10	7 3 9	1 4 6	-2.67	0.67	10
11	1 4 6	7 3 9	2.67	-0.67	11
12	1 4 3	7 6 9	4.67	-0.67	12
13	1 4 9	7 6 3	0.67	-0.67	13
14	1 6 3	7 4 9	3.33	-1.33	14
15	1 6 9	7 4 3	-0.67	-1.33	15
16	1 3 9	7 4 6	1.33	-2.00	16
17	4 6 3	7 1 9	1.33	-2.67	17
18	4 6 9	7 1 3	-2.67	-2.67	18
19	4 3 9	7 1 6	-0.67	-3.33	19
20	6 3 9	7 1 4	-2.00	-4.67	20

two-sample test

the **neutral permutation** is part of the set

perm	data	$T = \bar{x}_B - \bar{x}_A$	sorted	rank
1	7 1 4	6 3 9	2.00	
2	7 1 6	4 3 9	0.67	
3	7 1 3	4 6 9	2.67	
4	7 1 9	4 6 3	-1.33	
5	7 4 6	1 3 9	-1.33	
6	7 4 3	1 6 9	0.67	
7	7 4 9	1 6 3	-3.33	
8	7 6 3	1 4 9	-0.67	
9	7 6 9	1 4 3	-4.67	
10	7 3 9	1 4 6	-2.67	
11	1 4 6	7 3 9	2.67	
12	1 4 3	7 6 9	4.67	
13	1 4 9	7 6 3	0.67	
14	1 6 3	7 4 9	3.33	
15	1 6 9	7 4 3	-0.67	
16	1 3 9	7 4 6	1.33	
17	4 6 3	7 1 9	1.33	
18	4 6 9	7 1 3	-2.67	
19	4 3 9	7 1 6	-0.67	
20	6 3 9	7 1 4	-2.00	

Diagram illustrating the mapping of sorted values to ranks. The sorted values are: 4.67, 3.33, 2.67, 2.67, 2.00, 1.33, 1.33, 0.67, 0.67, 0.67, -0.67, -0.67, -0.67, -1.33, -1.33, -2.00, -2.67, -2.67, -3.33, -4.67. The ranks are: 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20. The value 2.00 is highlighted with a box and a rank of 5.

two-sample test

the rank of the original value determines **significance**

perm	data	$T = \bar{x}_B - \bar{x}_A$	sorted	rank	
1	7 1 4	6 3 9	2.00	4.67	1 significant
2	7 1 6	4 3 9	0.67	3.33	2
3	7 1 3	4 6 9	2.67	2.67	3
4	7 1 9	4 6 3	-1.33	2.67	4
5	7 4 6	1 3 9	-1.33	2.00	5
6	7 4 3	1 6 9	0.67	1.33	6
7	7 4 9	1 6 3	-3.33	1.33	7
8	7 6 3	1 4 9	-0.67	0.67	8
9	7 6 9	1 4 3	-4.67	0.67	9
10	7 3 9	1 4 6	-2.67	0.67	10
11	1 4 6	7 3 9	2.67	-0.67	11
12	1 4 3	7 6 9	4.67	-0.67	12
13	1 4 9	7 6 3	0.67	-0.67	13
14	1 6 3	7 4 9	3.33	-1.33	14
15	1 6 9	7 4 3	-0.67	-1.33	15
16	1 3 9	7 4 6	1.33	-2.00	16
17	4 6 3	7 1 9	1.33	-2.67	17
18	4 6 9	7 1 3	-2.67	-2.67	18
19	4 3 9	7 1 6	-0.67	-3.33	19
20	6 3 9	7 1 4	-2.00	-4.67	20

two-sample test

equivalently, a **p-value** can be computed

perm	data	$T = \bar{x}_B - \bar{x}_A$	sorted	rank		
1	7 1 4	6 3 9	2.00	4.67	1	significant
2	7 1 6	4 3 9	0.67	3.33	2	
3	7 1 3	4 6 9	2.67	2.67	3	
4	7 1 9	4 6 3	-1.33	2.67	4	
5	7 4 6	1 3 9	-1.33	2.00	5	$p = \frac{5}{20} = 0.25$
6	7 4 3	1 6 9	0.67	1.33	6	
7	7 4 9	1 6 3	-3.33	1.33	7	
8	7 6 3	1 4 9	-0.67	0.67	8	
9	7 6 9	1 4 3	-4.67	0.67	9	
10	7 3 9	1 4 6	-2.67	0.67	10	
11	1 4 6	7 3 9	2.67	-0.67	11	
12	1 4 3	7 6 9	4.67	-0.67	12	
13	1 4 9	7 6 3	0.67	-0.67	13	
14	1 6 3	7 4 9	3.33	-1.33	14	
15	1 6 9	7 4 3	-0.67	-1.33	15	
16	1 3 9	7 4 6	1.33	-2.00	16	
17	4 6 3	7 1 9	1.33	-2.67	17	
18	4 6 9	7 1 3	-2.67	-2.67	18	
19	4 3 9	7 1 6	-0.67	-3.33	19	
20	6 3 9	7 1 4	-2.00	-4.67	20	

two-sample test

test for **decrease**: use the negative test statistic

perm	data	$T = \bar{x}_A - \bar{x}_B$	sorted	rank		
1	7 1 4	6 3 9	-2.00	4.67	1	significant
2	7 1 6	4 3 9	-0.67	3.33	2	
3	7 1 3	4 6 9	-2.67	2.67	3	
4	7 1 9	4 6 3	1.33	2.67	4	
5	7 4 6	1 3 9	1.33	2.00	5	
6	7 4 3	1 6 9	-0.67	1.33	6	
7	7 4 9	1 6 3	3.33	1.33	7	
8	7 6 3	1 4 9	0.67	0.67	8	
9	7 6 9	1 4 3	4.67	0.67	9	
10	7 3 9	1 4 6	2.67	0.67	10	
11	1 4 6	7 3 9	-2.67	-0.67	11	
12	1 4 3	7 6 9	-4.67	-0.67	12	
13	1 4 9	7 6 3	-0.67	-0.67	13	
14	1 6 3	7 4 9	-3.33	-1.33	14	
15	1 6 9	7 4 3	0.67	-1.33	15	
16	1 3 9	7 4 6	-1.33	-2.00	16	$p = \frac{16}{20} = 0.80$
17	4 6 3	7 1 9	-1.33	-2.67	17	
18	4 6 9	7 1 3	2.67	-2.67	18	
19	4 3 9	7 1 6	0.67	-3.33	19	
20	6 3 9	7 1 4	2.00	-4.67	20	

two-sample test

two-sided test: use the absolute value of the test statistic

perm	data	$T = \bar{x}_B - \bar{x}_A $	sorted	rank		
1	7 1 4	6 3 9	2.00	4.67	1	significant
2	7 1 6	4 3 9	0.67	4.67	2	
3	7 1 3	4 6 9	2.67	3.33	3	
4	7 1 9	4 6 3	1.33	3.33	4	
5	7 4 6	1 3 9	1.33	2.67	5	
6	7 4 3	1 6 9	0.67	2.67	6	
7	7 4 9	1 6 3	3.33	2.67	7	
8	7 6 3	1 4 9	0.67	2.67	8	
9	7 6 9	1 4 3	4.67	2.00	9	
10	7 3 9	1 4 6	2.67	2.00	10	$p = \frac{10}{20} = 0.50$
11	1 4 6	7 3 9	2.67	1.33	11	
12	1 4 3	7 6 9	4.67	1.33	12	
13	1 4 9	7 6 3	0.67	1.33	13	
14	1 6 3	7 4 9	3.33	1.33	14	
15	1 6 9	7 4 3	0.67	0.67	15	
16	1 3 9	7 4 6	1.33	0.67	16	
17	4 6 3	7 1 9	1.33	0.67	17	
18	4 6 9	7 1 3	2.67	0.67	18	
19	4 3 9	7 1 6	0.67	0.67	19	
20	6 3 9	7 1 4	2.00	0.67	20	

formal procedure for a permutation test

given: data, test statistic, sig. level α , **possible exchanges**

- ▶ compute test statistic T_1 for original data
- ▶ compute test statistic T_i for all permutations $i = 2 \dots n_P$
(or a random selection \rightarrow 'Monte-Carlo')
 $i = 1$ is the 'neutral permutation'

- ▶ determine the rank of T_1 :
$$r = \sum_{i=1}^{n_P} [T_i \geq T_1]$$

where [true] = 1 and [false] = 0

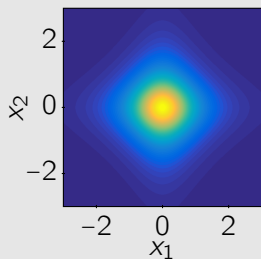
- ▶ determine the p -value:
$$p = \frac{r}{n_P}$$
- ▶ if $p \leq \alpha$, reject H_0

the test is exact if α is a multiple of $\frac{1}{n_P}$ and there are no ties, otherwise it is conservative

Exchangeability

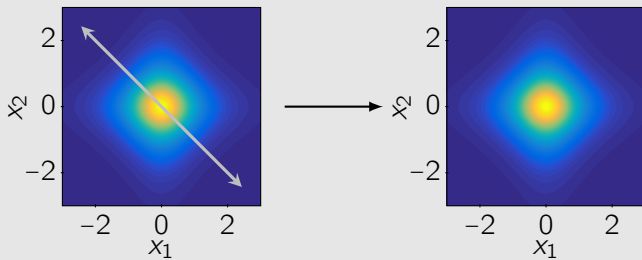
exchangeability

independent samples from the **same distribution** (i.i.d.) ...



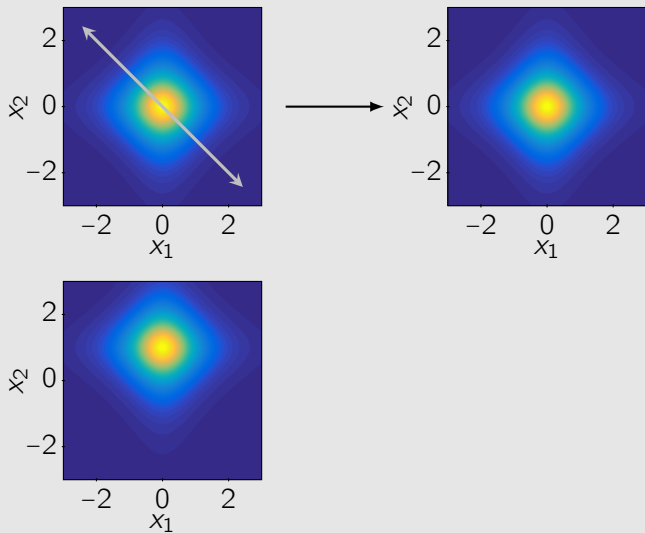
exchangeability

... can be **exchanged**: $x_1 \leftrightarrow x_2$



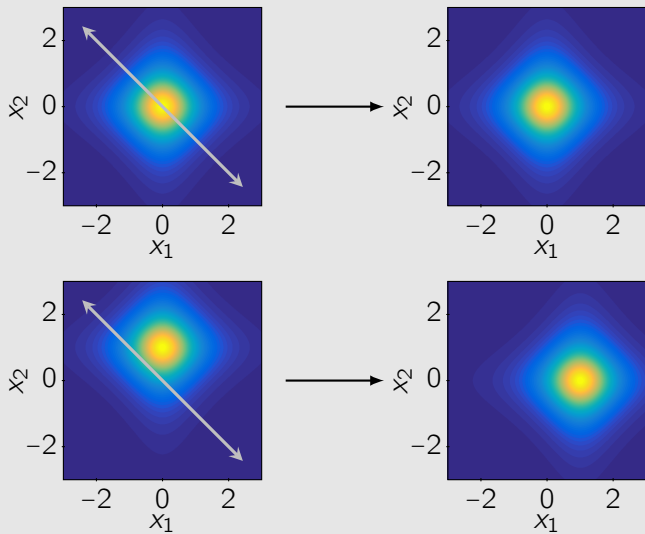
exchangeability

independent samples from **different distributions** . . .



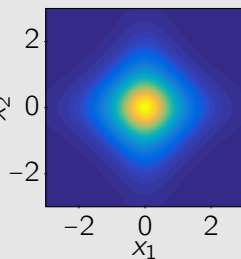
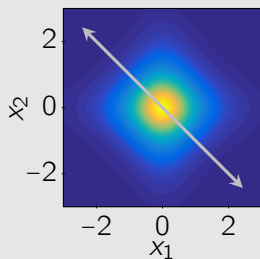
exchangeability

... **cannot** be **exchanged**

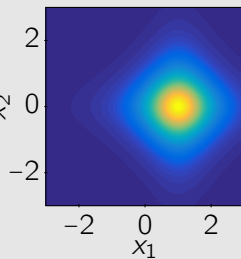
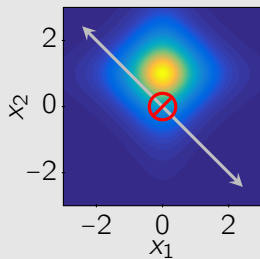


exchangeability

exchangeability refers to the **null hypothesis!**



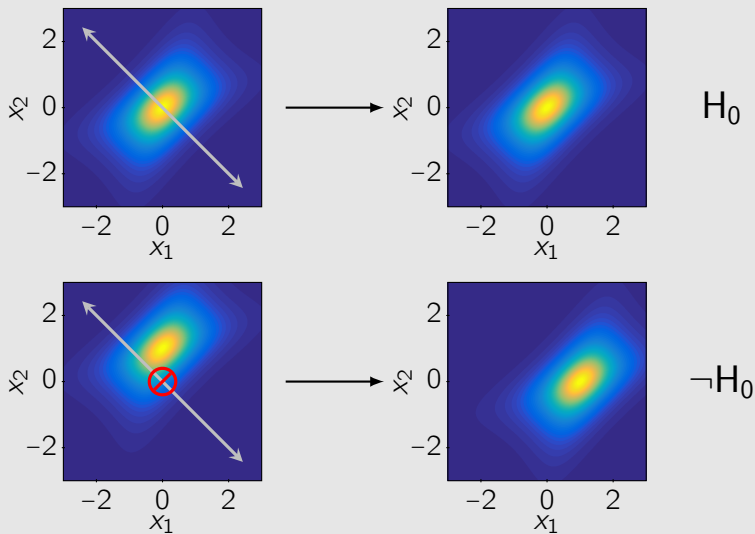
H_0



$\neg H_0$

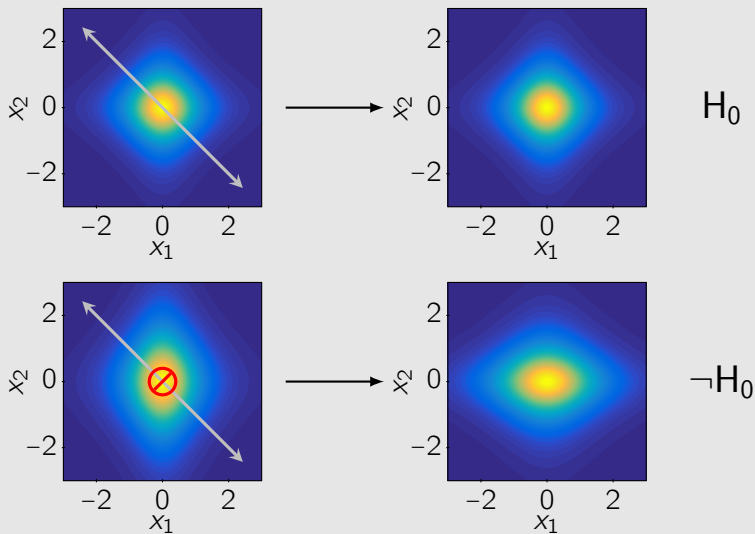
exchangeability

exchangeable also if there is dependency but **symmetric**



exchangeability

caution: null hypothesis may be wrong **in unexpected ways**



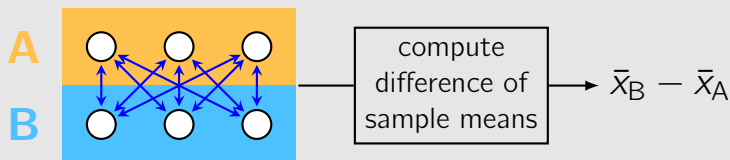
Some test examples

tests: univariate two-sample

independent values sampled from two univariate distributions

H_0 : two samples come from the **same distribution**

→ i.i.d. case: values can be exchanged freely



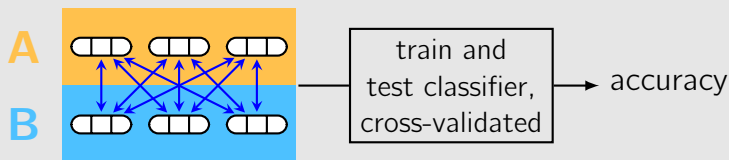
test statistic is sensitive to difference in means:
permutation analogue of the **two-sample t-test**

tests: multivariate two-sample

independent vectors from two multivariate distributions

H_0 : two samples from the **same multivariate distribution**

→ i.i.d. case: **vector values** can be exchanged freely



Problem with accuracies: few possible values → ties!

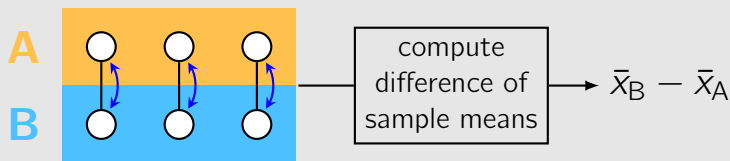
Example: classification of subject-specific patterns,
e.g. of patients based on structural MRI

tests: univariate paired

paired values sampled from two univariate distributions

H_0 : two samples come from the same distribution

→ partial dependency: values can be exchanged **within pairs**



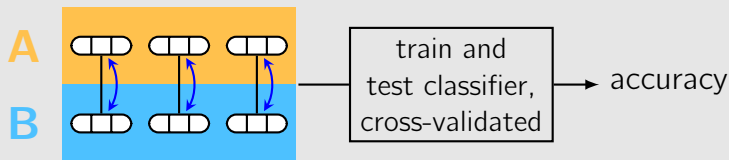
test statistic is sensitive to difference in means:
permutation analogue of the **paired t-test**

tests: multivariate paired

paired vectors sampled from two multivariate distributions

H_0 : two samples from the same multivariate distribution

→ partial dependency: vectors can be exchanged within pairs



Example:

- classification of condition-specific patterns across subjects
- classification of patterns across runs, within subject

alternative to accuracy: multivariate test

multivariate samples can also be compared using significance tests → potentially **more powerful**

NeuroImage 146 (2017) 113–120



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What's in a pattern? Examining the type of signal multivariate analysis uncovers at the group level[☆]



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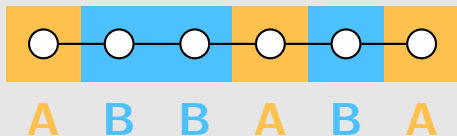
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^e Department of Psychology, Stanford University, Stanford, CA, United States

Limited exchangeability

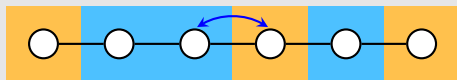
time series

fMRI trials occur in a time series with **serial dependency**

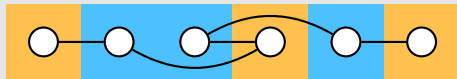


time series

problem: exchanges change the **dependency structure**



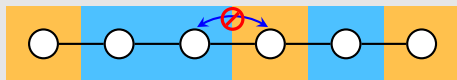
A B B A B A



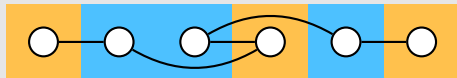
A B B A B A

time series

no exchangeability of values in fMRI time series!



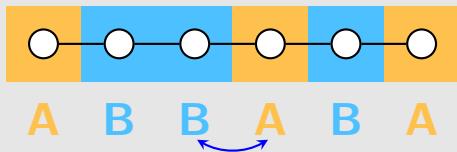
A B B A B A



A B B A B A

time series

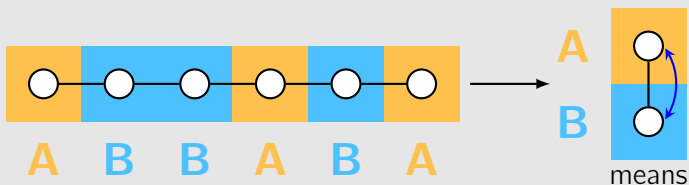
for a **randomized** trial sequence, **labels** can be exchanged



→ **randomization test**, \neq permutation test
relies on the distribution of the labels, not the data
disadvantage: not exact for given randomized design

time series

run-wise **means** or GLM estimates are often exchangeable



→ run-wise classification may be a better alternative

alternative to classification: cvMANOVA

cross-validated multivariate ANOVA allows for arbitrary time series designs → **more flexible** than decoding

NeuroImage 89 (2014) 345–357

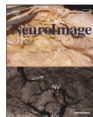


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Searchlight-based multi-voxel pattern analysis of fMRI by cross-validated MANOVA



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Further topics

multiple comparisons

several tests in parallel (voxels): control for **family-wise error**

precise correction depends on dependency between tests,
accounted for by using the **same permutation** across tests

test statistic T_{ij} , permutations $i = 1 \dots n_P$, tests $j = 1 \dots n_T$

- ▶ test whether there is an effect somewhere (omnibus H_0):
use the **maximum statistic**

$$M_i = \max_{j=1}^{n_T} T_{ij}, \quad p_{\text{omnibus}} = \frac{1}{n_P} \sum_{i=1}^{n_P} [M_i \geq M_1]$$

- ▶ test whether there is an effect at j , corrected

$$p_j = \frac{1}{n_P} \sum_{i=1}^{n_P} [M_i \geq T_{1j}]$$

beyond permutations

the principle underlying permutation tests:

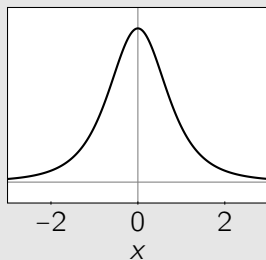
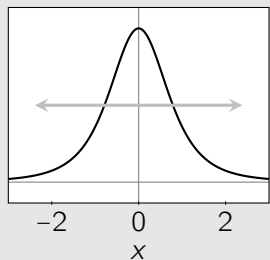
- ▶ find a group of **transformations** under which the H_0 -distribution is **invariant** (includes the neutral transformation)
- ▶ compute the test statistic from data after **each transformation** has been applied
- ▶ ...

permutation is a special case of transformation,
exchangeability is a special case of invariance

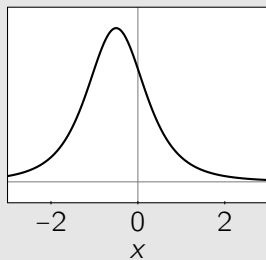
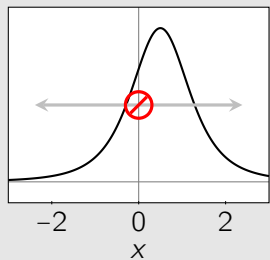
“group”: the combination of two transformations is another transformation (is also in the group)

beyond permutations: mirror symmetry

'sign-flip test': $x \leftrightarrow -x$



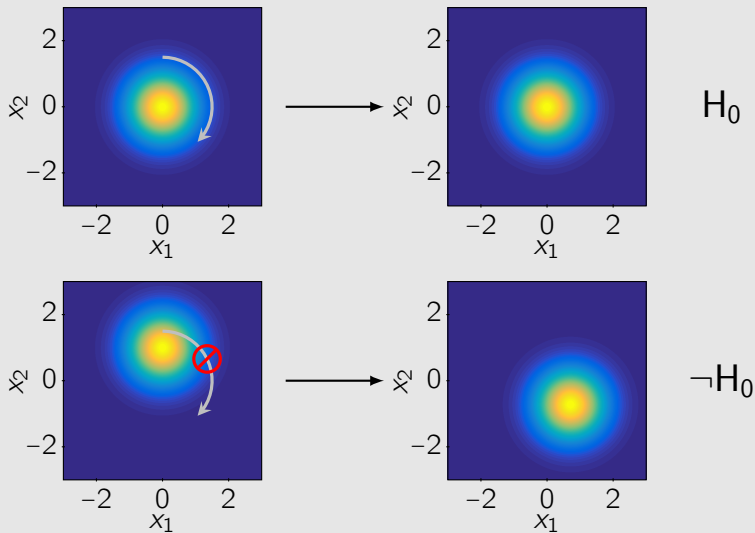
H_0



$\neg H_0$

beyond permutations: sphericity

'rotation test': <https://f1000research.com/posters/6-1085>



further reading

permutation tests, general

- Ernst, Permutation methods: A basis for exact inference, *Statistical Science* 2004
- Good, *Permutation, Parametric, and Bootstrap Tests of Hypotheses*, Springer 2005
- Lehmann & Romano, *Testing Statistical Hypotheses*, Springer 2005, [Sec. 5.8](#)

neuroimaging

- Nichols & Holmes, Nonparametric permutation tests for functional neuroimaging: A primer with examples, *Human Brain Mapping* 2001
- Winkler et al., Permutation inference for the general linear model, *NeuroImage* 2014
- Winkler et al., Non-parametric combination and related permutation tests for neuroimaging, *Human Brain Mapping* 2016

MVPA (caution!)

- Golland & Fischl, Permutation tests for classification: Towards statistical significance in image-based studies, *Information processing in medical imaging* 2003
- Etzel & Braver, MVPA permutation schemes: Permutation testing in the land of cross-validation, *PRNI* 2013
- Schreiber & Krekelberg, The statistical analysis of multi-voxel patterns in functional imaging, *PLOS ONE* 2013
- Stelzer et al., Statistical inference and multiple testing correction in classification-based multi-voxel pattern analysis (MVPA), *NeuroImage* 2013
- Allefeld et al., Valid population inference for information-based imaging: From the second-level t -test to prevalence inference, *NeuroImage* 2016

randomization tests

- Lehmann & Romano, *Testing Statistical Hypotheses*, Springer 2005, [Sec. 5.10](#)
- Edgington & Onghena, *Randomization Tests*, Chapman & Hall 2007