




Slides


Day 1, Session 1
Introduction



Workshop 7-8th July 2014
Ghent University

Goals


- Conduct SPM analyses
- Interpret SPM's results
- Report SPM methods and results



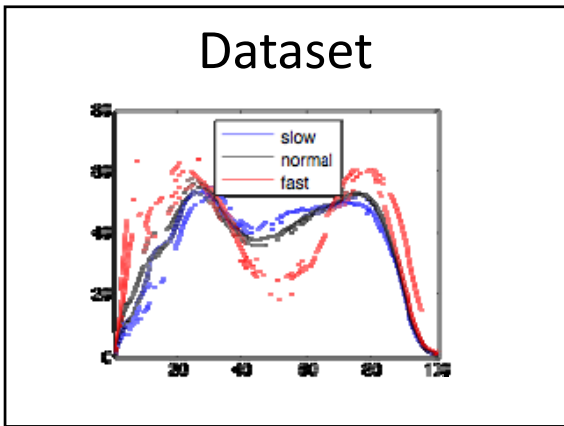
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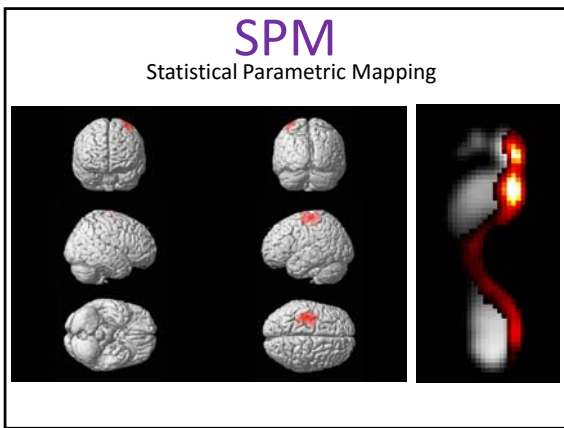
Not-Goals

- Become **highly proficient** in Matlab
- Understand **all aspects** of SPM
- Understand **all aspects** of conventional statistics
- Focus on **structuring** and **inputting** data



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Suitable for what biomechanical data?

a) Spatiotemporally smooth
 Sampled above Nyquist frequency
 Biological tissue viscoelasticity

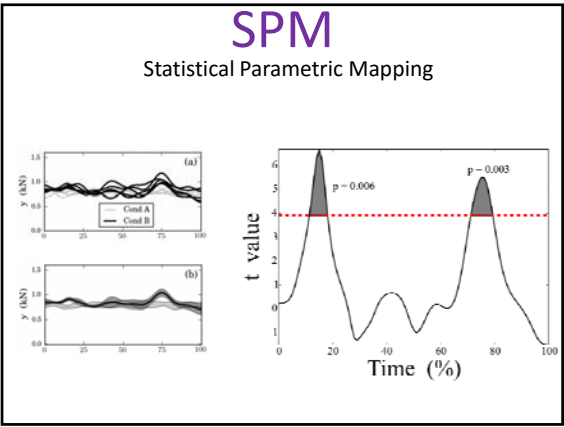
b) Bounded

Time or space

1D


2D

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Questions


- Have you heard of Matlab?
- Have you processed data in Matlab?
- Have you conducted statistical tests in Matlab?
- Have you validated probability computations in Matlab?



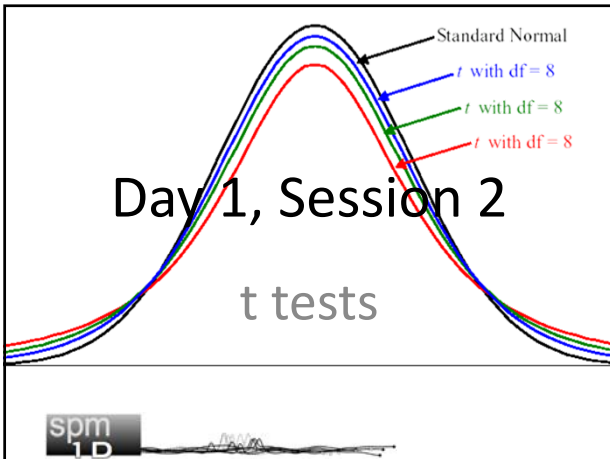
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MATLAB

- “MATrix LABORatory”
- Numerical array computations
- Useful for data analysis
- Broad functionality:
 - Statistics, optimization, signal processing, dynamic simulation, etc.



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Goals

- t-test recap
- Univariate vs 1DSPM calculations
- Run t-tests in Matlab
- Interpret output variables

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T-tests

- one-sample t-test
 - Used to test the difference between a single dataset and a constant (usually zero)
- paired t-test
 - Used to test for a differences in repeated measurements of a sample
- two-sample t-test
 - Used to test for a difference between two samples

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Experimental scenarios

1. One subject, multiple trials, two tasks
2. 100 subjects, one body mass value each, compared against national average
3. Surgical Procedure A vs. Surgical Procedure B on 40 subjects per procedure
4. 100 subjects, tested at ages 20 and 30
5. one subject, measured multiple times, compared to a "normal" database
6. Controls vs. Patients
7. Multiple subjects performing two different tasks once each

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First, we need to set up our null hypothesis.

Null Hypothesis (H_0)

– Predicts no significant difference

e.g. There is no significant difference in vertical ground reaction forces in injured versus uninjured runners

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Univariate two-sample t-test calculation

$$t = \frac{\bar{y}_B - \bar{y}_A}{\sqrt{\frac{1}{J}(s_A^2 + s_B^2)}}$$

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Univariate two-sample t -test

- 1. Compute mean values y_A and y_B
- 2. Compute the standard deviations s_A and s_B
- 3. Compute the t test statistic
- 4. Conduct statistical inference. Use α and the t distribution to compute the critical threshold. If $t > t_{\text{critical}}$ reject null hypothesis
- 5. Compute exact p -value using t and the univariate t -distribution.



Univariate ex_ttest_twosample.m

```
%(1) Conduct 0D test using spm1d:  
spm = spm1d.stats.ttest2(yA, yB);  
spmi = spm.inference(0.05, 'two_tailed', false);
```

yA = 10 x 1 double
yB = 10 x 1 double



Univariate reporting

- For example:
There was no significant difference between the male and female reaction times, $t(9) = 0.52$, $p = 0.15$.



SPM1D

two-sample t -test

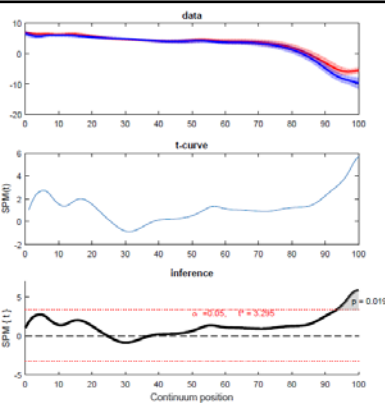
1. Compute mean fields $\bar{y}_A(q)$ and $\bar{y}_B(q)$
2. Compute the st. dev. of fields $s_A(q)$ and $s_B(q)$
3. Compute the t test statistic field
4. Conduct statistical inference.

Use α and the RFT t distribution to compute the critical threshold. If $SPM\{t\} > t$ critical reject null hypothesis for suprathreshold clusters.

5. Compute exact p -value for each cluster using cluster size and RFT distribution(s) for $SPM\{t\}$ topology.

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1D



SPM1D

calculation comparison

$$t = \frac{\bar{y}_B - \bar{y}_A}{\sqrt{\frac{1}{J}(s_A^2 + s_B^2)}}$$

$$SPM\{t\} \equiv t(q) = \frac{\bar{y}_B(q) - \bar{y}_A(q)}{\sqrt{\frac{1}{J}(s_A^2(q) + s_B^2(q))}}$$

The calculation of $t(q)$ does not mean that we are "doing 'q' t-tests".

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SPM1D

t-test Matlab function names

spm1d.stats.ttest(...)	One-sample t test
spm1d.stats.ttest_paired(...)	Paired t test
spm1d.stats.ttest2(...)	Two-sample test



SPM1D

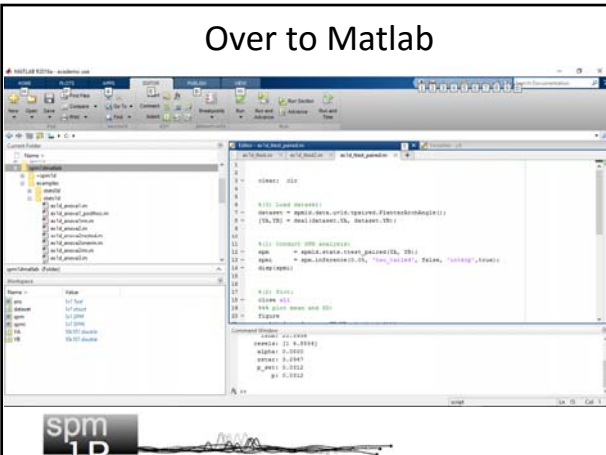
ex1d_ttest2.m

```
%(1) Conduct SPM analysis:  
spm = spm1d.stats.ttest2(YA, YB);  
spmi = spm.inference(0.05, 'two_tailed', true);
```

yA = 10 x 101 double
yB = 10 x 101 double



Over to Matlab



Questions resulting from the analysis
of these datasets?

Helpful tips?

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Day 1, Session 3

Probability and Random Field Theory

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Goals

- Describe the meaning of p values for 1D analyses
- Simulate experiments using smooth, random 1D data to validate p values

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1D

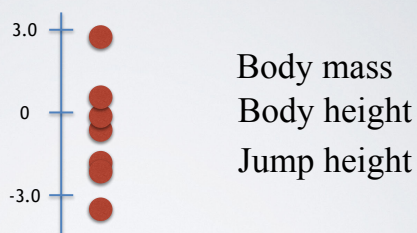






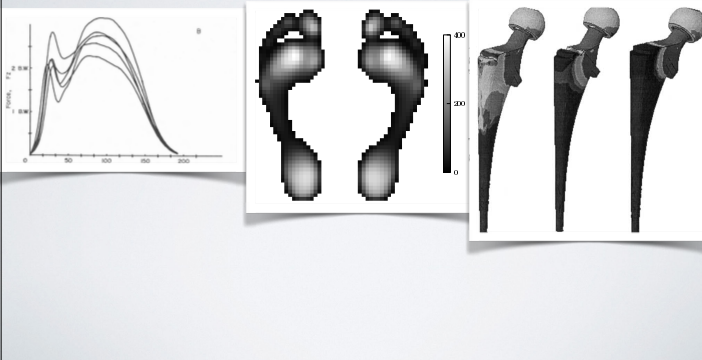
One can't use one's knowledge about randomness in oranges to make probabilistic conclusions regarding apples.

Zero-dimensional data



Variables which do not change in space or time

n -dimensional data



One can't use one's knowledge about randomness in **0D data** to make probabilistic conclusions regarding **n D data**.

Goals

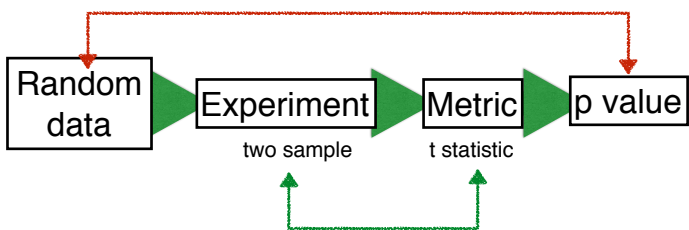
- Describe the meaning of p values for 1D analyses
- Simulate experiments using smooth, random 1D data to validate p values



What is a p value?

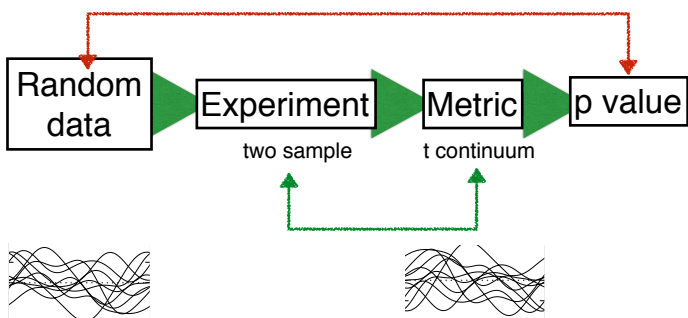
Demo

Summary



Demo 1D

Summary 1D



What is a p value?

The probability that a completely random nD process will yield a particular result.

Goals

- Describe the meaning of p values for 1D analyses
- Simulate experiments using smooth, random 1D data to validate p values

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Summary

t values describe experimental data

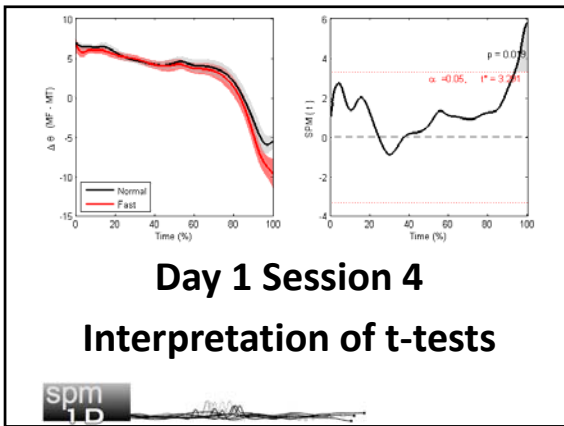
p values describe random data

Use an nD model of randomness to make probabilistic conclusions regarding nD data

p is not:

- x The probability that the null hypothesis is true
- x The probability that the alternative hypothesis is false
- x The probability that the observed result is random

✓ $P(\text{data}|H_0)$

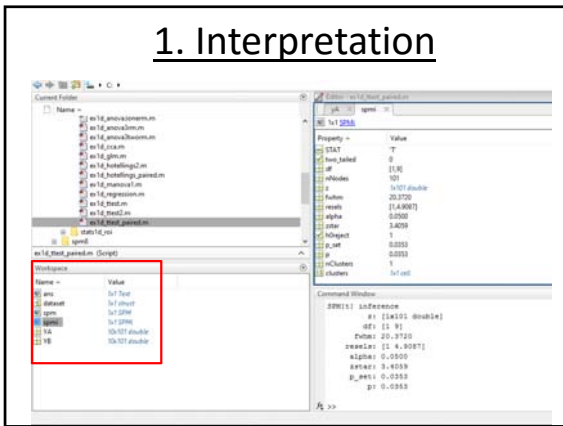


“Is it significant?”

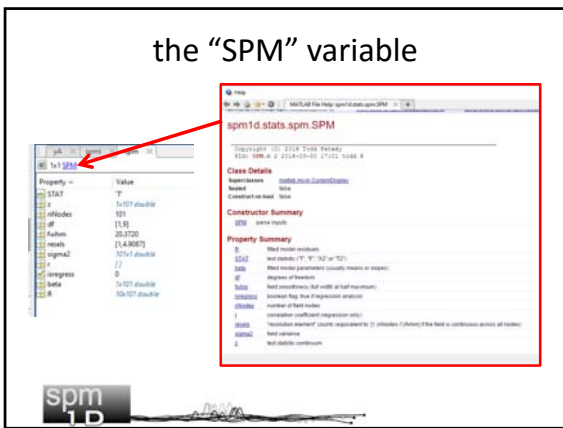
spm 1D

- Plan**
1. Interpret t-test results
 2. Describe the methods used
 3. Present the results
 4. Reviewer hints and tips
- spm 1D

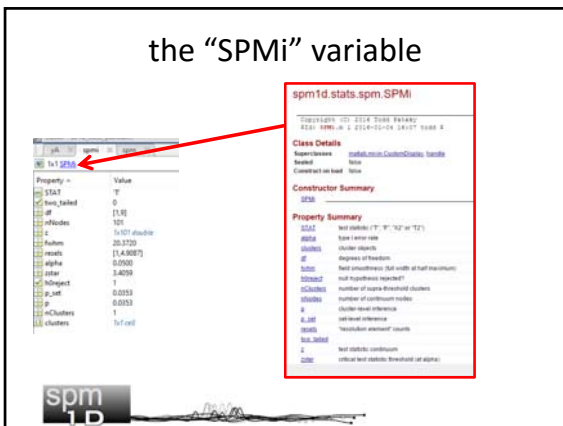
1. Interpretation



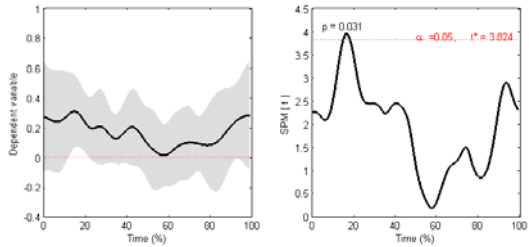
the "SPM" variable



the "SPMi" variable



Interpretation ex1d_ttest.m



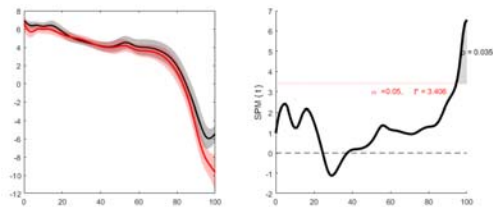
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```
>> disp(spmi)
```

```
SPM{t} inference  
z: [1x100 double]  
df: [1 9]  
fwhm: 11.2615  
resels: [1 8.7910]  
alpha: 0.0500  
zstar: 3.8213  
p_set: 0.0310  
p: 0.0310
```

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Key aspects – Interpretation ex1d_ttest_paired.m



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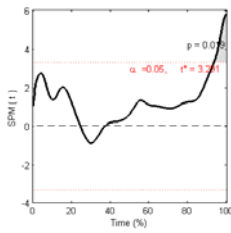
```
>> disp(spmi)
```

```
SPM{t} inference  
z: [1x101 double]  
df: [1 9]  
fwhm: 20.3720  
resels: [1 4.9087]  
alpha: 0.0500  
zstar: 3.4059  
p_set: 0.0353  
p: 0.0353
```

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Cluster specific p-values



Identically smooth random
1D data would produce a
cluster of this breadth with a
probability of $p=0.019$

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Clusters

```
spm1d.clusters{1}.details
```

```
Command Window  
>> spm1d.clusters{1}.details  
  
spm1d.geom.Cluster with properties:  
  endpoints: [94 4568 100]  
  origin: 1  
  levelapped: 0  
  extentR: 5.5632  
  extentL: 0.2732  
  N: 3.4059  
  WZ: [97.0453 5.0059]  
  P: 0.0353  
  
Methods, Superclasses
```

spm1d.geom.Cluster

Copyright (c) 2014 Todd Pataky
File: Cluster.m 1 2014-02-04 14:07:0000 8

Class Details

Superclasses: [spm1d.geom.Cluster](#)
Sealed: No
Construct on load: No

Constructor Summary

Cluster

Property Summary

E	probability value (based on random field theory)
level	cluster sign (+1 above -1, below -1)
extentR	starting and ending points (interpolated to threshold)
extentL	breadth (units: resels)
width	breadth (units: resels)
z	minimum height above threshold (actually zstar)
wraparound	true if the cluster wraps around the origin of a circular field
centroid	centroid

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2. Methods

What are the contents of a good methods section?

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SPM Methods

- a) Statistical tests used
- b) SPM code & analysis software
- c) Refer to key SPM/RFT literature
- d) Define terminology
- e) Specify alpha – correction?
- f) How results will be interpreted

[also relevant: data treatment, smoothing, averaging]

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a) Statistical tests used

- e.g. SPM One sample t -test
 - One or two-tailed?
 - Dependent variable tested
 - Independent variable

"A SPM two-tailed two-sample t -test was used to compare male versus female knee angles."

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b) SPM code

- Can refer to <http://www.spm1d.org/> for open source code
- The current Python version of **spm1d** is:
- The current Matlab version of **spm1d** is:

Software

Python 2.7.2; Enthought Python Distribution, Austin, TX.
Matlab R2016a (8.3.0.532), The Mathworks Inc, Natick, MA.
[Home → Help → About Matlab]



c) Key SPM/RFT literature

SPM literature (Neuroimaging)

- Friston KJ, Ashburner JT, Kiebel SJ, Nichols TE, Penny WD. (Eds.) Statistical parametric mapping: the analysis of functional brain images. London: Elsevier; 2007.
 - A book that describes the SPM analysis concepts with specific application to brain images.
- Friston KJ, Holmes AP, Worsley KJ, Poline JB, Frith CD, Frackowiak RSJ (1995). Statistical parametric maps in functional imaging: a general linear approach. *Human Brain Mapping* 2, 189–210.
- SPM documentation repository, Wellcome Trust Centre for Neuroimaging: <http://www.fil.ion.ucl.ac.uk/spm/doc/>

SPM/RFT Literature (Biomechanics)

- Pataky, T. C. . (2012). One-dimensional statistical parametric mapping in python. *Computer Methods in Biomechanics and Biomedical Engineering*, 15(3):295–301.
- Pataky, T. C. (2016). Rft1d: Smooth one-dimensional random field upcrossing probabilities in python. *Journal of Statistical Software*, page in press.



d) Terminology –

- SPM vs SPM{t}

*SPM refers to the overall methodological approach
SPM{t} to the scalar trajectory variable*

- Suprathreshold cluster

“adjacent points of the SPM{t} curve often exceed the critical threshold, we therefore call these “supra-threshold clusters”.

- Critical threshold

“Value at which only α % (5%) of smooth random curves would be expected to cross”



e) Specify alpha

- You may wish to correct alpha for multiple comparisons / dependent variables.
- Remember this would have to be entered manually in the code.

$$p_{critical} = 1 - (1 - \alpha)^{1/N}$$

"To retain a family-wise Type 1 error rate of $\alpha = 0.05$ we adopted a Sidák corrected threshold of 0.012 for four comparisons"

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f) How results will be interpreted

- What does a high value of SPM{t} mean?
- What is the consequence of crossing the critical threshold (or not)?

"if the SPM{t} trajectory crosses the critical threshold at any time node, the null hypothesis is rejected"

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An example method

- Refer to [handout](#)
"Day1Session4 - Example method write-up.docx"
- Task – Using the colour-coding in the bullet-points, colour the text in the example write-up to match the relevant bullet-point.
The first sentence has been started as an example.
What could be more concise?

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Example methods to refer to

- Pataky TC (2010). Generalized n-dimensional biomechanical field analysis using statistical parametric mapping. *Journal of Biomechanics* 43, 1976-1982.
- Pataky TC (2012) One-dimensional statistical parametric mapping in Python. *Computer Methods in Biomechanics and Biomedical Engineering*. 15, 295-301.
- Pataky TC, Robinson MA, Vanrenterghem J (2013). Vector field statistical analysis of kinematic and force trajectories. *Journal of Biomechanics* 46 (14): 2394-2401.

Applications

- Vanrenterghem, J., Venables, E., Pataky, T., Robinson, M. (2012). The effect of running speed on knee mechanical loading in females during side cutting. *Journal of Biomechanics*, 45, 2444-2449.
- De Ridder, R., Willems, T., Vanrenterghem, J., Robinson, M., Pataky, T., Roosen, P. (2013). Gait kinematics of subjects with chronic ankle instability using a multi-segmented foot model. *Medicine and Science in Sports and Exercise*, 45, 2129-2136.
- Robinson, M.A., Donnelly, C.J., Tsao, J., Vanrenterghem, J. (2014). Impact of knee modelling approach on indicators and classification of ACL injury risk. *Medicine & Science in Sports & Exercise*, 46 (7), 1269-1276.



3. Presentation of results

- Key information to present:
 - a) Was the critical threshold exceeded?
 - b) Direction of effect
 - c) Consequence for the null hypothesis
 - d) Descriptive data:
 - critical threshold, p-value/s, number of supra-threshold clusters, extent of clusters, degrees of freedom.



Figure caption

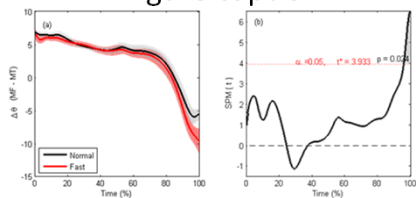


Figure 1 a) Mean trajectories for longitudinal arch angles during normal (black) and fast (red) walking. b) The paired samples t-test statistic SPM {t}. The critical threshold of 3.933 (red dashed line) was exceeded at time = 96% with a supra-threshold cluster probability of $p=0.024$ indicating a significantly more negative angle in the fast condition.



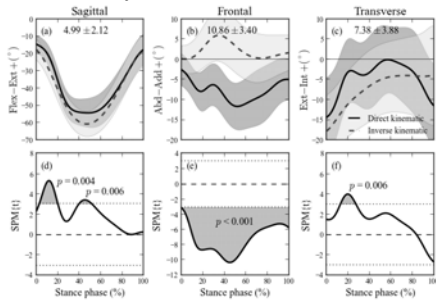
Results section example

Mean longitudinal arch angles during normal and fast walking were highly similar for the majority of time (figure 1a). However one supra-threshold cluster (96-100%) exceeded the critical threshold of 3.933 as the $\Delta\Theta(\text{MF-MT})$ angle during fast walking was significantly more negative¹ than during normal walking (figure 1b). The precise probability that a supra-threshold cluster of this size would be observed in repeated random samplings was $p=0.024$. The null hypothesis was therefore rejected.

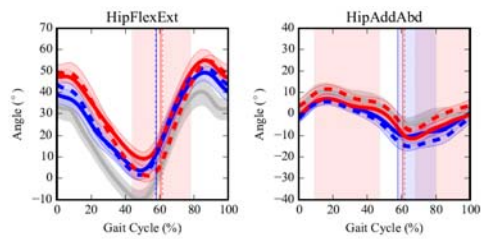
~~$\text{SPM}(t)_{\text{max}} = 6.55, p = 0.024, \text{d.f.} = 1,9$~~

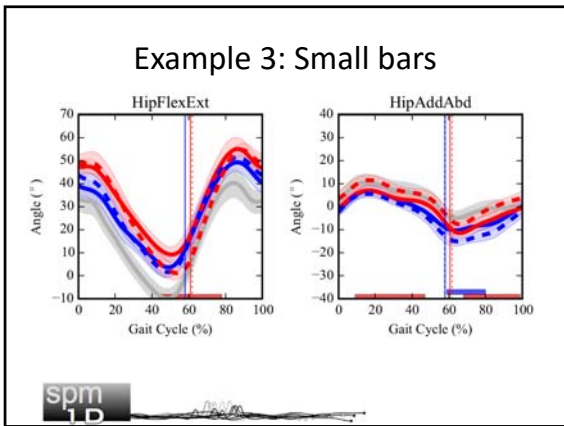


Example 1: Data and stats



Example 2: Shading



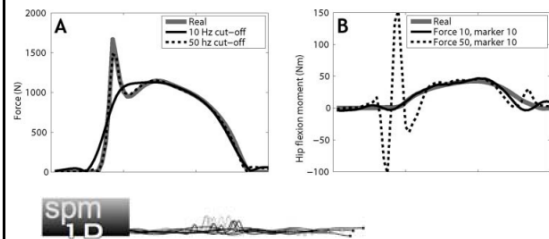


- 4. Reviewers – Hints and tips**
- Anticipate some resistance
 - Be consistent throughout
 - Present statistical results with original data
 - Highlight similarity to univariate interpretation
 - Refer to papers that use similar analyses
 - Benefits of SPM vs PCA, FDA etc
 - Use supplementary material where appropriate
- spm 1D**

- Tasks – Analyse and report...**
- Analyse and report the results of the data
[paired_data.mat](#)
1. Write the methods for the scenario below
 - Knee flexion angle data during running
 - 34 male participants
 - Two speed conditions
 - Data are normalised to the stance phase
 2. Write a figure caption
 3. Write a results section
- spm 1D**

Day 2, Session 1

Registration and smoothing



Goals

- Refresh smoothing and registration principles of biomechanical data
- Identify how smoothness and registration can affect SPM outcome

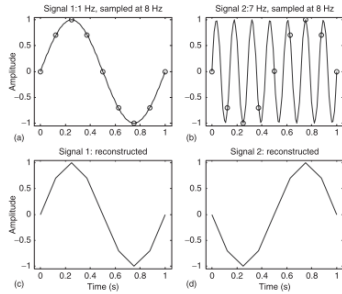


Overview

- Sampling of biomechanical signals
- Smoothing of biomechanical signals
- Registration of biomechanical signals
- Exercise on effect smoothing on SPM



Sampling biomechanical data



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Smoothing biomechanical data

Remove artefact \longleftrightarrow Retain signal

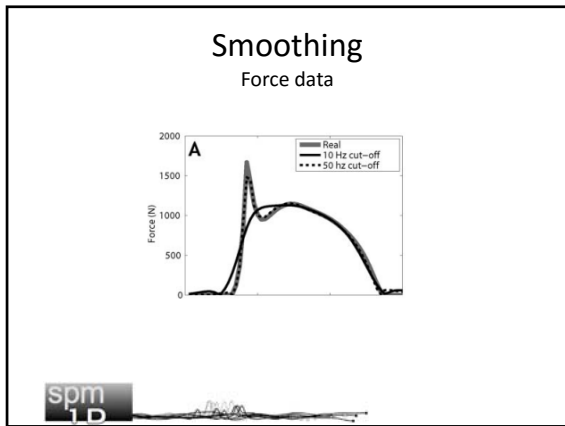
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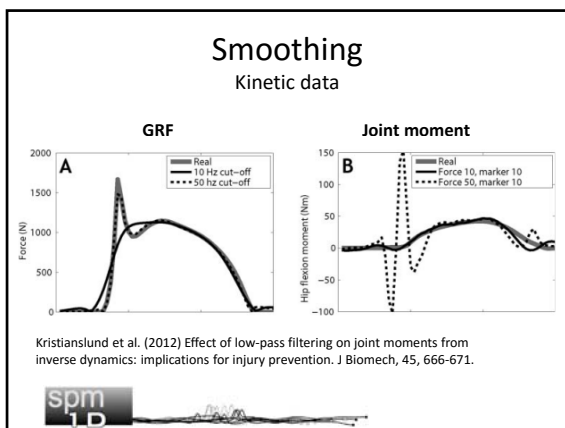
Smoothing Kinematic data

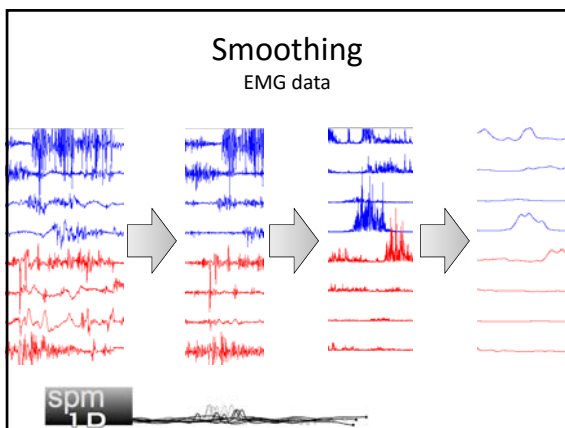


Displacement: $x(t) = \sin t$; Noise $(t) = 0.001 \sin 50t$; Signal to noise ratio: 1:0.001.
 Velocity: $v(t) = \cos t$; Noise $(t) = 0.05 \cos 50t$; Signal to noise ratio: 1:0.05.
 Acceleration: $a(t) = -\sin t$; Noise $(t) = -2.5 \sin 50t$; Signal to noise ratio: 1:2.5.

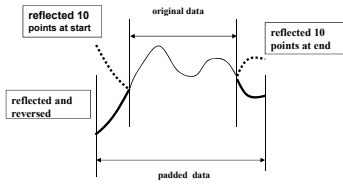
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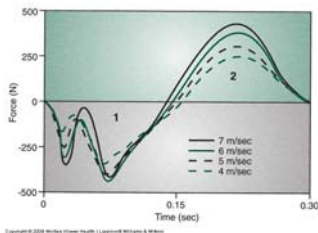


End point effects



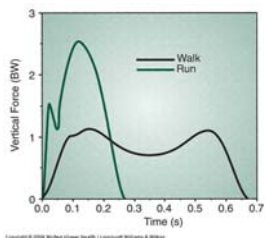
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Registration

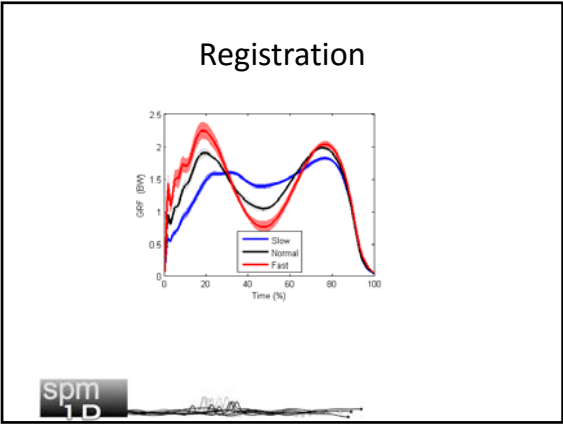


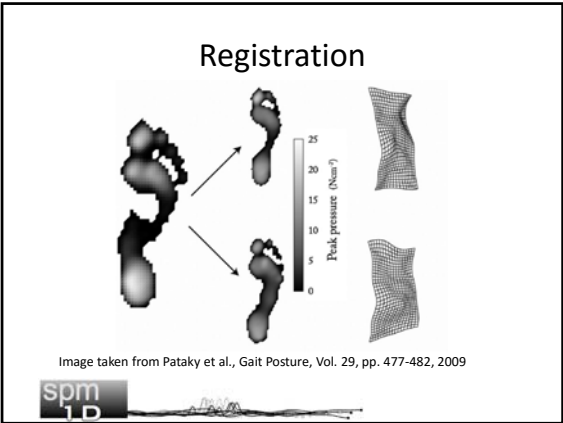
spm
1D

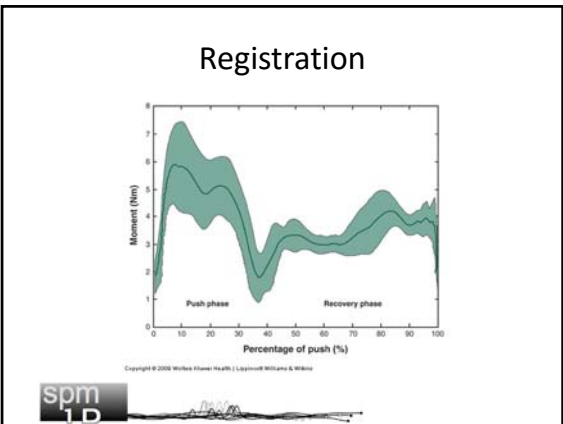
Registration



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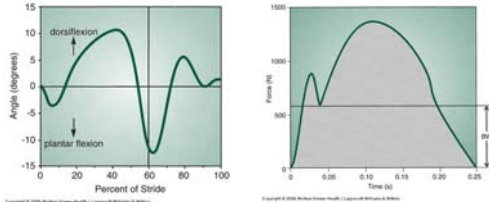






Linear registration

- Temporal context is retained uniformly



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Relevance to SPM

- Sampling affects SPM when Nyquist is violated
- Smoothing affects dependence 'nodes'

$$x'_i = a_0 x_i + a_1 x_{i-1} + a_2 x_{i-2} - b_1 x'_{i-1} - b_2 x'_{i-2}$$

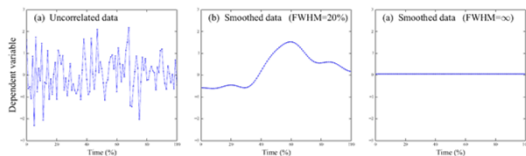
=> Affects SPMi (see next slide)

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Inference

RFT handles the issue of multiple comparisons and statistical inference .

(Adler, 1981)



If neighbouring data are completely uncorrelated we'd conduct 100 independent t tests. If data are correlated then we have <100 independent tests.

The temporal (or spatial) gradient estimates # independent processes

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Relevance to SPM

- Sampling affects SPM when Nyquist is violated
- Smoothing affects dependence 'nodes'

$$x'_i = a_0 x_i + a_1 x_{i-1} + a_2 x_{i-2} - b_1 x'_{i-1} - b_2 x'_{i-2}$$

=> Affects SPMi

- Registration affects t-curve

=> Affects SPM{t}

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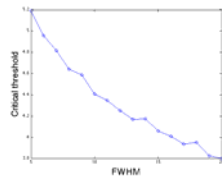


Exercise

Smoothness of group trajectories affects outcome SPM by lowering threshold

– Can we investigate relationship smoothness and threshold?

```
z = randn1d(J, Q, fwhm);
spm = spm1d.stats.tttest(z);
spm_i = spm.inference(0.05);
```

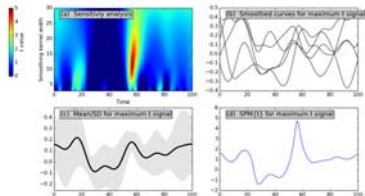


spm
1D



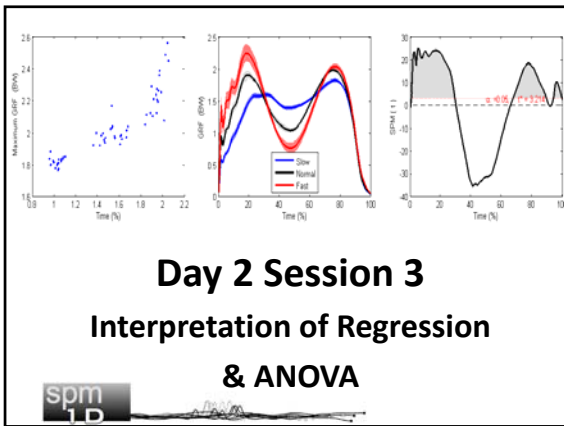
Sensitivity of data processing

e.g. treat smoothness as a sensitivity measure



spm
1D





Plan

- Regression example
 - Code
 - Interpretation
- ANOVA example
 - Code
 - Interpretation

spm
1D

Regression

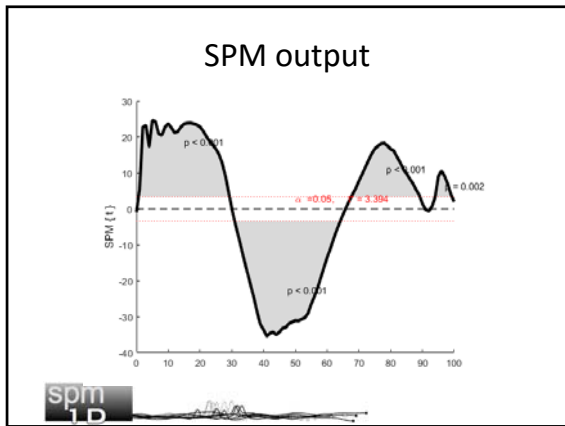
```

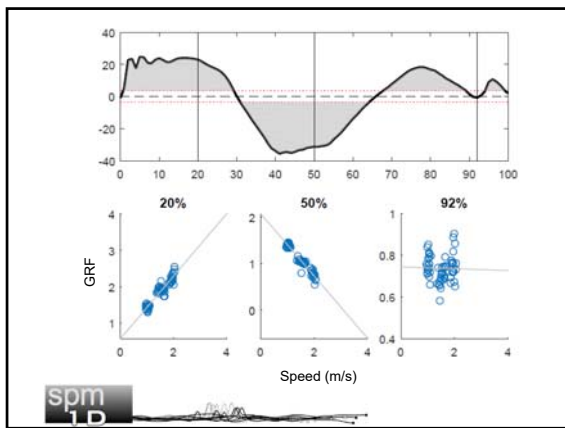
%(1) Conduct SPM analysis:
spm = spm1d.stats.regress(Y, x);
spm_i = spm.inference(0.05, 'two_tailed', true);
  
```

"x" independent variable
walking speed

"Y" dependent variable
vertical ground
reaction force

spm
1D





Why a SPM{t} value?

The t value is the statistic upon which statistical inferences (p values) are based.

The regression coefficient r and t value are alternative expressions of effect magnitude, and map directly to each other given the sample size.

spm.r is the correlation coefficient

spm 1D

Regression interpretation

There was a significant relationship between walking speed and vGRF. A greater walking speed significantly increased the vGRF during the first and last 30% stance but significantly reduced GRF from ~30-70% stance. As random data would produce this effect <5% time. The null hypothesis was therefore rejected.



Figure caption

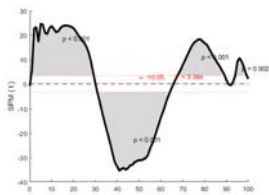
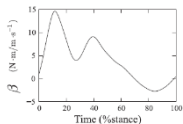
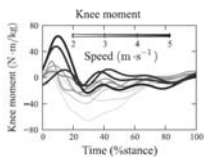


Figure 1. A statistical inference curve indicating significant relationships between walking speed and vertical ground reaction force. There were significant positive relationships during early and late stance and a significant negative relationship during mid-stance.



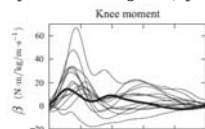
Random effects analysis

One subject's data

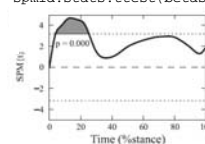


All subjects

`spm1d.stats.regress(speed,mom);`



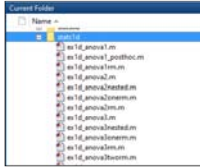
`spm1d.stats.ttest(betas);`



Vanrenterghem et al. (2012) JoB

ANOVA

- Many ANOVA types



spm
1D

SPM code

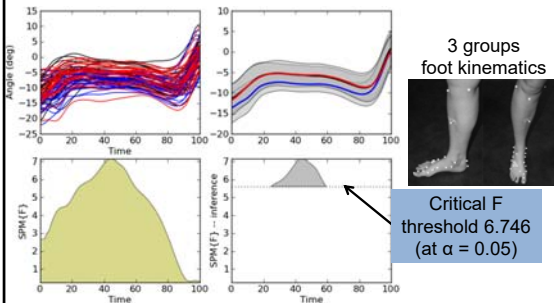
```
%(1) Conduct SPM analysis:  
spm      = spm1d.stats.anova1(Y, A);  
spm_i    = spm.inference(0.05);
```

```
%(1) Conduct SPM analysis:  
spm_bs   = spm1d.stats.anova1(Y, A);  
          %between-subjects model  
spm      = spm1d.stats.anova1rm(Y, A, SUBJ);  
          %within-subjects model  
spm_i    = spm.inference(0.05);
```

spm
1D

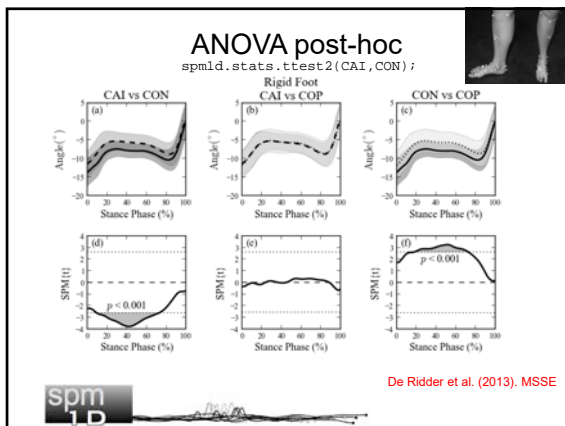
One-way ANOVA

```
spm1d.stats.anova(CAI, COP, CON);
```



spm
1D

De Ridder et al. (2013). MSSE



ANOVA interpretation

- There was a significant difference in angle between the three groups between ~20-60% stance. Identically smooth random 1D data would produce a cluster of this breadth with a probability of $p < 0.05$. Post-hoc independent t -tests showed that CAI and COP were both significantly greater than the CON group.

One way RM ANOVA

```

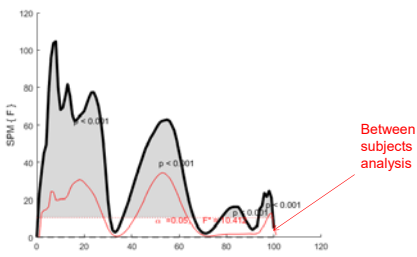
% (1) Conduct SPM analysis:
spm_bs = spm1d.stats.anova(Y, A);
        % between-subjects model
spm     = spm1d.stats.anova1rm(Y, A, SUBJ);
        % within-subjects model
spm_i   = spm.inference(0.05);
    
```

Warning: Only one observation per subject found. Residuals and inference will be approximate. To avoid approximate residuals: (a) Add multiple observations per subject and per condition, and (b) ensure that all subjects and conditions have the same number of observations.

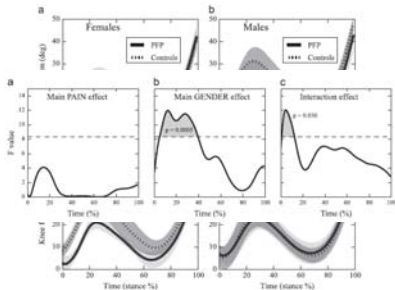
```
> In spm1d.stats.anova.designs.ANOVA1rm/check_for_single_responses (line 31)  
In spm1d.stats.anova1rm (line 9)
```



SPM output



Two-way ANOVA



Pataky et al. (2015). JoB

Tasks – Analyse and report...

- Finish ANOVA worksheet
- Analyse the example datasets 1-4
 1. Write the analysis code
 2. Plot the results
 3. Write a figure caption
 4. Write up the results

spm
1.0



Day 2, Session 4

Introduction to Vector Analysis with SPM

spm
1D



Topics

- Overview
- False positives
- Vector field statistics
- Reminder of SPM limitations
- Concluding remarks

spm
1D



Topics

- Overview
- False positives
- Vector field statistics
- Reminder of SPM limitations
- Concluding remarks

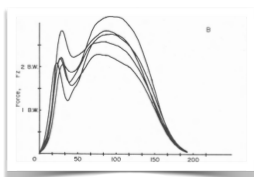
spm
1D





Scalar 1D

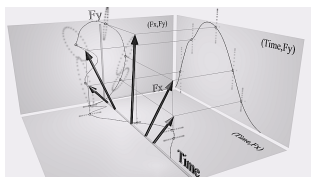
Univariate 1D



t tests
regression
ANOVA

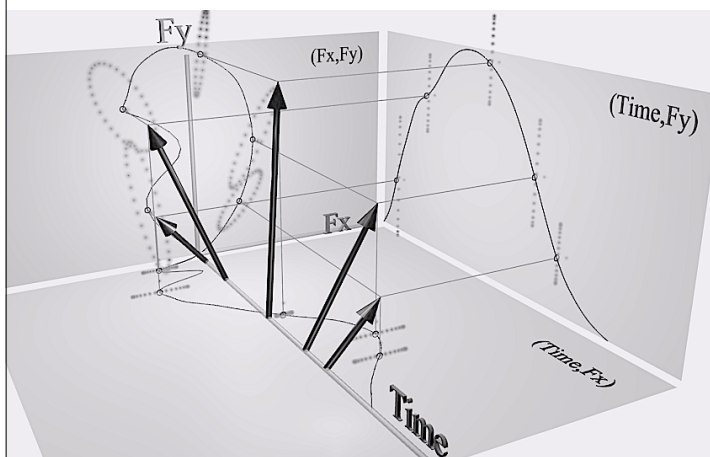
Vector 1D

Multivariate 1D

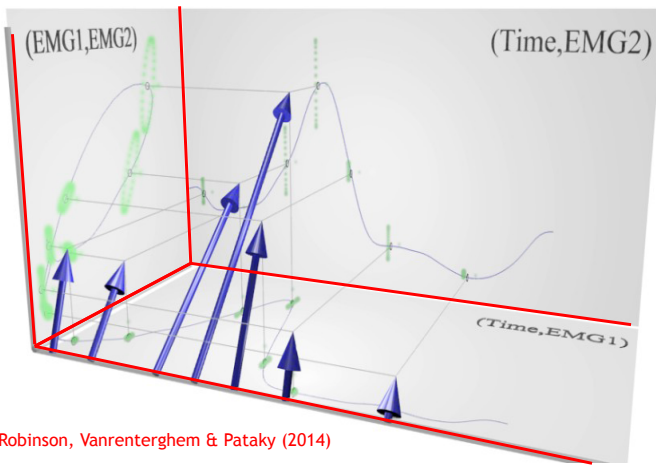


Hotelling's T^2 tests
CCA
MANOVA

Vector data: forces



Vector data: EMG



Topics

- Overview
- False positives
- Vector field statistics
- Reminder of SPM limitations
- Concluding remarks



False positives

Demo



Methods

- Assembled ~1000 vector trajectories from public datasets
- Estimated the median smoothness for different classes of data
 - Kinematics, dynamics, EMG
- Computed false positive rates for OD thresholds
 - Analytically
 - Validated using simulation

Results

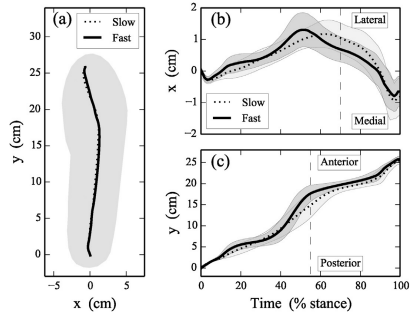
- Probability of a false positive:
 - One scalar trajectory
 - $p = 0.382$
 - One 3-component vector trajectory
 - $p = 0.764$
 - Two 3-component vector trajectories
 - $p = 0.945$

Topics

- Overview
- False positives
- Vector field statistics
- Reminder of SPM limitations
- Concluding remarks



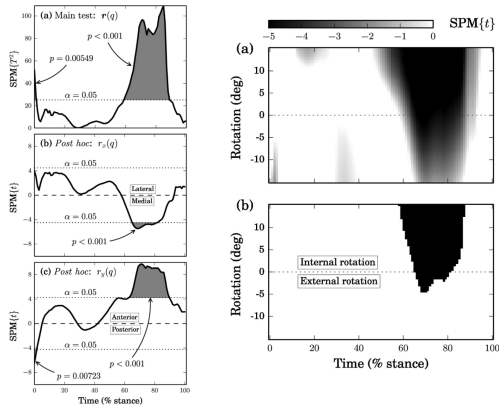
Example 1: COP



Pataky et al. (2014) JoB

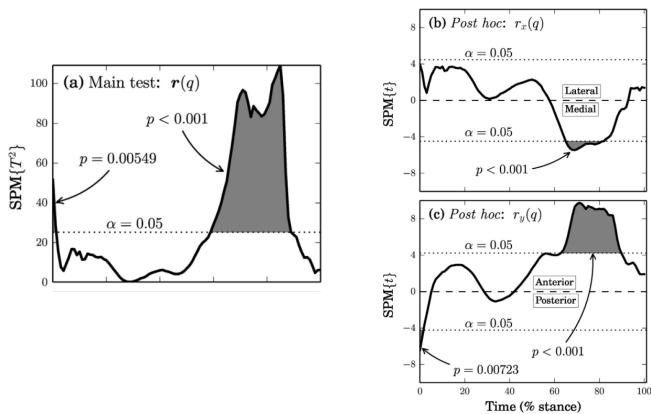
spm
1D

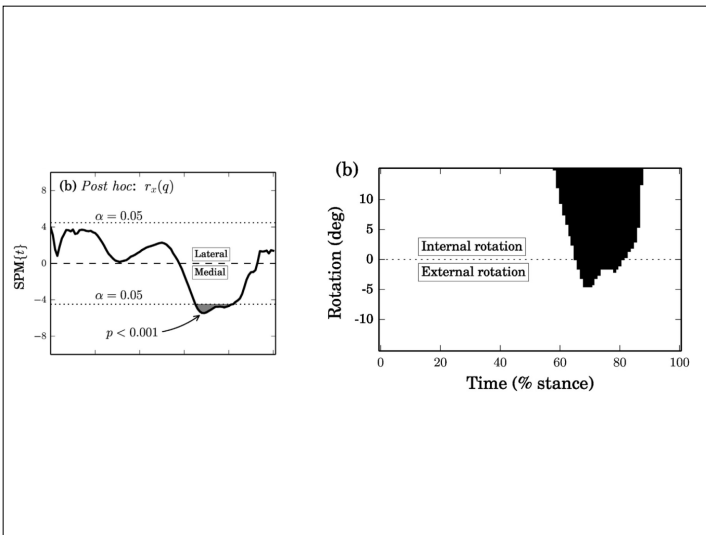
Hotelling's T^2 tests



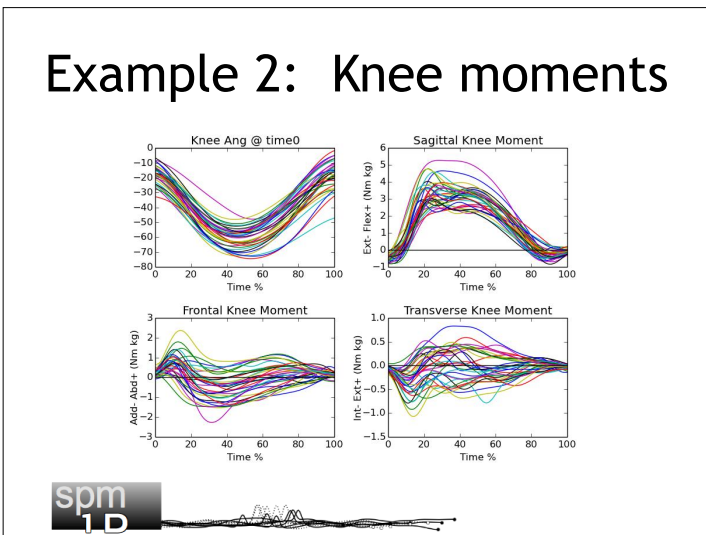
spm
1D

Hotelling's T^2 test

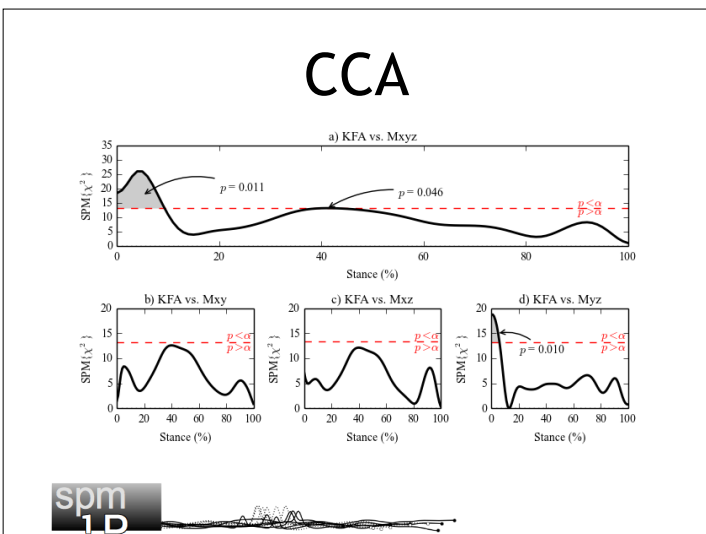




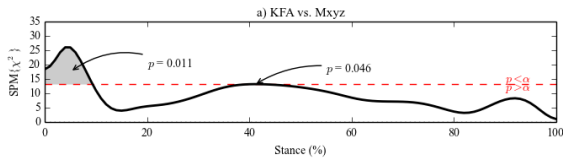
Example 2: Knee moments



CCA



CCA



Topics

- Overview
- False positives
- Vector field statistics
- Reminder of SPM limitations
- Concluding remarks

spm
1D

SPM limitations

- SPM has same limitations as other methods e.g. non-random sampling, non-blind experimentation, non-homologous data....
- Requires data registration (temporal normalization)
- SPM's procedures are more complex, especially RFT-based inference
- Accessibility (Python/Matlab programming)

spm
1D

SPM limitations



spm
1D



Topics

- Overview
- False positives
- Vector field statistics
- **Reminder of SPM limitations**
- Concluding remarks

spm
1D



Concluding remarks

- Support:
 - Website WWW.SPM1D.ORG
 - Handouts + notes + worksheets workshop
 - Todd, Mark, Jos
- Certificate
- Feedback

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1D