

EEG Spectral Fingerprints

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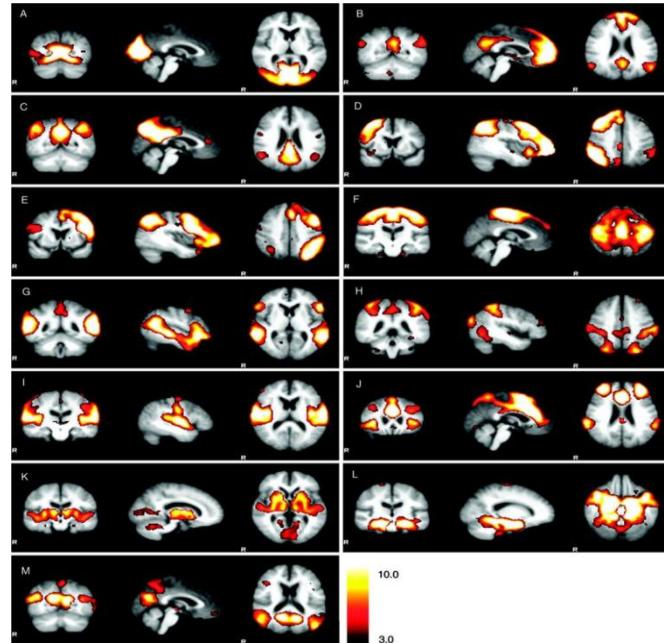
20.06.2018
58th Cracow School of Theoretical Physics, Zakopane

Goal of the study

- ▶ We want to develop a **reliable method** which allow us to see activity of **deep sources** of human brain and **dynamics of the networks** from EEG:
 - ▶ **optimal representation of EEG signal** that expose information about brain networks and deep sources reliably
 - ▶ discovering **specific and stable features** of each brain structure
 - ▶ ability to model brain functions as **dynamic systems**
(thanks to high time resolution of the EEG signal)

Support

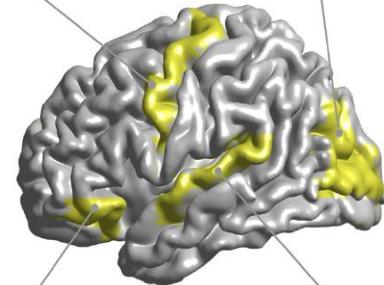
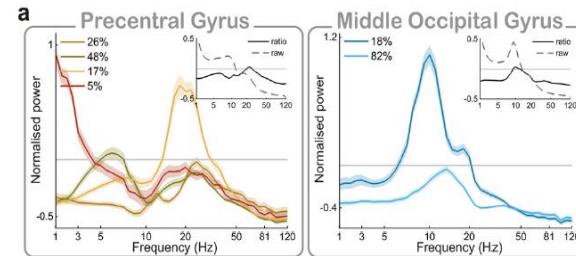
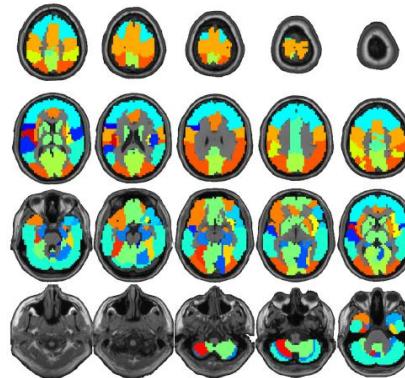
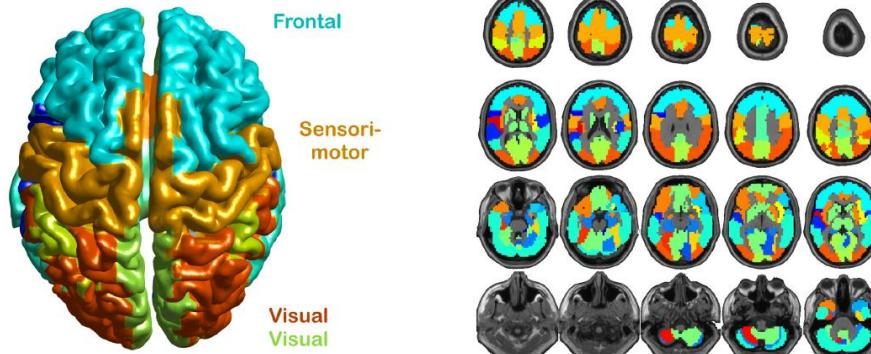
- ▶ Different stable **resting state networks** can be found in the human brain
 - ▶ Damoiseaux et al. 2008 (*Cerebral Cortex*)
- ▶ **Natural frequencies** exist for different brain regions
 - ▶ Rosanova et al. 2009 (*J. Neuroscience*)
- ▶ Patterns are stable and gathered across **core patterns**
 - ▶ Krienen et al. 2014 (*Phil. Trans. of the Royal Society B*)
- ▶ Feedforward and feedback **loops** in primates visual cortex (delta, alpha)
 - ▶ van Kerkoerle et al. 2014 (*PNAS Neuroscience*)
- ▶ Sensory or lower-to-higher order **hierarchy** in primates
 - ▶ Murray et al. 2014 (*Nature Neuroscience*)
- ▶ Human brain **does not follow** that hierarchy
 - ▶ Mellem et al. 2017 (*J. Neurophysiology: Neural Circuits*)
- ▶ **Gene expression** is correlated with functional brain networks
 - ▶ Richiardi et al. 2015 (*Science: Brain Networks*)

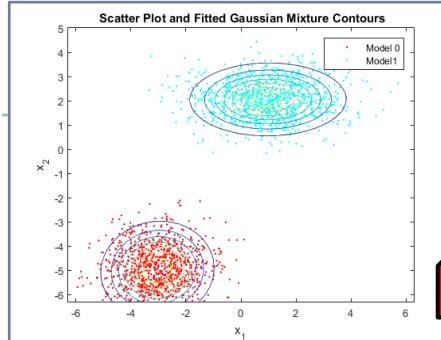
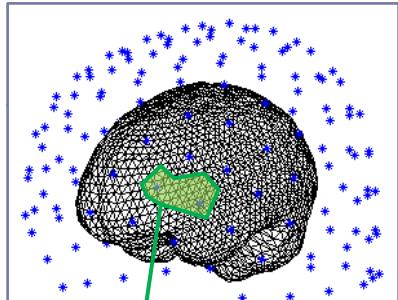


(Damoiseaux et al. 2008)

Brain Fingerprinting

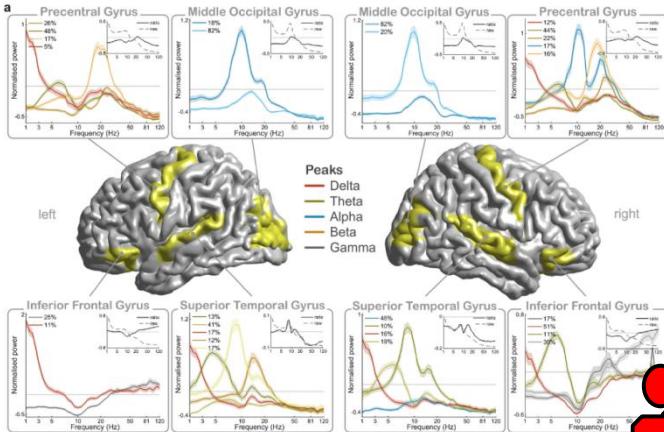
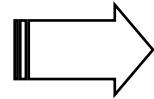
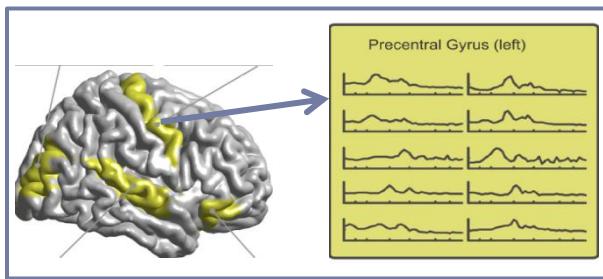
- ▶ Brain fingerprinting: spectral fingerprints based on approach in paper of A. Keitel i J. Gross, „**Individual human brain areas can be identified from their characteristic spectral activation fingerprints**”, *PLoS Biol*, t. 14, nr 6, s. e1002498, 2016.
- ▶ Model brain activity in each ROI as a **Gaussian Mixture Model** based on **spectral** content of MEG signal





Single
subject

$d \in \text{ROI}$



Group model



The Data

Datasets

	N	Spectral Fingerprints	Ranks	Opt. Number of modes	Networks
MEG reproduction	22	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
MEG - gamma	22	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
MEG – reduced number of trials (on average 279/466)	22	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
EEG Age-ility (RIKEN/worse coregister)	22	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
EEG Age-ility (fixed forward model)	22	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
EEG N=12 (data from Toruń)	12	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

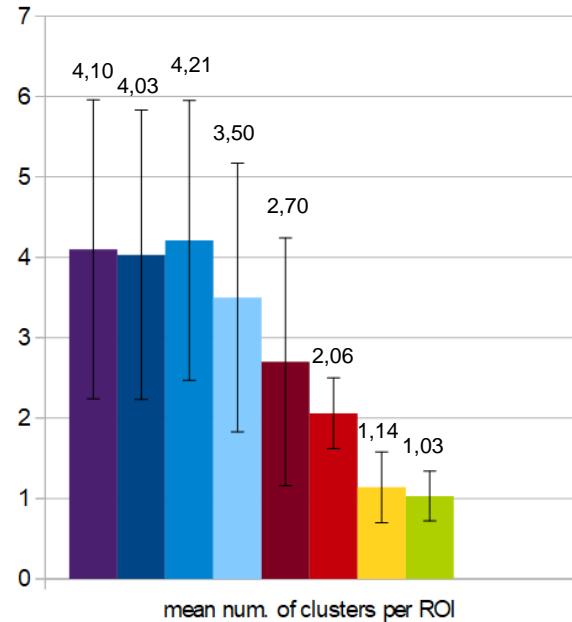
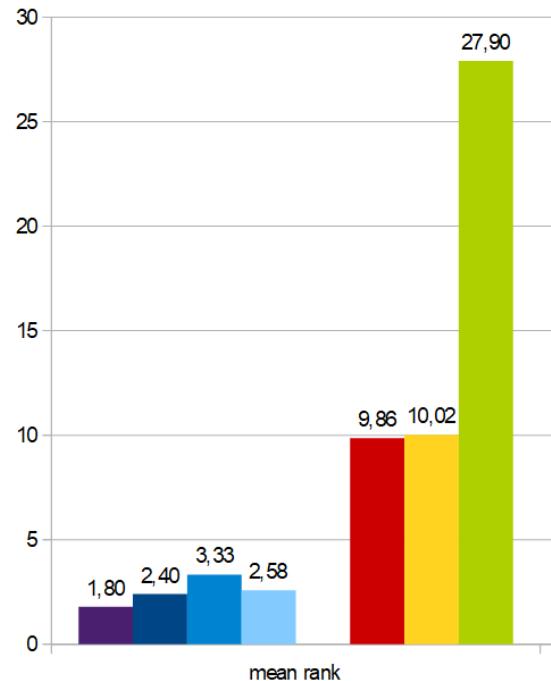
MEG : rest, 466 ± 20 , eyes open, N=22, perfect coregistration

EEG Age-ility: rest, 118 ± 7 s, eyes closed, N=22, poor (manual) coregistration of EEG to MRI

EEG N=12: rest, 460 ± 32 s, eyes open, N=22, fair (< 1cm per electrode) coregistration of EEG to MRI

Comparison between main results

- MEG (orig)
- MEG (reprod.)
- MEG (-gamma)
- MEG (reduced trials)
- EEG N=7
- EEG N=12
- EEG Agility N=22
- EEG Agility N=22 (worse coregister)

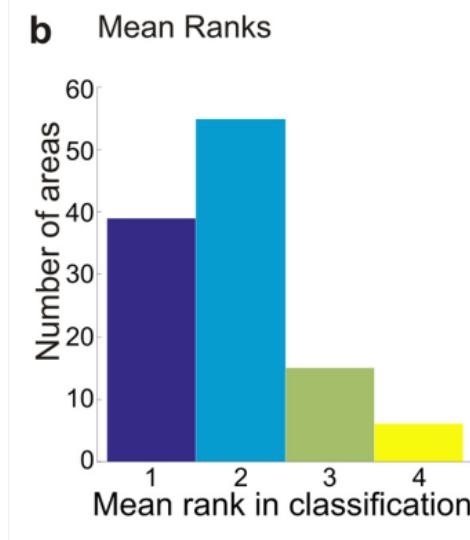




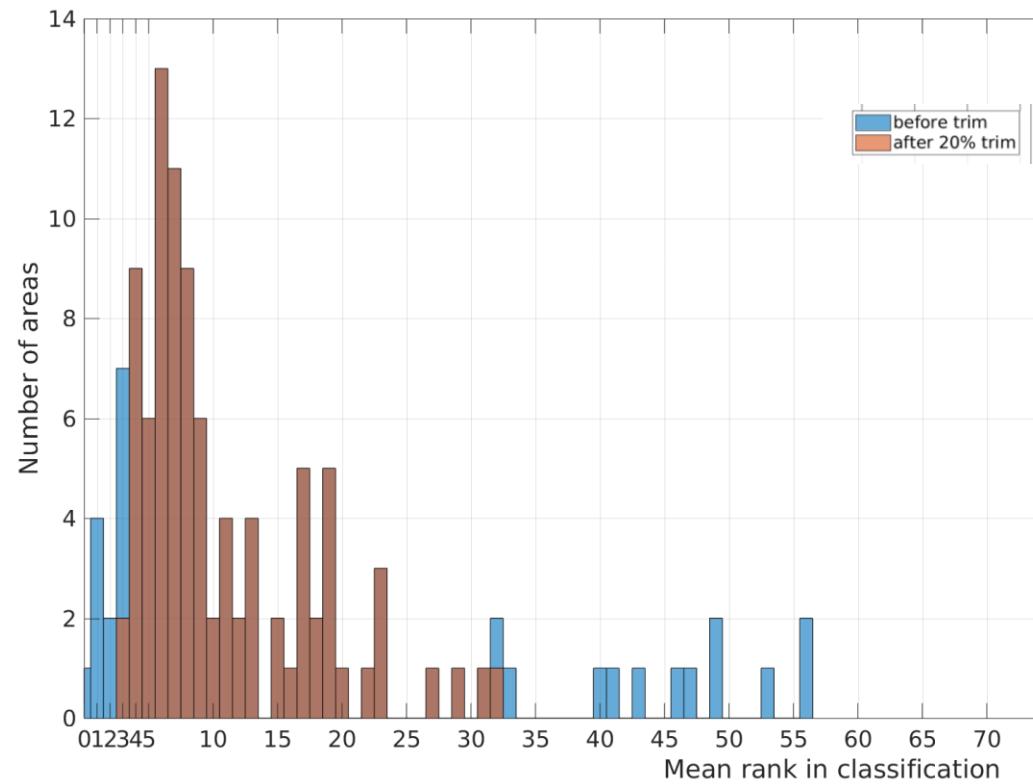
EEG Age-ility (N=22, fixed forward model)

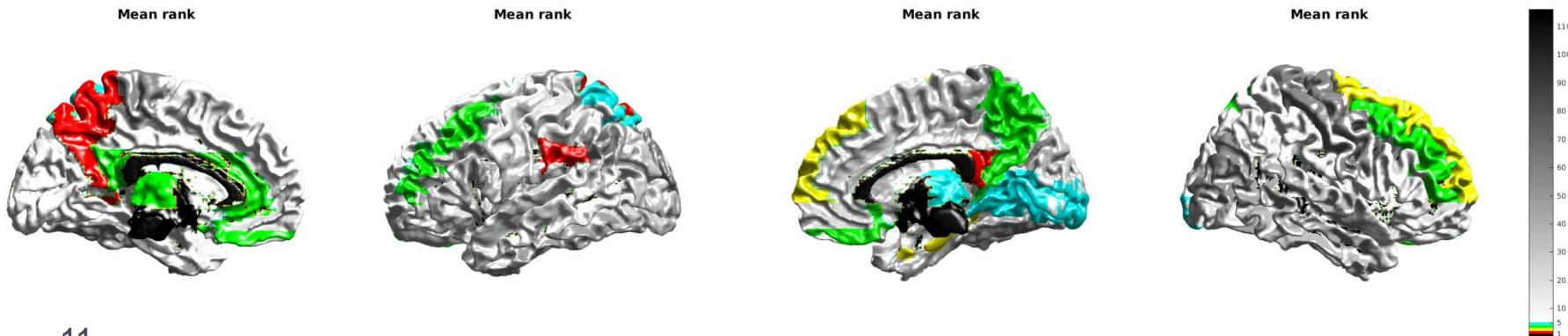
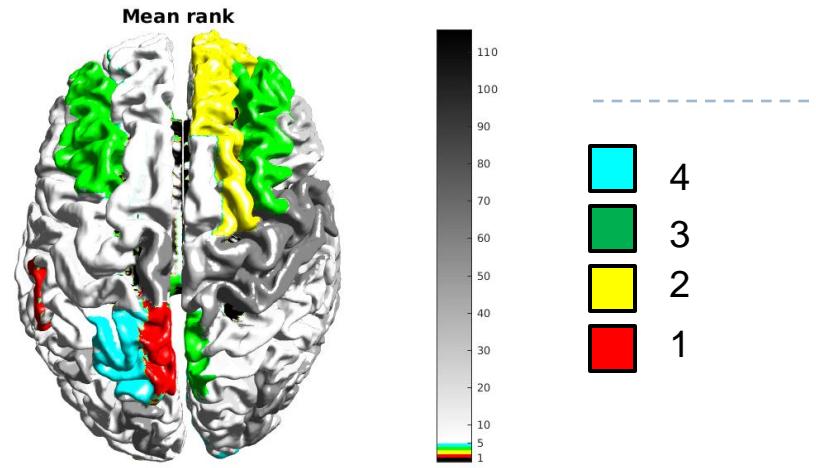
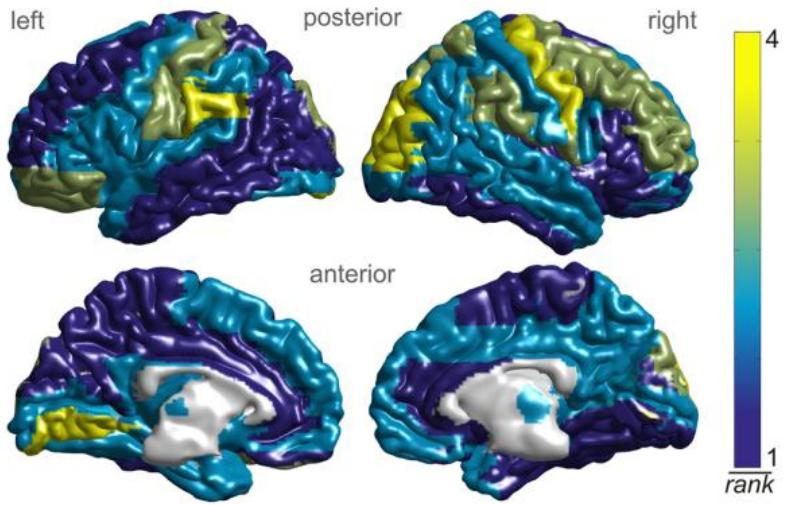


Left: Keitel & Gross
(GoodGrid, GoodFixCINum).
Mean rank = 1.8

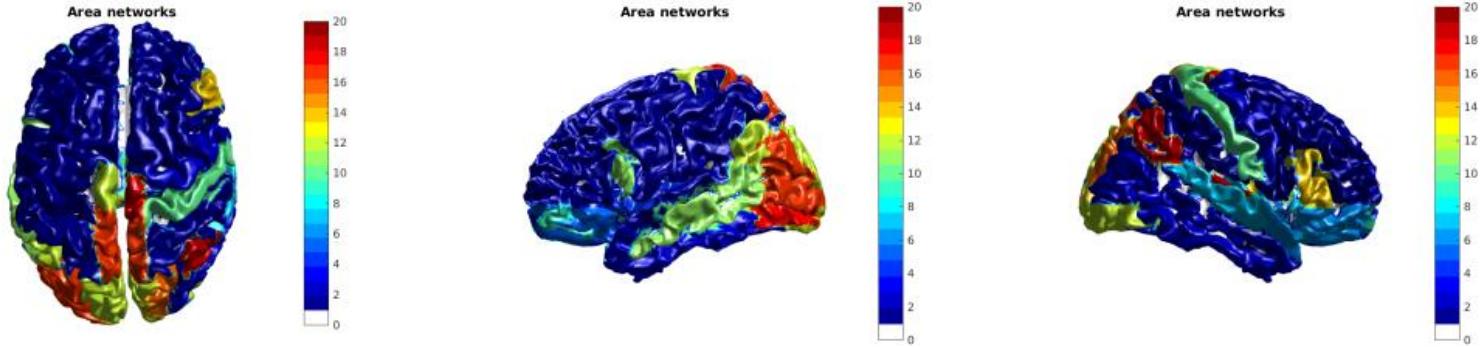
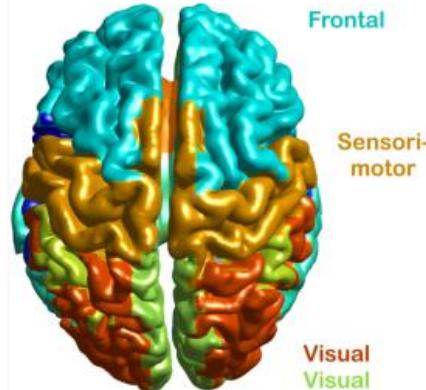


Right: EEG N=22 Agility
Mean rank = 10.02





Left-up: Keitel & Gross, other figures: EEG N=22 Agility (GoodGrid, GoodFixCINum).



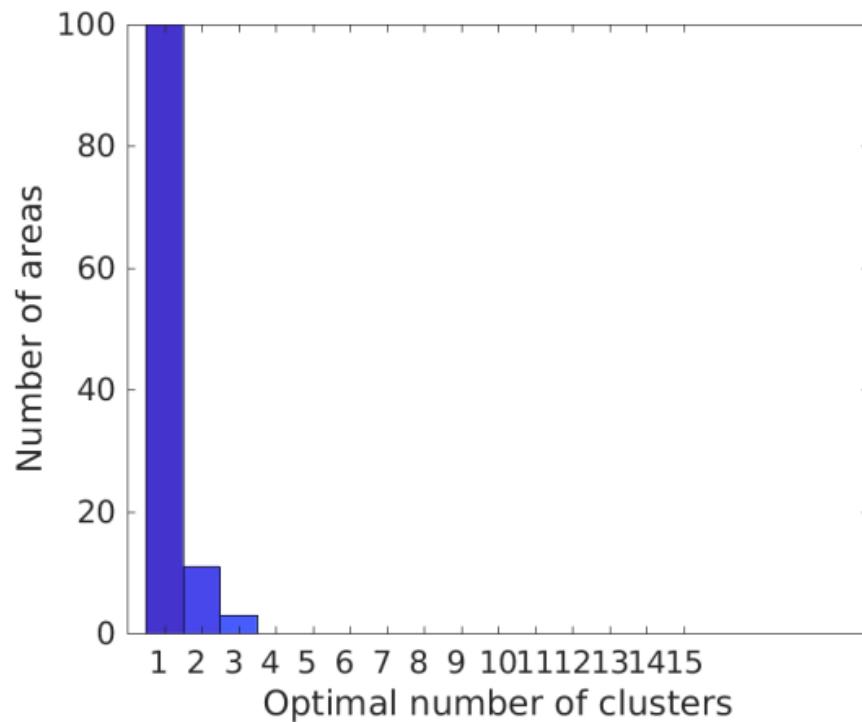
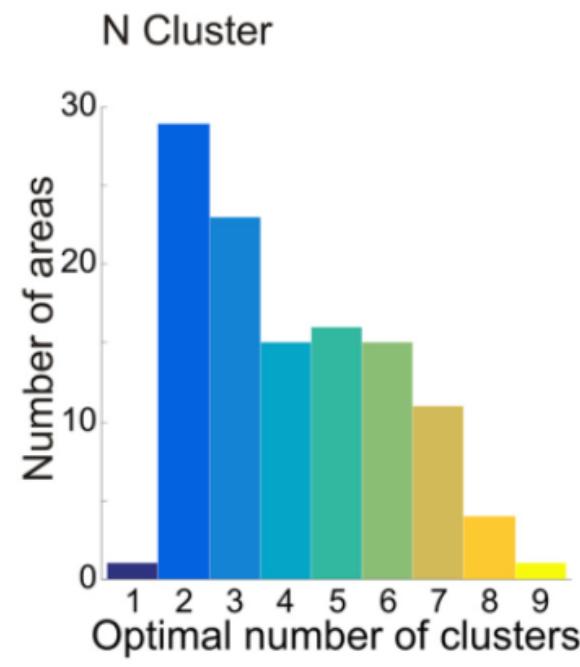
Left: Keitel & Gross

Agility(GoodGrid, GoodFixCINum).

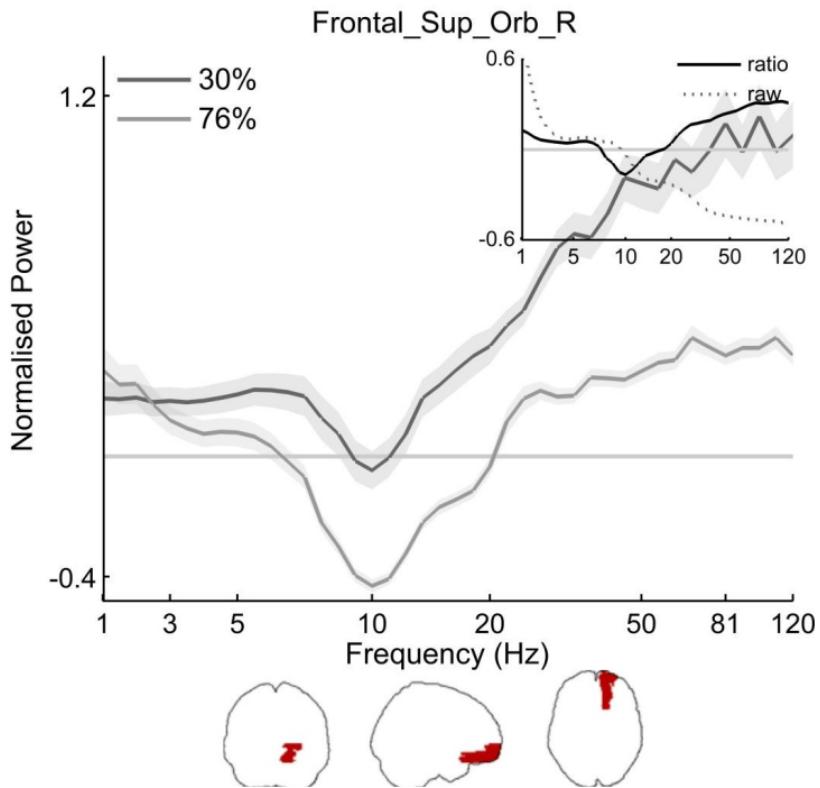
Avg. num. of clusters = 4.10 ± 1.86

Right: EEG N=22

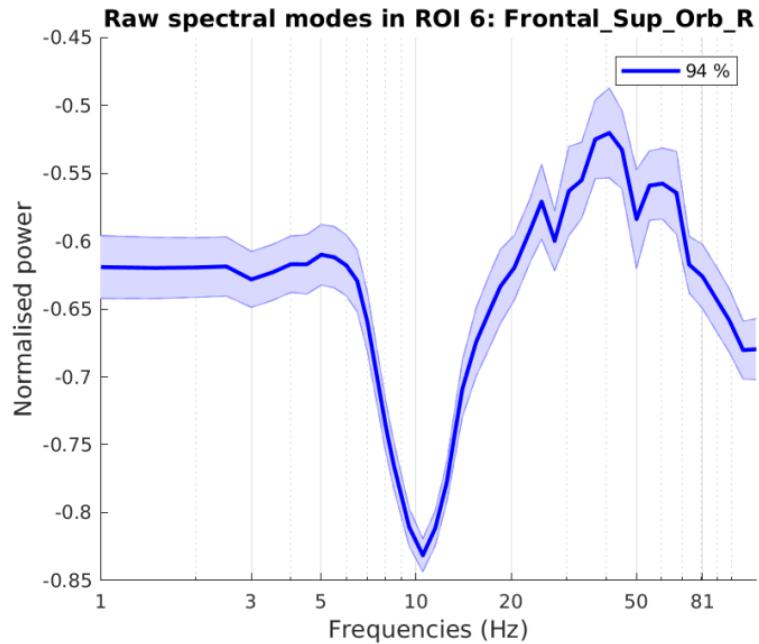
Avg. num. of clusters = 1.14 ± 0.44



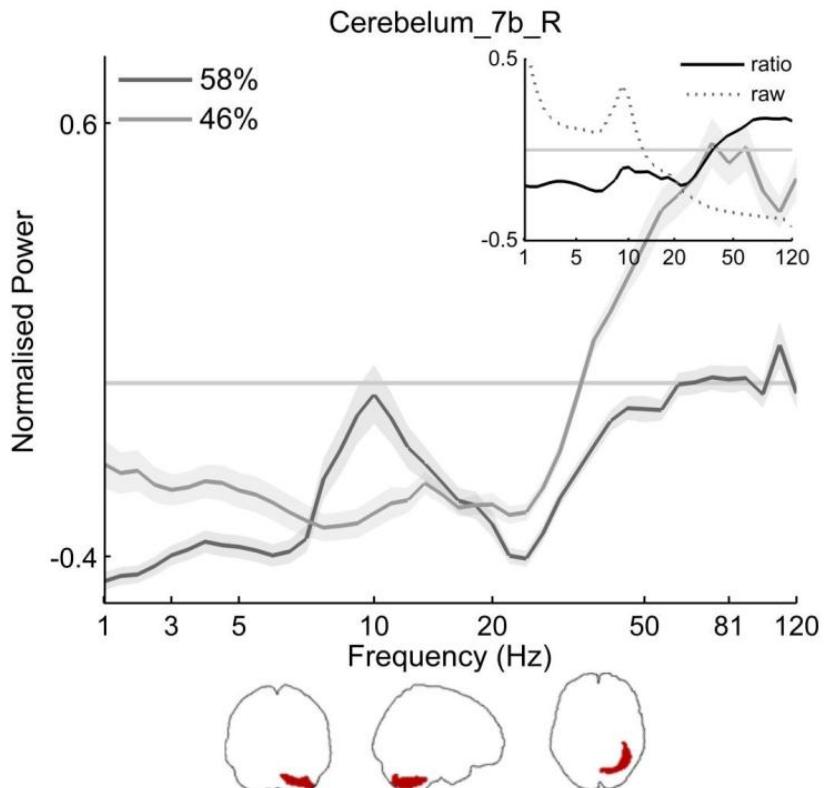
Keitel & Gross, 2016



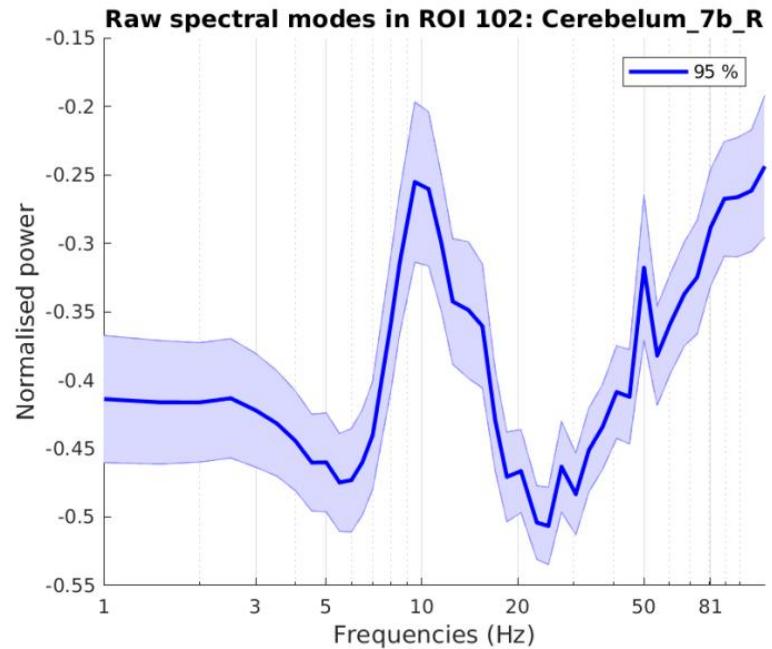
EEG Age-ility



Keitel & Gross, 2016

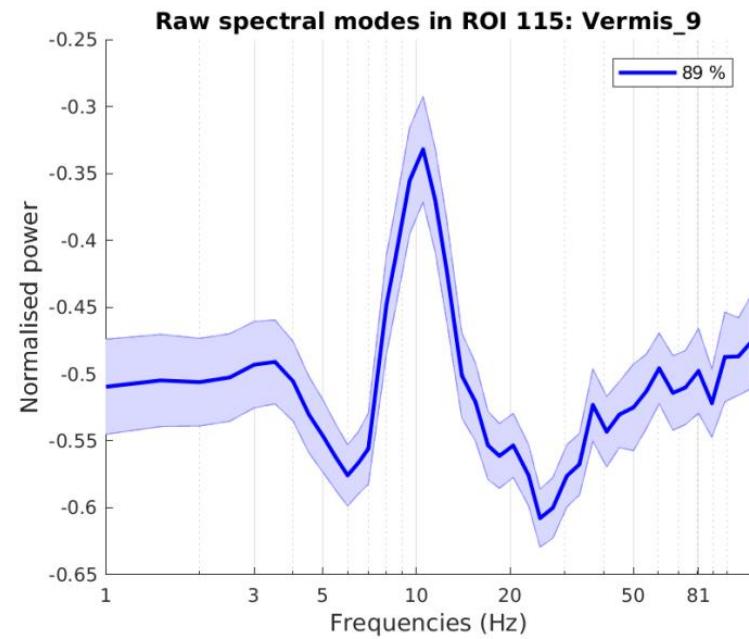
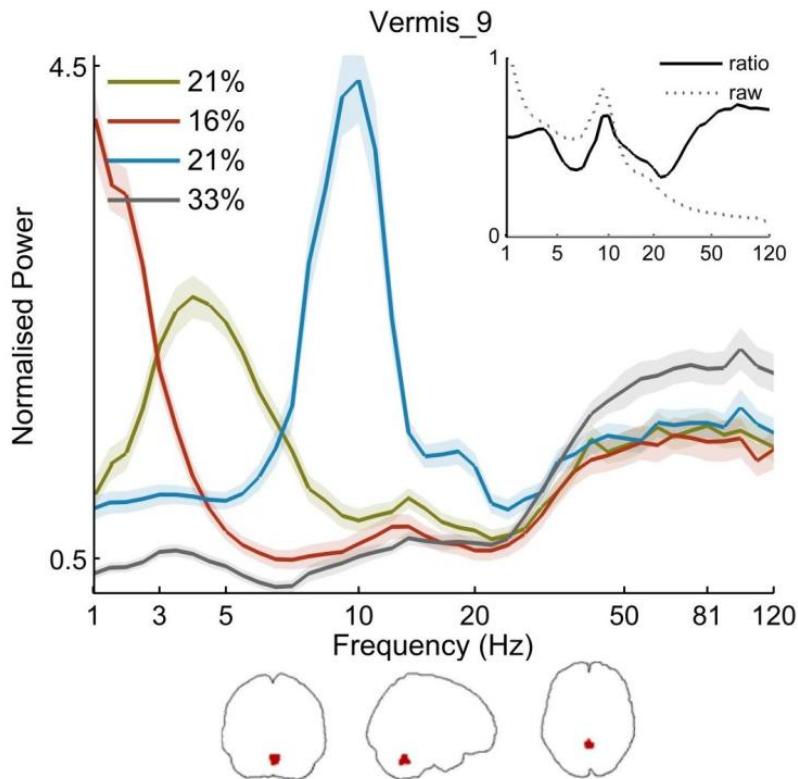


EEG Age-ility



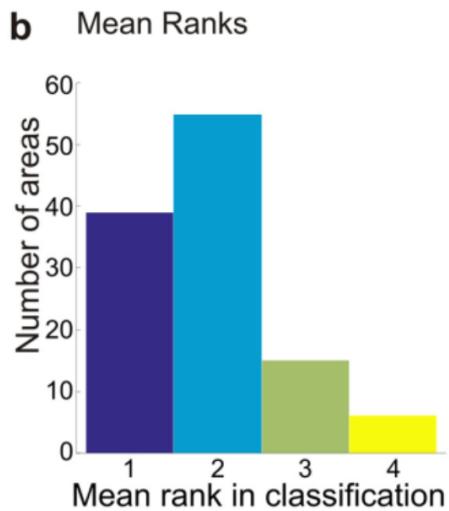
Keitel & Gross, 2016

EEG Age-ility

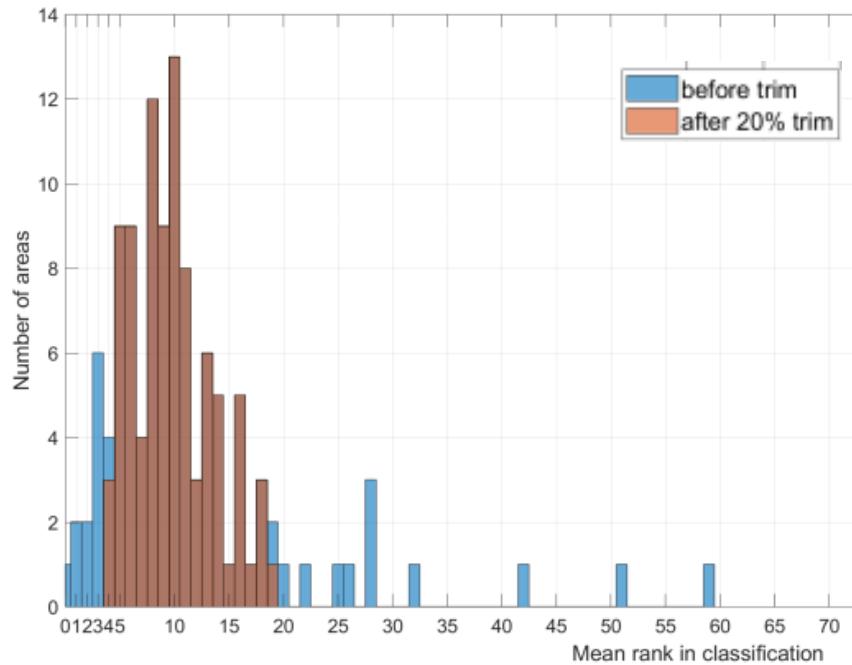


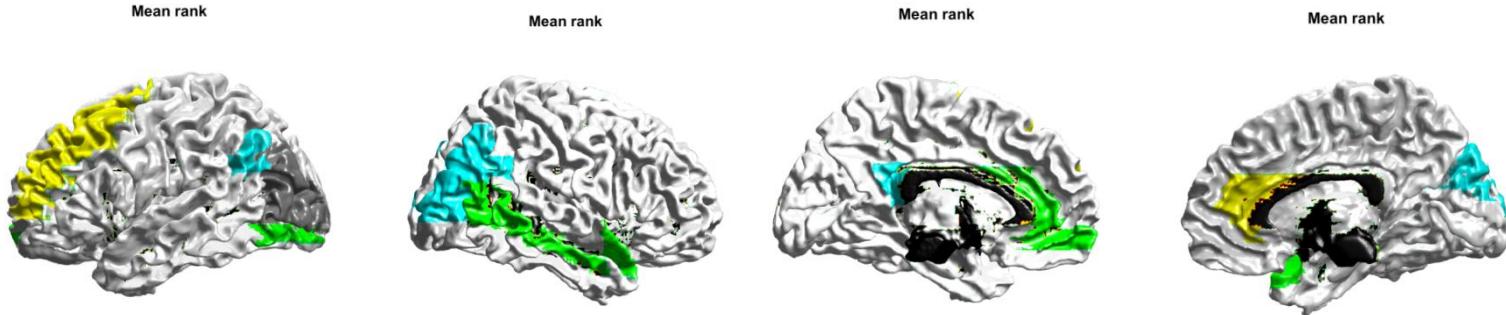
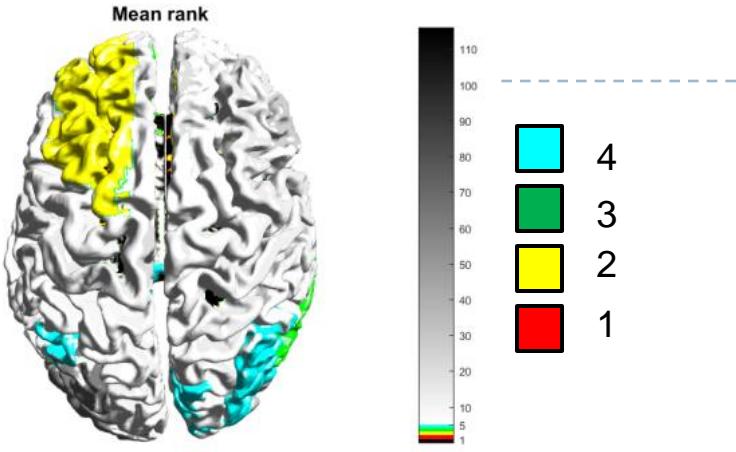
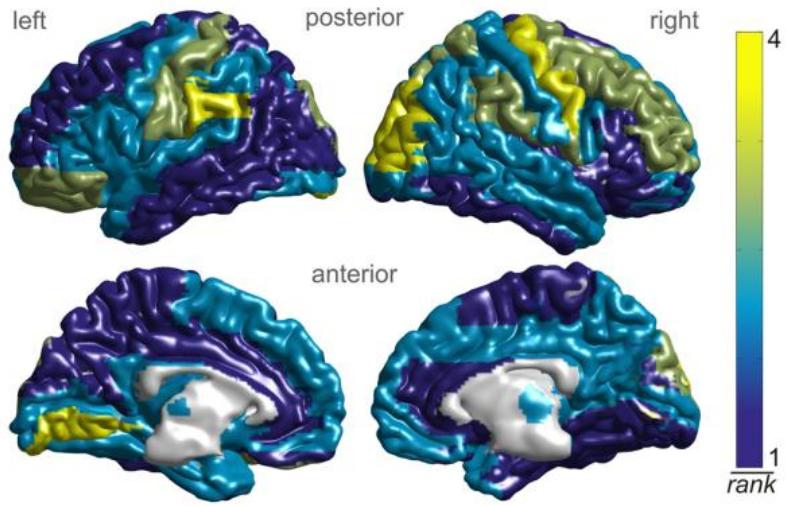
EEG N=12 (Toruń)

Left: Keitel & Gross
Mean rank = 1.8

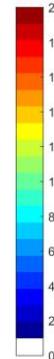
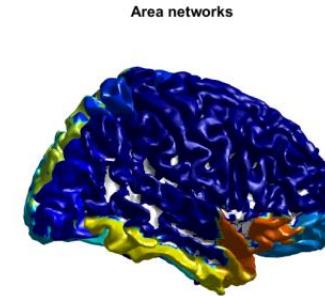
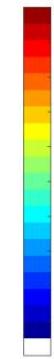
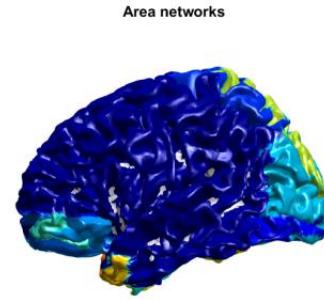
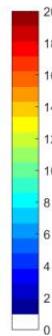
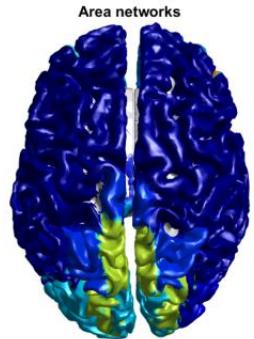
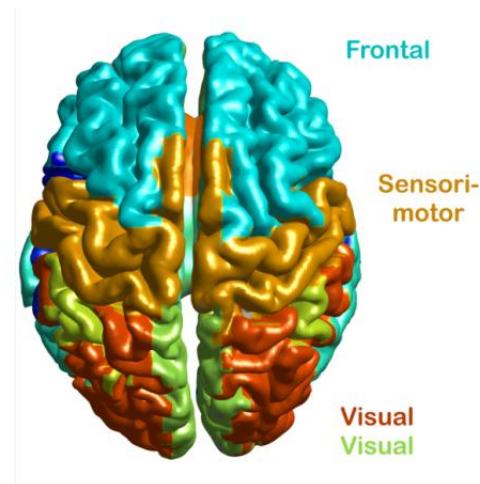


Right: EEG N=12 Torun.
Mean rank = 9.86



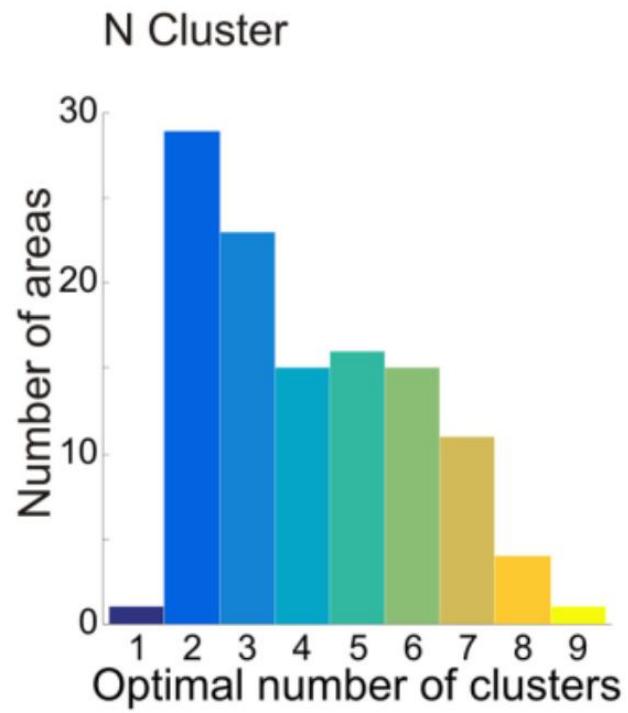


Left-up: Keitel & Gross, other figures: EEG N=12 Torun



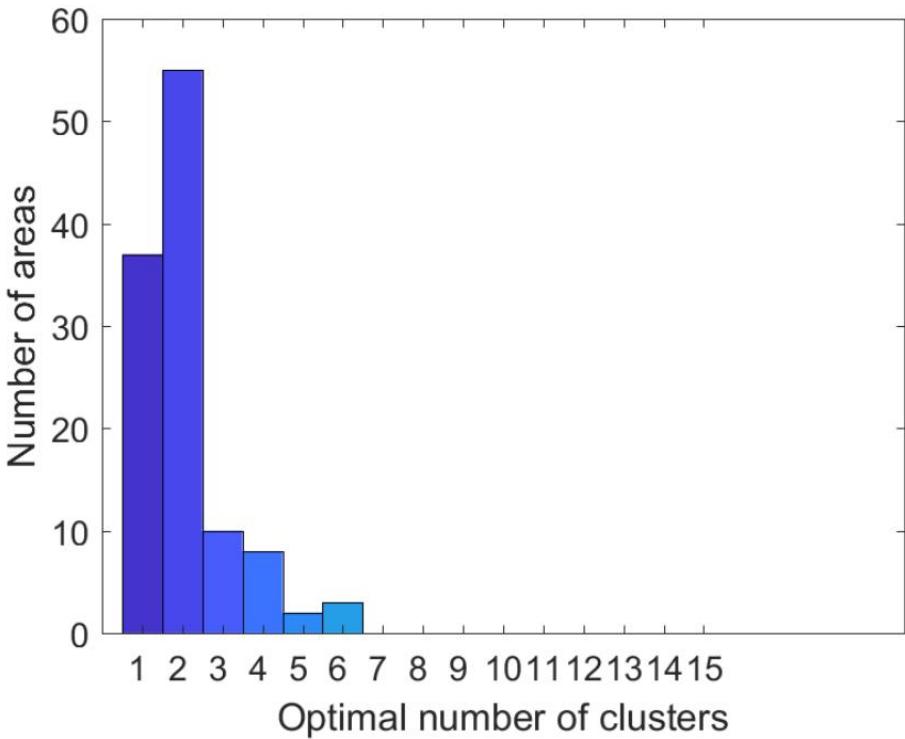
Left: Keitel & Gross

Avg. num. of clusters = 4.10 ± 1.86

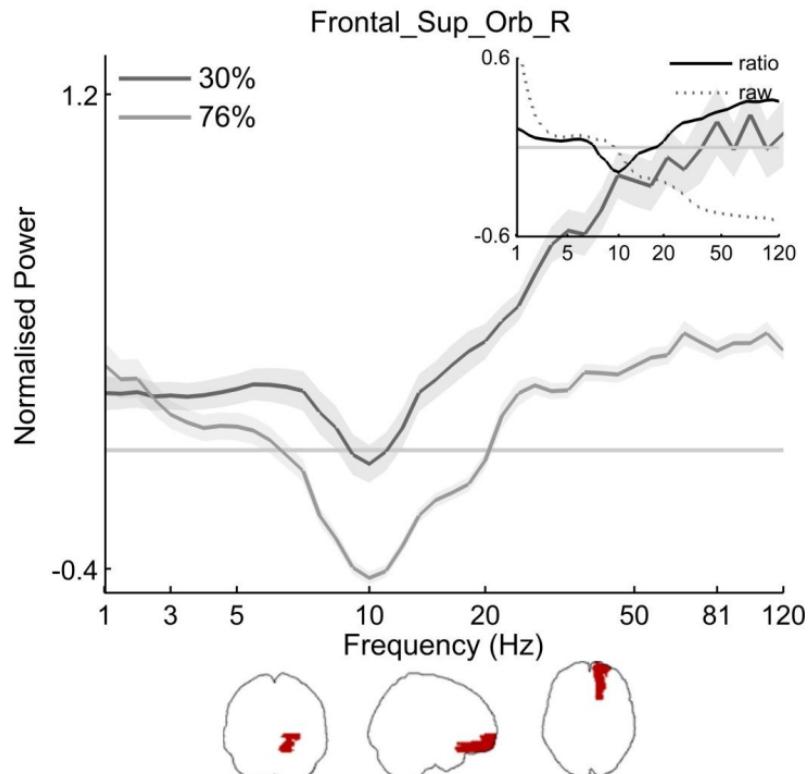


Right: EEG N=12 Torun.

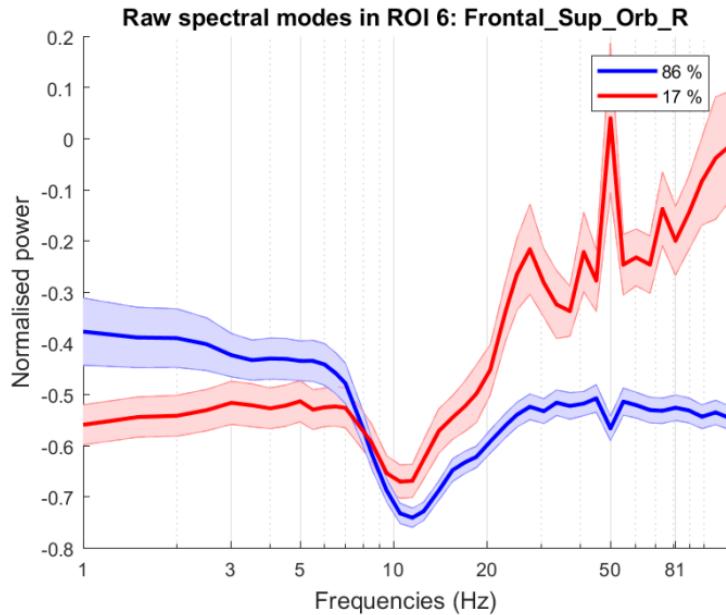
Avg. num. of clusters = 2.06 ± 1.13

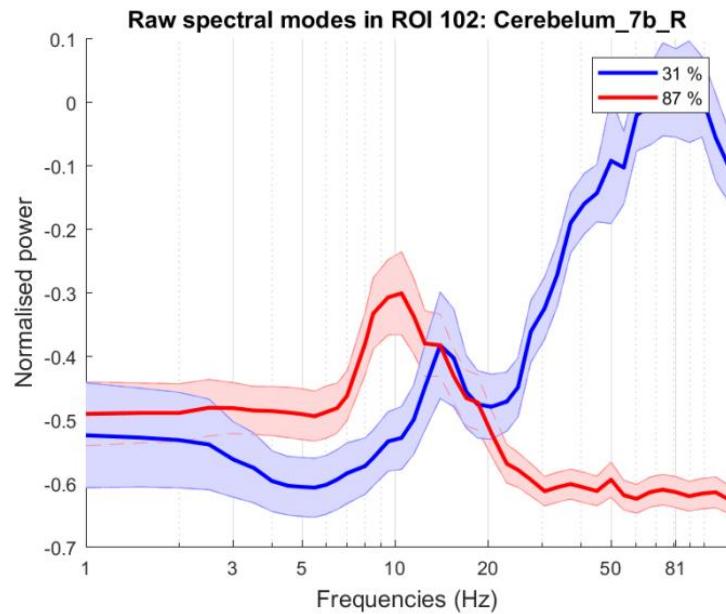
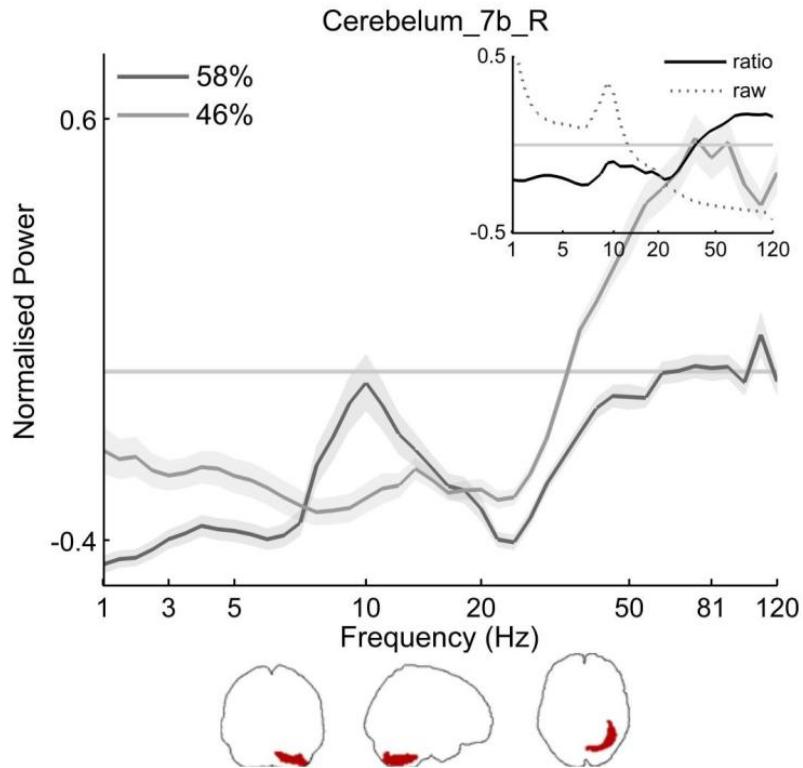


Keitel & Gross, 2016

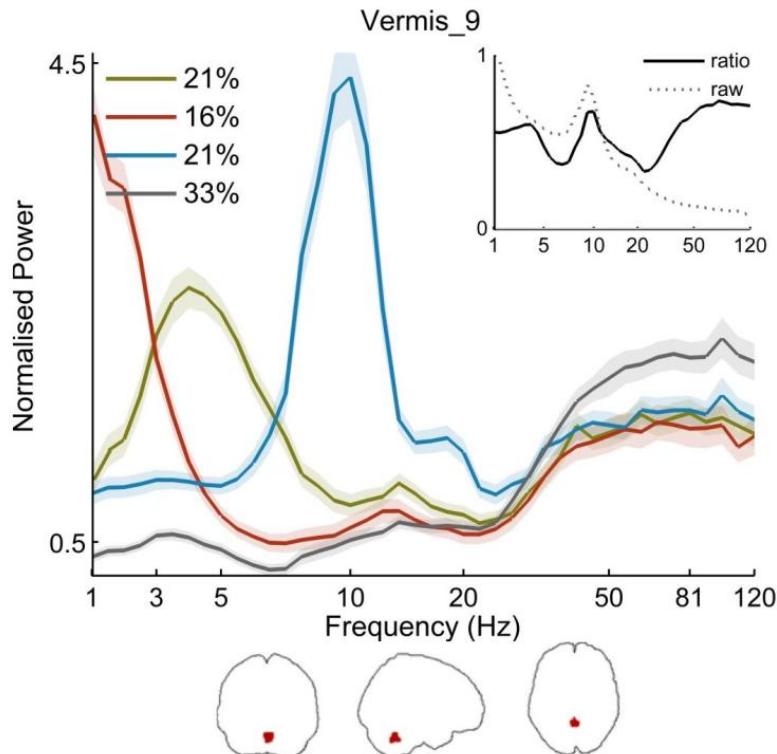


EEG (Toruń)





Keitel & Gross, 2016



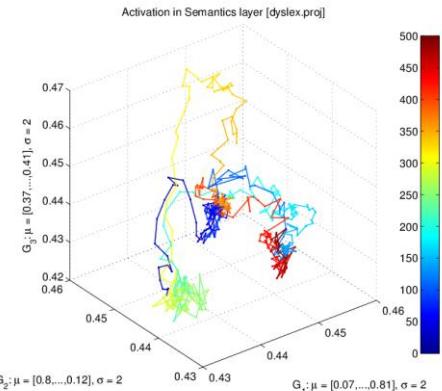
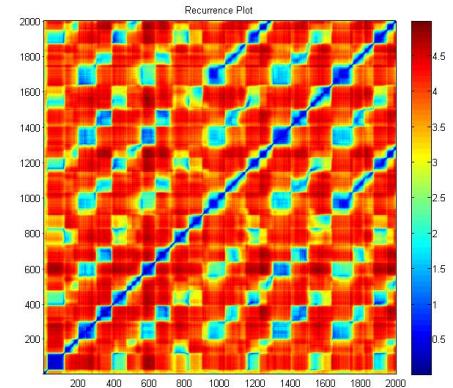
EEG (Toruń)

Conclusions

- ▶ **Spectral fingerprinting** is a procedure that allows to discover spectral dynamics of each ROI from functional data (MEG, EEG)
- ▶ Distant **homologue areas** are connected together
(similarity is not an artifact caused by short distance similarity and source mixing)
- ▶ **MEG**: ROI spectral fingerprints are specific so we can classify them
- ▶ **EEG**: ROI recognition accuracy basing on its spectral fingerprint is well above chance level (but still some improvement is needed)

Plans for the future

- ▶ Prove that EEG spectral fingerprints :
 - ▶ Are specific for each brain region and stable across participants
 - ▶ Can be used to differentiate ROIs (classification, ranks)
- ▶ Calculate spectral fingerprints on **resting-state EEG N=22 or more**
- ▶ Spectral fingerprints from EEG recording with **128 electrodes or more**
- ▶ Improve inverse problem solution with MV-PURE approach
(Piotrowski, Nikadon, Gutierrez [arXiv:1712.02997 \[eess.SP\]](https://arxiv.org/abs/1712.02997))
- ▶ Check whether spectral fingerprints exhibit **attractor structure**
- ▶ Study **trajectories in higher-dimensional spaces**



Contributions

- ▶ **prof. Włodzisław Duch, dr Joanna Dreszer**
 - ▶ for finding an idea and setting Spectral Fingerprints as one of the direction of research
- ▶ **Jakub Wojciechowski**
 - ▶ ongoing help with analysis of the methods and results from Keitel & Gross 2016 paper, preprocessing of EEG datasets
- ▶ **Jan Nikadon and dr Tomasz Piotrowski**
 - ▶ help with generating forward model for source reconstruction
- ▶ **prof. Andrzej Cichocki**
 - ▶ substantial part of work was done during the internship in RIKEN, BSI, Japan under professor's supervision and inspiration
- ▶ **Dr Anne Keitel & prof. Joachim Gross**
 - ▶ For providing access to the data and advices on know-how



- **Nicolaus Copernicus University** in Poland
 - Faculty of Physics, Astronomy and Informatics
 - Department of Informatics
 - (pol. Katedra Informatyki Stosowanej)



NICOLAUS COPERNICUS
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and Informatics

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Informatyki
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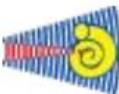
Neurocognitive Laboratory



FACULTY OF PHYSICS,
ASTRONOMY AND INFORMATICS



CENTRE FOR MODERN
INTERDISCIPLINARY
TECHNOLOGIES



INSTITUTE OF PHYSIOLOGY
AND PATHOLOGY OF HEARING



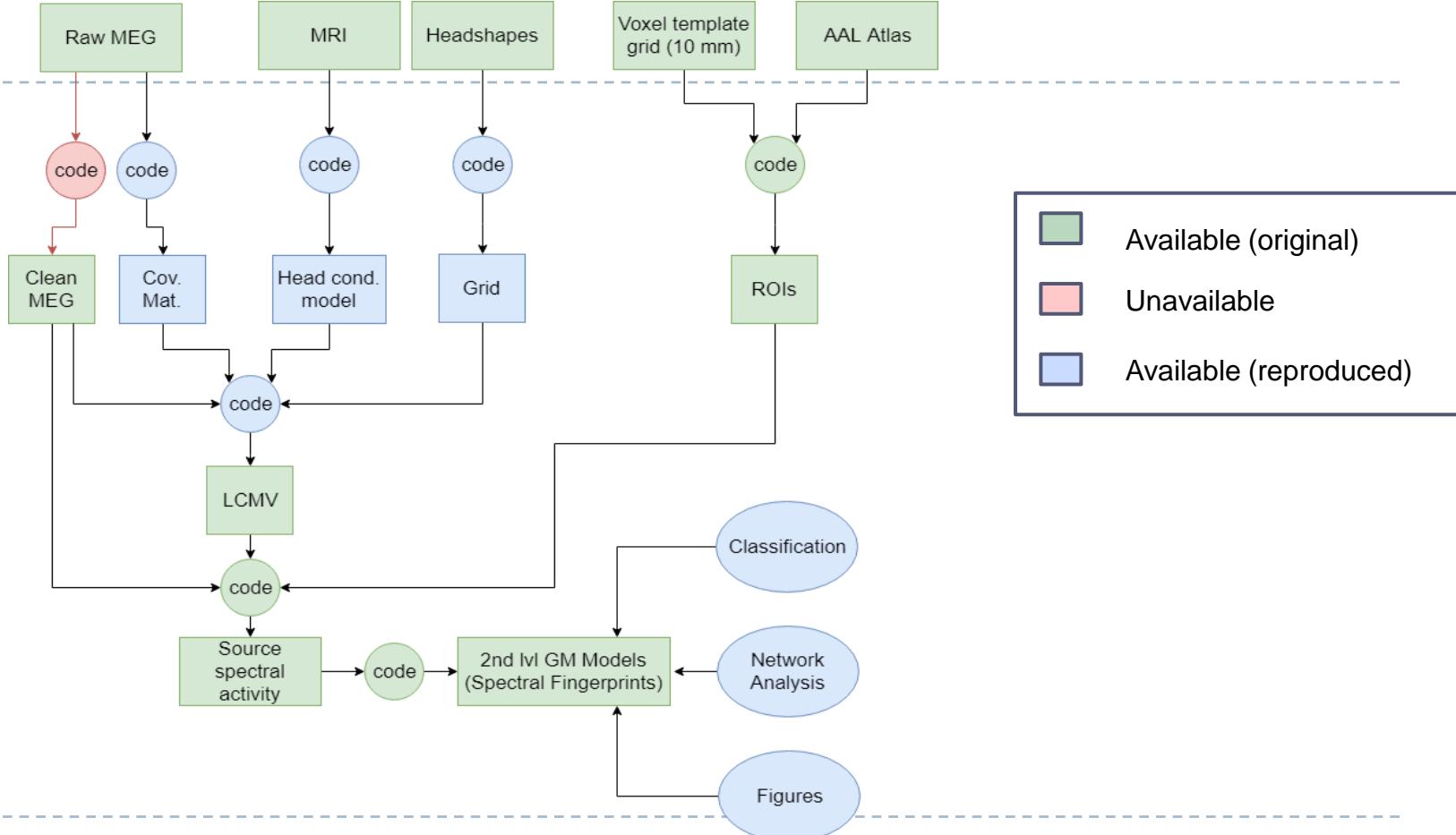
„In search of the sources of brain's cognitive activity”.

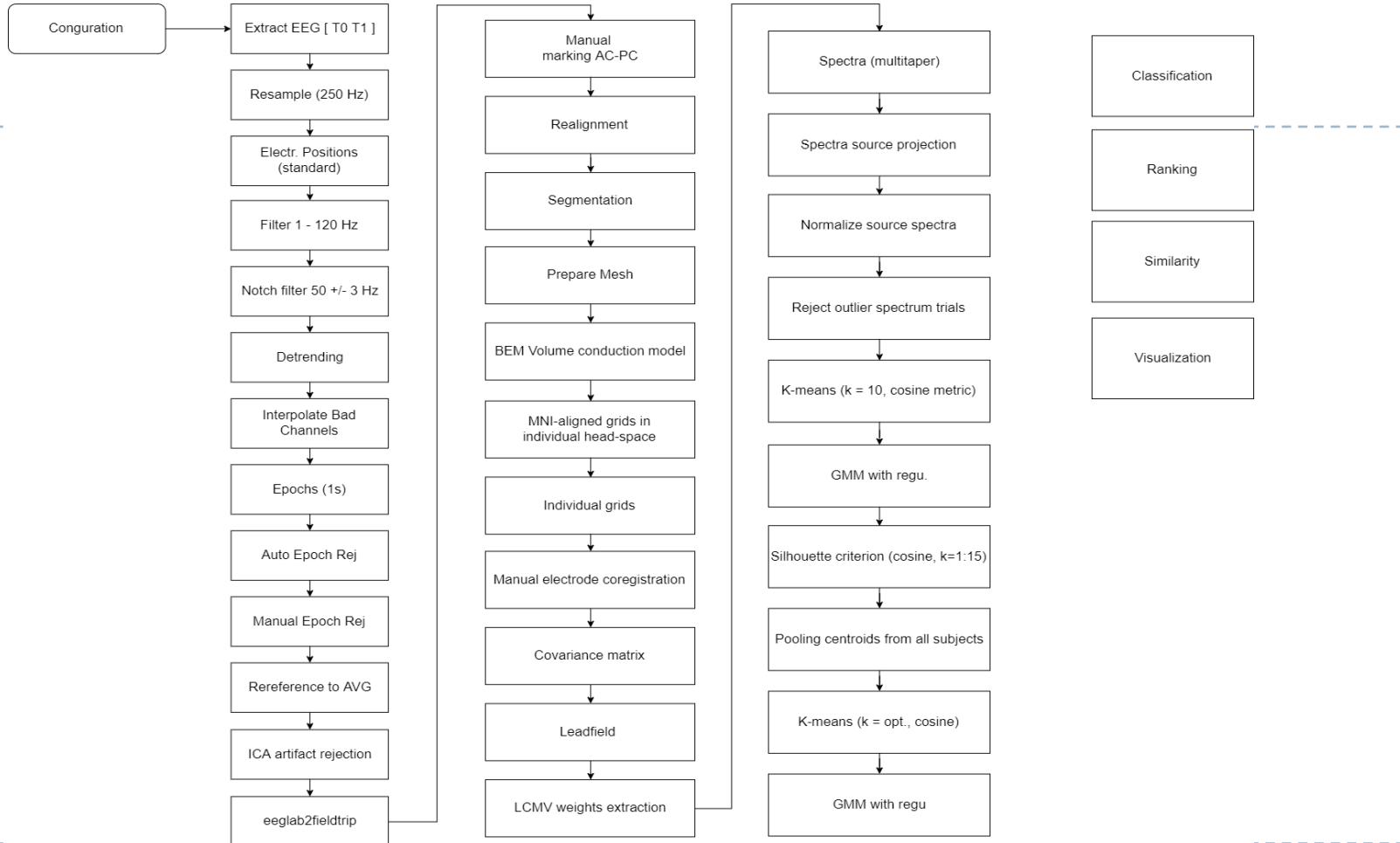
This study was supported by the National Science Center Poland
(UMO-2016/20/W/NZ4/00354) grant awarded to Prof. dr hab. Andrzej Cichocki.



Thank you !

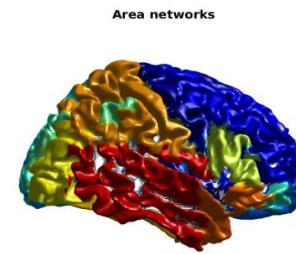
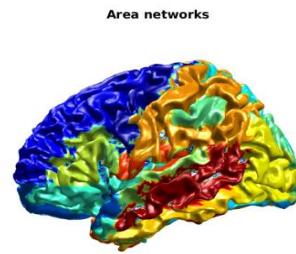
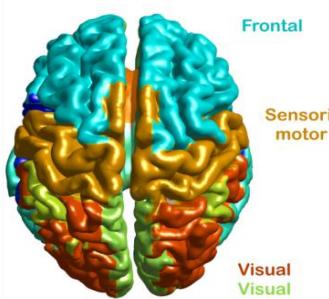
Diagram of code availability (updated 24.01.2018)





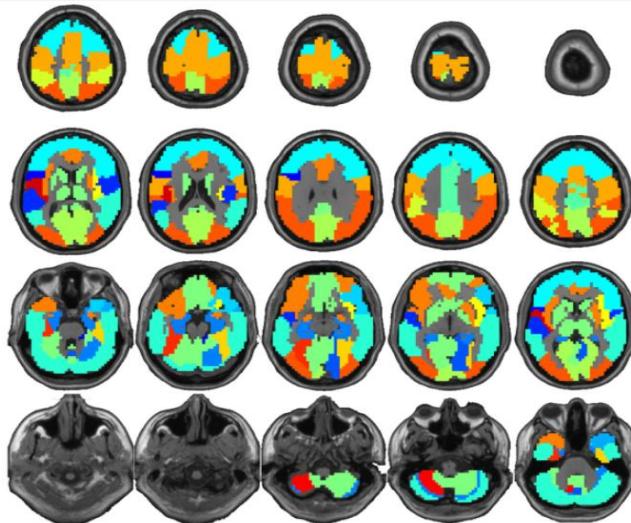
MEG reproduction results – ROI similarity (network analysis)

Left-up: Keitel & Gross, other figures: Reproduction.

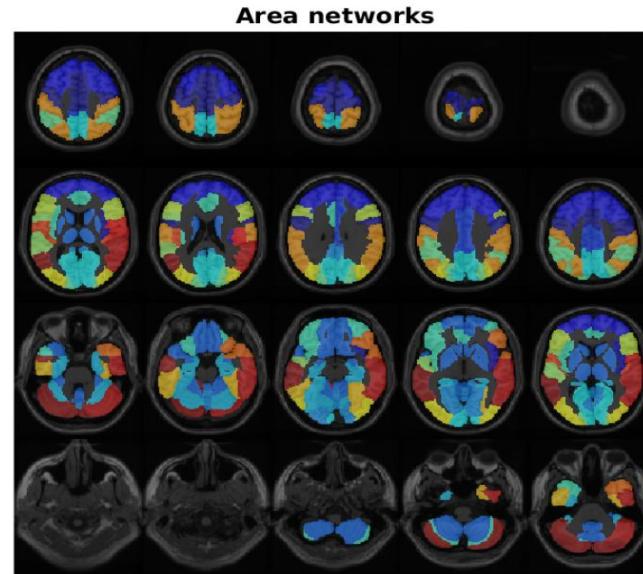


MEG reproduction results – ROI similarity (network analysis)

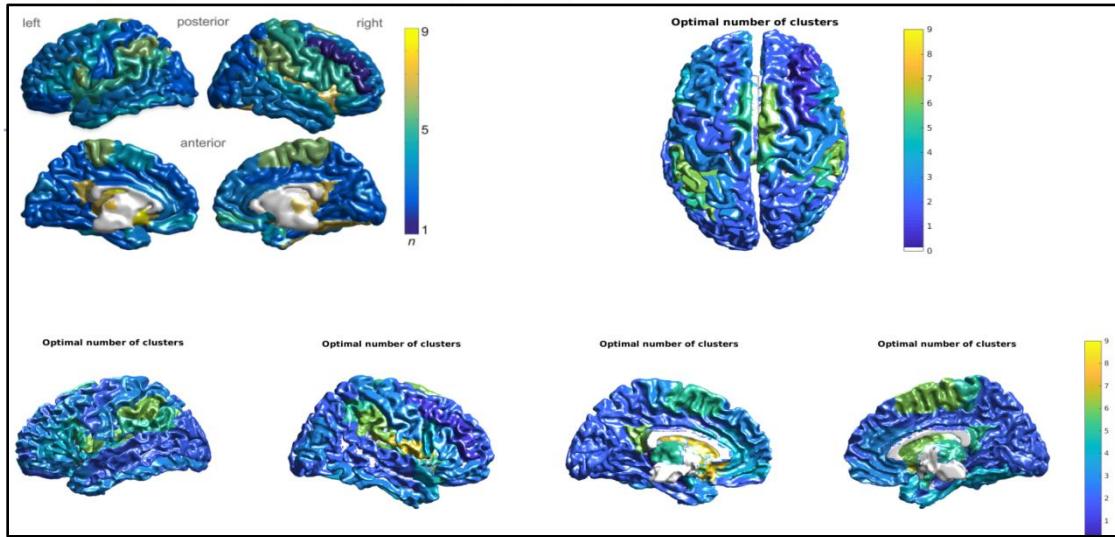
Left: Keitel & Gross



Right: Reproduction.



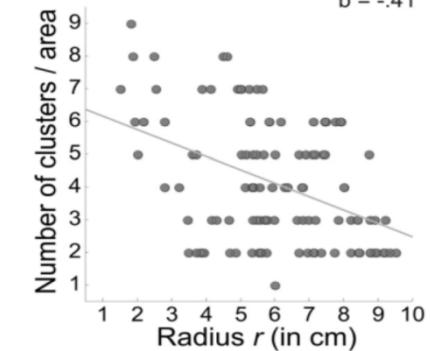
Left-up: Keitel & Gross, other figures: Reproduction.



Left: Keitel & Gross

$$R^2 = 0.18, F = 24.15, p < 0.001$$

$$b = -.41$$

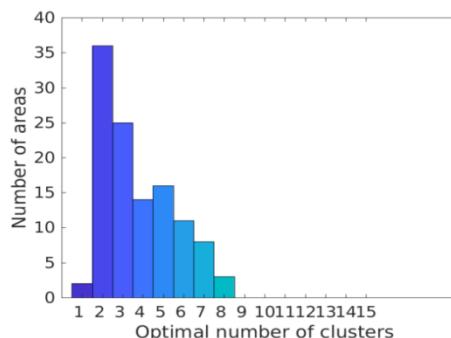
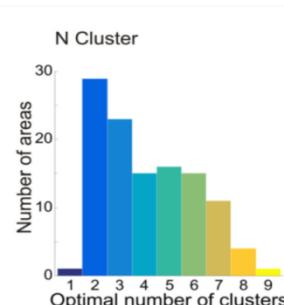


Left: Keitel & Gross

$$\text{Avg. num. of clusters} = 4.10 \pm 1.86$$

Right: Reproduction.

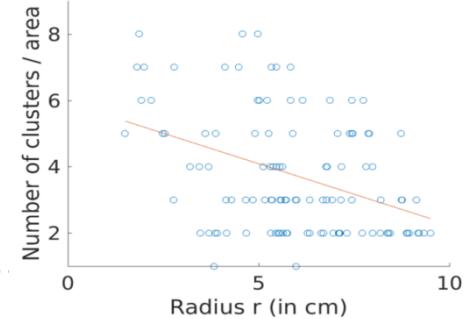
$$\text{Avg. num. of clusters} = 3.75 \pm 1.78$$



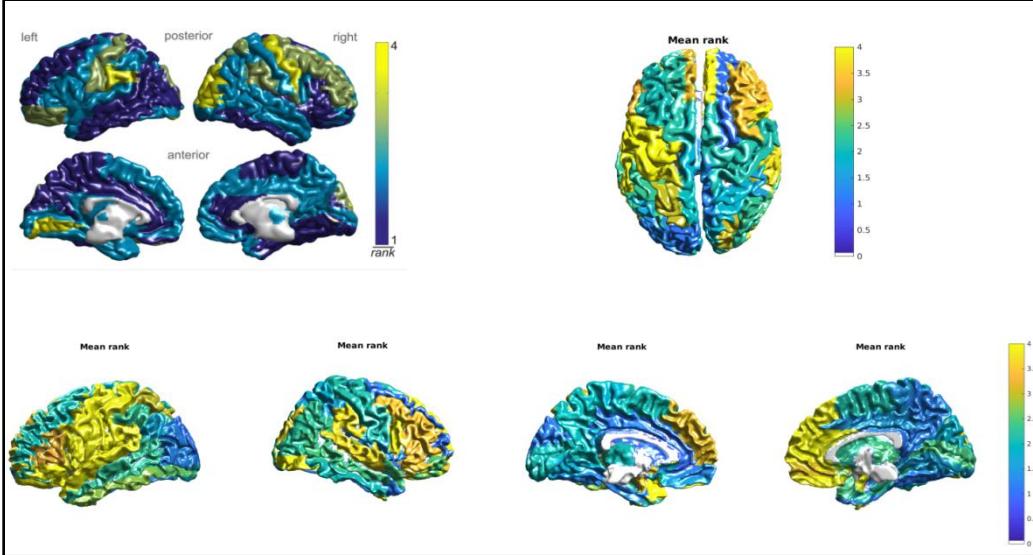
t: Reproduction.

$$R^2 = 0.16, F = 21.75, p < 0.001$$

$$y = (5.9279) + (-0.3677) * x$$

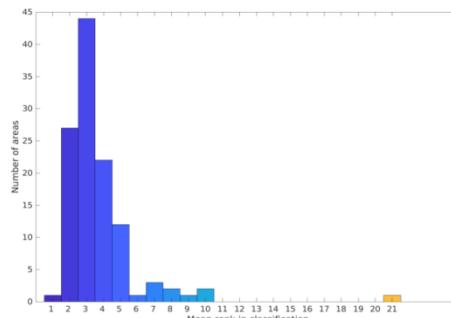
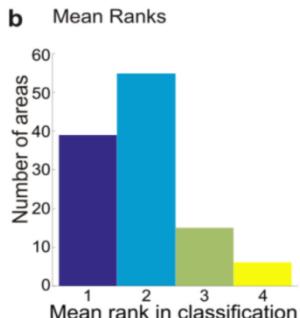


Left-up: Keitel & Gross, other figures: Reproduction.



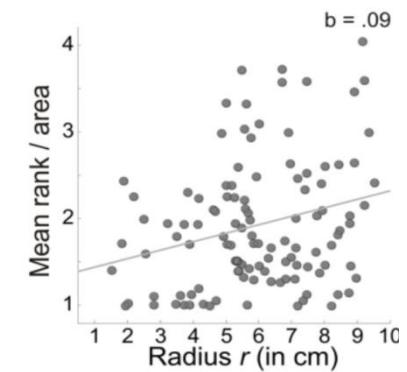
Left: Keitel & Gross
Mean rank = 1.8

Right: Reproduction.
Mean rank = 2.28



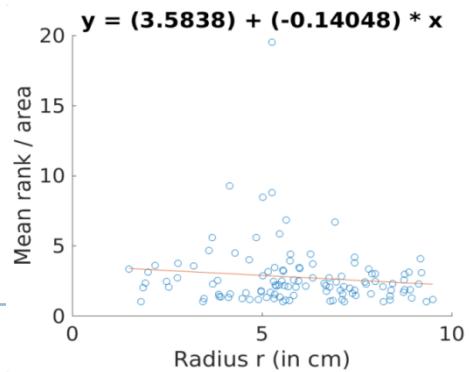
Left: Keitel & Gross

$R^2 = 0.07, F = 8.15, p = 0.04$



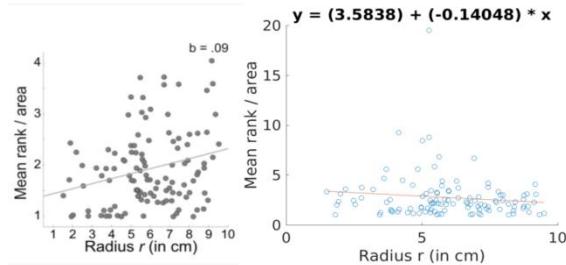
: Reproduction.

$R^2 = 0.01, F = 1.71, p = 0.19$



Left: Keitel & Gross
 $R^2 = 0.07, F = 8.15, p = 0.04$

Right: Reproduction.
 $R^2 = 0.01, F = 1.71, p = 0.19$

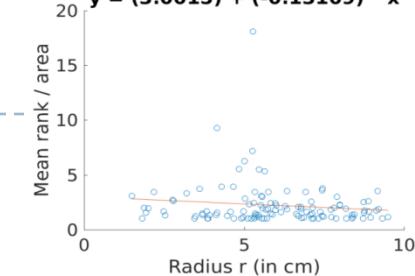
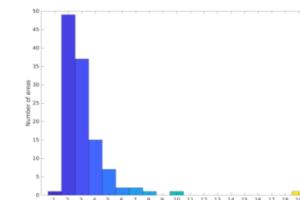


With homologues

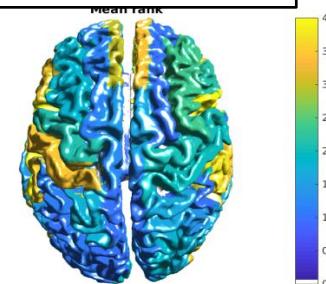
Mean rank (K & G) = 1.4

Mean rank (reprod.) = 1.79

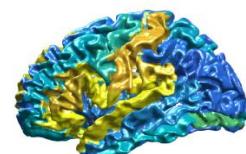
$$y = (3.0015) + (-0.13109) * x$$



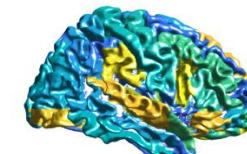
With homologues



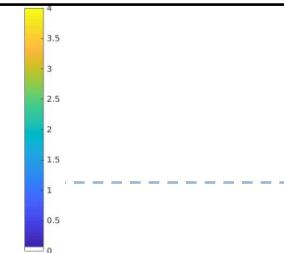
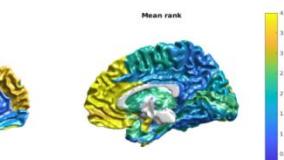
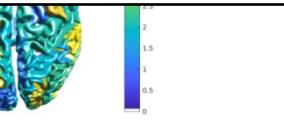
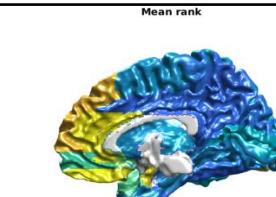
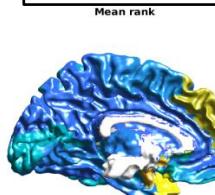
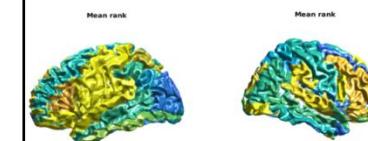
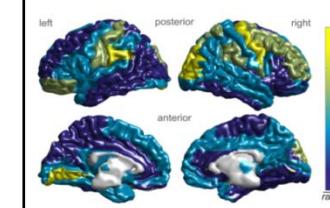
Mean rank



Mean rank

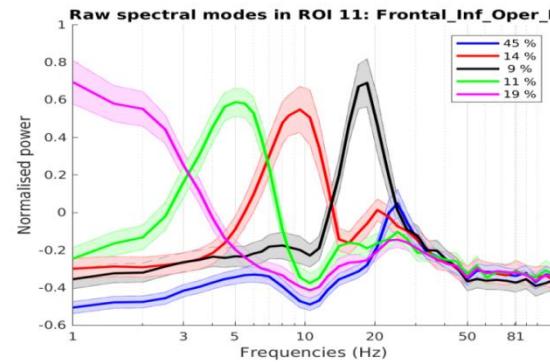
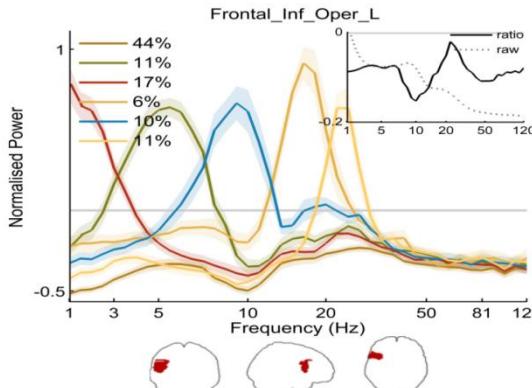
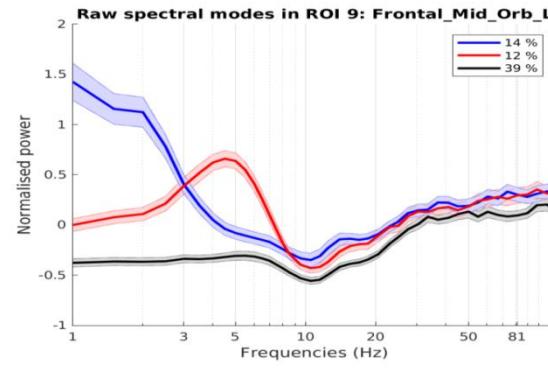
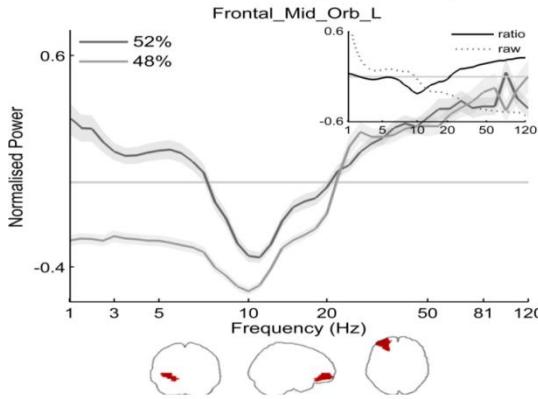


No homologues (previous slide)

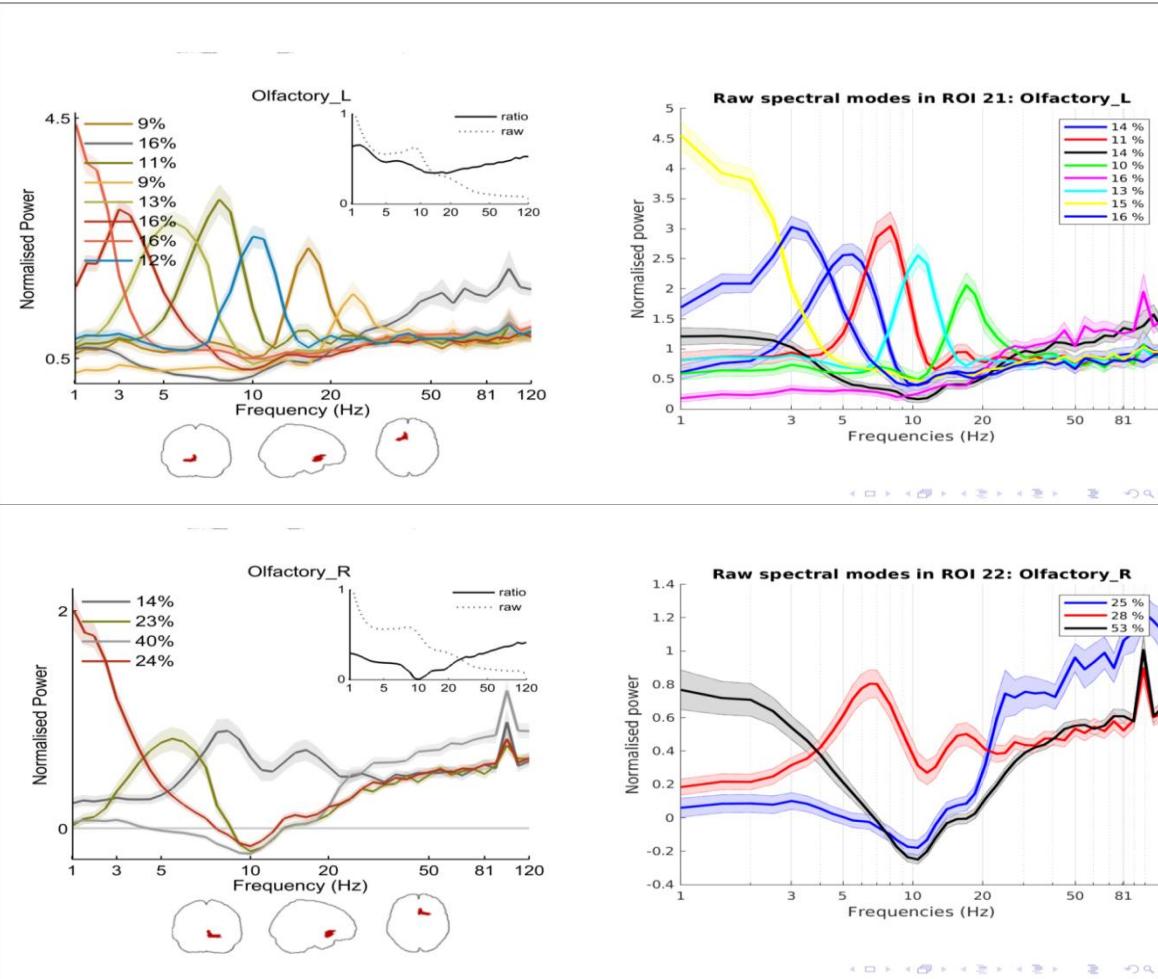


MEG reproduction results – ROI spectral fingerprints

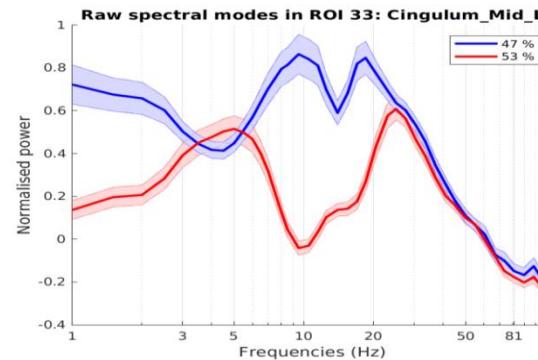
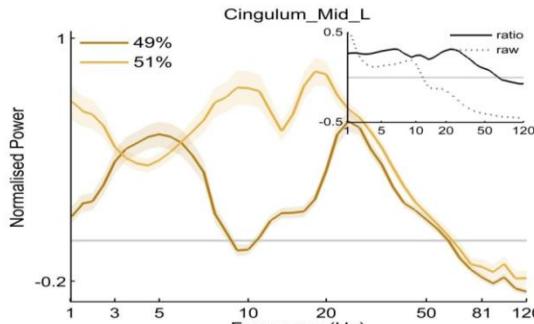
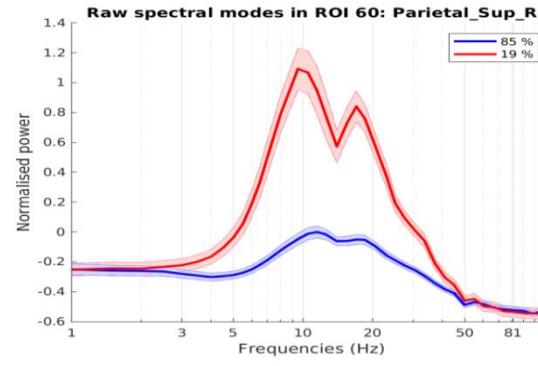
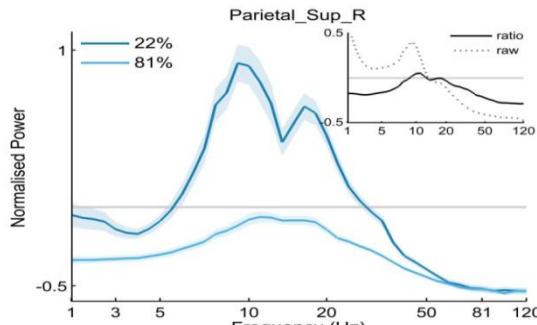
Frontal areas (inferior)



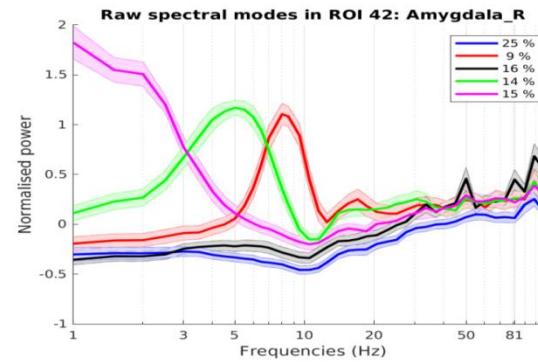
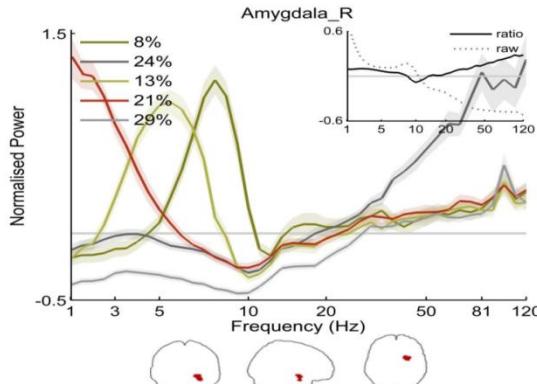
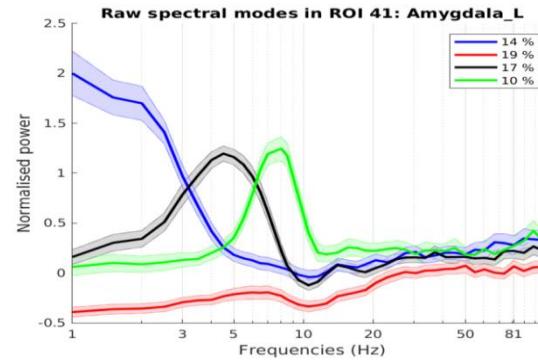
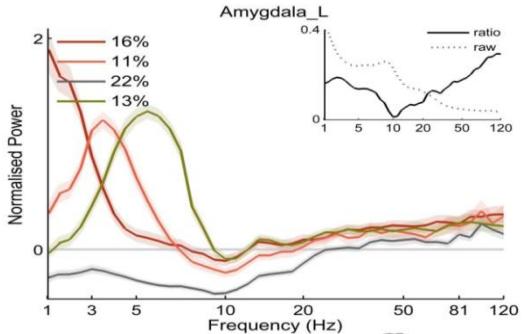
MEG reproduction results – ROI spectral fingerprints



MEG reproduction results – ROI spectral fingerprints

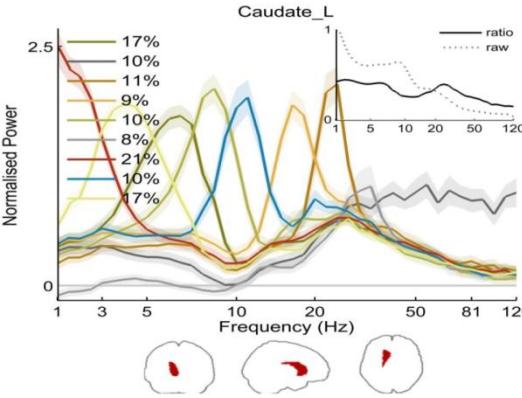


MEG reproduction results – ROI spectral fingerprints

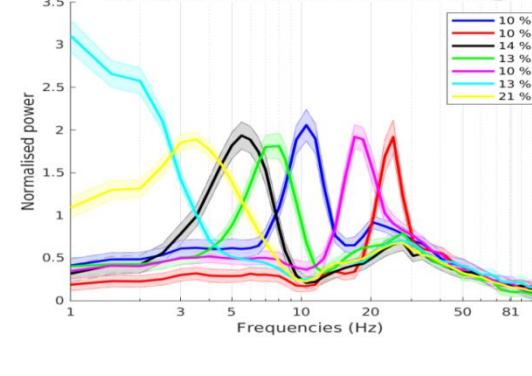


MEG reproduction results – ROI spectral fingerprints

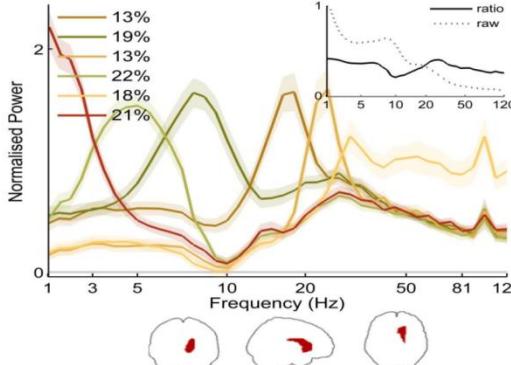
Basal ganglia



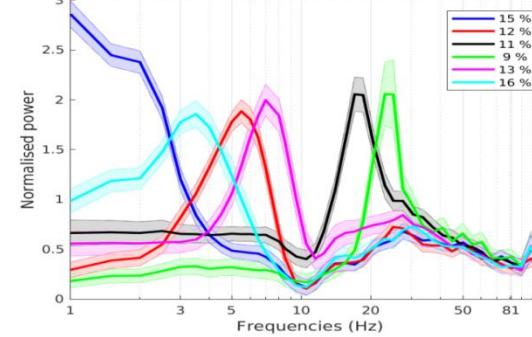
Raw spectral modes in ROI 71: Caudate_L



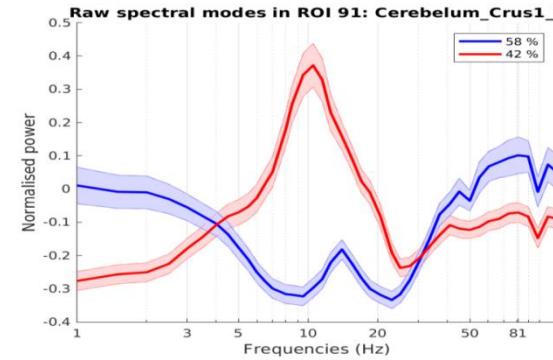
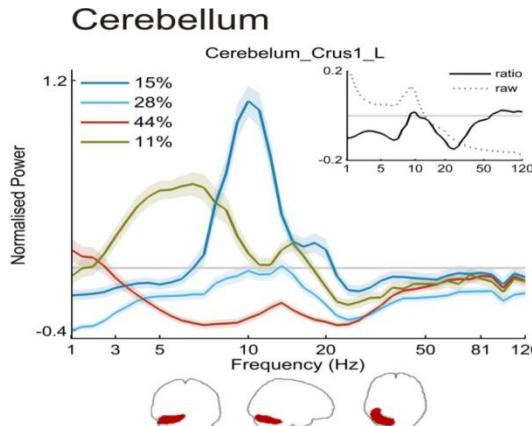
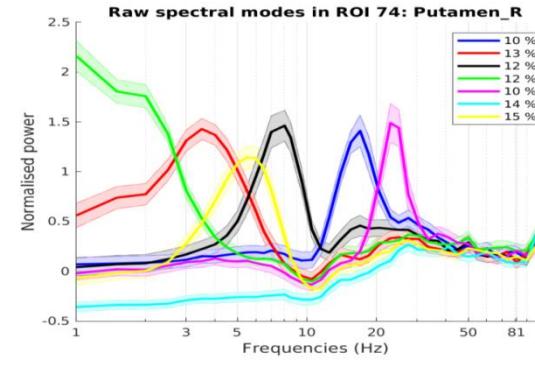
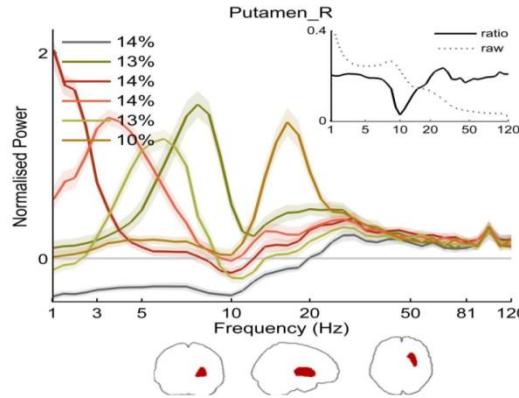
Caudate_R



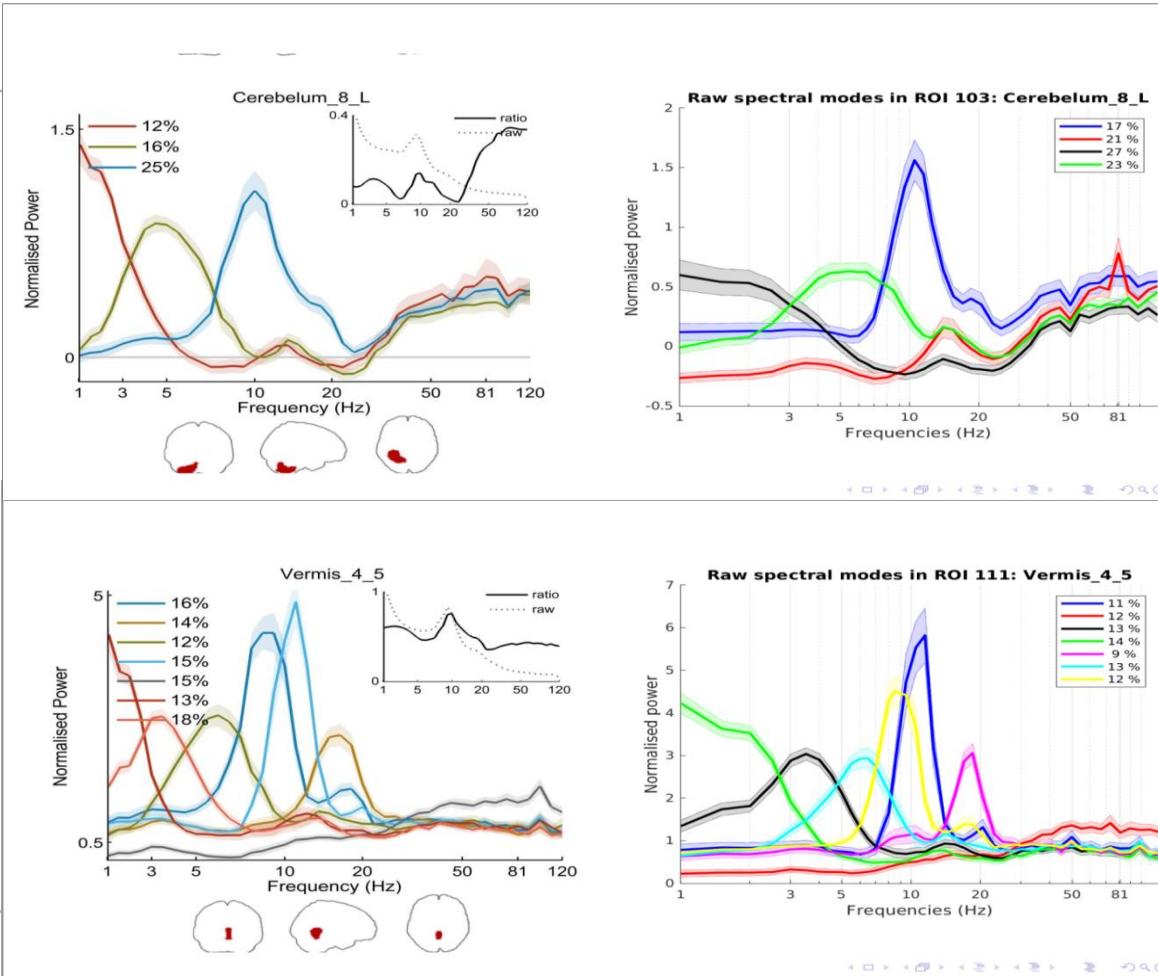
Raw spectral modes in ROI 72: Caudate_R



MEG reproduction results – ROI spectral fingerprints



MEG reproduction results – ROI spectral fingerprints



MEG reproduction results – ROI spectral fingerprints

