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# Comparing the SPIJN algorithm with conventional NNLS for myelin water fraction mapping

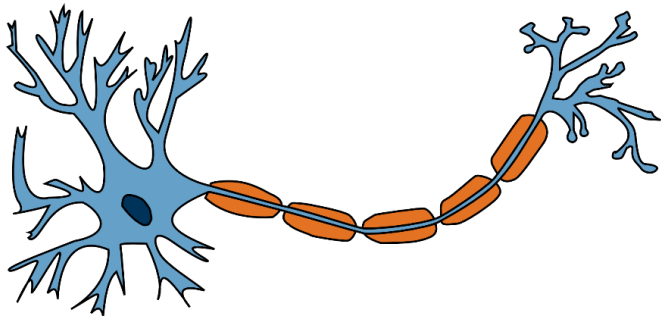
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Session **AI in Action**  
Date **01 October 2020**  
Time **14:30**  
Program # **S03.07**  
Abstract ID **A-1118**

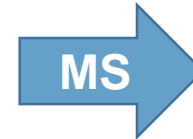
<sup>1</sup> Technical University of Munich, School of Medicine, Department of Neuroradiology, Munich, Germany  
<sup>2</sup> Philips Research Europe, Hamburg, Germany



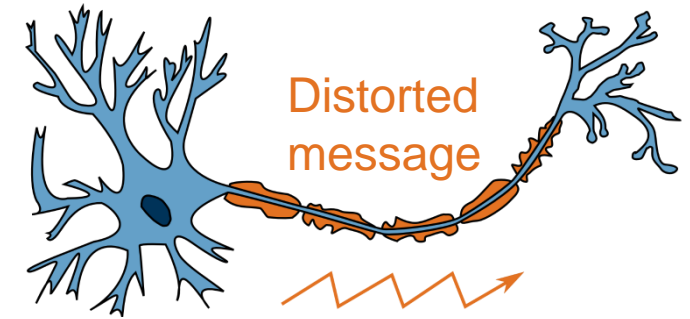
### Myelin



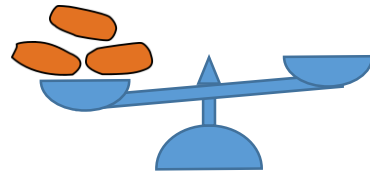
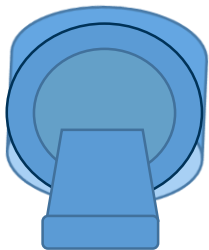
- Insulation
- Fast signal conduction



### Demyelination



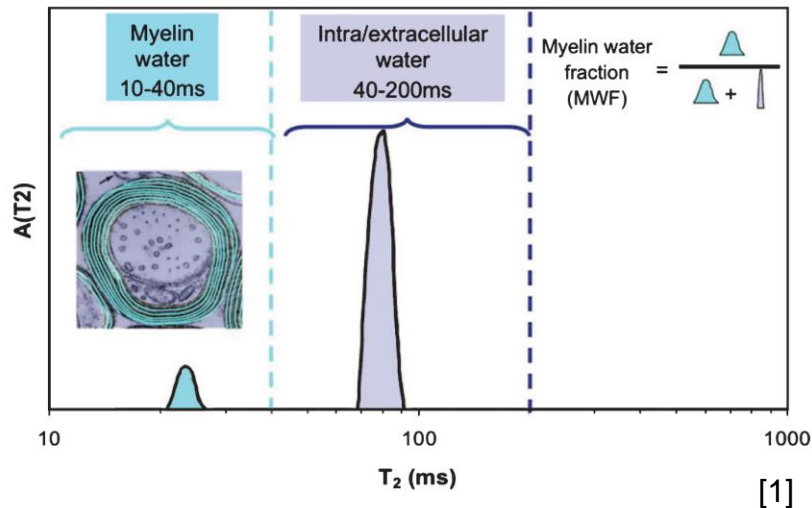
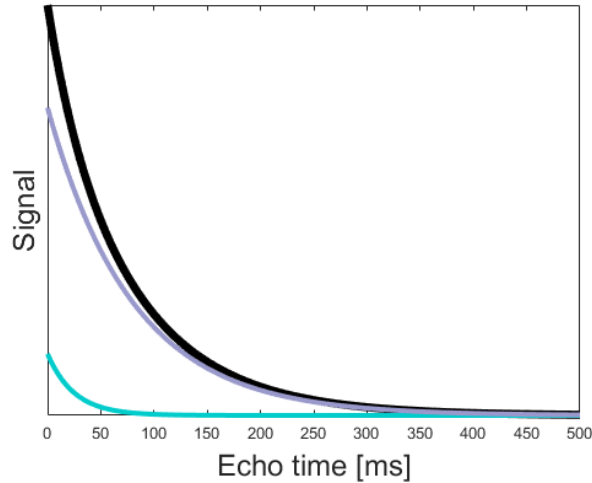
### Myelin imaging



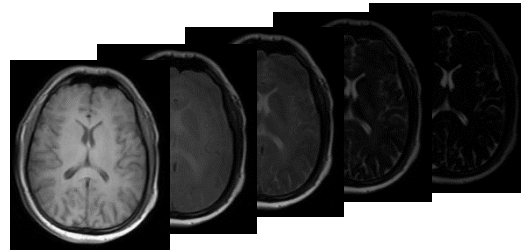
- Integrity of nerve tissue
- Monitor demyelinating diseases

MS nerve fiber: modified from <https://commons.wikimedia.org/wiki/File:Neuron.jpg>

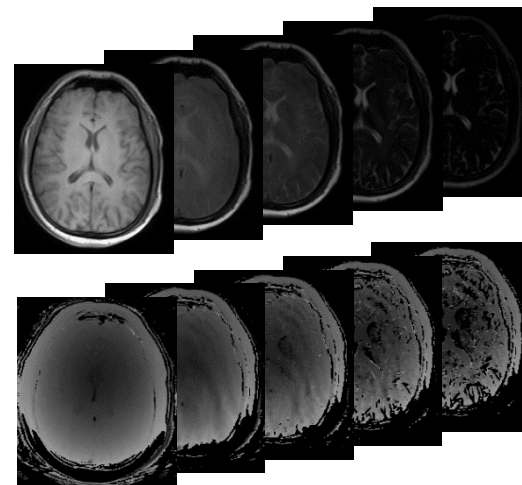
## Myelin Water Imaging



## Calculating T2 distributions



- Magnitude data



- Complex data

**NNLS**

[2]

- Non-Negative Least Squares

**SPIJN magn-based**

[3]

- Sparsity Promoting Iterative Joint NNLS
- MR Fingerprinting based
- Joint sparsity constraint
- New and faster method

**SPIJN comp-based**

[3]

- SPIJN can be improved by including complex data

**Aim** →

Compare NNLS and SPIJN (based on magnitude and complex data) for MWF mapping



## Cohort



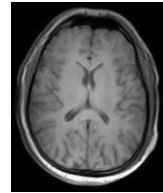
- n = 8 (3♀ / 5♂)
- Ø 33 years (29-49)
- healthy

## Sequence

- 3D GRASE
- TE1/ΔTE = 8/8 ms
- TR = 1120 ms
- 48 echoes
- 1x2x5mm<sup>3</sup>
- 20 slices
- α = 90°
- Philips Engenia Elition 3T

## Comparing NNLS & SPIJN

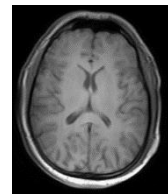
NNLS



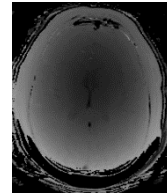
SPIJN magn

- Only magnitude data
- n = 8 (3♀ / 5♂)

NNLS



SPIJN magn

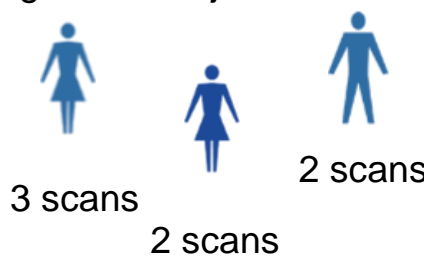


SPIJN comp

- Complex data available
- n = 4 (2♀ / 2♂)

## Reproducibility analysis

- Rescanning same subjects
- n = 3

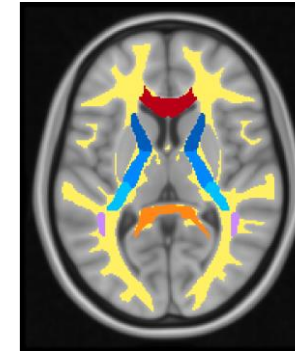


NNLS

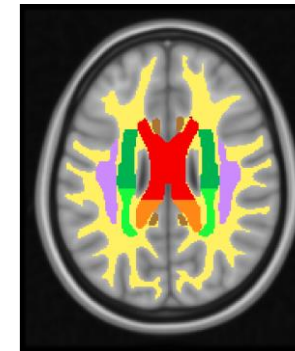
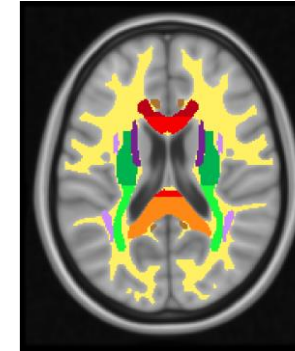
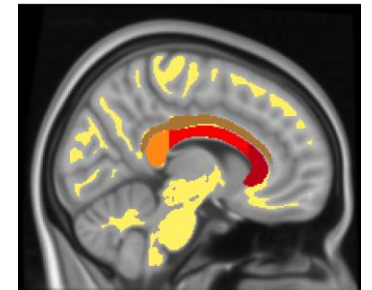
SPIJN magn

## Quantitative analysis

12 WM regions: Whole brain WM

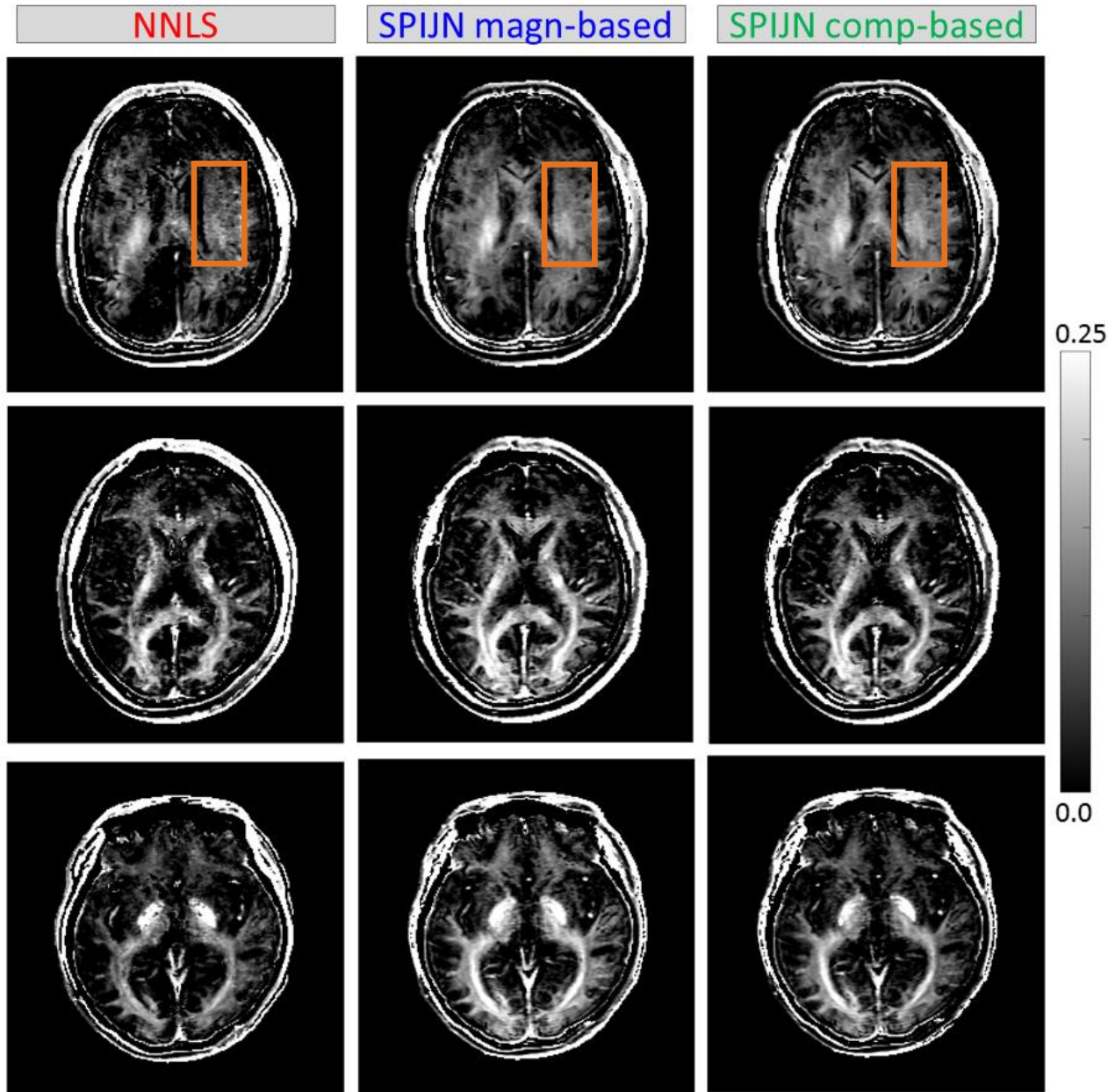


11 labels from the JHU ICBM-DTI-81 White-Matter Atlas



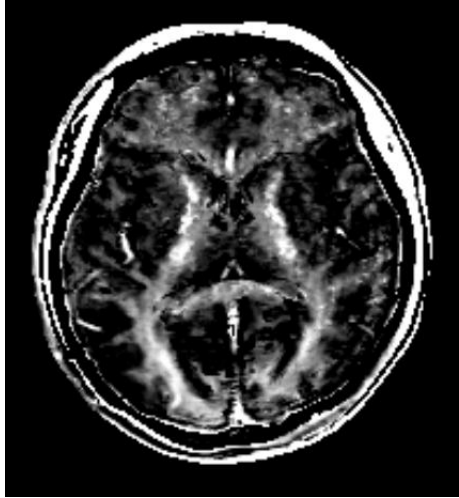
- Genu of CC
- Body of CC
- Splenium of CC
- Ant. limb of IC
- Post. limb of IC
- Retrolent. of IC
- Sup. corona rad.
- Post. corona rad
- Cingulate gyrus
- Sup. long. fasc.
- Sup. fronto-occ. facs.

JHU atlas: Mori, S., et al., MRI atlas of human WM, 2005.

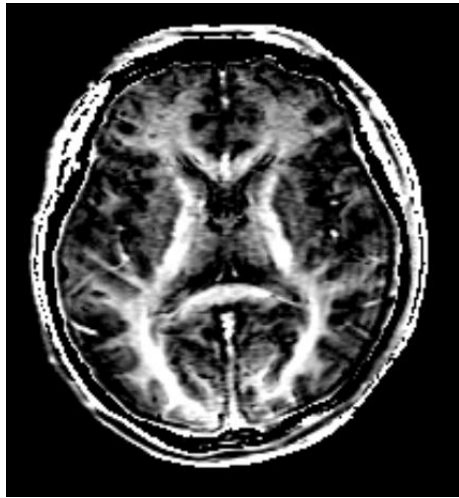


- Good agreement of areas with high MWF
- NNLS MWF maps appear a bit more noisy and porous
- SPIJN MWF maps appear relatively smooth
  - probably due to stronger regularization

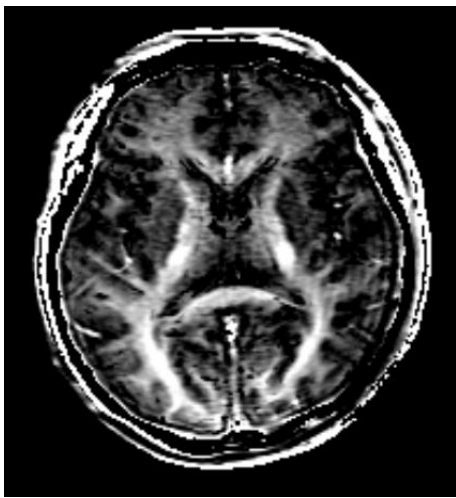
NNLS



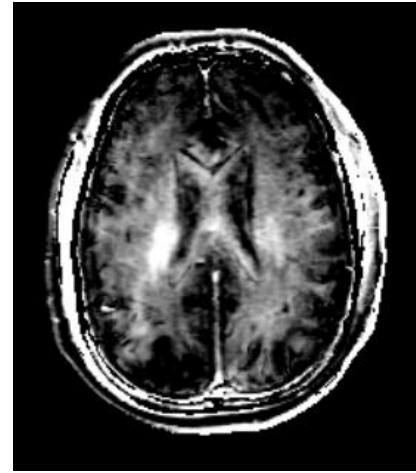
SPIJN magn-based



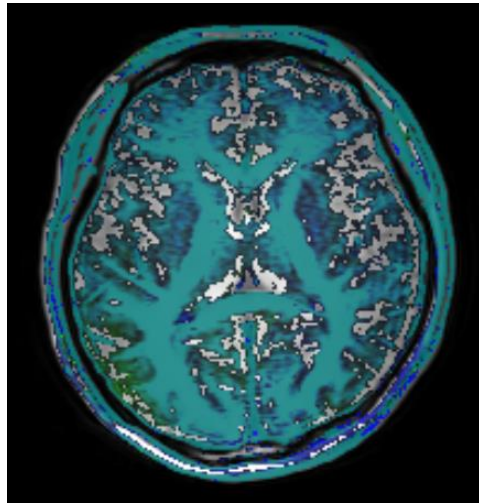
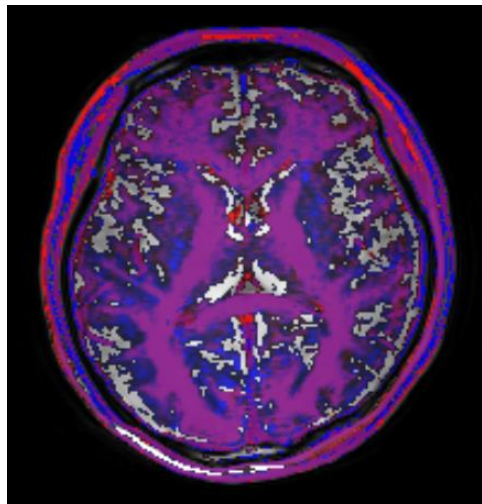
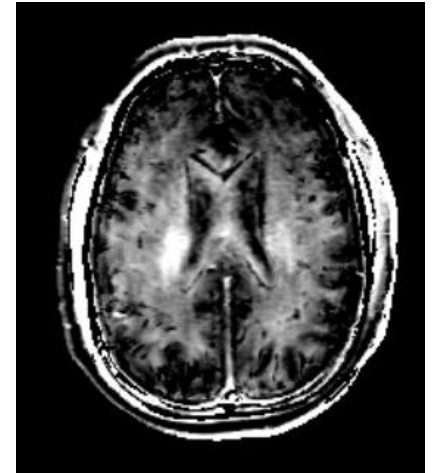
SPIJN comp-based



SPIJN magn-based



SPIJN comp-based



NNLS

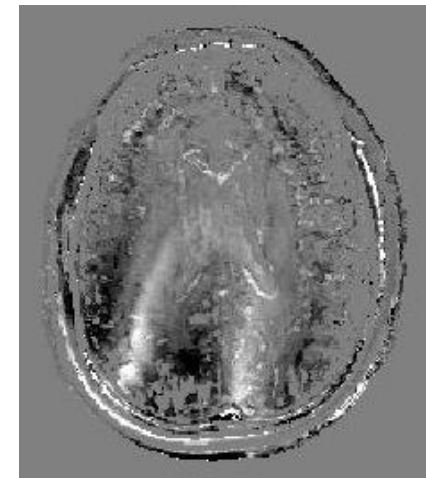


SPIJN magn



SPIJN comp

SPIJN magn – SPIJN comp



-0.05

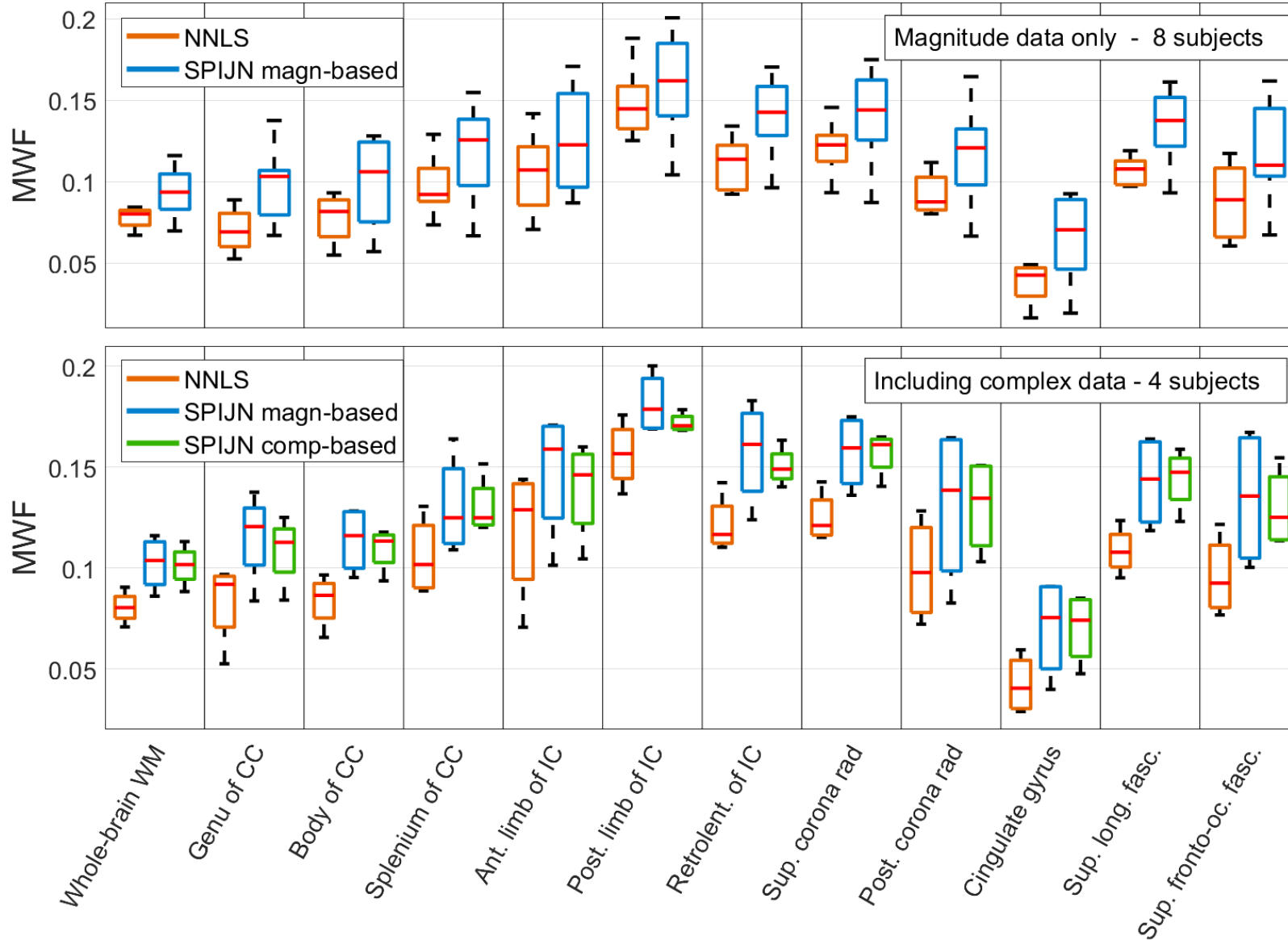
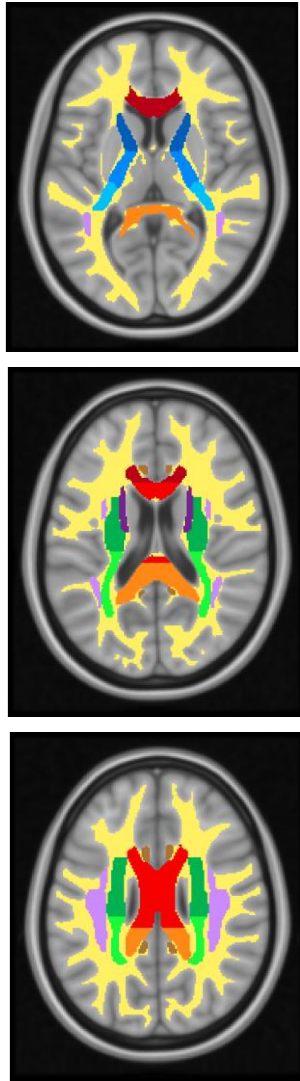


0.05

- SPIJN magn and SPIJN comp very comparable
- Small phase-like patterns in the subtraction image

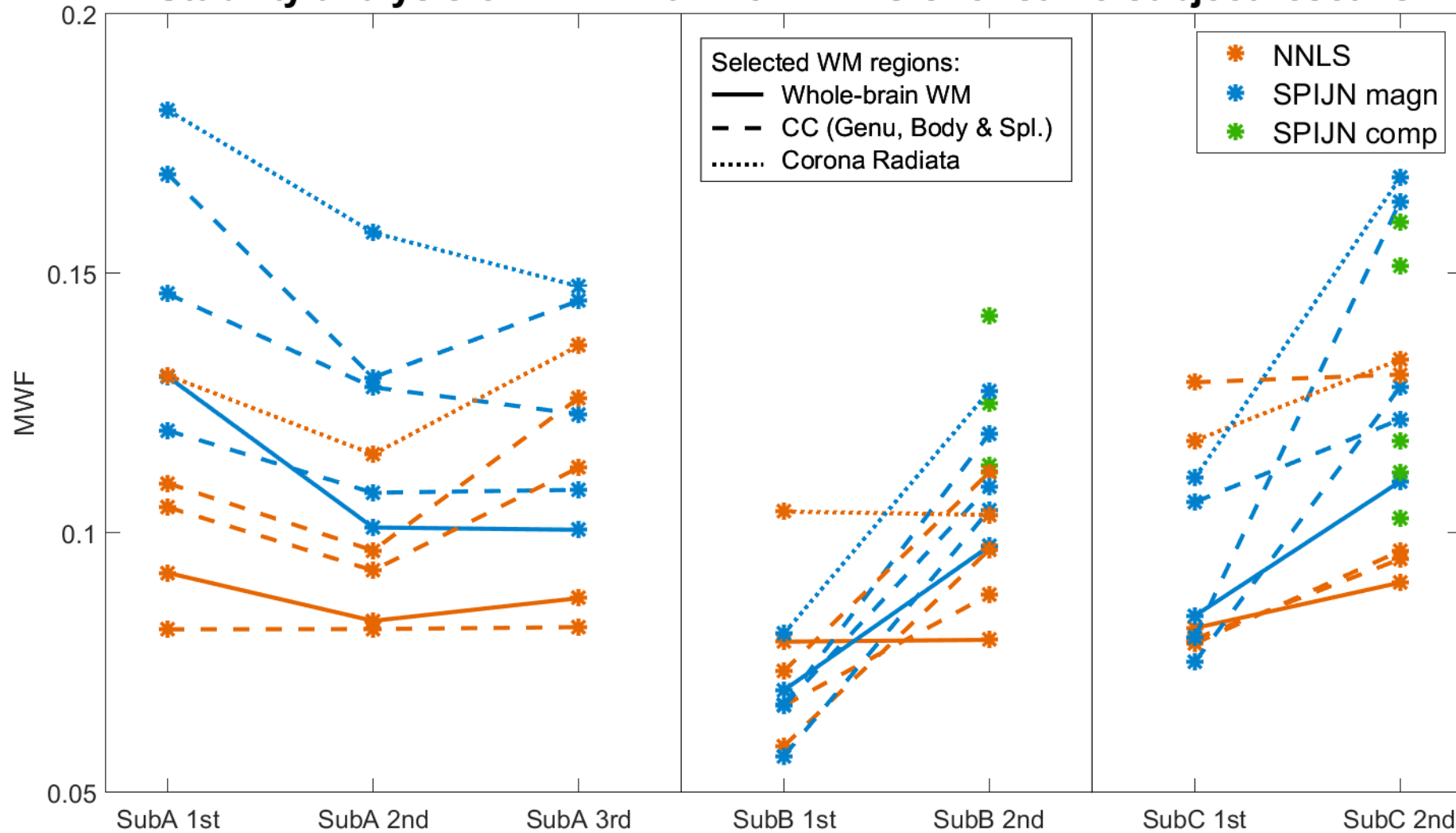


Comparison of MWF values obtained by NNLS and SPIJN in 12 WM segments



- Similar mean MWF values of NNLS and SPIJN
- SPIJN systematically higher and varies more strongly between subjects
- Slightly lower MWF and smaller variance of mean MWF across subjects with complex-data based SPIJN

## Stability analysis of MWF within 5 WM VOIs for same-subject rescans



- Rescanning of the same subjects
- Complex data only for 2nd scan of Sub B & C
- Comparison of several WM regions
- NNLS MWF values tend to be more stable



- SPIJN yielded visually smoother MWF maps
- NNLS algorithm seemed to provide more stable MWF values
- Complex-data based SPIJN processing provided better agreement with NNLS MWF
- Future studies:
  - Reproducibility analysis for complex-data based SPIJN
  - Effect of SPIJN regularization



Thank you for your attention!

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