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# SCAIFIELD a multi-center 7T study on Ataxia

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## Outline

Spinocerebellar ataxias (SCA)



- The SCAIFIELD project
- <u>Whole</u>-brain qMRI at 7T
- Initial Results
- Summary and outlook



- SCA: most common autosomal dominantly inherited progressive ataxia disorders worldwide
- First gene-therapies with a gene silencing approach have started. They might be preventive.
- Early (imaging) biomarkers would enable trials in the pre-ataxic stage!







# The SCAIFIELD project



**Goal:** Develop <u>quantitative MRI</u> methods for <u>whole-brain</u> imaging at <u>ultra-high field (</u>7T) and apply the novel technology in a <u>multi-center study</u> on <u>spinocerebellar ataxia</u> (SCA) patients

Challenge: Cerebellum imaging at 7T

#### Pls

Tony Stöcker, DZNE, Bonn, Germany (Coordinator) Pal Erik Goa, NTNU, Trondheim, Norway Pierre Maquet, GIGA, Liège University, Belgium Thomas Klockgether, DZNE, Bonn, Germany Ergin Atalar, Bilkent University, Ankara, Turkey



#### **Advisory Board**

Gülin Öz, CMRR, University of Minnesota, Minneapolis Robin Heidemann, UHF Group, Siemens Healthineers Cathalijne van Doorne and Julie Greenfield, Euro-Ataxia



PPI activities

- (i) Incorporate patient perspective in study design
- (ii) Providing easy-to-read information about research outcomes
- (iii) Explain possibilities to participate in clinical enrolment



# The SCAIFIELD project



I. Method development: UHF-Neuroimaging including cerebellum and brainstem





## Ataxia Imaging @ $7T \Rightarrow$ mitigating B1 inhomogeneity (transmit field)



... without lengthy subjectspecific calibration !

## Universal Pulses<sup>1</sup>

- pTx pulses, which homogenize B1 for data base of subjects
- UPs generalize, i.e. they also perform on new subjects



<sup>&</sup>lt;sup>1</sup> Gras et al, MRM 2017



parallel transmission (pTx)





Acquire B1 maps per channel Acquire B0 map Calculate X RF pulses  $(\Sigma \approx 5 \cdot 10 \text{ min})$ Apply pulses in sequence

## SCAIFIELD method development: rapid whole-brain 7 Tesla MRI

#### Custom sequences with tailored Universal Pulses (UPs) and ultra-fast imaging readouts

- UP calculation <sup>1,2,3,4</sup> with BLT database (Bonn-Liège-Trondheim field-map database collection > 70 subjects)
- Fast readouts <sup>5,6,7,8</sup> combining 2D parallel imaging, partial Fourier, elliptical sampling, EPI factor, and compressed sensing



<sup>1</sup> Löwen et al, *MRM* 2022, 8(86): 2564–72; <sup>2</sup> Gras et al,, MRM 2018, 80:53–65,; <sup>3</sup> Gras, et al, MRM 2019, 81(5): 3202–8; <sup>4</sup> Löwen et al, MRM, 2024. <sup>5</sup> Brenner et al, *Magma* 2014, 27(5):455–62. <sup>6</sup> Pracht et al, MRM 2018, 79(5): 2620–28; <sup>7</sup> Stirnberg and Stöcker, MRM 2022, 85:1540-1551; <sup>8</sup> Stirnberg et al, MRM 2024

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Custom sequences:

- $pTx \Rightarrow$  homogenous whole-brain MRI
- UPs  $\Rightarrow$  calibration-free and robust
- Advanced readouts  $\Rightarrow$  high SNR/CNR
- MP-RAGE, 3D TSE, 3D FLAIR

<

 $\checkmark$ 

<sup>1</sup> Löwen et al, *MRM* 2022, 8(86): 2564–72; <sup>2</sup> Gras et al,, MRM 2018, 80:53–65,; <sup>3</sup> Gras, et al, MRM 2019, 81(5): 3202–8; <sup>4</sup> Löwen et al, MRM, 2024.

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## rapid quantitative and molecular whole-brain MRI @ 7T

#### Chemical Exchange Saturation Transfer<sup>1</sup> (CEST) - 1.6 mm iso

Multi-Parametric Mapping<sup>2</sup> (MPM) - 0.6 mm iso



<sup>1</sup> Völzke et al. "PUSHUP-CEST: Calibration-free whole-brain ultra-high field CEST imaging using universal parallel transmission ", MRM 2024
<sup>2</sup> Wang, Ehses, Stöcker, Stirnberg. "Reproducibility of Rapid MPM at 3T and 7T with Highly Segmented and Accelerated 3D-EPI." MRM 2022 88(5): 2217–32.

## SCAIFIELD: 7T-qMRI Acquisition Protocol

#	Contrast (Method)	Resolution / mm <sup>3</sup>	Coverage	Duration / min	Quantitative Analysis	Hypothesis
1	B0 & B1 map (3DREAM)	5.0 x 5.0 x 5.0	whole brain	02:38	Calibration: flip angle maps and field maps	$(\rightarrow qMRI analysis)$
2	MPRAGE	0.6 x 0.6 x 0.6	whole brain	07:24	segmentation / volumetry	volume loss <sup>1</sup>
3	MPM + QSM (segm. 3D-EPI)	0.6 x 0.6 x 0.6	whole brain	13:41	T1, T2*, PD, MT,QSM / volumetry	T1, T2* reduction <sup>2</sup> (PD,MT?) / volume loss / iron load <sup>2</sup>
4	CEST (segm. 3D-EPI)	1.6 x 1.6 x 1.6	whole brain	07:37	Amides, NOE (?)	decreased?
5	DWI (SMS-SE-EPI, 2-shell HARDI)	1.5 x 1.5 x 1.5	whole brain	10:44	Microstructure & connectivity: e.g. DTI, NODDI, tractography	reduced connectivity / cerebellar demyelination <sup>3,4</sup>
6	MRSI (3D-CRT)	5.0 x 5.0 x 5.0	cerebellum	08:56	Neurochemical profile: Cr, Glu, Gln, NAA,	MRS Score: SCA classification <sup>5</sup>

≈ 52 min



Reetz K et al., *Brain* 136, no. 3 (2013)
Deistung A et al., *Brain Comm.* 4, 2022
Mascalchi M, et al. PLoS One (2018)

4. Piccinin CC, et al. Mov Disord. (2020)

5. Joers JM et al., Annals Neurology 83, (2018)



## **Example Patient Data**





DZNE

## **Example Patient Data**

Diffusion: brain microstructure (1.5 mm iso)





## <u>DTI</u>

FA: Fractional anisotropyMD: Mean diffusivityRD: Radial diffusivityAD: Axial diffusivity

#### <u>NODDI</u>

ICVF: Intracellular Volume Fraction

ISOVF: Isotropic Volume Fraction

OD: Orientation Dispersion



1.00

0.75

0.50

0.25

0.00



## **Tissue parameter mapping: molecular information**

Multi-Parametric Mapping (MPM) - 0.6 mm iso

Chemical Exchange Saturation Transfer (CEST) - 1.6 mm iso





HC

## **SCAIFIELD: summary and next steps**

## **SCAIFIELD** Achievements

- Fast sequences for homogeneous whole-brain quantitative MRI at 7T with high resolution and image quality
- Tailored analysis pipelines with focus on cerebellum for brain segmentation, MPM, QSM, CEST, DWI, CSI
- Sequence role out at all SCAIFIELD sites (Bonn, Liège, Trondheim, Essen) / patient scanning started Q3 2024

## Next steps

- Ataxia patient study is ongoing and will finish in Q2 2025
- Automated global analysis pipeline and group analyses
- Backtranslation to 3T
- qMRI as early imaging biomarker for Ataxia ( $\Rightarrow$  in future, therapy monitoring in pre-ataxic stage?)

## Impact beyond the project

- Imaging protocol ready for clinical research, paving the way for increased routine use of 7T MRI
- Sequences will be made freely available in Q1 2025 (for Siemens 7T Terra and 7T Plus, WIP: Terra.X)





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#### **SCAIFIELD Team**

**Bonn** 

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# **Thank You**

**D**JPND research



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