

Impact of deep learning reconstruction on the accuracy of automated hippocampal subfield segmentation in high-resolution T2-weighted turbo spin echo imaging

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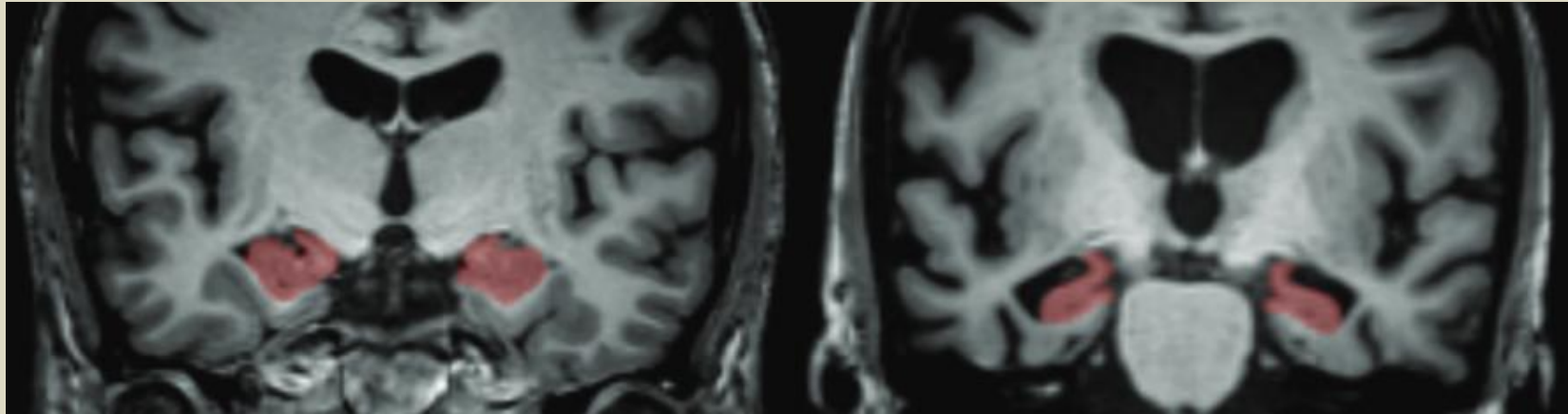
Hippocampus



- Major site of neuronal loss in neuropsychiatric disorders

Healthy control

Patients with Alzheimer's disease



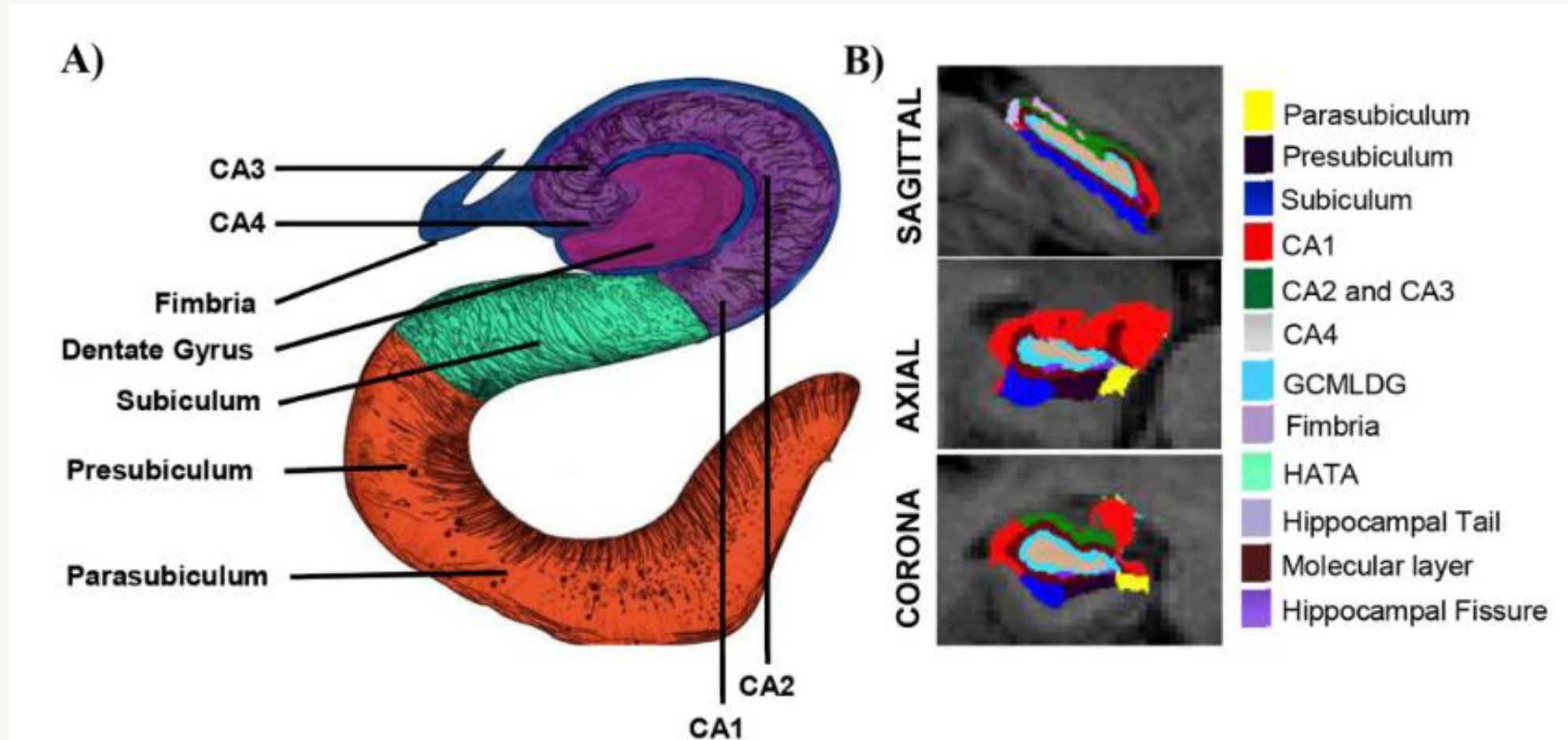
Teipel S, et al. Lancet Neurol. 2015

- Hippocampal subfield evaluation
 - More sensitive for early Alzheimer's disease than whole atrophy*
 - Demonstrates high disease specificity**

* Apostolova LG, et al. Neurobiology of Aging. 2010

** Blanken AE, et al. NeuroImage: Clinical. 2017

Automated segmentation of high-resolution T2WI



Kannappan B, et al. PLoS One. 2022

High-quality images require long acquisition times

Deep learning reconstruction (DLR)



➤ Deep Resolve Boost

- Reduces noise caused by high acceleration factors

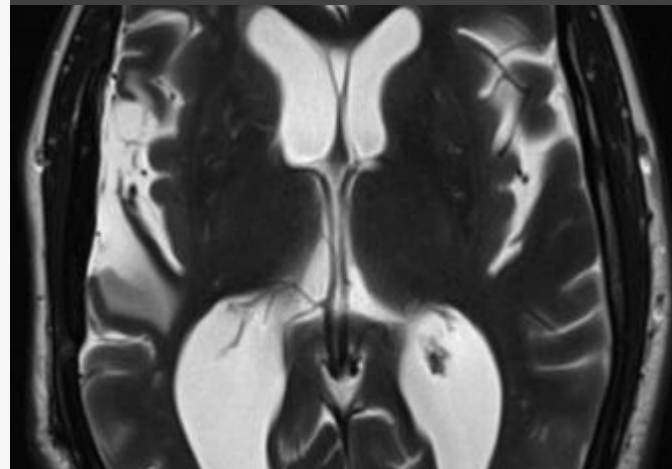
➤ Deep Resolve Sharp

- Enables high-resolution imaging via deep learning

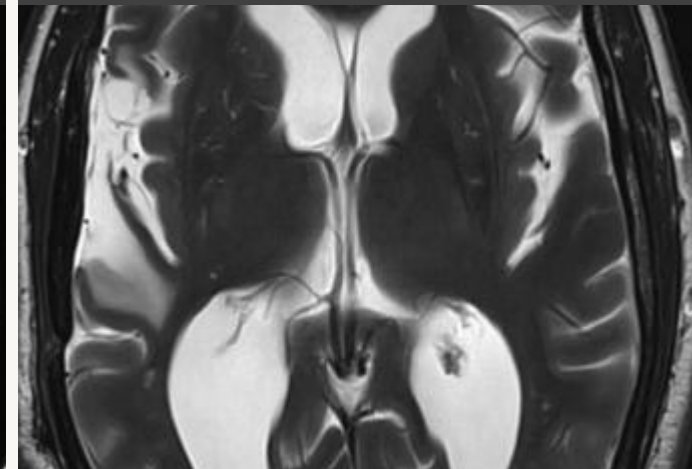
Reduce acquisition time while maintaining image quality

Lyo SK, et al. Radiol Adv. 2025

Conventional: 1 min 36 s



DLR: 1 min 4 s



Impact on hippocampal subfield segmentation is unclear



To evaluate whether accurate hippocampal subfield segmentation can be maintained using deep learning reconstruction, even with reduced acquisition time.

Equipment and Subjects



➤ Subjects

- **10 healthy volunteers (Total: 20 bilateral hippocampi)**
7 males, 3 females; mean age, 29.8 ± 6.5 years (range: 23–47 years)

➤ MRI system

- MAGNETOM Vida 3.0 T (Siemens Healthineers, Germany)
- 20-channel head/neck coil

Imaging sequence



Parameters	3D T1WI		2D T2WI		
Acquisition plane	Sagittal		Coronal		
TR/TE (ms)	2300/2.98		8020/50		
Matrix (phase × frequency)	240 × 256		448 × 448		
Field of view (mm)	256		175		
Slice thickness (mm)	1		2		
Flip angle (degrees)	9		122		
Acceleration factor (GRAPPA)	2	2	2	3	4
DLR setting*	off	off	on		
Acquisition time (min:s)	5:21	4:17	4:17	2:56	2:24

*Deep Resolve Boost (medium) and Sharp (on)

Imaging sequence



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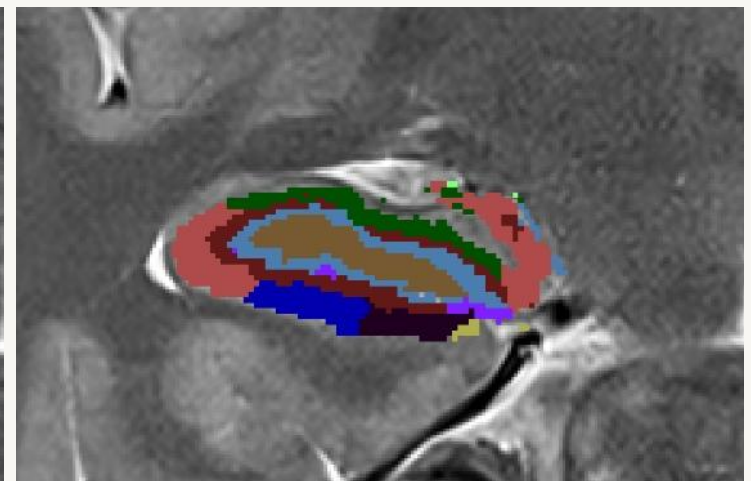
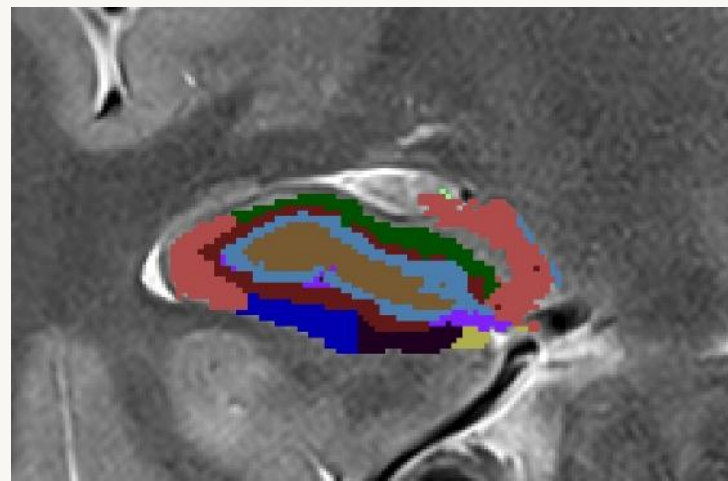
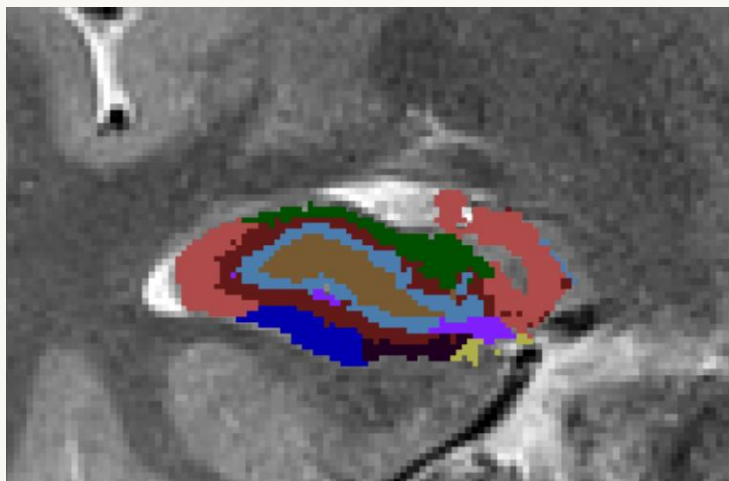
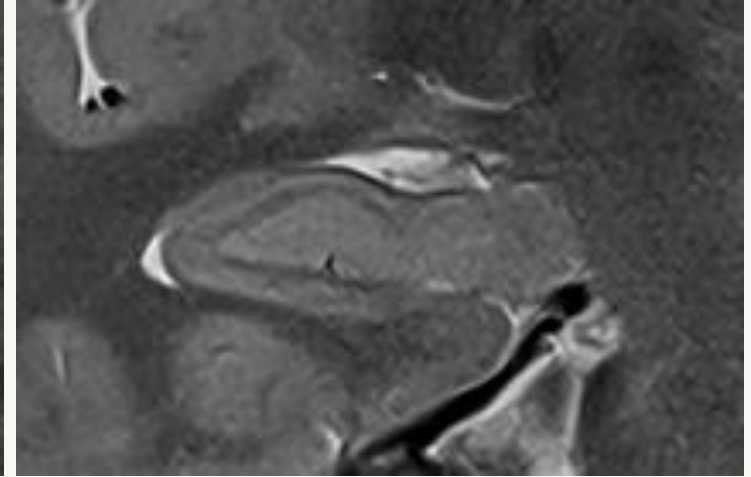
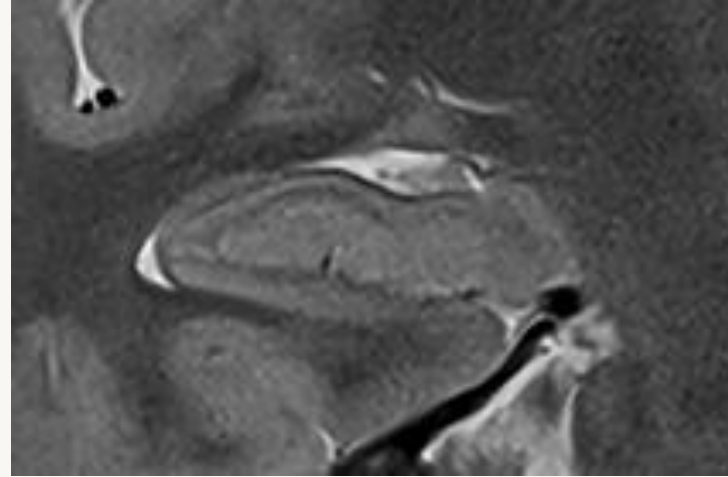
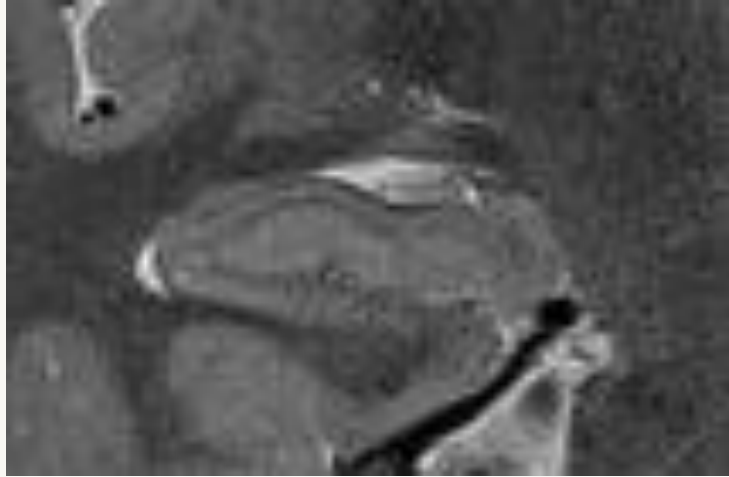
*Deep Resolve Boost (medium) and Sharp (on)

Analysis methods



- Hippocampal segmentation using FreeSurfer (version 7.4.1)
 - CA1, CA3, DG+CA4, Subiculum, Tail and Whole hippocampal
- Statistical analysis (R version 4.4.0)
 - Comparison of volume differences
 - Assessment of volume change rate
 - Paired t-test (Holm correction)
 - Evaluation of Reliability: ICC (2,1), Bland-Altman analysis

Acquired images



GRAPPA 2
4:17

GRAPPA 3+DLR
2:56

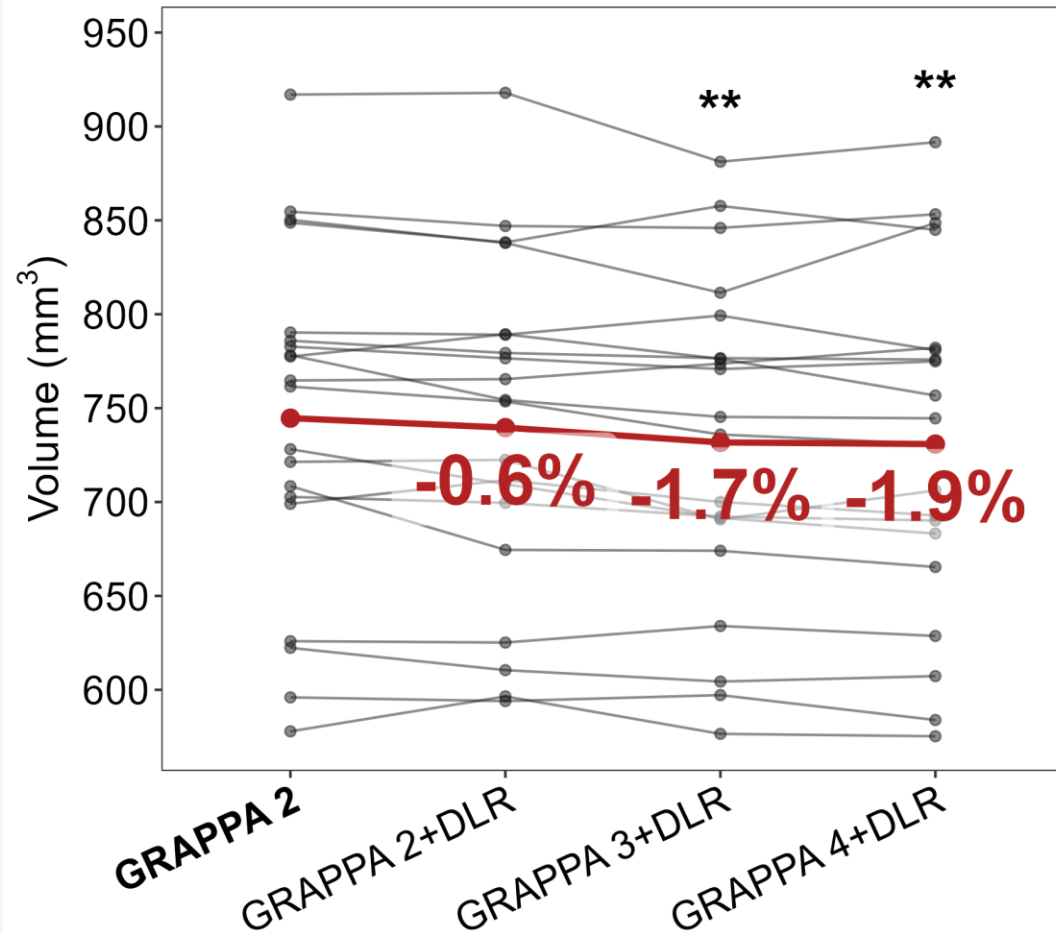
GRAPPA 4+DLR
2:24

Results of volume change rates



CA1

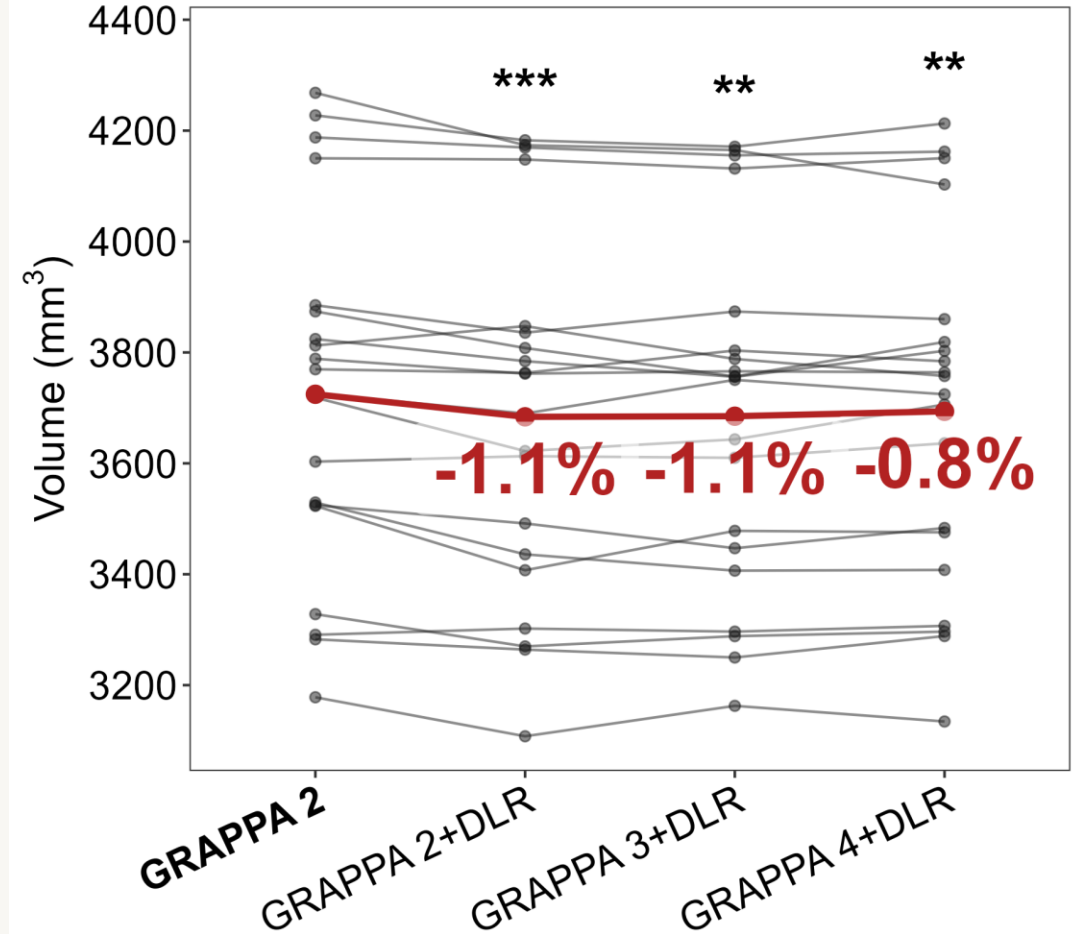
ANOVA p.adj = < 0.001 | ICC(2,1) = 0.982



* p < 0.05, ** p < 0.01, *** p < 0.001 (vs GRAPPA 2)

Whole hippocampus

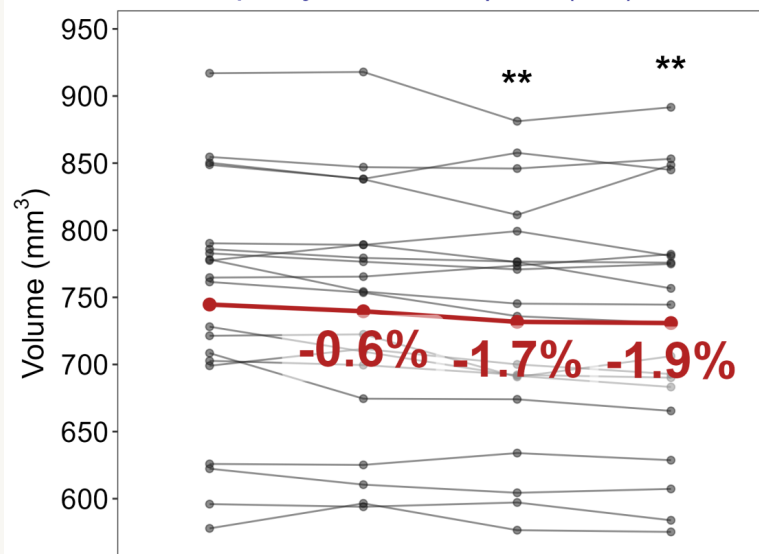
ANOVA p.adj = < 0.001 | ICC(2,1) = 0.989



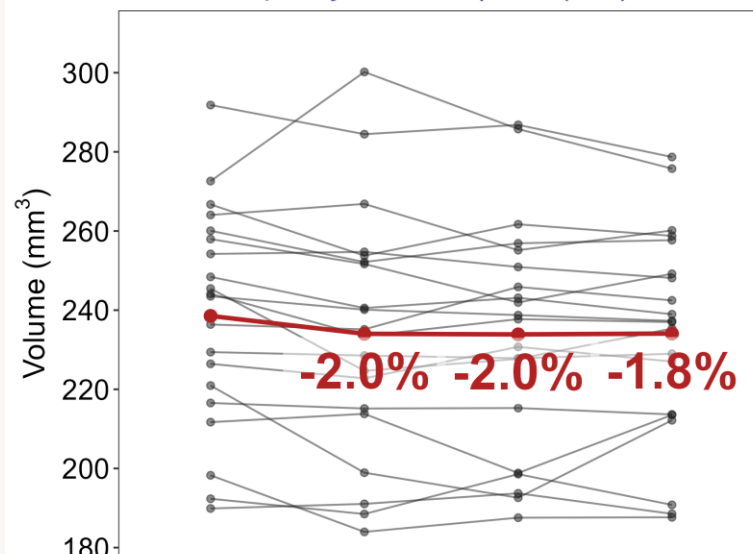
* p < 0.05, ** p < 0.01, *** p < 0.001 (vs GRAPPA 2)

CA1

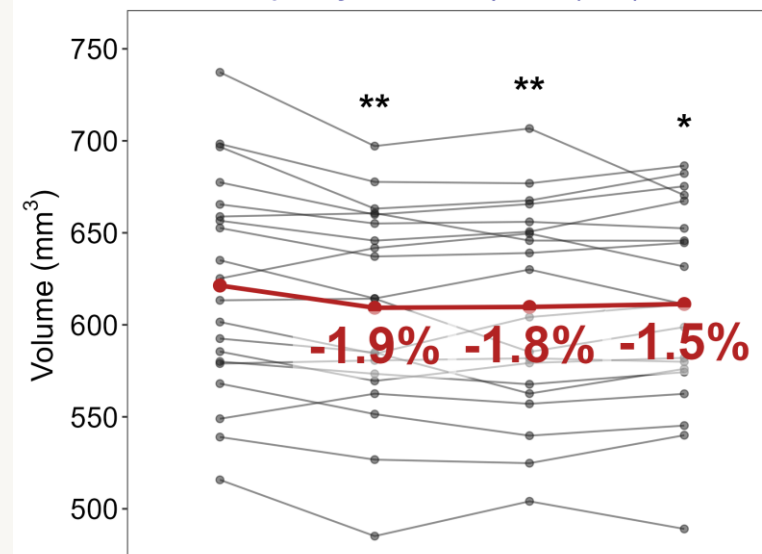
ANOVA p.adj = < 0.001 | ICC(2,1) = 0.982

**CA3**

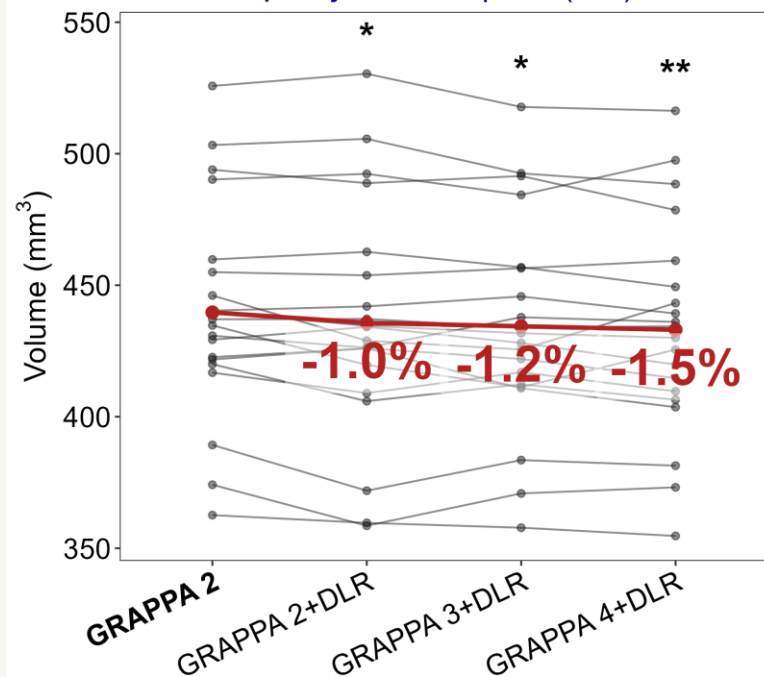
ANOVA p.adj = 0.112 | ICC(2,1) = 0.955

**DG+CA4**

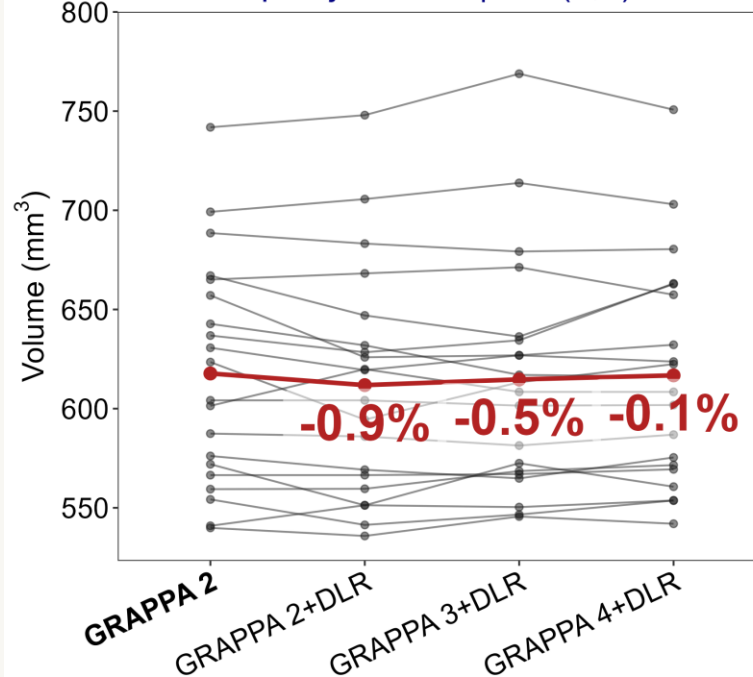
ANOVA p.adj = 0.004 | ICC(2,1) = 0.959

**Subiculum**

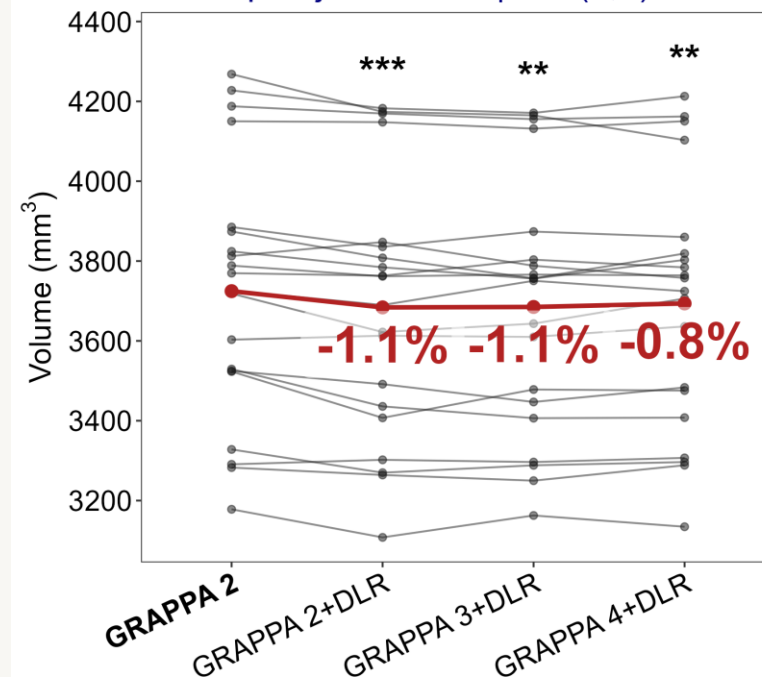
ANOVA p.adj = 0.018 | ICC(2,1) = 0.977

**Hippocampal Tail**

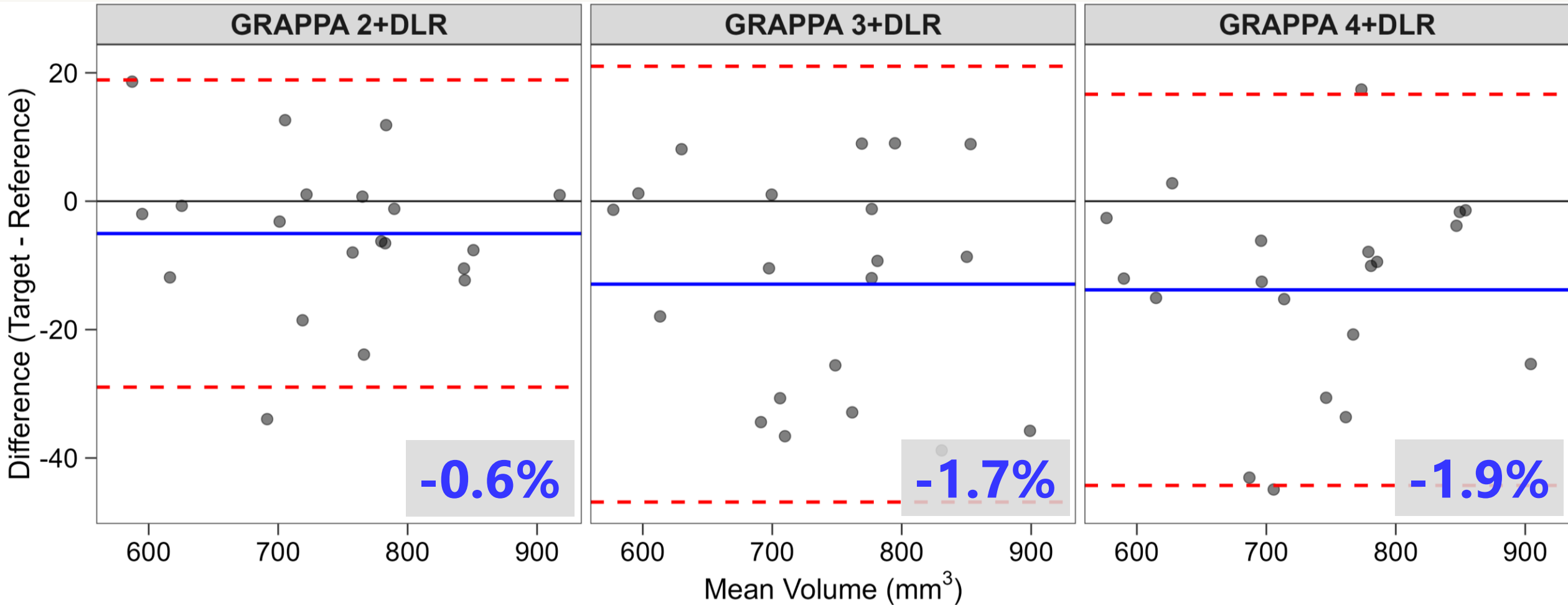
ANOVA p.adj = 0.172 | ICC(2,1) = 0.975

**Whole hippocampus**

ANOVA p.adj = < 0.001 | ICC(2,1) = 0.989



Bland-Altman analysis (CA1)

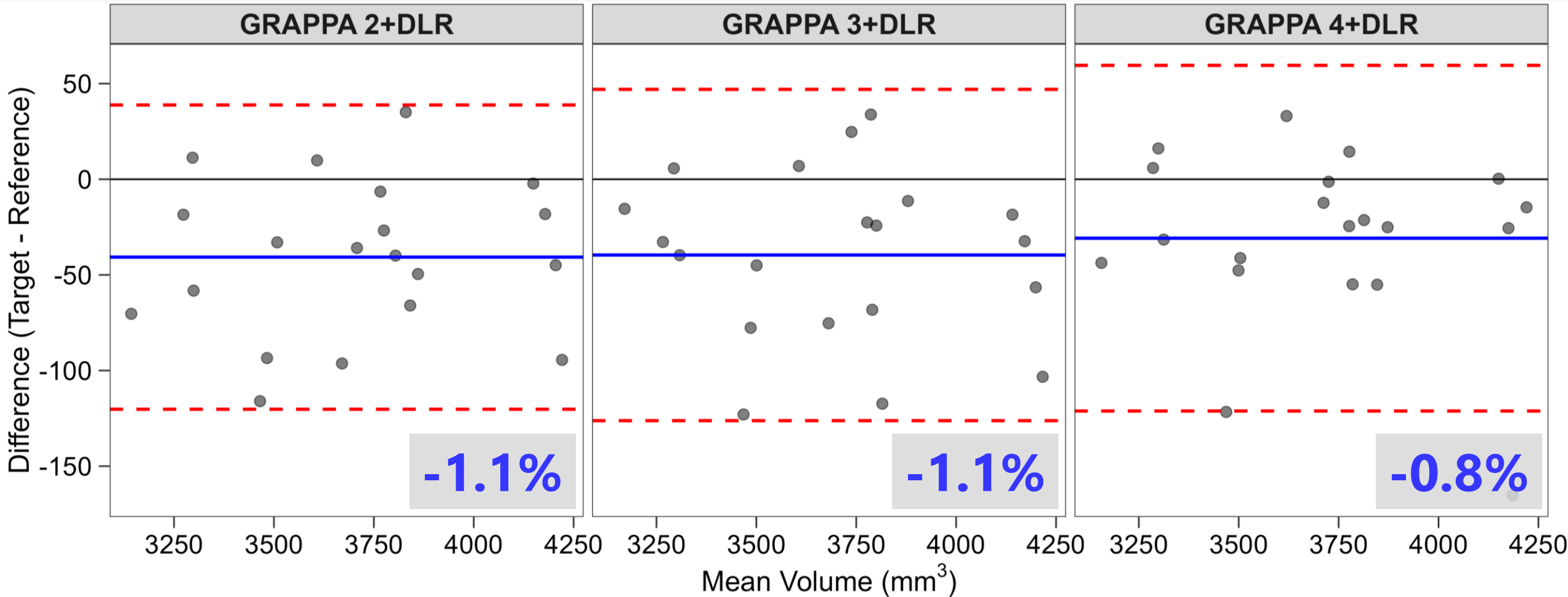


Bias [95% LoA] (mm³)
-5.0 [-29.0, 18.9]

Bias [95% LoA] (mm³)
-12.9 [-46.9, 21.0]

Bias [95% LoA] (mm³)
-13.8 [-44.3, 16.7]

Bland-Altman analysis (Whole hippocampus)



Bias [95% LoA] (mm³)
-40.7 [-120.2, 38.8]

Bias [95% LoA] (mm³)
-39.6 [-126.2, 47.0]

Bias [95% LoA] (mm³)
-30.8 [-121.2, 59.5]

Discussion



➤ Volume change rates

- **~2% reduction** in CA1, DG+CA4, Subiculum, and whole hippocampus
- **ICC > 0.90** for each subfield across different acceleration factors

High reproducibility despite high acceleration factors

➤ Is this 2% change acceptable?

- Longitudinal reproducibility error in FreeSurfer: **~5%***
- Hippocampal atrophy in Alzheimer's disease can reach up to **34.2%****

* Brown EM, et al. NeuroImage. 2020

** Blanken AE, et al. NeuroImage: Clinical. 2017

Statistically significant, but clinically insignificant



- Volume reduction trends with DLR
 - Advanced **denoising** and **high-resolution** processing
Deep Resolve Boost and Deep Resolve Sharp
 - Exclusion of "pseudo-structures" caused by noise**
- Clinical impact of accelerated imaging
 - Scan time reduction (GRAPPA 4): 4 min 17 s to **2 min 24 s**
 - Shorter scan time: Better feasibility and less motion artifact**

Approximately 45% scan time reduction is possible via DLR



Hippocampal subfield segmentation using DLR-processed images maintains **high segmentation accuracy**, even when acquisition time is **reduced by 45%**.