



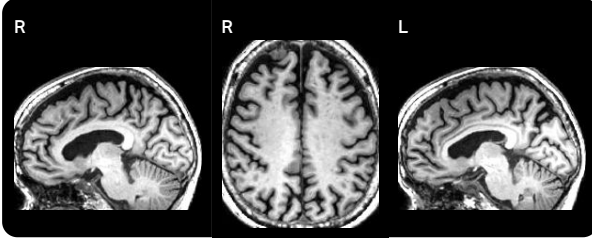
PATIENT ID	NAME	SEX	DATE OF BIRTH	STUDY DATE	SEQUENCE(S)
052_S_4807		F	-(75Y)	04.08.2015 <i>3 previous studies considered.</i>	Accelerated Sag IR-FSPGR (T1) ... (+3)

Clinically Relevant Slices

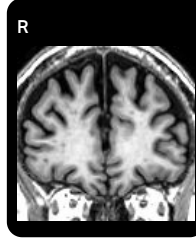
Mesiotemporal



Fronto-parietal



Fronto-polar



Temporo-polar



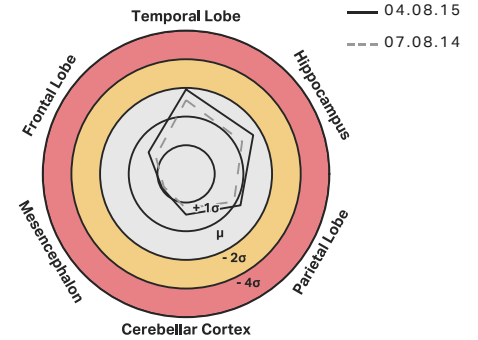
Mesencephalon/pons



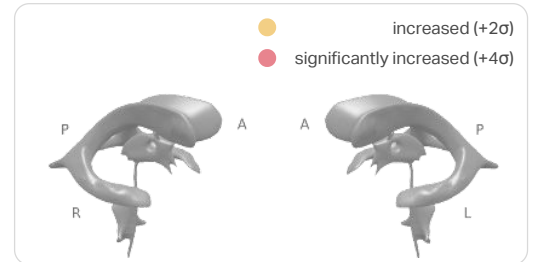
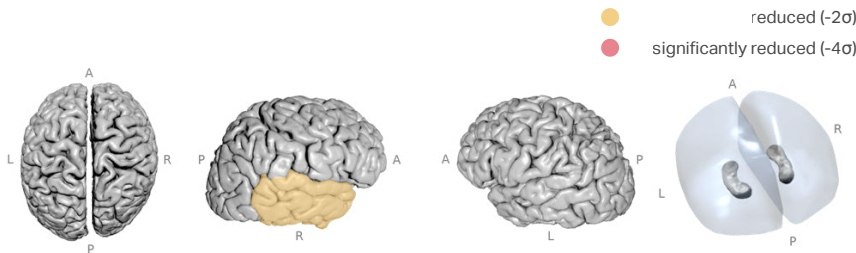
Total Volumes

REGION	PERCENTILE*		VOLUME [ml]		NORMAL RANGE [ml]
	07.08.14	04.08.15	07.08.14	04.08.15	04.08.15
WHOLE BRAIN	25.0 %	12.9 %	1,141.0	1,111.0	1,078.3–1,229.1
TOTAL WHITE MATTER	8.2 %	5.2 %	481.0	469.6	461.3–550.5
TOTAL GREY MATTER	55.3 %	40.0 %	660.0	641.4	588.5–709.7
CEREBRAL CORTEX	52.4 %	36.1 %	452.6	437.4	397.7–494.5

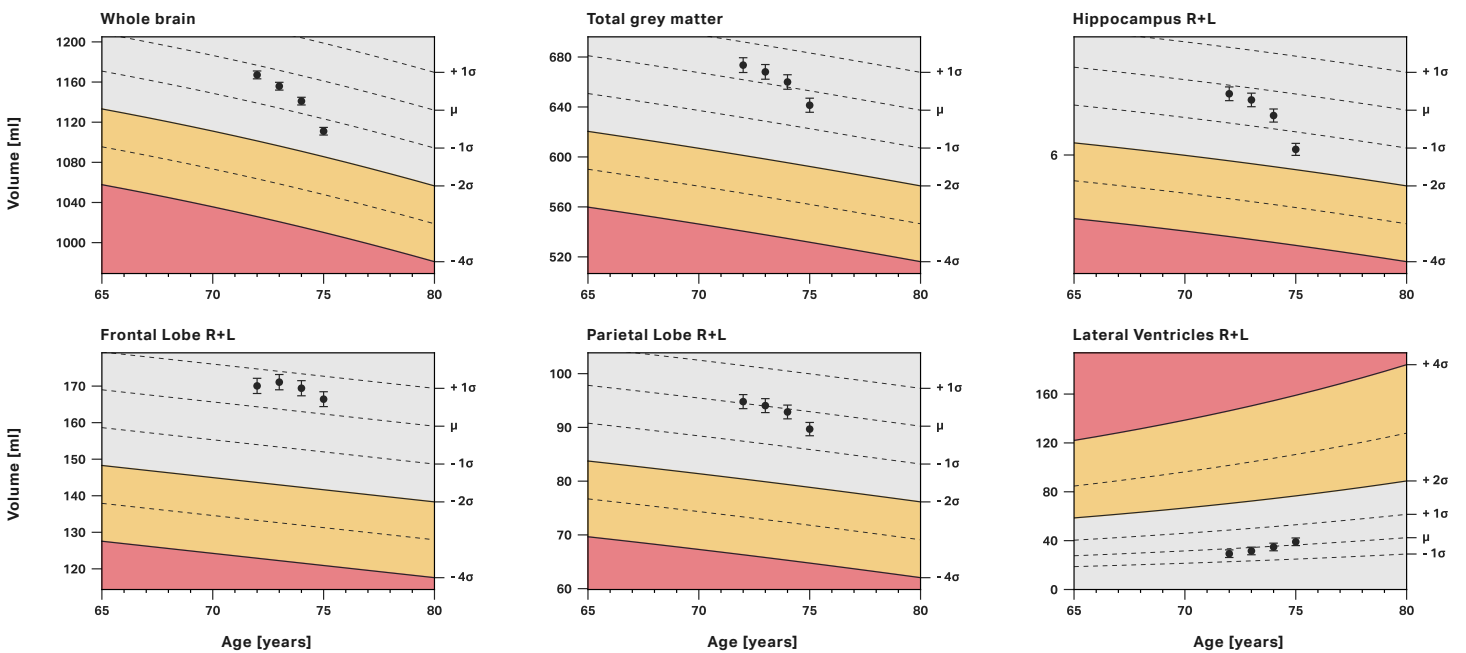
● reduced (-2σ)
● significantly reduced (-4σ)



Schematic Overview (04.08.2015)



Percentile Graphs (04.08.2015 + 3 previous studies considered)



Error bars indicate 90% confidence intervals based on a large cohort of longitudinal measurements where sequence parameters were kept constant.





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052_S_4807		F	-(75Y)	04.08.2015 <i>3 previous studies considered.</i>	Accelerated Sag IR-FSPGR (T1) ... (+3)

Supratentorial Volumes

● reduced (-2σ)
● significantly reduced (-4σ)

REGION	RIGHT					LEFT				
	PERCENTILE*		VOLUME [ml]		NORMAL RANGE [ml]	PERCENTILE*		VOLUME [ml]		NORMAL RANGE [ml]
	07.08.14	04.08.15	07.08.14	04.08.15	04.08.15	07.08.14	04.08.15	07.08.14	04.08.15	04.08.15
FRONTAL LOBE	74.0 %	69.8 %	85.6	84.1	70.9–91.9	72.4 %	67.7 %	83.8	82.3	69.8–90.1
PARIETAL LOBE	42.5 %	28.0 %	45.6	43.7	38.7–52.7	50.9 %	42.6 %	47.3	46.0	39.6–53.7
PRECUNEUS	11.8 %	5.5 %	9.9	9.4	9.0–13.0	16.1 %	7.4 %	9.8	9.3	8.7–12.8
OCCIPITAL LOBE	86.9 %	73.0 %	36.6	34.9	27.8–38.6	65.3 %	52.5 %	37.4	36.0	30.0–41.7
TEMPORAL LOBE	4.2 %	1.1 %	58.1	55.1	56.2–72.3	17.3 %	6.6 %	58.3	55.3	53.4–69.1
HIPPOCAMPUS	20.0 %	7.9 %	3.4	3.2	3.0–4.3	33.4 %	7.9 %	3.2	2.9	2.7–3.8
PARAHIPPOCAMPAL GYRUS	7.4 %	1.7 %	2.7	2.5	2.5–3.5	82.0 %	56.3 %	3.4	3.2	2.7–3.7
ENTORHINAL CORTEX	2.8 %	0.7 %	1.8	1.6	1.8–2.7	10.2 %	4.6 %	1.9	1.8	1.7–2.7
CAUDATE	88.5 %	87.3 %	3.9	3.8	2.6–4.2	75.6 %	78.9 %	3.5	3.5	2.4–3.9
PUTAMEN	79.1 %	70.9 %	4.5	4.4	3.5–4.9	67.6 %	59.7 %	4.5	4.4	3.6–5.0
PALLIDUM	77.9 %	81.9 %	1.5	1.5	1.2–1.7	94.8 %	84.7 %	1.6	1.6	1.2–1.7
THALAMUS	39.1 %	29.3 %	7.7	7.5	6.9–8.6	53.6 %	48.2 %	7.9	7.8	7.0–8.7

Infratentorial Volumes

● reduced (-2σ)
● significantly reduced (-4σ)

REGION	PERCENTILE*			VOLUME [ml]		NORMAL RANGE [ml]
	07.08.14	04.08.15	07.08.14	04.08.15	04.08.15	
BRAINSTEM	98.7 %	98.6 %	31.3	31.1	24.1–30.8	
MESENCEPHALON	92.4 %	87.6 %	8.3	8.2	6.9–8.5	
PONS	99.6 %	99.5 %	18.2	18.1	12.8–17.4	
CEREBELLAR CORTEX	75.2 %	71.8 %	104.1	102.4	82.8–113.3	

Ventricle Volumes

● increased (+2σ)
● significantly increased (+4σ)

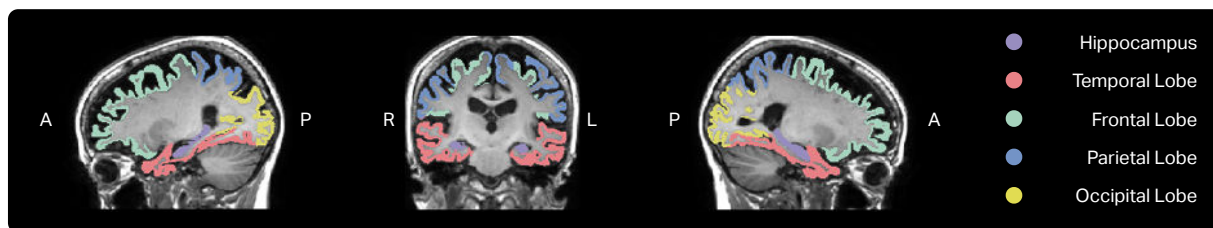
REGION	PERCENTILE*			VOLUME [ml]		NORMAL RANGE [ml]
	07.08.14	04.08.15	07.08.14	04.08.15	04.08.15	
RIGHT LATERAL VENTRICLE	36.0 %	46.5 %	14.7	16.6	7.9–36.4	
LEFT LATERAL VENTRICLE	58.7 %	68.1 %	20.1	22.4	8.7–39.4	
THIRD VENTRICLE	82.1 %	85.3 %	1.5	1.6	0.7–1.9	
FOURTH VENTRICLE	73.2 %	79.4 %	1.7	1.7	0.8–2.1	

*The percentile gives the expected proportion of the normal population with the same covariates having a lower volume than the one measured (based on the measurements of a normal collective).

Quality Control (04.08.2015)

IMAGE QUALITY	Good
NOTES	See appendix

Automatic Segmentation (04.08.2015)





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Appendix

The automated quality control yielded a good image quality. However, the following points need attention:

1. The slice thickness of the T1 sequences (04/08/15, 07/08/14, 13/08/13 and 15/02/13) could be further improved. For optimal results, a max. slice thickness of 1.1 mm is recommended.
2. The spacing between slices of the T1 sequences (04/08/15, 07/08/14, 13/08/13 and 15/02/13) could be further improved. For optimal results, a max. spacing between slices of 1.1 mm is recommended.

	04.08.2015 SCAN	07.08.2014 SCAN
DICOM FIELD	T1	T1
SEQUENCE DESCRIPTION	Accelerated Sag IR-FSPGR	Accelerated Sag IR-FSPGR
SCANNER MANUFACTURER	GE MEDICAL SYSTEMS	GE MEDICAL SYSTEMS
SCANNER TYPE	DISCOVERY MR750	DISCOVERY MR750
FIELD STRENGTH	3.0 T	3.0 T
REPETITION TIME	7.32 ms	7.32 ms
ECHO TIME	3.02 ms	3.02 ms
INVERSION TIME	400.00 ms	400.00 ms
PIXEL SPACING	1.1 mm × 1.1 mm	1.1 mm × 1.1 mm
SLICE THICKNESS	1.2 mm	1.2 mm
SPACING BETWEEN SLICES	1.2 mm	1.2 mm
ACQUISITION MATRIX	[0, 256, 256, 0]	[0, 256, 256, 0]
NUMBER OF SLICES	196	196

	13.08.2013 SCAN	15.02.2013 SCAN
DICOM FIELD	T1	T1
SEQUENCE DESCRIPTION	Accelerated Sag IR-FSPGR	Accelerated Sag IR-FSPGR
SCANNER MANUFACTURER	GE MEDICAL SYSTEMS	GE MEDICAL SYSTEMS
SCANNER TYPE	DISCOVERY MR750	DISCOVERY MR750
FIELD STRENGTH	3.0 T	3.0 T
REPETITION TIME	7.32 ms	7.32 ms
ECHO TIME	3.02 ms	3.02 ms
INVERSION TIME	400.00 ms	400.00 ms
PIXEL SPACING	1.1 mm × 1.1 mm	1.1 mm × 1.1 mm
SLICE THICKNESS	1.2 mm	1.2 mm
SPACING BETWEEN SLICES	1.2 mm	1.2 mm
ACQUISITION MATRIX	[0, 256, 256, 0]	[0, 256, 256, 0]
NUMBER OF SLICES	196	196

