

# Multi-contrast MRI Atlas of the Cynomolgus Macaque Brain

# INTRODUCTION

- Species-specific anatomical brain templates are required for accurate spatial normalization of single-subject MRI, automated brain tissue segmentation, and surface generation. Templates facilitate the comparison across subjects, studies, and laboratories.
- Multiple templates are available for rhesus macaques, but only a few for cynomolgus macaques, which differ in brain morphology and endocranial volume<sup>3,10</sup>.
- Most templates are based on T1-weighted images, but other MRIcontrasts may provide essential additional information.
- In this work, we present high-quality multi-contrast symmetric and asymmetric cynomolgus macaque brain templates along with wholebrain parcellations and surface reconstruction.

## METHODS

• 14 healthy female cynomolgus macaques (Macaca fascicularis) within an age range of 8 to 20 years (mean = 15 years) were included.

	$T_1w$	T <sub>2</sub> w	GRE	MTw	PDw	$T_1w$
sequence	3D MPRAGE	FSE	3D ME-GRE	3D FLASH	3D FLASH	3D FLASI
resolution (mm)	0.5x0.5x0.5	0.4x0.4x0.9	0.31x0.31x0.31	0.5x0.5x0.5	0.5x0.5x0.5	0.5x0.5x0
repetition time (ms)	2700	8000	57	30	25	10
echo time (ms)	2.7	17	3.7/4.9/48	3.2	3.2	3.2
flip angle (deg)	8	180	20	5	5	15
acquisition time (ms)	) 14.3	5.2	24	15.3	6.3	5.1
3T Siemens MAGNETOM Prisma: 7 cm single loop coil						

31 Siemens MAGNETOM Prisma; 7 cm single loop coll



- T1-weighted (T1w) template was generated using ANTs, and the produced warps were further used to transform weighted contrasts and quantitative parametric maps into the same space<sup>2,11</sup>.
- QSM reconstruction was carried out using a multi-scale dipole inversion approach1. R2\* maps were estimated by fitting the magnitude of signal decay over ten echoes.
- Maps of magnetization transfer saturation (MTsat) and apparent T1 relaxation time (**T1app**) were calculated using the method described by Helms et al., 2008.
- Multi-contrast symmetric and asymmetric templates were generated with an isotropic resolution of 0.25 mm and 0.5 mm, respectively.

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T1w: T1-weighted; MTw: Magnetization Transfer-weighted; mGRE: mean of multiecho gradient-echo magnitude across echo times; T2w: T2-weighted; QSM: Quantitative Susceptibility Mapping; R2\*: effective transverse relaxation rate; Mtsat: Magnetization Transfer saturation; T1app: apparent T1 relaxation time.

- The T1w template provides an excellent gray and white matter contrast.
- QSM and R2\* templates make subcortical gray matter nuclei clearly distinguishable from other parts of the brain.
- MTsat template reveals differences in the degree of myelination.
- In contrast to T1w, the T1app template provides quantitative values of the longitudinal relaxation time.



### **Tissue segmentation**

- used tissue
- correctly.

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RESULTS



The T1w template was three-class segmentation providing cerebrospinal fluid (CSF), gray matter (GM), and white matter (WM) probability masks.

The GM segmentation failed in the case of subcortical gray matter nuclei. Only caudate and putamen were classified

QSM and R2\* templates have been used to overcome this limitation (see talk #1248).





## CONCLUSION

- This study presents high-resolution multi-contrast MRI symmetric and asymmetric brain templates of cynomolgus macaque in stereotaxic space.
- Provided templates include both weighted MRI as well as quantitative parametric maps.
- The availability of multiple MRI contrasts allowed for better and more detailed visualization of brain structures and may be used for further anatomical parcellation.
- Transformation to commonly used rhesus macaque templates will support comparative studies across species.

#### Surfaces

- Nonlinear alignment warps to the NMT v2 atlas were used to propagate the Cortical Hierarchy Atlas of the Rhesus Macaque (CHARM) and Subcortical Atlas of the Rhesus Macaque (SARM) labels<sup>4,6</sup>.
- Similarly, nonlinear warps to the D99 atlas and INIA19 atlas were used to propagate D99 and NeuroMaps labels into the template space<sup>8,9</sup>.

#### References

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