

A4 / LEARN White Matter Hyperintensity Segmentation Methods

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We segmented white matter hyperintensity (WMH) from FLAIR MRI for each MR imaging session independently. To follow is the step-by-step processing procedure for accurate segmentation of WMH using an automated tool (hypermapp3r.readthedocs.io).

1. We identified FLAIR and MPRAGE MRI from the same session for all subjects
2. Using `spm_coreg`, we co-registered FLAIR MRI to MPRAGE MRI from the same session
3. Using `fsl_anat` (https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/fsl_anat), we performed bias field correction on the FLAIR image.
4. We used `ICVmapper` for brain extraction (i.e. skull removal) on MPRAGE
5. We resliced the brain-extracted MPRAGE to the subject space FLAIR
6. Using `HyperMapper` algorithm, we segmented WMH in the FLAIR space.
7. We upsampled WMH from `HyperMapper` to the MPRAGE space
8. We co-registered the baseline `FreeSurfer` segmentation mask (<https://surfer.nmr.mgh.harvard.edu/>) to the subject space MPRAGE to create a subject-specific white matter mask from the following aseg ROIs (2 41 43 44 77 4 5 11 50)
9. The WMH `HyperMapper` images were masked by the `FreeSurfer` white matter mask to remove unwanted voxels
10. We computed the WMH volume and intracranial volume in cubic mm.

References

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