

MS Journal Appendix for MRI methodology

| Hardware | |
|-----------------------------------|--|
| Field strength | |
| Manufacturer | |
| Model | |
| Coil type (e.g. head, surface) | |
| Number of coil channels | |

| Acquisition sequence | | |
|---|-----|----|
| Type (e.g. FLAIR, DIR, DTI, fMRI) | | |
| Acquisition time | | |
| Orientation | | |
| Alignment (e.g. anterior commissure/posterior commissure line) | | |
| Voxel size | | |
| TR | | |
| TE | | |
| TI | | |
| Flip angle | | |
| NEX | | |
| Field of view | | |
| Matrix size | | |
| Parallel imaging | Yes | No |
| If used, parallel imaging method: (e.g. SENSE, GRAPPA) | | |
| Cardiac gating | Yes | No |
| If used, cardiac gating method: (e.g. PPU or ECG) | | |
| Contrast enhancement | Yes | No |

Acquisition sequence

If used, provide name of contrast agent, dose and timing of scan post-contrast administration

Other parameters:

| Image analysis methods and outputs | |
|--|--|
| Lesions | |
| Type (e.g. Gd-enhancing, T2-hyperintense, T1-hypointense) | |
| Analysis method | |
| Analysis software | |
| Output measure (e.g. count or volume [ml]) | |
| Tissue volumes | |
| Type (e.g. whole brain, grey matter, white matter, spinal cord) | |
| Analysis method | |
| Analysis software | |
| Output measure (e.g. absolute tissue volume in ml, tissue volume as a fraction of intracranial volume, percentage change in tissue volumes) | |
| Tissue measures (e.g. MTR, DTI, T1-RT, T2-RT, T2*, T2', ¹H-MRS, perfusion, Na) | |
| Type (e.g. whole brain, grey matter, white matter, spinal cord, normal-appearing grey matter or white matter) | |
| Analysis method | |
| Analysis software | |
| Output measure | |
| Other MRI measures (e.g. functional MRI) | |
| Type (e.g. whole brain, grey matter, white matter, spinal cord, normal-appearing grey matter or white matter) | |
| Analysis method | |
| Analysis software | |
| Output measure | |

Other analysis details: