Table S1

| GE MR imaging protocol |  |  |  |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- |
|  | Plane | TR (ms) | TE (ms) | FOV (mm) | Slice thickness (mm) |
| T2 FSE | sag | 3997.0 | 91.0 | 24.0 | 3 |
| T2 FSE | ax | 3214.0 | 91.0 | 24.0 | 4 |
| T2 FSE | cor | 3615.0 | 91.0 | 24.0 | 4 |
| DWI b1000 | ax | 10265.0 | minimum | 24.0 | 4 |
| 3D SWAN | ax | minimum | 40.6 | 25.0 | 1.5 (isotropic) |
| T1 FSE | ax | 512.0 | Min Full | 24.0 | 4 |
| FLAIR | ax | 9745.0 | 133.1 | 24.0 | 4 |
| DSC Perfusion | ax | 1500.0 | 35.0 | 24.0 | 4 |
| T1 fs FSPGR (+C) | ax |  | out of Phace | 25.6 | 1.2 (isotropic) |
| T1 FSE (+C) | ax | 590.0 | Min Full | 24.0 | 4 |

Table S2

| Philips MR imaging protocol |  |  |  |  |
| :--- | :--- | :--- | :--- | :--- |
|  | Plane | TR (ms) | TE (ms) | FOV (mm) |
|  | sag | $3000-5500$ | 100 | 24.0 |
| T2 TSE | ax | $3000-5500$ | 100 | 24.0 |
| T2 TSE | cor | $3000-5500$ | 100 | 3 |
| T2 TSE | ax | 4927 | 90 | 24.0 |
| DWI b1000 | ax | 51 | 12 | 24.0 |
| 3D SWI | ax | $400-680$ | 12 | 24.0 |
| T1 TSE | ax | 10000 | 140 | 24.0 |
| FLAIR | ax | 1677 | 40 | 24.0 |
| DSC Perfusion | ax | 13 | 6 | 24.0 |
| T1 fs WATS + +C) | ax | $400-680$ | 12 | 24.0 |
| T1 TSE (+C) | ax |  | 24.0 | 4 |



Fig. S1 Representative flow cytometry histograms of CD14/SSc-gated monocytes (left) and the CD14+ subsets analysed.


Fig S 2
Fig. S2 M2-monocyte subsets representation according to ependymal enhancement. Graphs were generated using Prism GraphPad 7.0a Macintosh. Linear regression and Pearson correlation coefficient did not measure any linear relationship between the variables.


Fig. S3 M2-monocyte subsets representation according to corpus callosum infiltration. Graphs were generated using Prism GraphPad 7.0a Macintosh. Linear regression and Pearson correlation coefficient did not measure any linear relationship between the variables.


Fig. S4 M2-monocyte subsets representation according to midline shift. Graphs were generated using Prism GraphPad 7.0a Macintosh. Linear regression and Pearson correlation coefficient did not measure any linear relationship between the variables.


Fig S 5
Fig. S5 M2-monocyte subsets representation according to macroscopic hemorrhage. Graphs were generated using Prism GraphPad 7.0a Macintosh. Linear regression and Pearson correlation coefficient did not measure any linear relationship between the variables.


Fig S6
Fig. S6 M2-monocyte subsets representation according to the edema score. Graphs were generated using Prism GraphPad 7.0a Macintosh. Linear regression and Pearson correlation coefficient did not measure any linear relationship between the variables.


Fig. S7 M2-monocyte subsets representation according to the score of intratumoral susceptibility signal intensity (ITSS). Graphs were generated using Prism GraphPad 7.0a Macintosh. Linear regression and Pearson correlation coefficient did not measure any linear relationship between the variables.

