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MR-based Parenchymal CSF water fraction (CSFF) is a marker of glymphatic clearance function and relates to beta-amyloid deposition on ¹¹C-PiB PET

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Synopsis

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Glymphatic clearance has recently gained increasing attention as a major driver of neurodegenerative diseases like Alzheimer's disease. Perivascular spaces is believed to reflect glymphatic clearance such that enlargement of the PVS is believed to reflect blocked glymphatic clearance. We present preliminary work on a MR-based FAST-T2 CSF water fraction map to study glymphatic clearance. Our results show a quadratic relationship between CSFF and PiB PET SUVR in AD-specific regions. This indicates that CSFF could be a biomarker of dysfunctional glymphatic clearance, leading to early beta amyloid accumulation, since early on, CSFF increases without a significant change in PiB PET SUVR.

Introduction

Glymphatic clearance has recently gained increasing attention as a major driver of neurodegenerative diseases like Alzheimer's disease (AD)¹. Perivascular space (PVS) is of the key route of the glymphatic system, facilitates the water exchange between CSF and ISF, and is believed to reflect glymphatic function. The enlargement of and fluid stasis in the PVS is believed to reflect blocked glymphatic clearance². This dysfunctional clearance can lead to the deposition of extracellular amyloid beta and intracellular tau protein, which can be quantified using positron emission tomography (PET)³. Therefore, early detection of the clearance dysfunction is crucial to the prevention and diagnosis of neurodegenerative disease⁴.

In this study, we assessed the relationship between CSF water fraction (CSFF) mapping and beta-amyloid deposition using Pittsburgh compound B (11C-PiB) PET. We further assessed CSF water fraction mapping at low (5mm thickness) and high (2mm thickness) resolution⁵ to exclude the possible concerns about the partial volume effect (PVE).

Material and Methods

Subjects

A total of 45 subjects (21 normal controls, 11 MCI/AD, and 13 unknown diagnoses) were scanned on a 3T Siemens Prisma scanner with a 64-channel head-neck coil. The six echoes (TE=0, 7.5, 17.5, 67.5, 147.5, 307.5 ms) of MR FAST-T2 data were acquired with both 2mm (highRes) and 5mm (lowRes) thickness with common in-plane resolution 0.93mmX0.93mm.⁶ T1 MPRAGE images (isotropic 0.5 mm3) were acquired for coregistration and ROI parcellation. A total of 18 subjects (11 controls and 7 MCI/AD) had concomitant PiB PET, all of which had low-resolution FAST-T2 data.

Processing

The 6 echoes FAST-T2 data were fitted into the three-water compartment model with regularized nonlinear least square⁷. Then the CSFF map was obtained by dividing the long T2 (T2>200 ms) compartment signal magnitude by the total signal magnitude. The CSFF was co-registered to the FreeSurfer T1 space for ROI value extraction⁸. The PiB PET data was processed with temporal realignment, summation of 60-90 min time frames, coregistration to FreeSurfer T1 space, and calculation of the SUVR with a cerebellum cortex reference⁹. All the ROIs used in this work were eroded by 1 voxel to reduce the PVE.

Analysis

First, the regression was examined between CSFF and the PiB SUVR in AD-mask¹⁰ to evaluate the linear and nonlinear association between beta-amyloid deposition and CSFF. Secondly, we qualitatively checked the low- and high-resolution CSFF map and overlayed them on the susceptibility image (SWI) to see the consistency of hyperintensity spots (CSF/free water in PVS) on the CSFF map and hypointensity spots (medullary vein) surrounding lateral ventricle. Thirdly, we compared the ROI values of CSFF from both low- and high-resolution maps. The linear regression and Bland-Altman analyses were done to check the consistency between the low- and high-resolution CSFF to further exclude the concerns about PVE.

Results

Figure 1 shows the quadratic relationship between the PiB SUVR and the low-resolution CSFF in ADmask (p₁=0.006 for the first-order term, p₂=0.001 for the second-order term, whole model R²=0.88). The healthy controls and MCI/AD subjects are presented with blue dots and red dots, respectively.

Figure 2 shows the high and low-resolution CSFF maps overlayed on the same subject's SWI images. (a) is high-resolution CSFF on SWI, and (b) is low-resolution CSFF on SWI. High CSFF signals around the lateral ventricle and along with medullary veins are observed.

Figure 3 shows the linear regression and Bland-Altman plots comparing low- and high-resolution CSFF. The data show for: (a) the global cerebral cortex (p<0.0001, R^2 =0.91); (b) the global cerebral white matter (p<0.0001, R^2 =0.72); (c) the parietal lobe (p<0.0001, R^2 =0.86); (d) the temporal lobe (p<0.0001, R^2 =0.91); (e) the occipital lobe (p<0.0001, R^2 =0.94); (f) the frontal lobe (p<0.0001, R^2 =0.77). The Bland-Altman plots show that 97% of points lie within 1.96 standard deviations (which corresponds to a 95% confidence interval).

Discussion and conclusion

We present preliminary work on an MR-based FAST-T2 CSF water fraction map to study glymphatic clearance. Our results show a quadratic relationship between CSFF and the PiB PET SUVR in AD-specific regions. This indicates that CSFF could be a biomarker of dysfunctional glymphatic clearance, leading to early beta-amyloid accumulation, since early on, CSFF increases without a significant change in PiB PET SUVR (Figure 1).

Figure 2 (a) and (b) show that the PVS surrounding medullary veins can be seen on both high- and low-resolution CSFF confirmed by cross-validation on SWI. Hyperintense CSFF foci with the medullary vein on SWI surrounding the lateral ventricle indicate a potential CSF clearance pathway, which is also corresponding to the findings of recent DTI-ALPS studies¹¹.

Figure 3 shows that the CSFF value is consistent in both low (5mm thickness) and high (2mm thickness) resolution mapping within 1 voxel-eroded ROIs. This will justify the low-resolution FAST-T2 data in the analysis when high-resolution (thin-slice) images are not required/available.

Enlarged PVS can be seen on both low and high-resolution CSFF but smaller ones are not visible on CSFF or conventional T1w or T2w. CSFF corresponding to the long T2 signal that represents free water could be a more accurate measure of total CSF in the glymphatic system than the total PVS load.

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Figure 1 Relationship between CSFF and PiB PET SUVR in AD mask. Blue dots represent healthy control and the red dots indicate MCI/AD subjects. It clearly shows that using both PiB SUVR and CSFF in AD-mask can distinguish normal controls from MCI/AD subjects.

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Figure 2 CSF fraction (CSFF) map at (a) high (0.93x0.93x5 mm3) and (b) low (0.93x0.93x2 mm3) resolution overlayed on the susceptibility-weighted image (SWI). Red arrows point to the medullary veins, at which CSFF shows a hyperintensity signal.



Figure 3 Linear regression and Bland-Altman plots for high and low-resolution CSFF in (a) GM, (b) WM, (c) parietal lobe, (d) temporal lobe, (e) occipital lobe, and (f) frontal lobe. The CSFF from high and low-resolution FAST-T2 data are of high consistency across all the considered global and local ROIs.

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