S5. Tissue and parcellated volumes for each cohort

We here show results from a repeat of the analysis performed for Table 2, using brains from cohort 1 (Table 3), mean (standard deviation) age 128.4 (7.4) days and cohort 2 (Table 4), age 471.5 (5.9) days separately. The pattern of mean volumes, Tc1 > WT (or vice versa) was the same, in each structure, in both cohorts, and the same as when both were combined (Table 2). The tendency when analysing the two cohorts separately was to decrease significance: p-values were generally greater and more likely to exceed the arbitrary significance threshold of p=0.05.

Table 3

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<td>mean</td>
<td>std</td>
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Cohort 1 mean absolute volumes (mm\(^3\)), by group, of (a) probabilistic tissues: \( BV = GM + WM \); \( TIV = BV + CSF \) and (b) parcellated regions via integration of Jacobian determinants, and their standard deviations. (Bonferroni-adjusted two-tailed p-values shown, omitted where >>0.05).

### Table 4

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<td>1.30x10(^{-5})</td>
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Cohort 2 mean absolute volumes (mm\(^3\)), by group, of (a) probabilistic tissues: \( BV = GM + WM \); \( TIV = BV + CSF \) and (b) parcellated regions via integration of Jacobian determinants, and their standard deviations. (Bonferroni-adjusted two-tailed p-values shown, omitted where >>0.05).