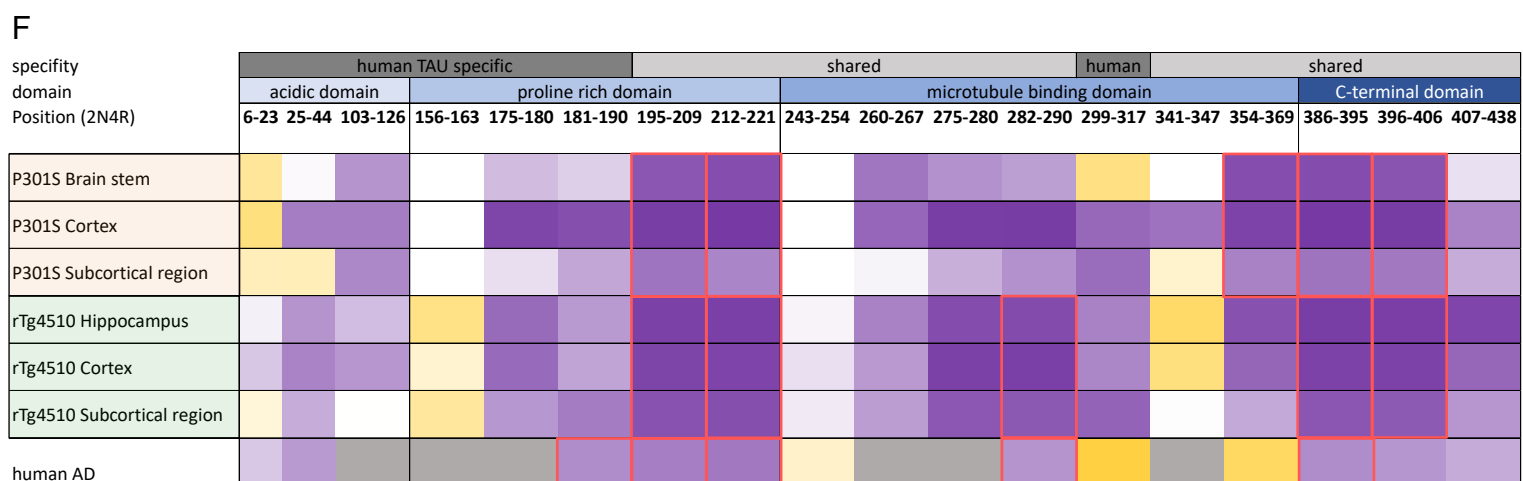
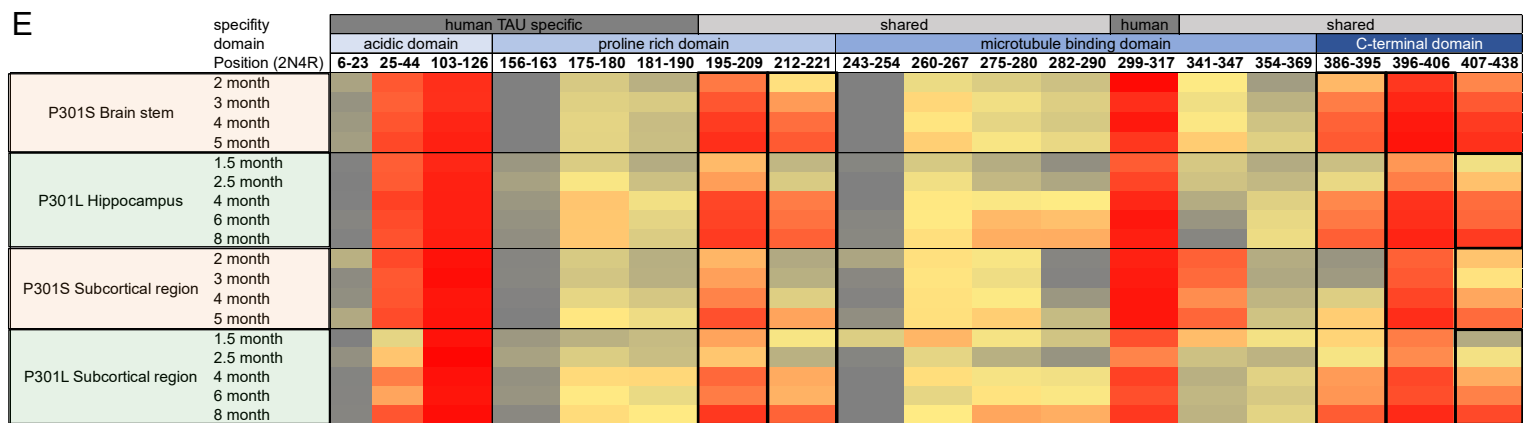


modification extent: 0% 50% 100% modified ■ peptides with highest progression



Pearson correlation: 1 no correlation -1

Average top 5 peptides

not monitored