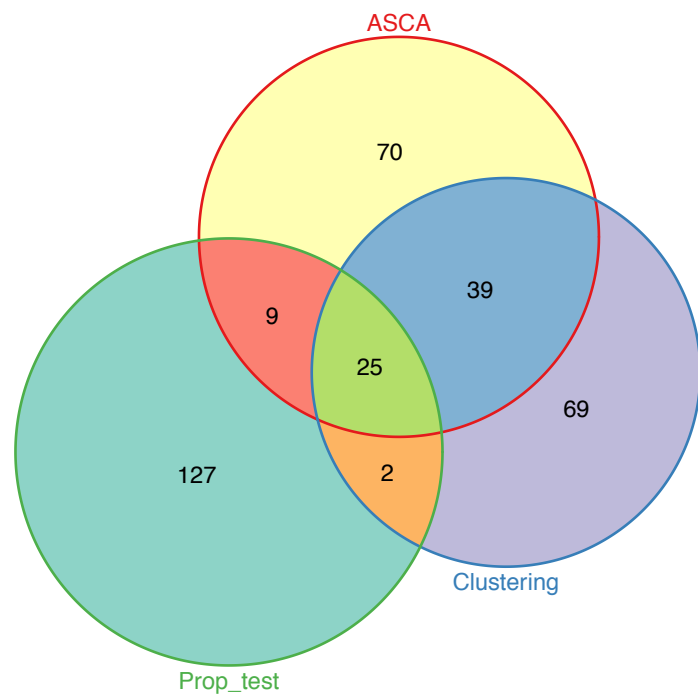
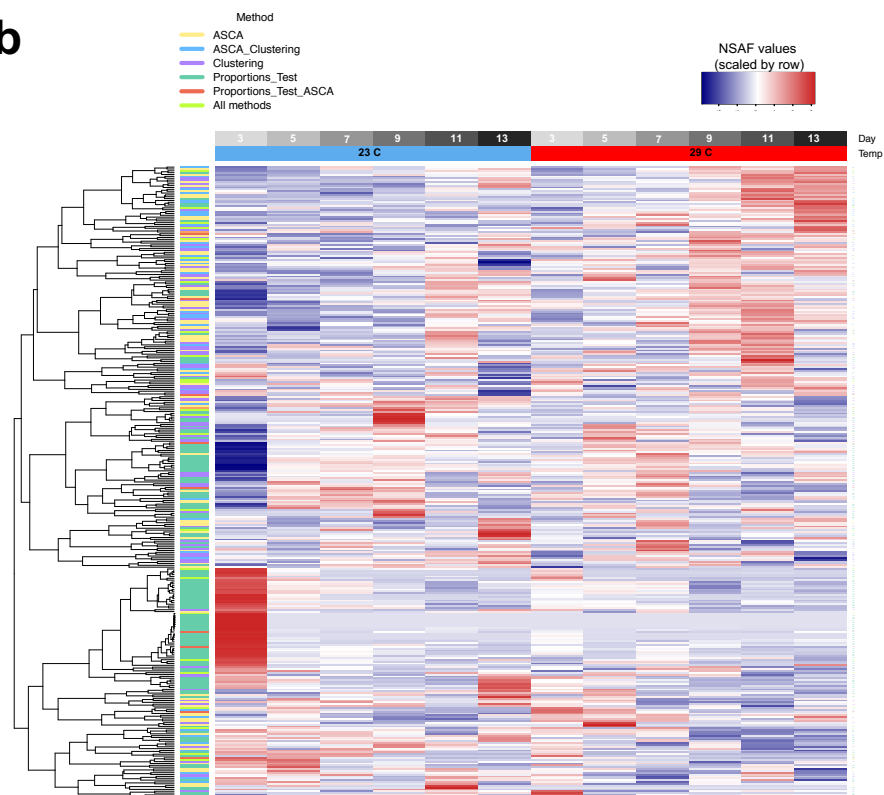
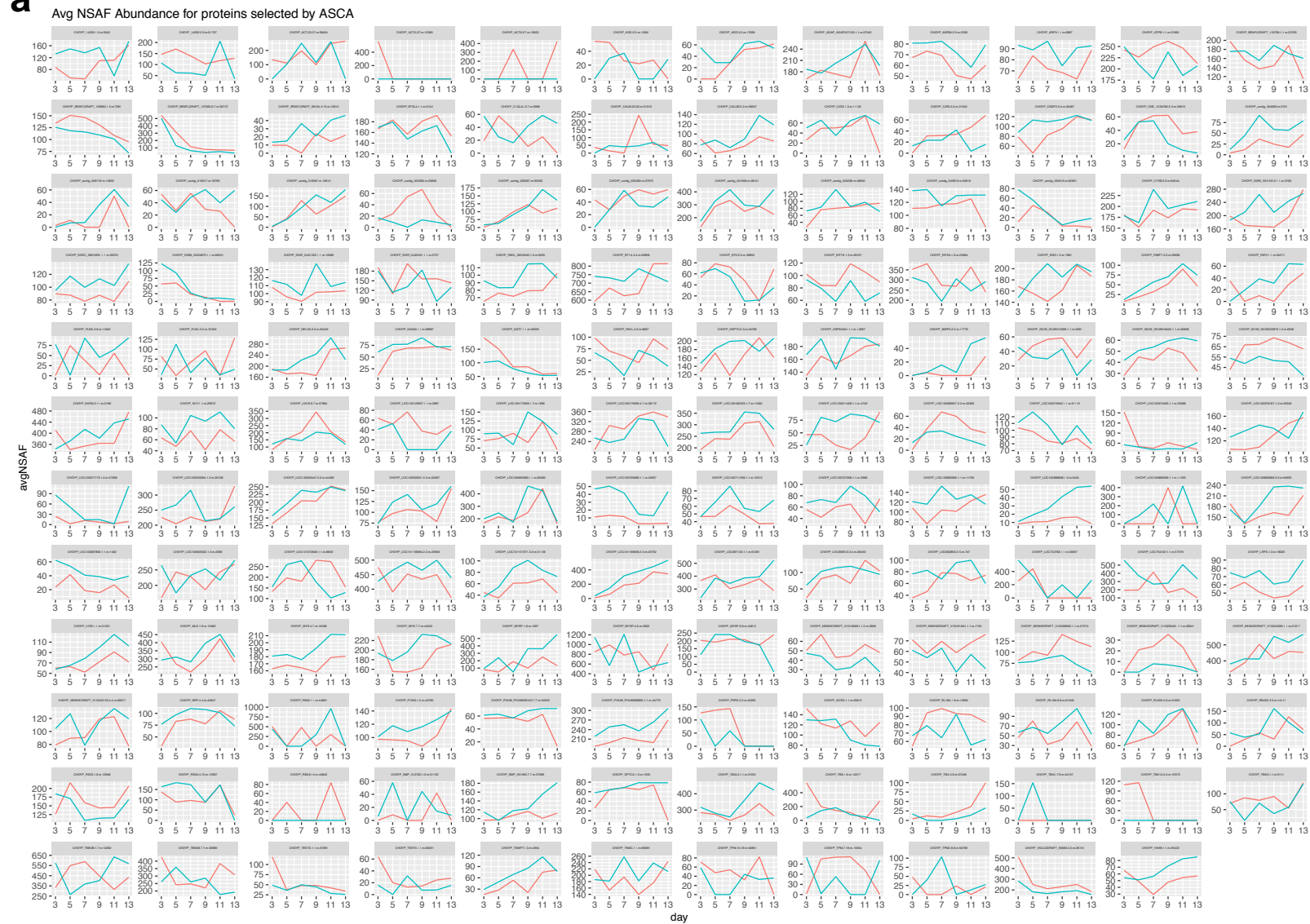
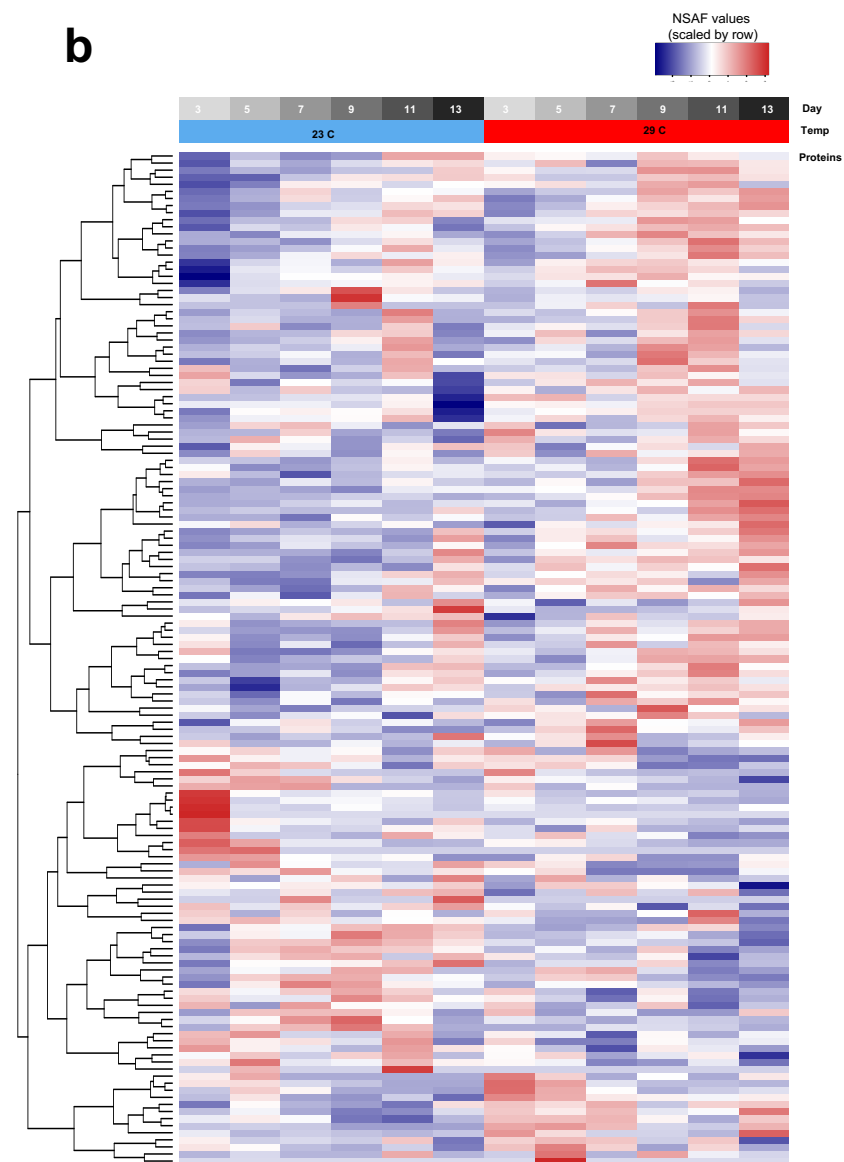


**a****b**

a

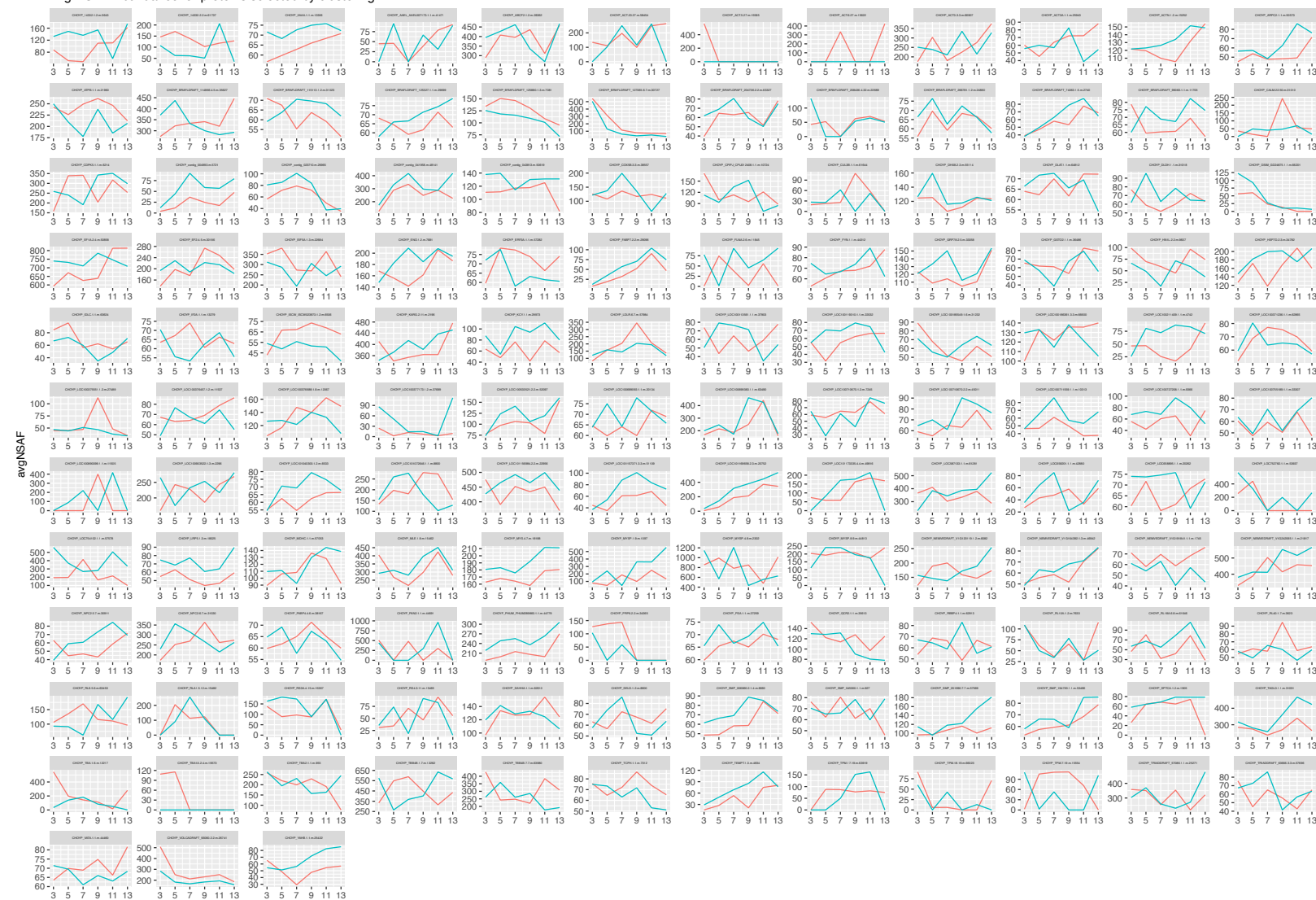


b

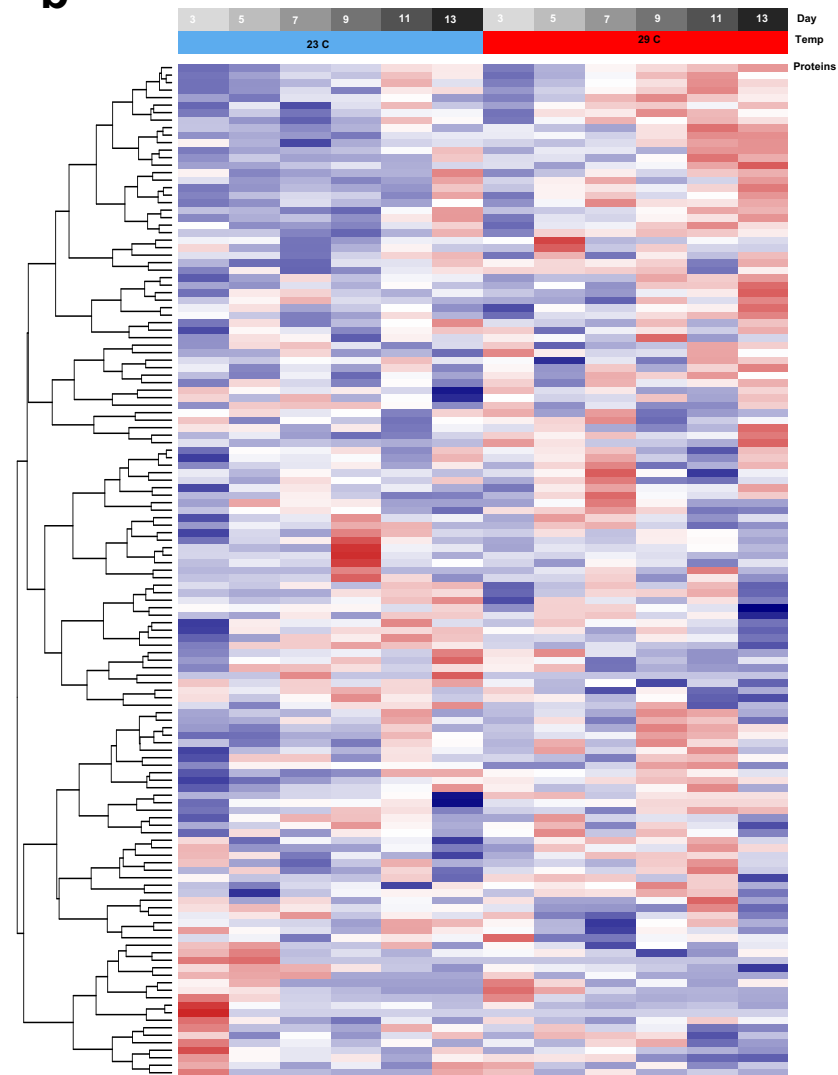


a

Avg NSAF Abundance for proteins selected by clustering

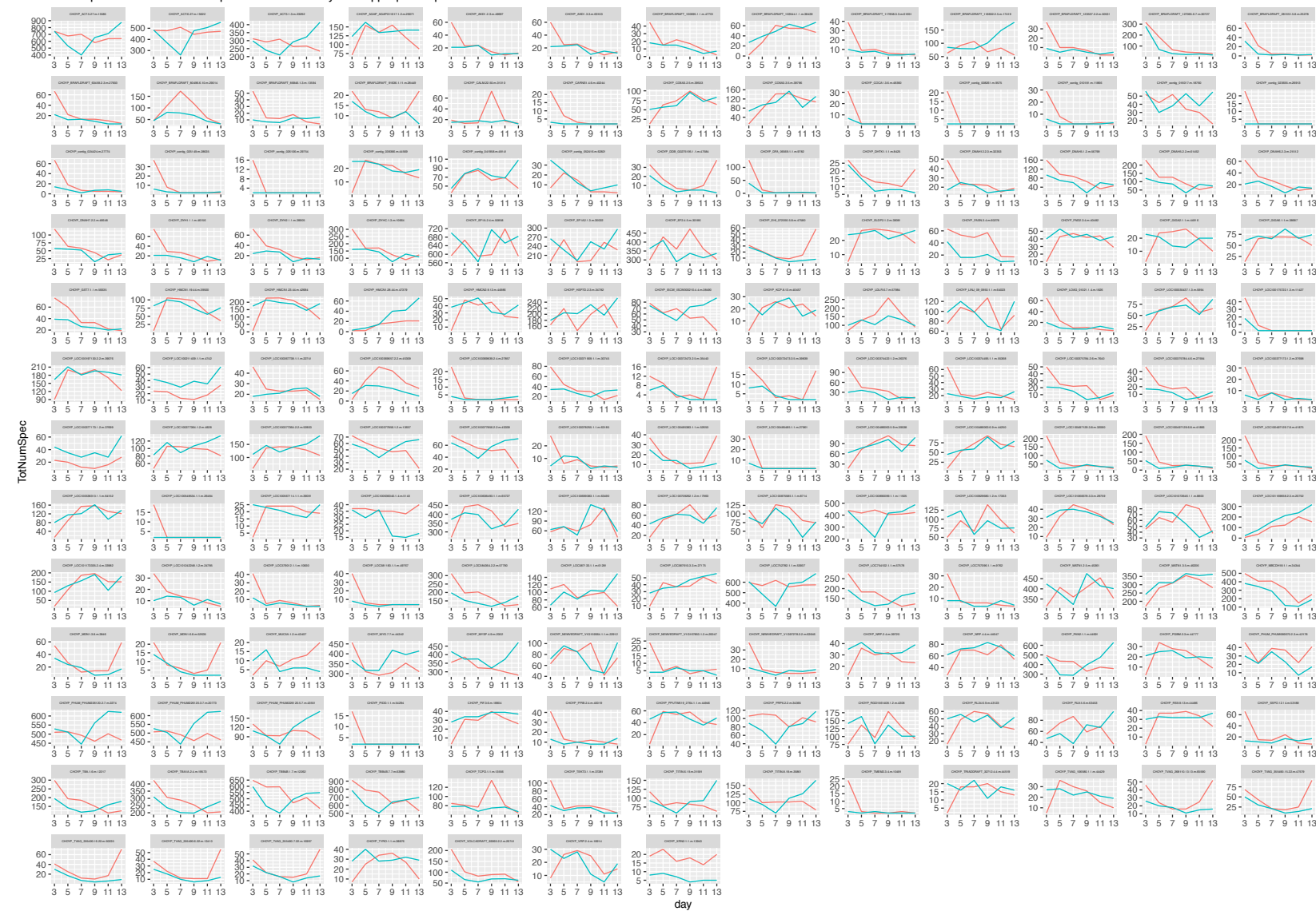


b



**a**

Total Spectral Abundance for proteins selected by ChiSq prop. test pval 0.1

**b**