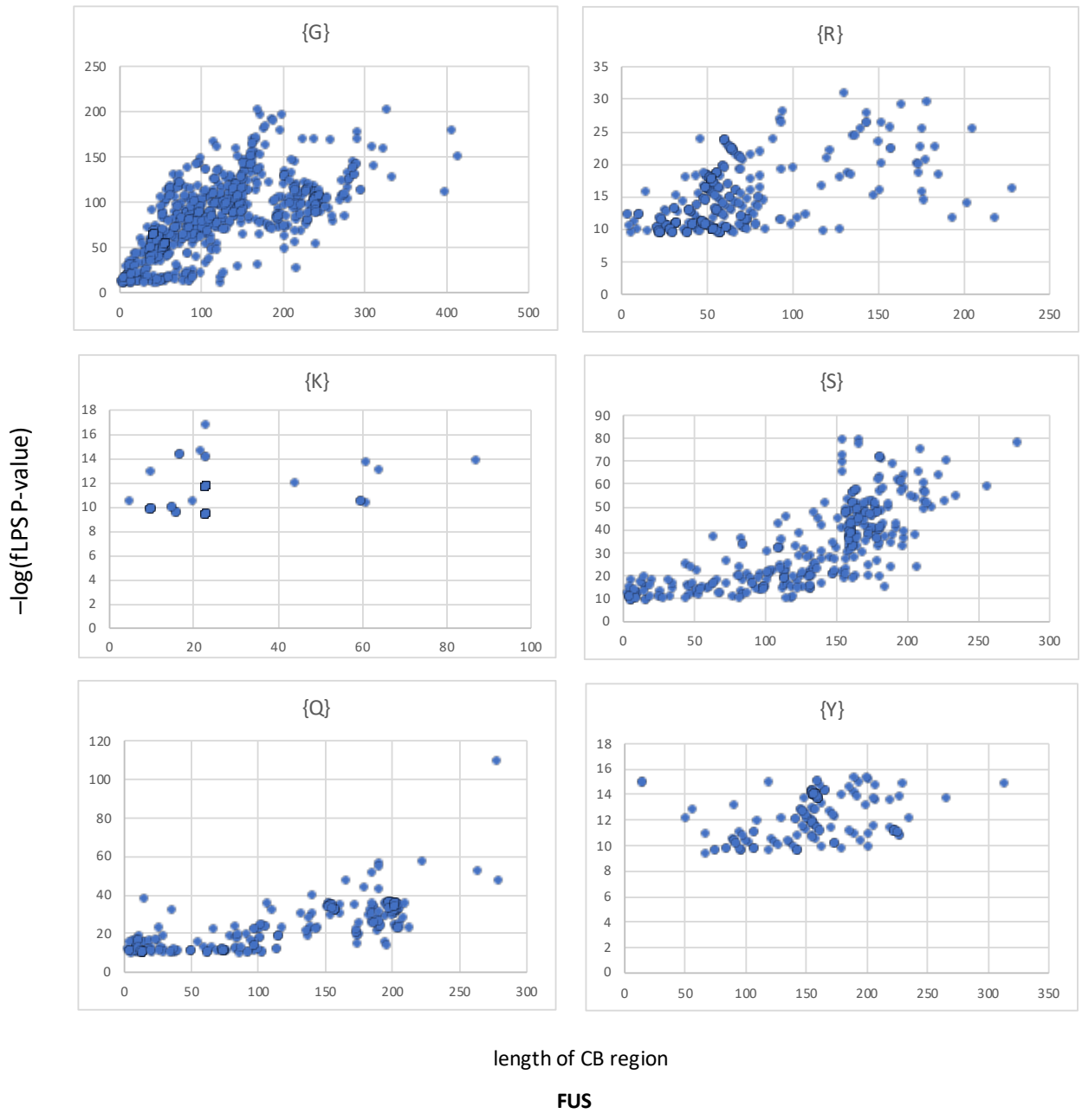
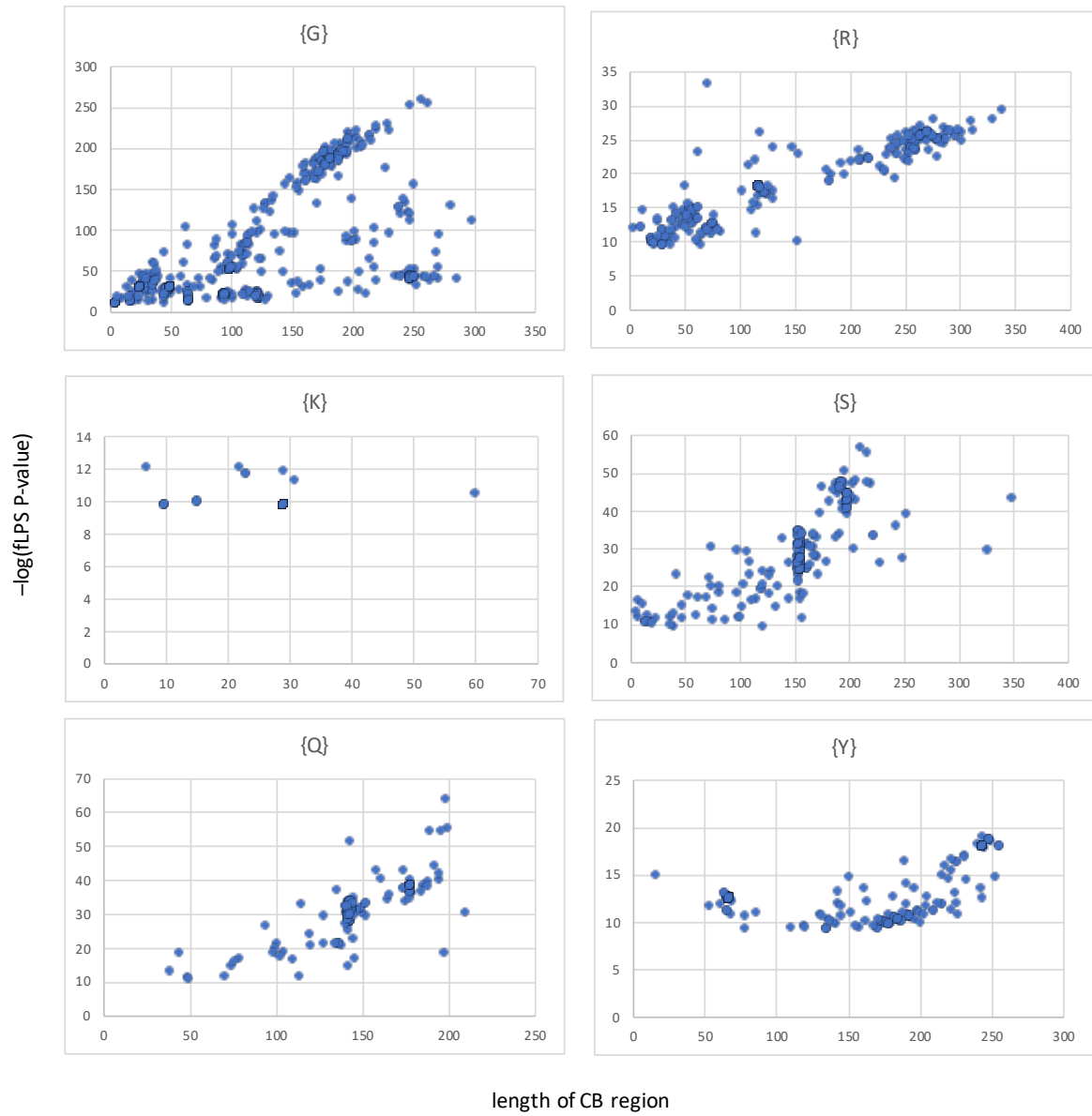


Figure Legend:

Plots of CB region length (in residues) versus $-\log(\text{P-value})$ for the six most common single-residue biases for each of FUS, TAF15 and EWSR1 respectively. The plots for each protein are arrayed on one page each. The P-values are from the fLPS program.

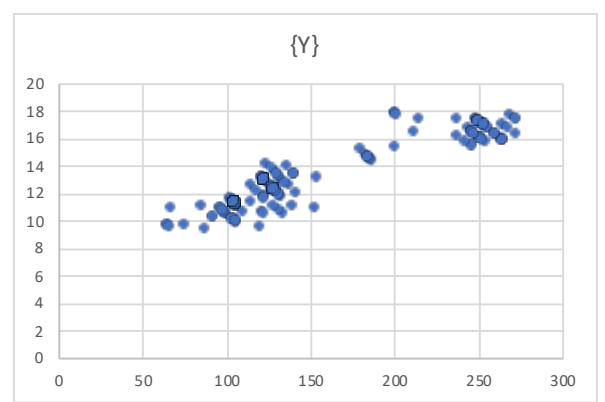
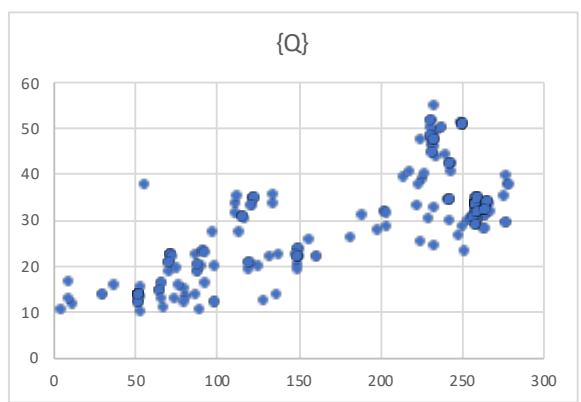
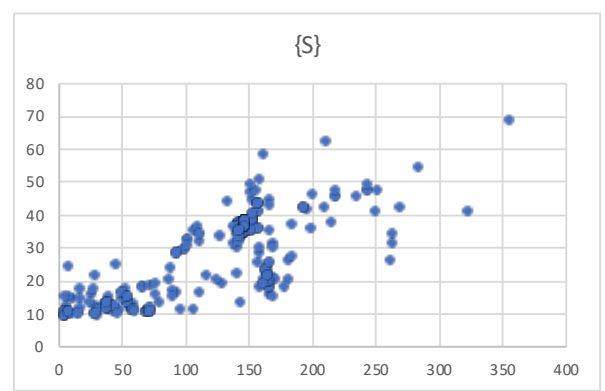
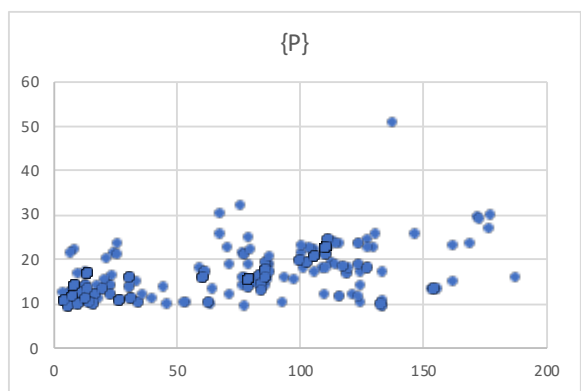
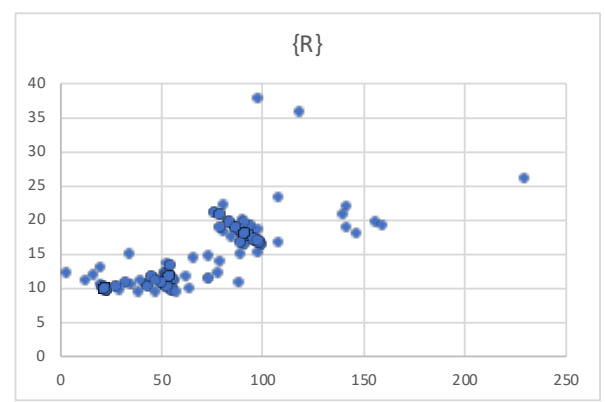
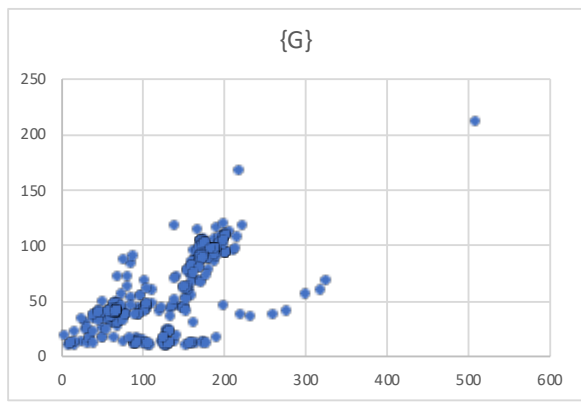




length of CB region

TAF15

length of CB



length of CB region

EWSR1