**Figure Legend:**
Total counts of sequences containing single-residue CB regions with P \( \leq 1 \times 10^{-4} \) for (A) FUS, (B) TAF15 and (C) EWSR1 orthologs. The CB regions are labelled \( \{x\} \), where \( x \) is the biasing residue. These are counts of sequences, not of CB regions.
(A) FUS (total 396)

(B) TAF15 (total 250)

(C) EWSR1 (total 280)