

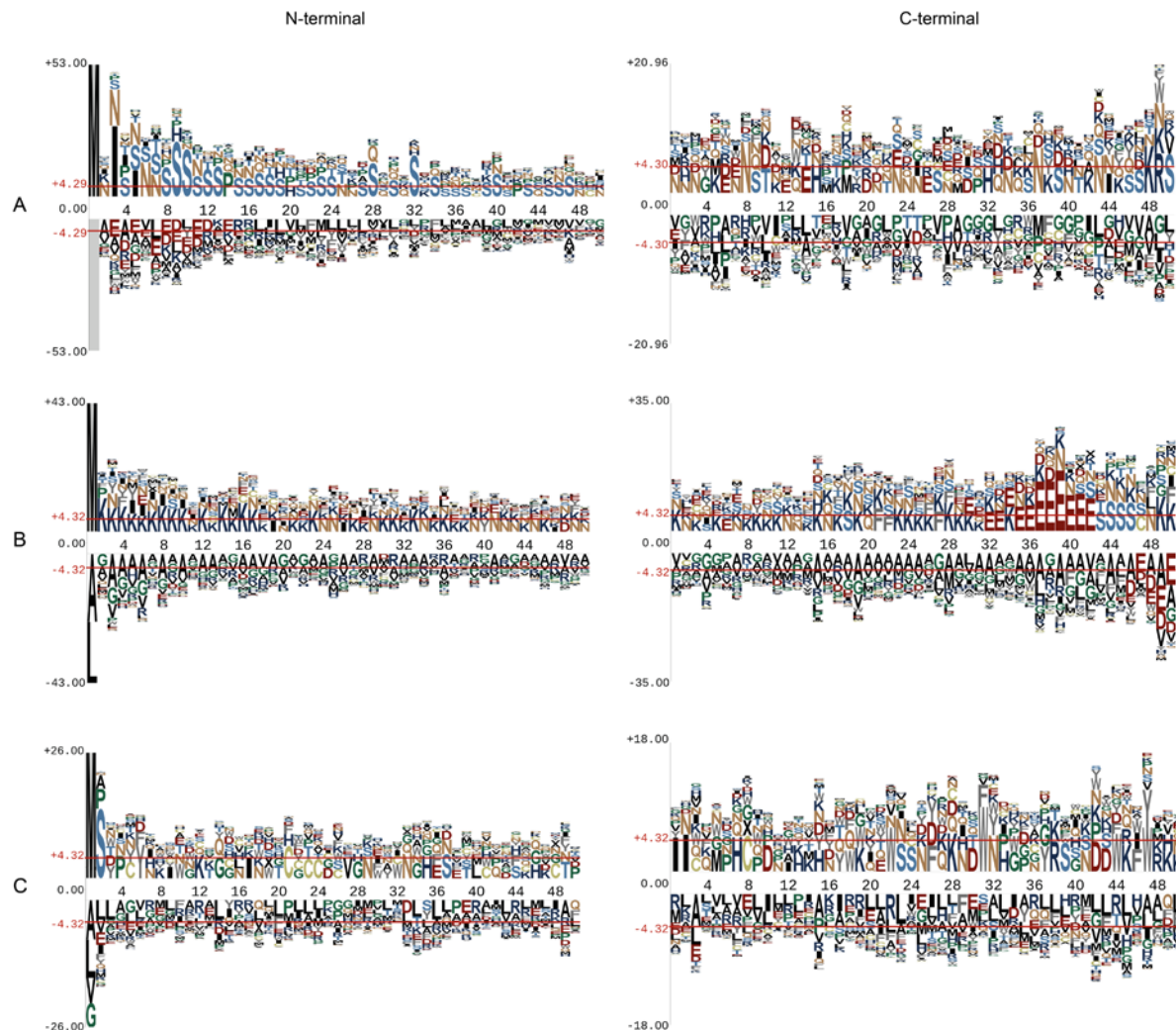
SecretEPDB: a comprehensive web-based resource for secreted effector proteins of the bacterial types III, IV and VI secretion systems

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Supplementary Figure S1. Sequence logo representations showing the amino acid conservation and preferences of N- and C-terminal motifs of (A) T3SEs, (B) T4SEs, and (C) T6SEs. In all panels the x-axis represents residue position: amino acids above the x-axis are favored, while those below the x-axis are disfavored at the corresponding positions. Note that these sequence logos are generated using the whole set of each type of secreted effectors, rather than species-specific subsets of sequence data.