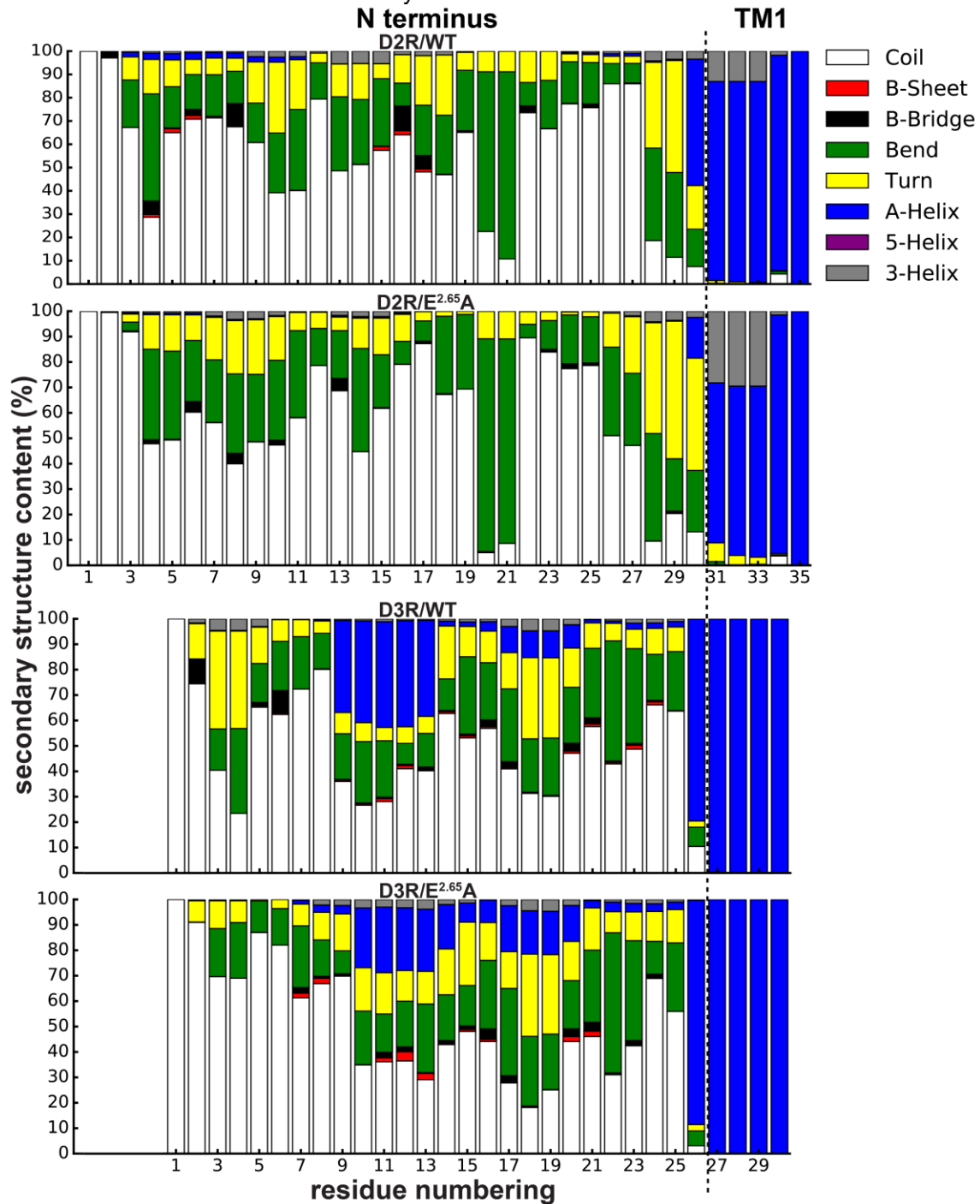


S4 Fig. Distribution of secondary structure content for each of the N terminal residues. The secondary structures were classified by DSSP v2.0.4 [1]. The boundary between NT and TM1 is indicated by the dotted line.



Reference

1. Kabsch W, Sander C. Dictionary of protein secondary structure: pattern recognition of hydrogen-bonded and geometrical features. *Biopolymers*. 1983;22(12):2577-637. doi: 10.1002/bip.360221211. PubMed PMID: 6667333.