

## Supplemental Figure 1

for Resilience of BST-2/Tetherin structure to single amino acid substitutions

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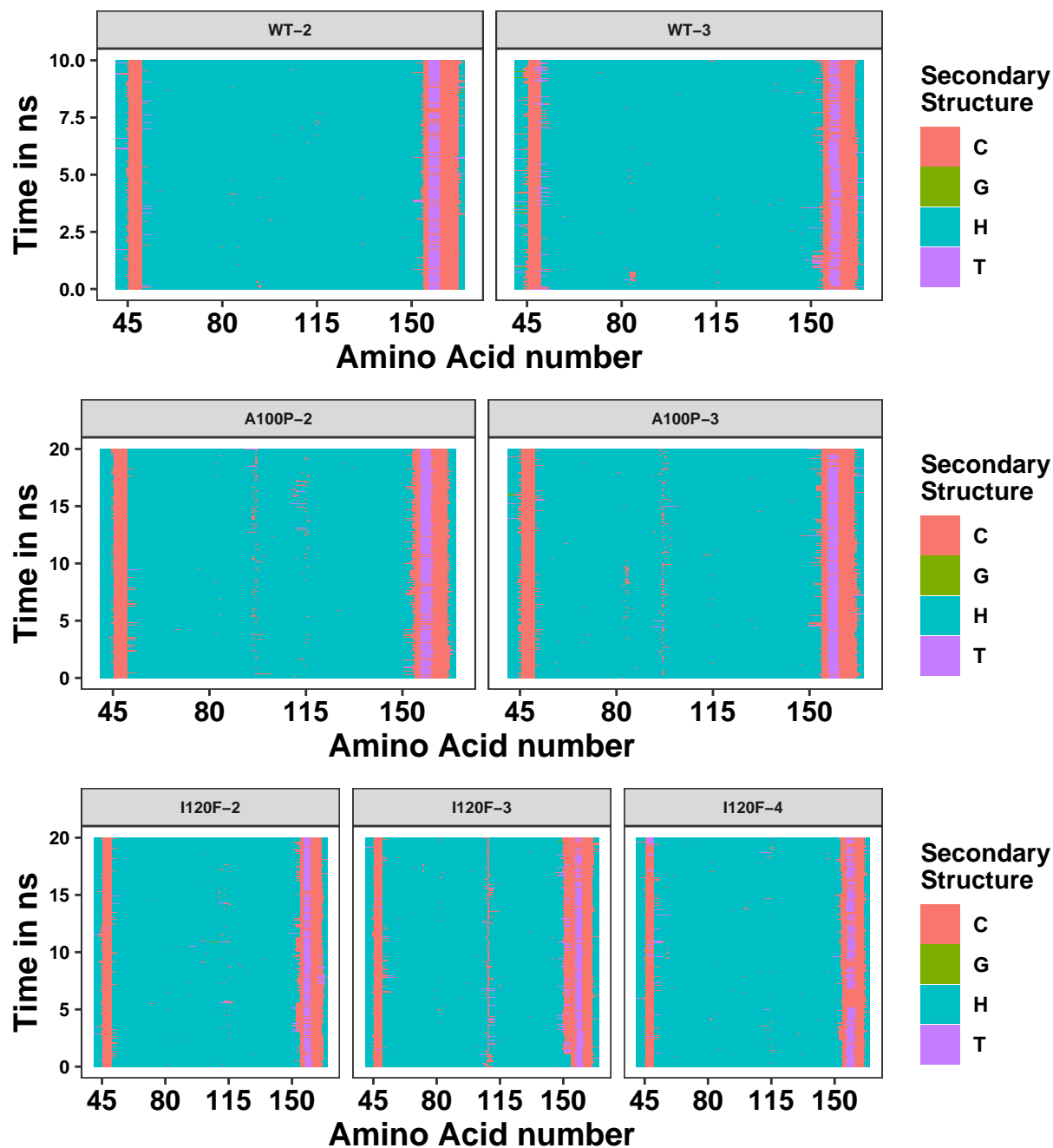


Figure 1: (A) Secondary structure over time plots from replicates of the WT, A100P and I120F simulations. Secondary structures were assigned by YASARA based on hydrogen bonding distances and psi angles at 25 picosecond intervals. ‘H’ stands for alpha helix, ‘T’ for turn, ‘C’ for coil, and ‘G’ for 3/10 helix.