

Electronic supporting information for:
Resistance to tyrosine kinase inhibitors can be mediated by changes to the conformation space of the kinase

1 Supplemental Figures

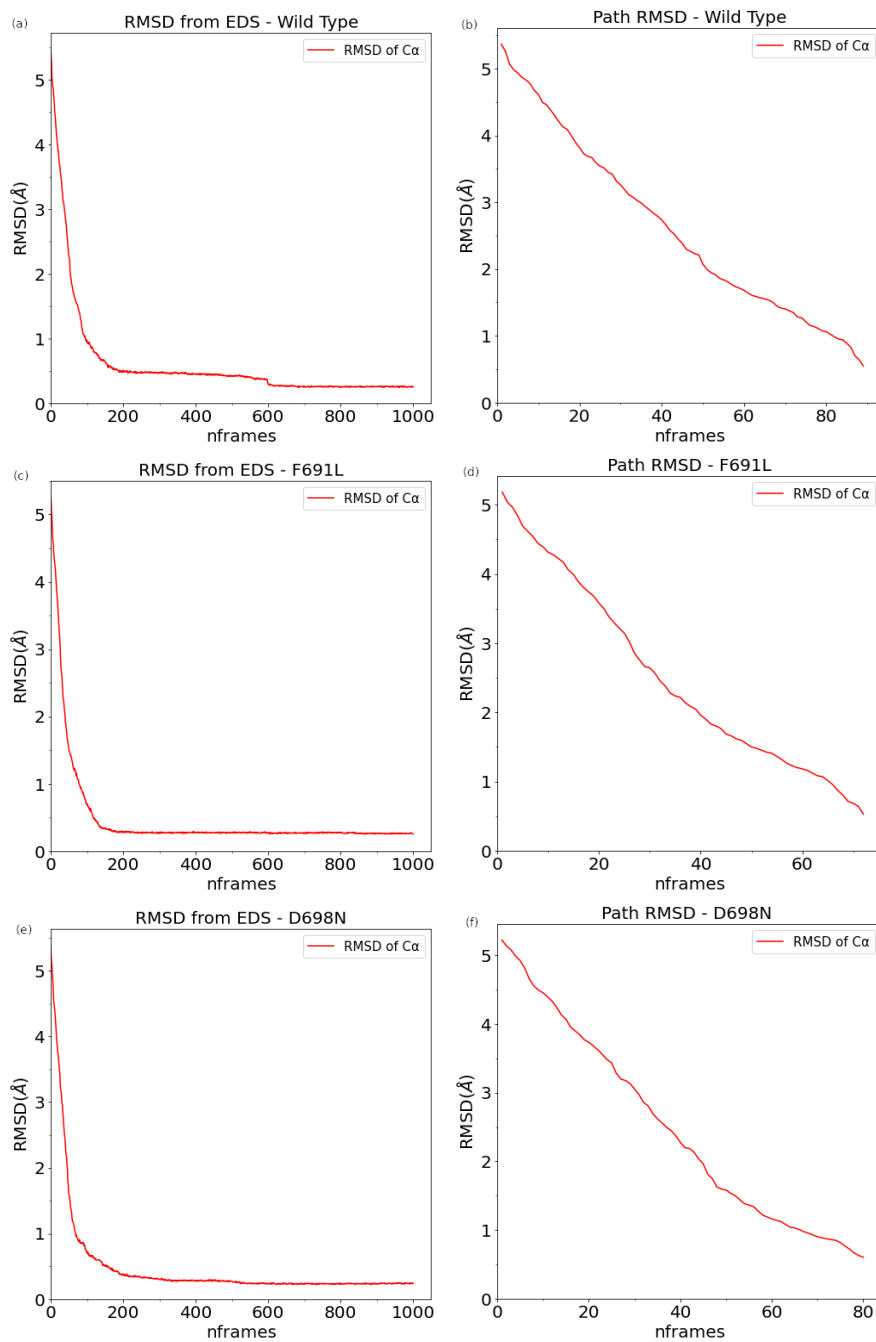


Figure S1: C α Root mean square deviation (RMSD) profiles with respect to the target structures, for trajectories calculated by EDS (a, c and e) and wT-metaD (b, d and f), shown for wt-FLT (a and b), and the mutants F691L (c and d), and D698N (e and f).