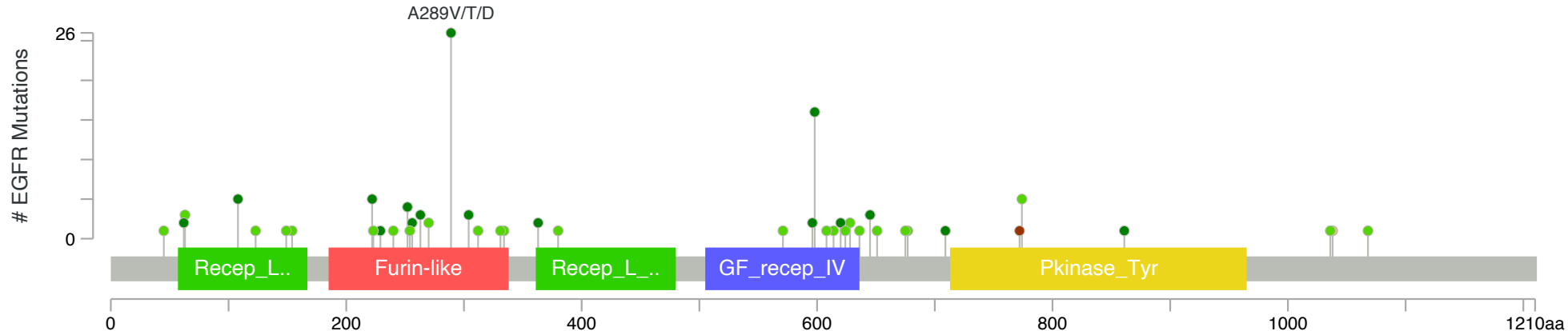


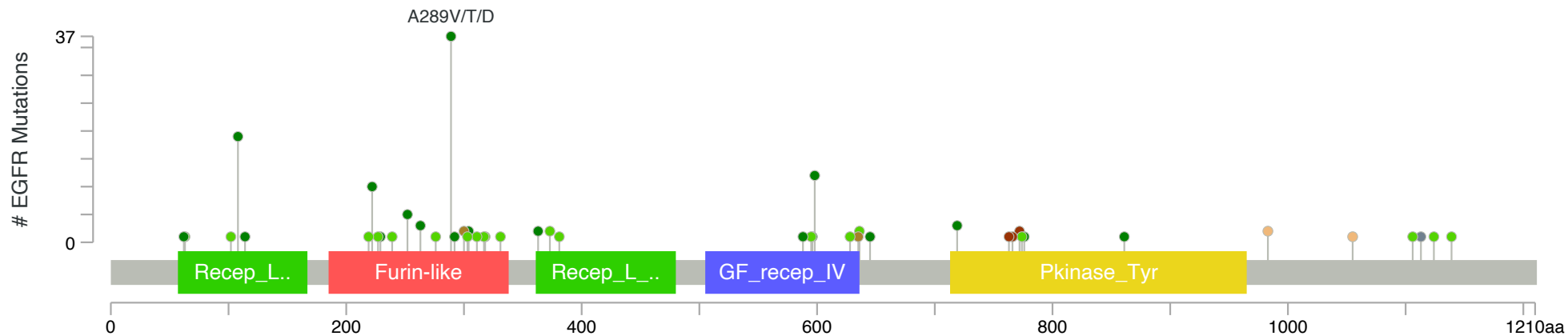
TCGA cohort

EGFR kinase domain: 7/118 mutation sites (5.9%)



MSKCC cohort

EGFR kinase domain: 10/245 mutation sites (4.1%)



Supplementary Figure S1. EGFR mutational distributions summarized from TCGA and MSKCC cohort, respectively.