

Figure S1: Fusions found in more than 1 histology



В

300

0

0



Figure S2: Distribution of spanningDelta for annoFuse prioritized fusions from TCGA and PBTA cohorts

50 100 150 200

change_spanning

0

50 100 150 200



Figure S3: Sensitivity of TCGA fusions retained by annoFuse



Figure S4: Distribution of kinase genes fused in 5' and 3' genes per histologies