

Description of Additional supplementary files

Supplementary Data S1: Selection estimates across copy number states

Supplementary Data S2: Selection estimates in diploid state

Supplementary Data S3: Selection change associated with the deletion of the gene.

Supplementary Data S4: Selection change associated with the gain of the gene.

Supplementary Data S5: Clusters of gene-tumor type pairs

Supplementary Data S6: Cancer types with largest amount of data