

Supplemental Material

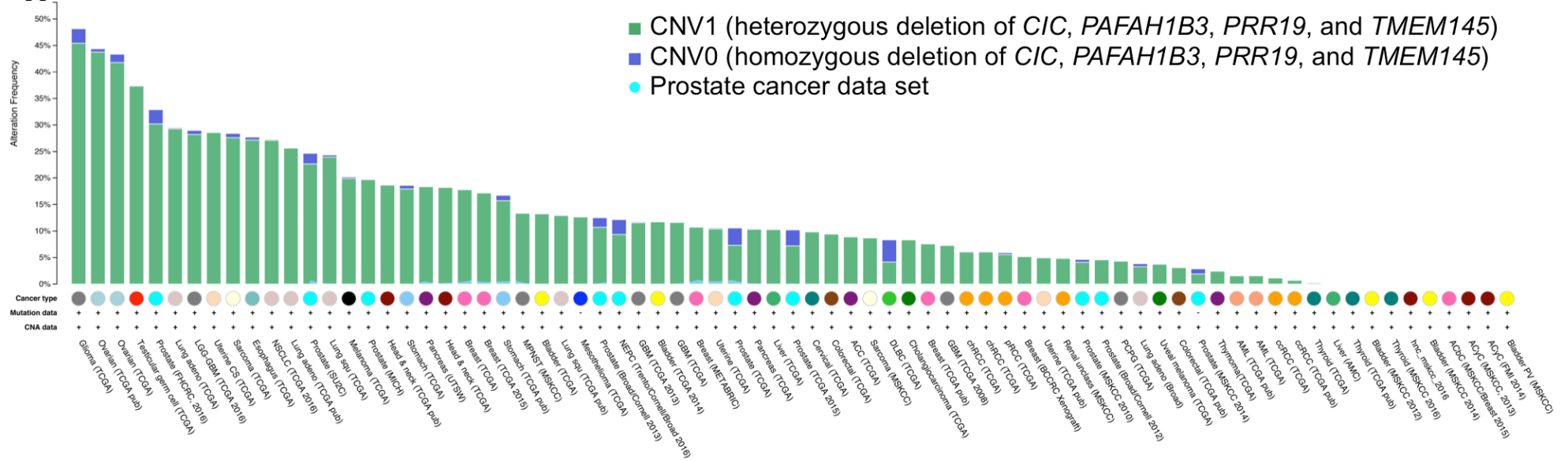


Figure S1. Histogram of homozygous (CNV0) and heterozygous (CNV1) gene deletion events, associated with four neighbouring genes (*CIC*, *PAFAH1B3*, *PRR19*, and *TMEM145*), in diverse cancers. The y-axis show the percentage of tumours that have an alteration in the four genes (CNV0 events shown in blue; CNV1 events in green), and the x-axis denote a cBioPortal data set (prostate cancer data sets are indicated in turquoise).