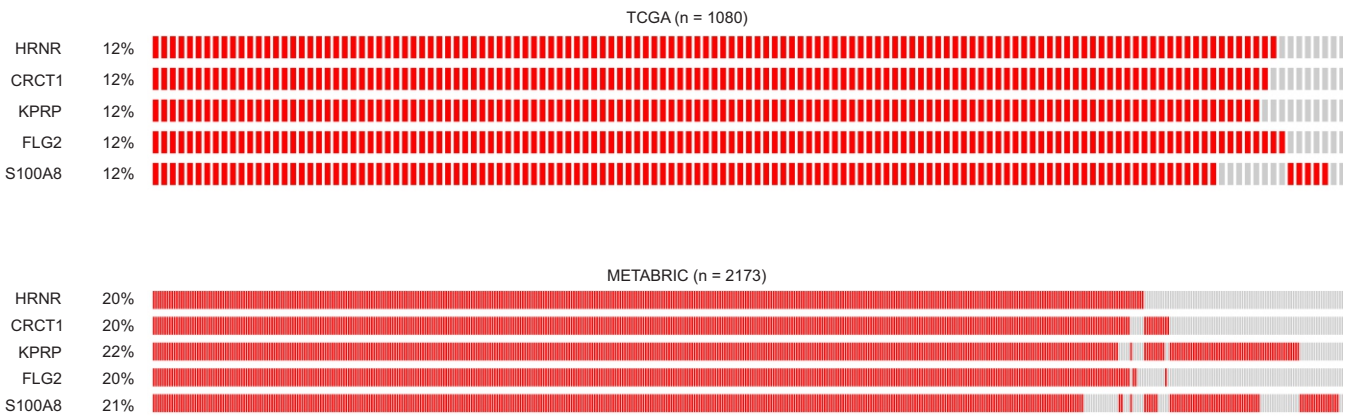


Figure S3



**Figure S3.** Copy number increases of the indicated genes in both TCGA and METABRIC datasets. The analyses were determined using tools provided by cBioPortal. Only the proportions of tumors with the indicated amplifications (red) are included.