Supplementary Figures and Tables

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Figure S1: CNA gene counts from TCGA rare and common cancer cohorts. Gene-level results produced by GISTIC were downloaded from the cBioPortal. Only high-level amplifications (red) and putative homozygous deletions (blue) were included. **A)**. All protein-coding genes. **B)**. Oncogenes. **C)**. Tumor suppressor genes.
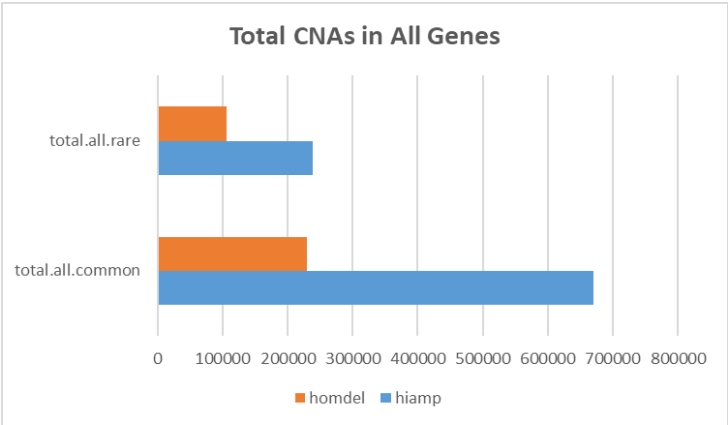
Figure S1A
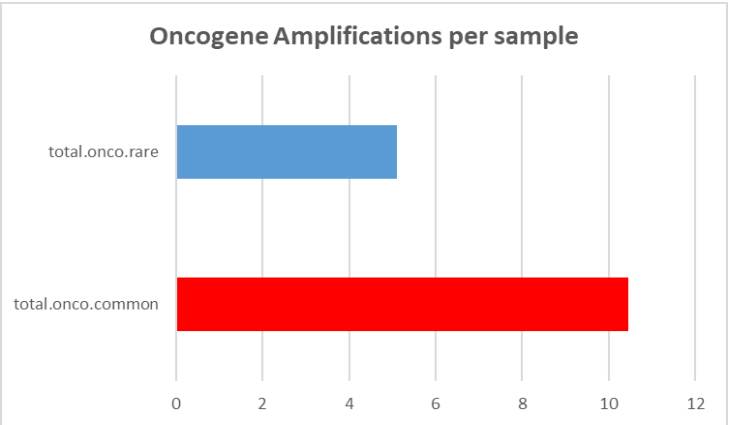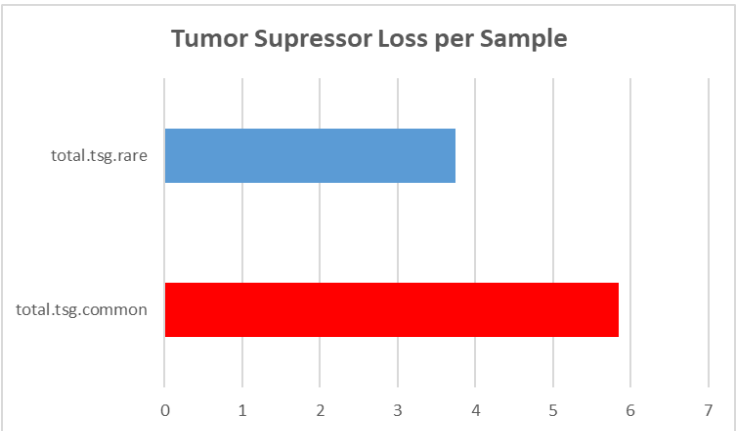

Figure S1B

Figure S1C

Figure S2. CNA gene counts from TCGA rare and common cancer cohorts, broken down by cancer type. **A)**. All protein-coding genes. **B)*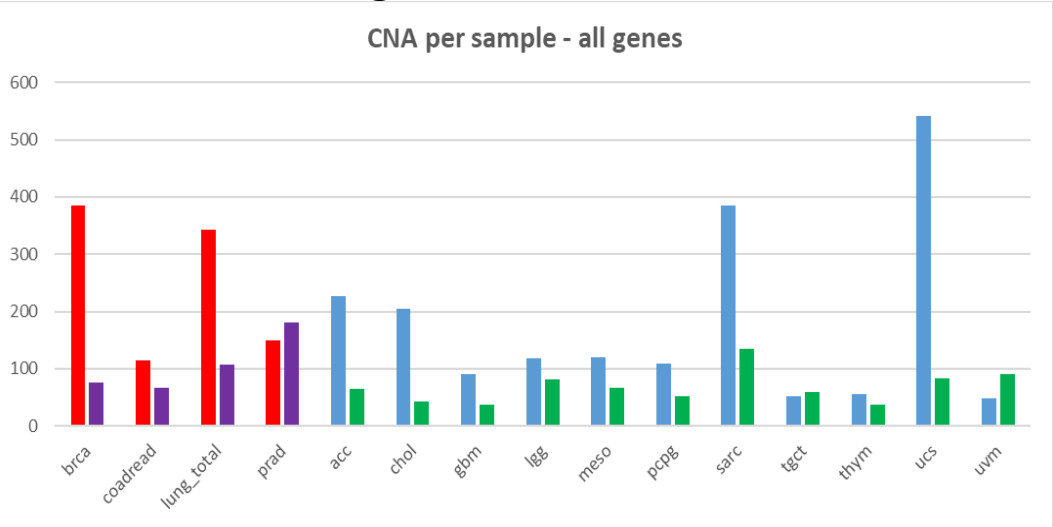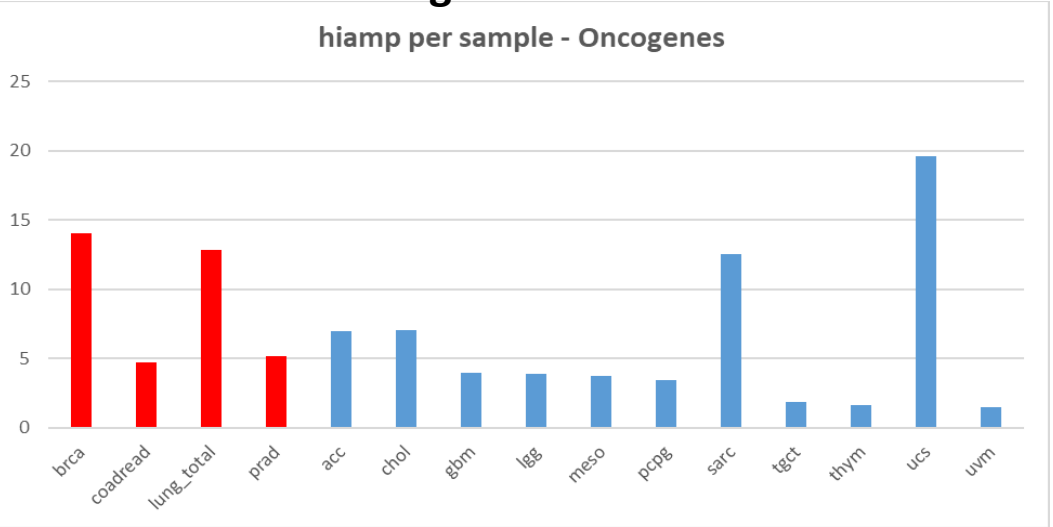*. Oncogenes. **C)**. Tumor suppressor genes.