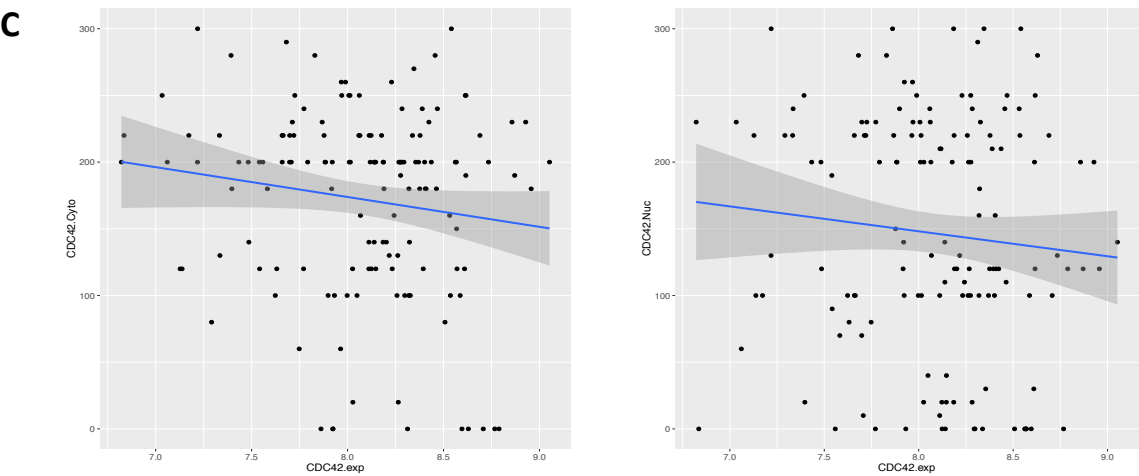
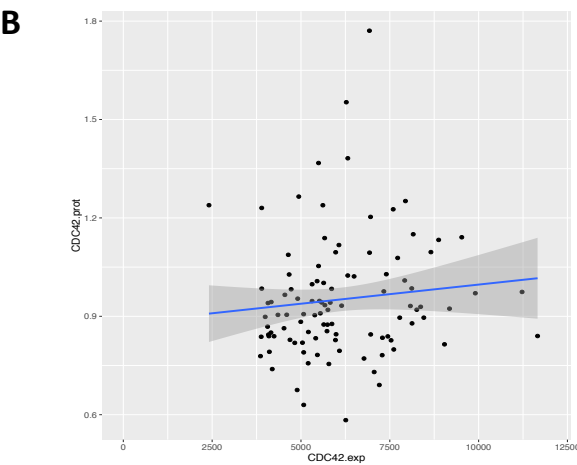
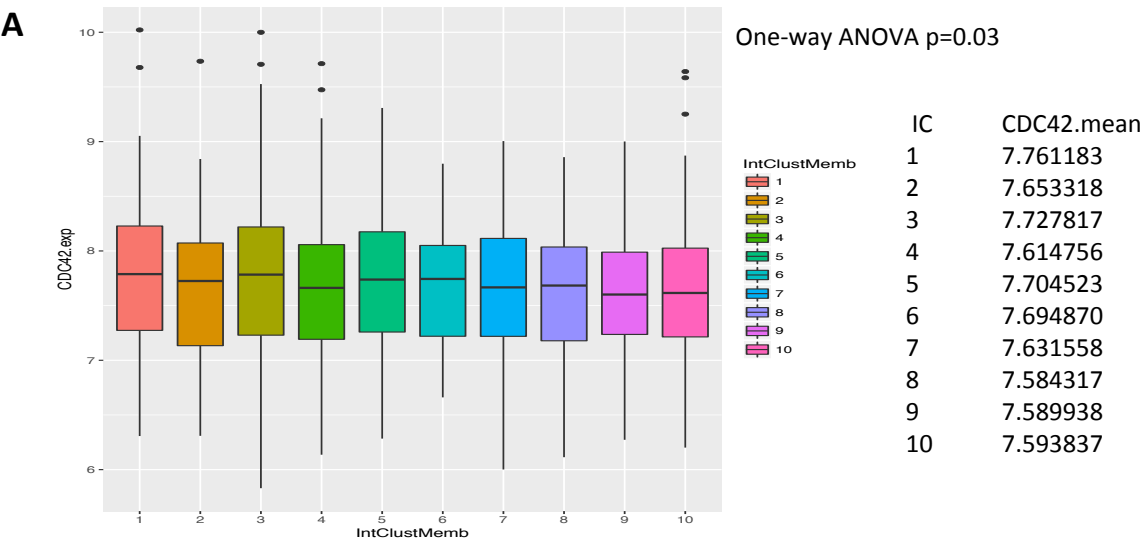


Supplementary Figure 1

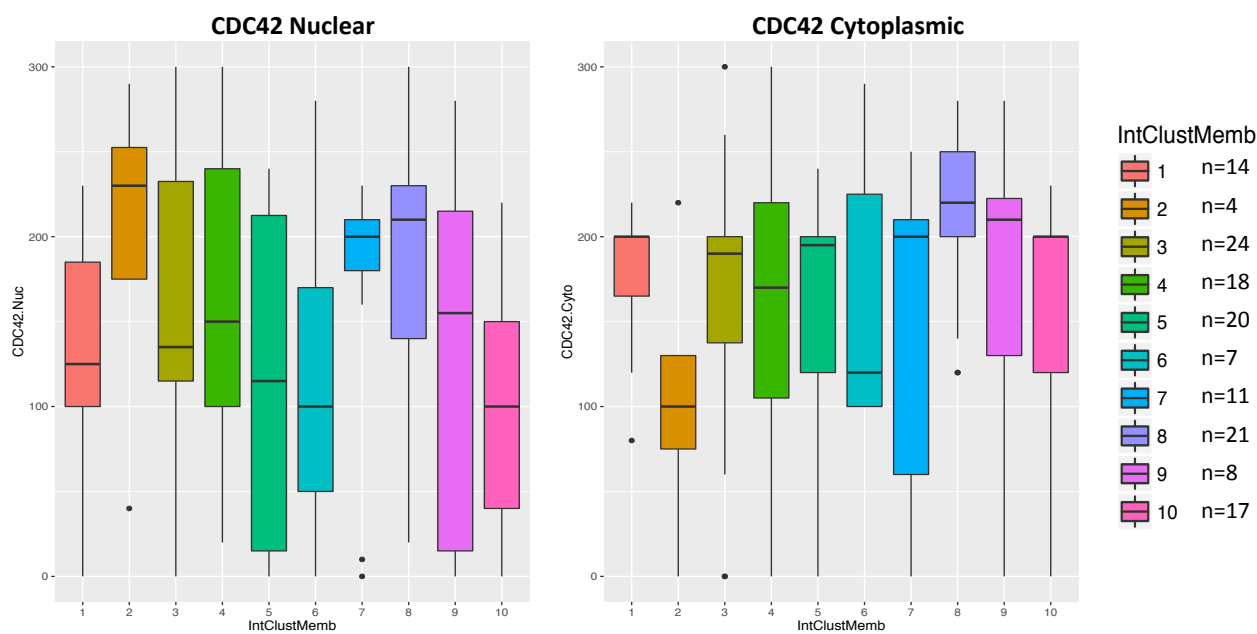


A. mRNA expression of *CDC42* in METABRIC data by integrated cluster membership (IntClustMemb). While the overall expression differed between clusters ($p=0.03$, one-way ANOVA), none of the Tukey post-tests were statistically significant. Also shown is the mean expression value for each of the ICs

B. In TCGA breast cancer data, *CDC42* mRNA expression (CDC42.exp) is not significantly correlated with mass spectrometry protein levels (Spearman $r=0.16$, $p=0.09$).

C. mRNA expression of *CDC42* from METABRIC cases did not correlate with either cytoplasmic (left) or nuclear (right) expression by immunohistochemistry (H-scores)

Supplementary Figure 2



Distribution of CDC42 H-scores by Integrated Cluster membership of cases within METABRIC cohort.