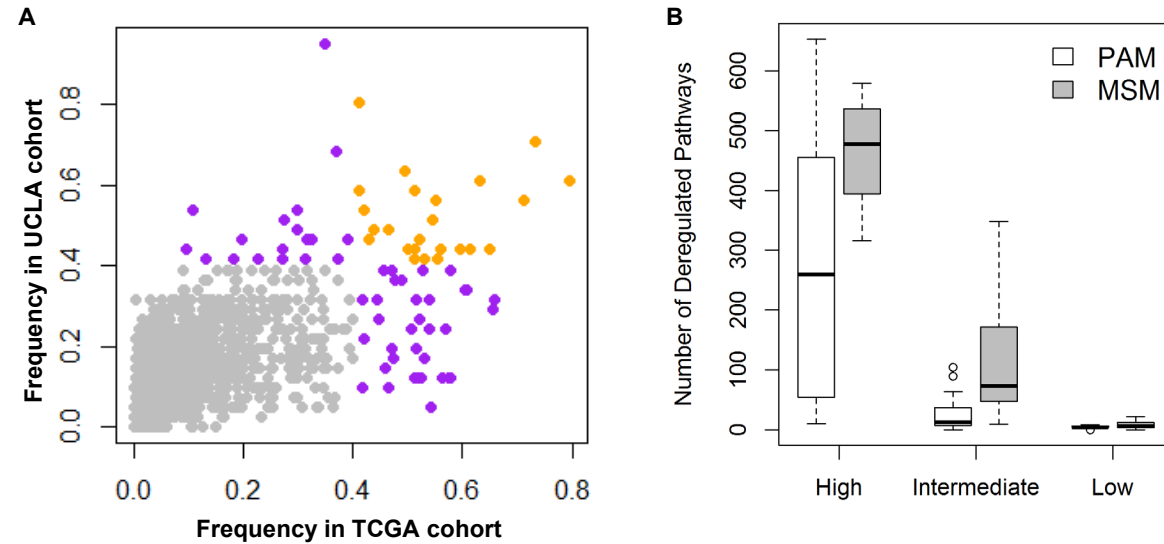


Supplementary Figure S3



Supplementary Figure S3. Pathways, frequently deregulated by the mutated genes in TCGA LUAD and UCLA cohorts.

A) Recurrence rate of 1341 pathways affected by genes mutated in ADC for TCGA and UCLA cohort. Each data point represents a pathway. Pathways, frequently affected in either or both cohorts are indicated in purple or orange color, respectively. **B)** Distributions of the number of pathways deregulated by PAM- (white bars) and MSM- (grey bars) bearing genes in three patient groups identified in **Figure 2D**.