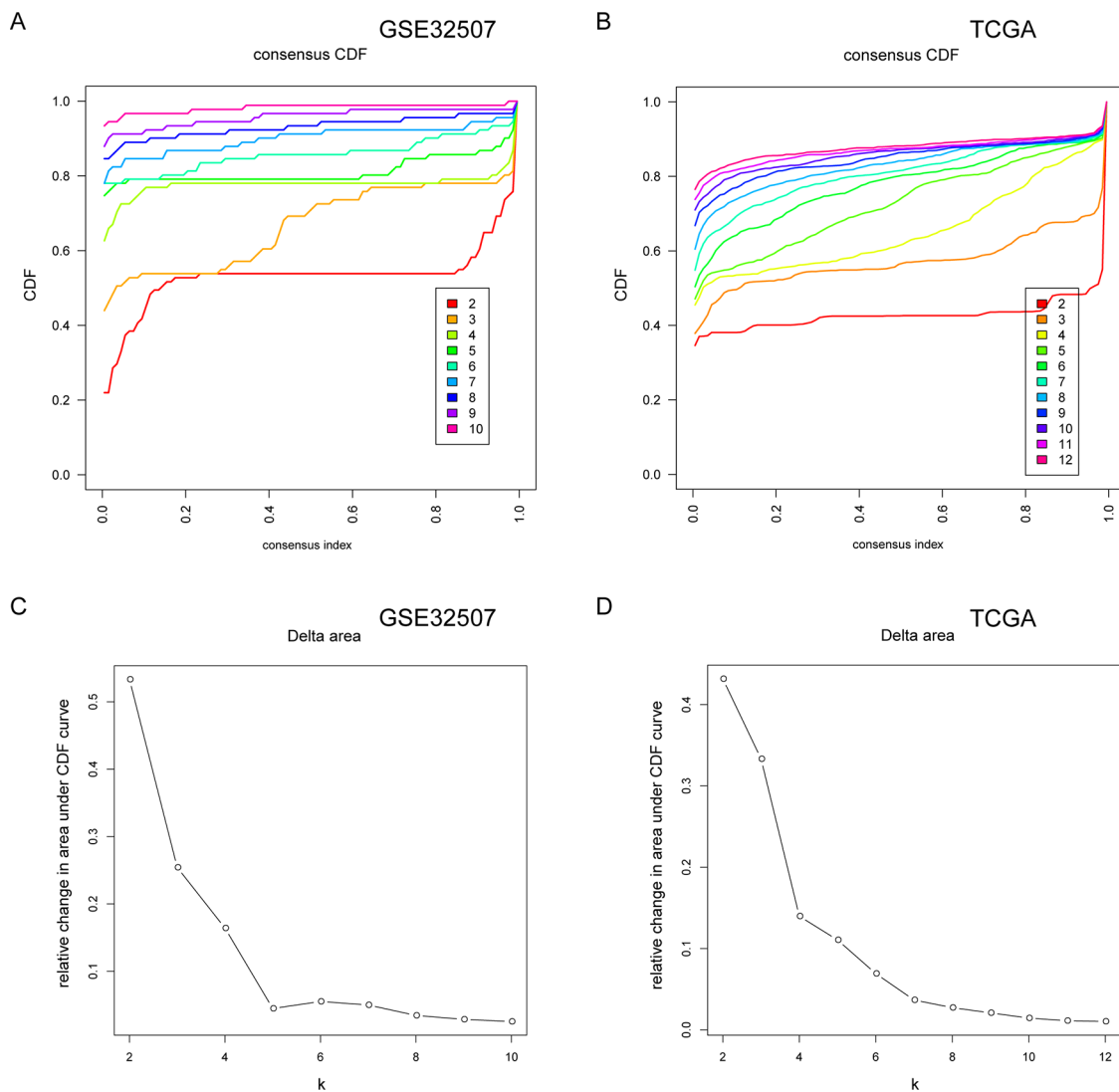
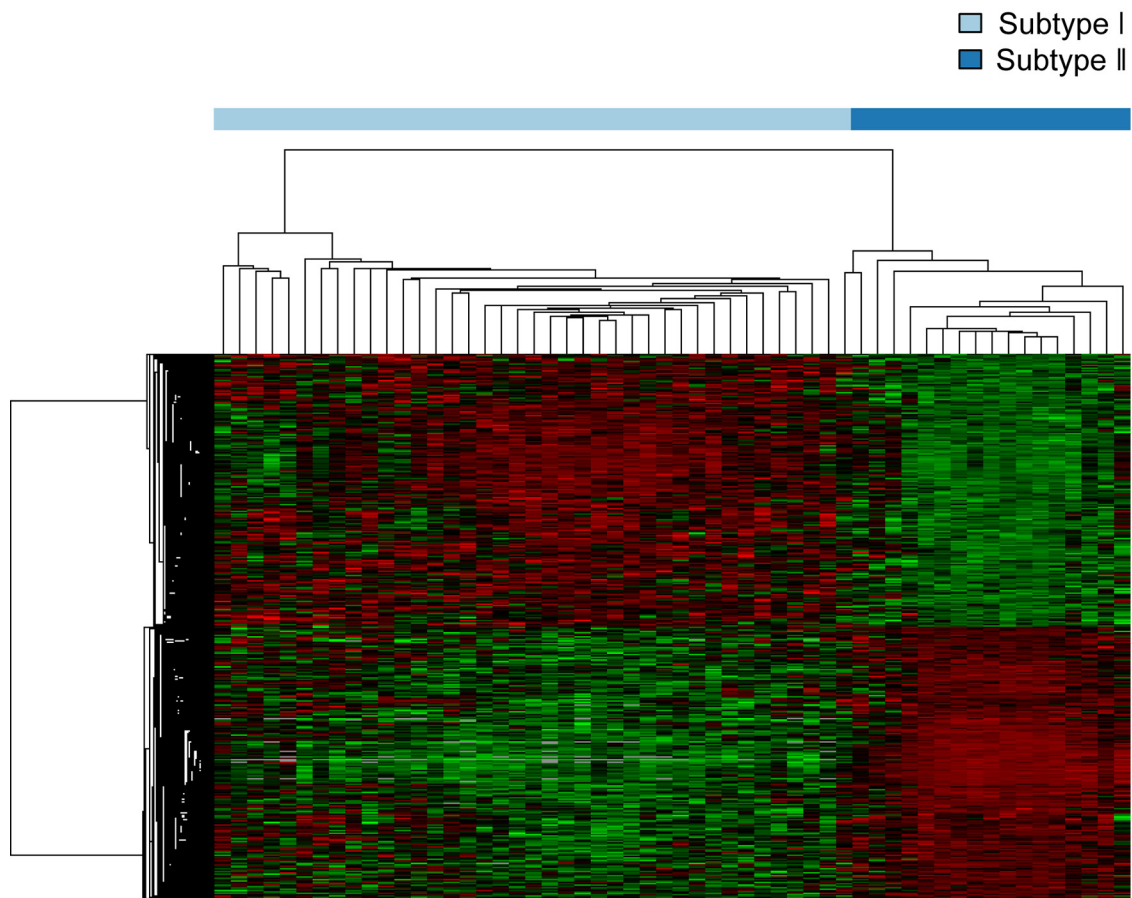


Identification of distinct molecular subtypes of uterine carcinosarcoma

SUPPLEMENTARY FIGURES AND TABLES



Supplementary Figure 1: Delineation of two molecular subtypes of UCS. A. and B. Empirical cumulative distribution plots for GSE32507 and TCGA dataset, respectively. C. and D. The increased area under the CDF curve along with increased number of molecular subtypes.



Supplementary Figure 2: Heatmap of top 500 genes over-expressed in distinct molecular subtypes. After SAM-seq, we picked top 500 genes over-expressed in each subtype to form TOP500. Hierarchical clustering of TOP500 was performed by Cluster 3.0 using centroid linkage method. Red, over-expressed genes; Green, down-expressed genes.

Supplementary Table 1: Clinicopathologic Characteristics (N = 57)

See Supplementary File 1

Supplementary Table 2: Pathways enriched in each molecular subtype

See Supplementary File 1