

# Uterine carcinosarcoma (UCS)

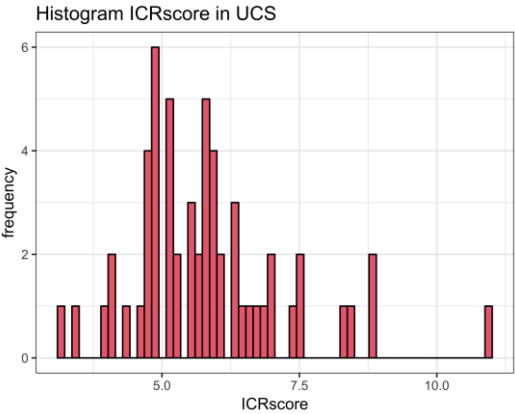
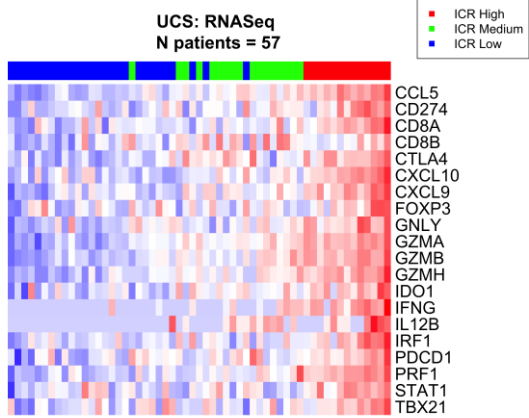
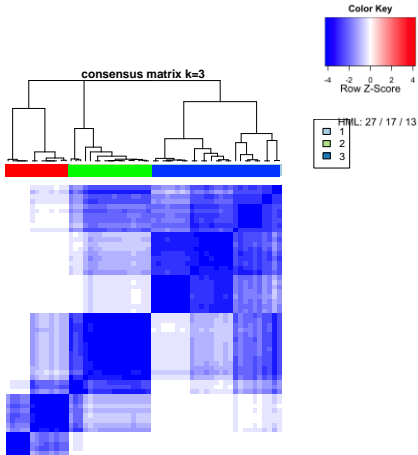
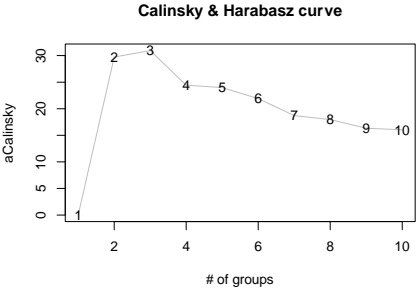


Figure: Gene expression data was obtained from TCGA, pancancer normalized (EDAseq) and log transformed. Samples are ordered by ICR score. Optimal k for clustering was 3. The combination action performed to reduce to 3 clusters is: none Resulting in mean ICR scores of 4.82 / 5.91 / 7.84 for Low / Medium / High resp.

## Deconvolution immune cell populations

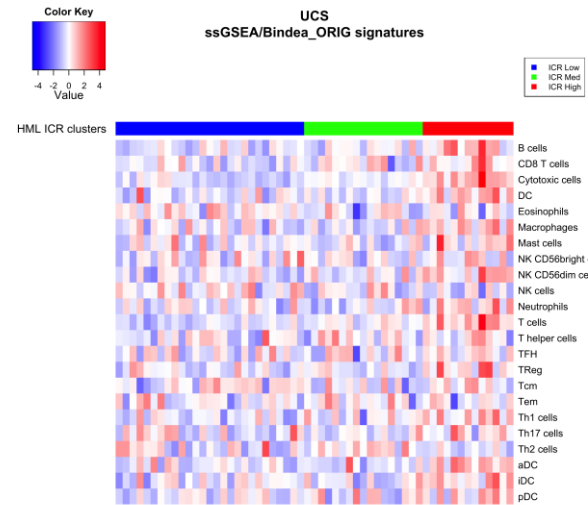
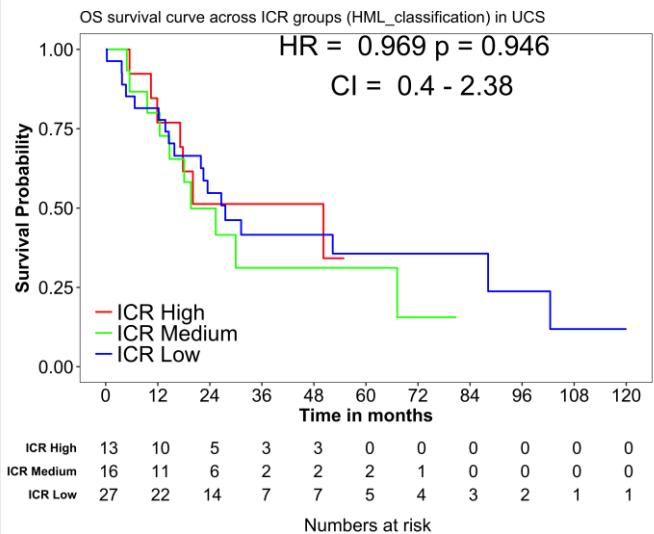


Figure: EDAseq normalized, log transformed gene expression data was obtained from TCGA, using Assembler\_Pance\_Normalized\_filtered. Bindea\_ORIG enrichment z-scores were used to generate this heatmap.

## Survival analysis



## ssGSEA oncogenic pathways

