

# Ovarian serous cystadenocarcinoma (OV)

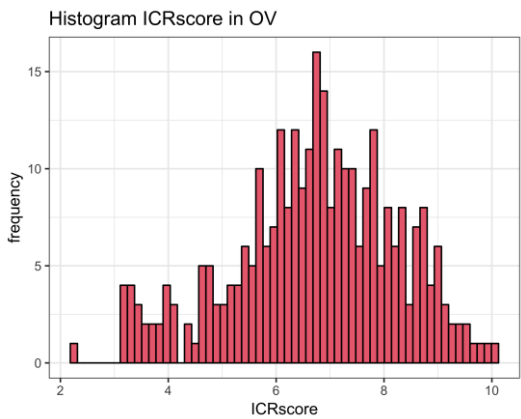
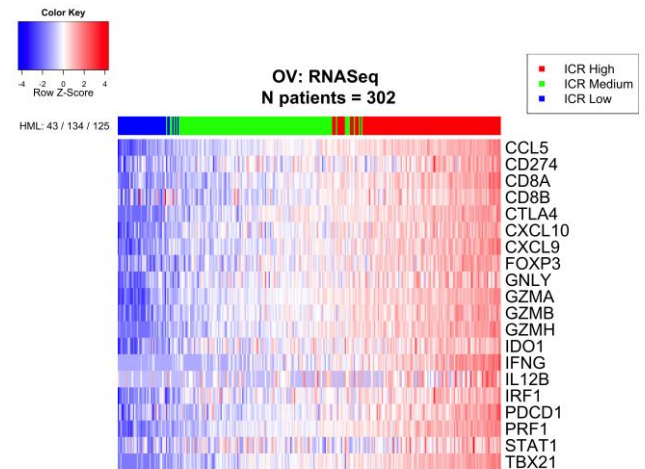
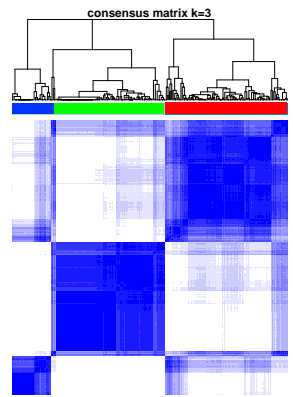
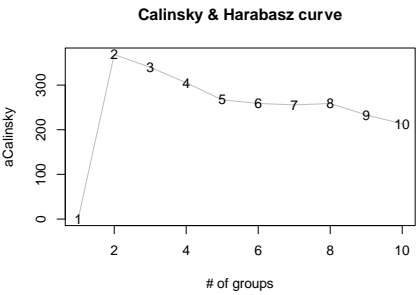


Figure: Gene expression data was obtained from TCGA, pancancer normalized (EDAseq) and log transformed. Samples are ordered by ICR score. Optimal k for clustering was 3. The combination action performed to reduce to 3 clusters is: none Resulting in mean ICR scores of 4.06 / 6.26 / 8.11 for Low / Medium / High resp.

## Deconvolution immune cell populations

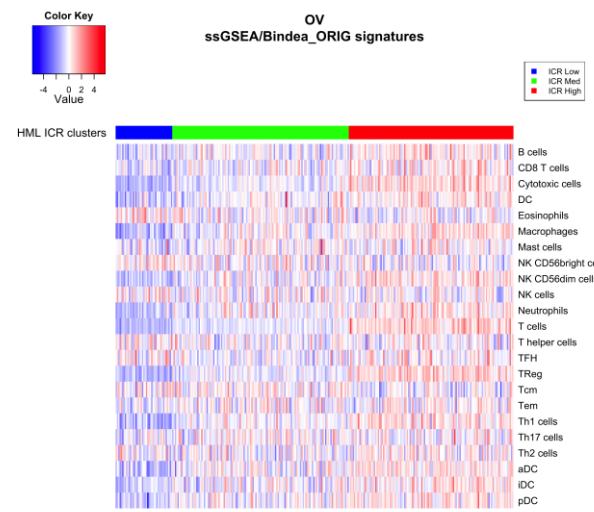
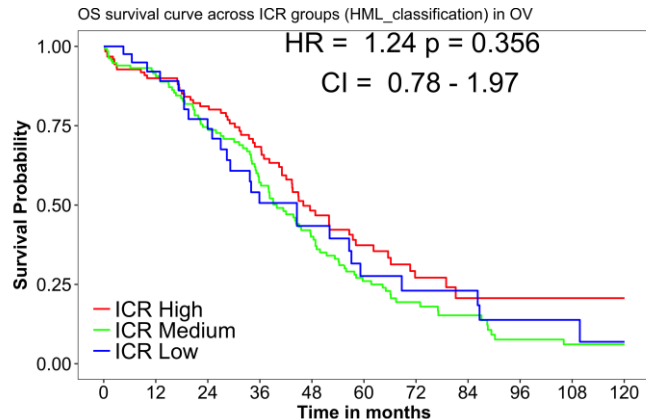


Figure: EDAseq normalized, log transformed gene expression data was obtained from TCGA, using Assembler\_Panca\_Normalized\_filtered. Bindea\_ORIG enrichment z-scores were used to generate this heatmap.

## Survival analysis



ICR High	124	96	79	54	32	21	12	6	4	3	2
ICR Medium	134	106	81	58	40	26	14	10	5	4	2
ICR Low	43	31	24	15	12	7	5	5	2	2	1

Numbers at risk

## ssGSEA oncogenic pathways

