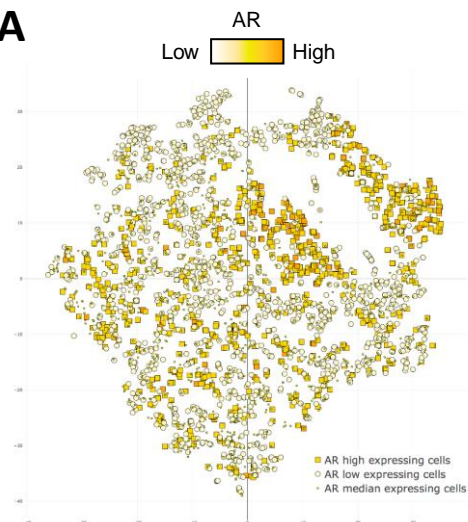


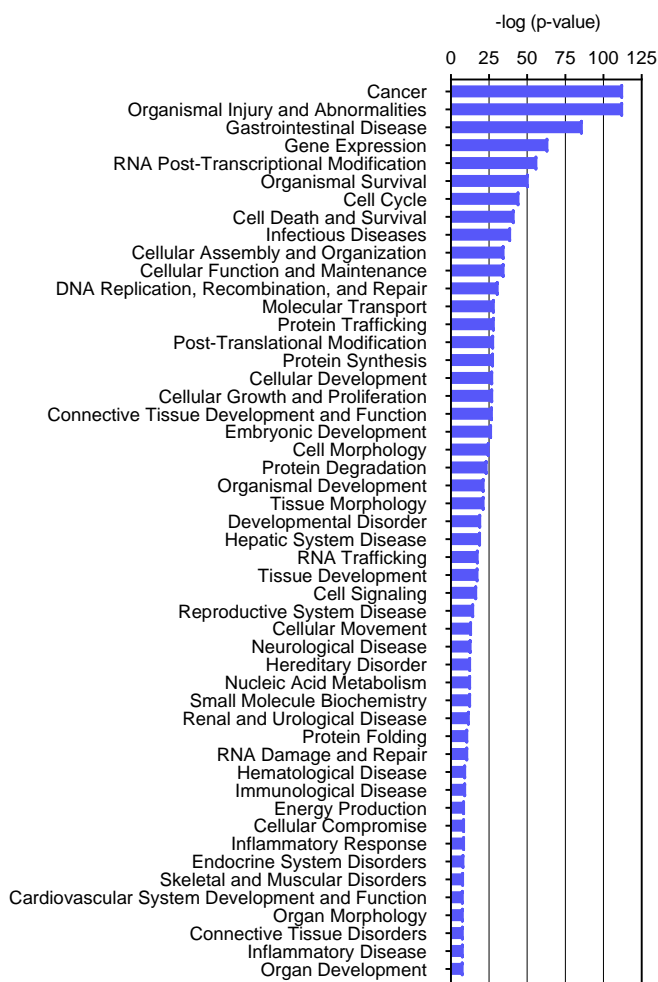
Supplemental Figure 3

A

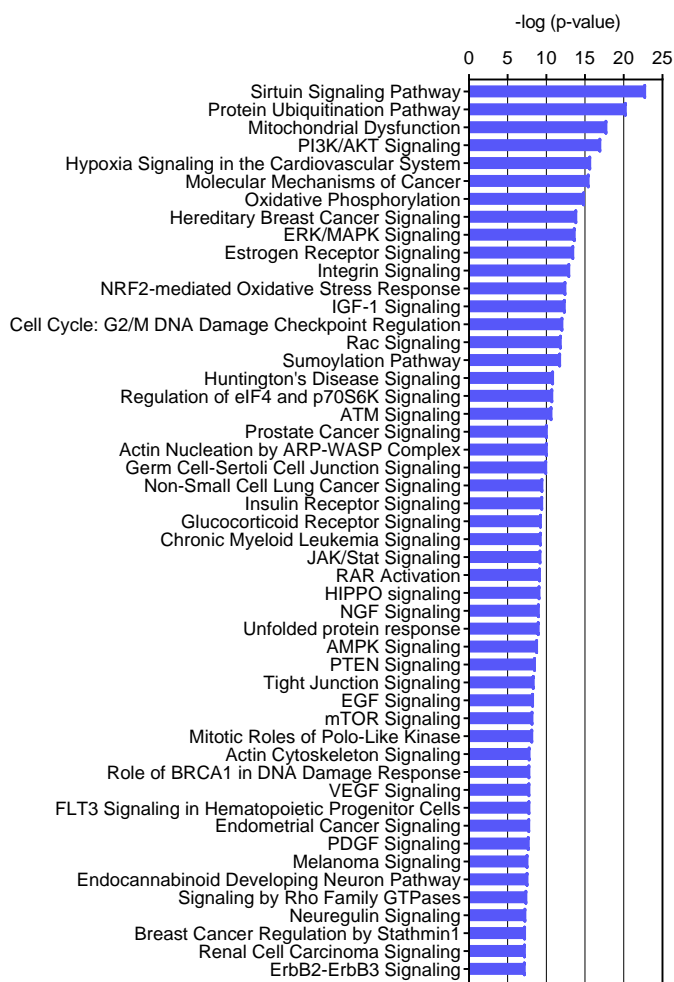


B

Diseases and Function Pathways



Canonical Pathways



Supplemental Figure 3. Single-cell RNAseq of AR expressing MDA-MB-453 TNBC cells. scRNAseq was performed on untreated MDA-MB-453 cells cultured in full serum media. Cells were divided in groups based on their expression of AR, AR^{Low} and AR^{High}. **(A)** tSNE plot of AR levels in cells clustered by gene expression. **(B)** IPA was performed on the genes differentially expressed between AR^{Low} and AR^{High} groups. Shown are the top 50 most significantly altered disease and function pathways and canonical pathways.