Enrichment of bottom 30% genes by correlation with purity (48 genes) HALLMARK EPITHELIAL MESENCHYMAL TRANSITION HALLMARK XENOBIOTIC METABOLISM HALLMARK INFLAMMATORY RESPONSE-HALLMARK MYOGENESIS-Biological processes Number of genes found HALLMARK ESTROGEN RESPONSE LATE-HALLMARK COMPLEMENT-HALLMARK UV RESPONSE DN HALLMARK ESTROGEN RESPONSE EARLY HALLMARK FATTY ACID METABOLISM HALLMARK NOTCH SIGNALING-0.5 1.5 -log10(B-H adjusted p-value)