

Enrichment of middle 30-70% genes by correlation with purity (62 genes)

Biological processes

HALLMARK EPITHELIAL MESENCHYMAL TRANSITION

HALLMARK HEDGEHOG SIGNALING

HALLMARK ANGIOGENESIS

HALLMARK ESTROGEN RESPONSE EARLY

HALLMARK IL2 STAT5 SIGNALING

HALLMARK KRAS SIGNALING DN

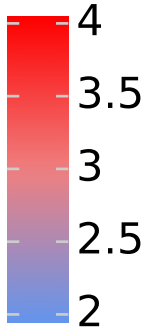
HALLMARK MTORC1 SIGNALING

HALLMARK COMPLEMENT

HALLMARK GLYCOLYSIS

HALLMARK HYPOXIA

Number of genes found



0 0.3 0.6 0.9 1.2

$-\log_{10}(\text{B-H adjusted p-value})$