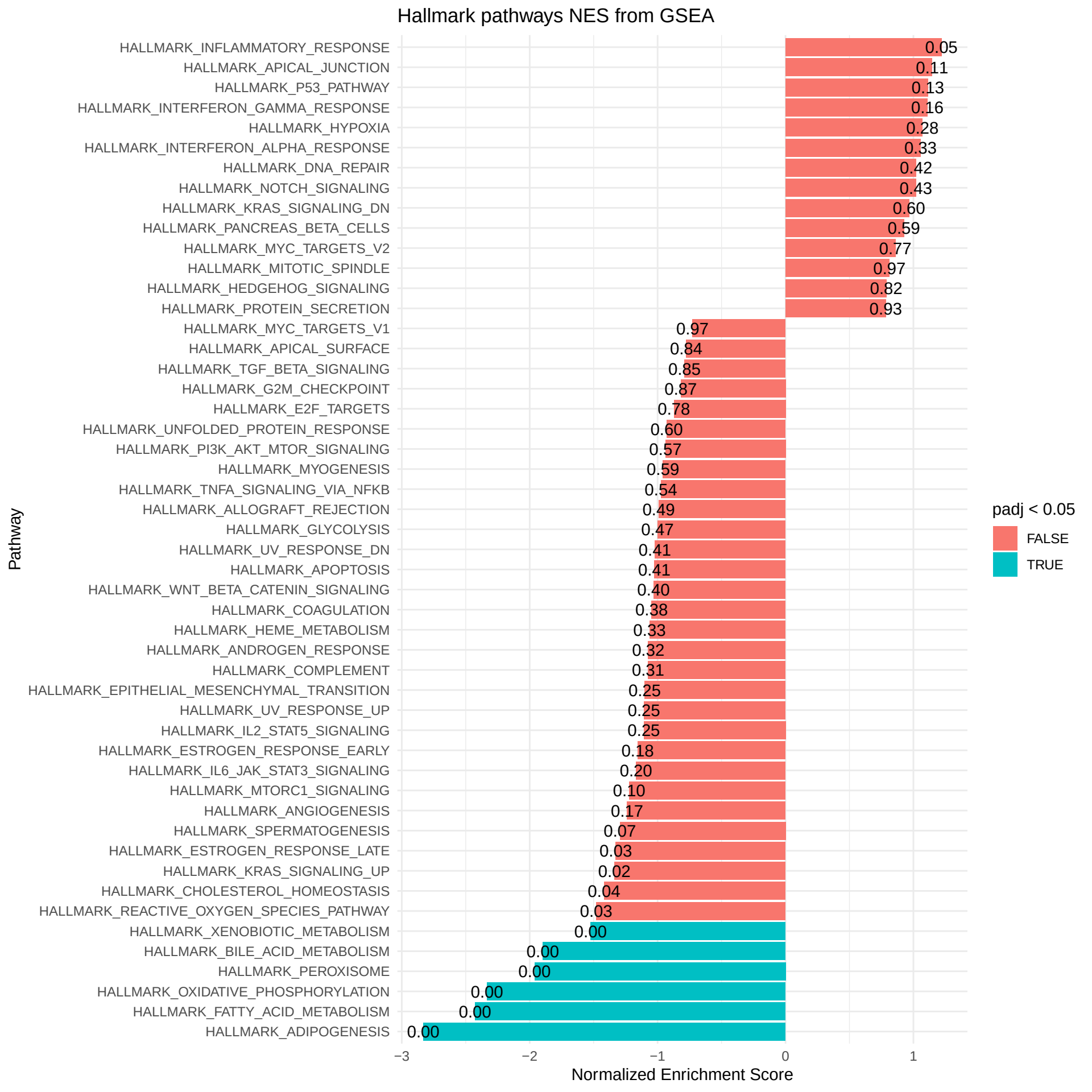
Hallmark pathways NES from GSEA
HALLMARK_INFLAMMATORY_RESPONSE 0.05
HALLMARK_APICAL_JUNCTION 0.11
HALLMARK_P53_PATHWAY 0.13
HALLMARK_INTERFERON_GAMMA_RESPONSE 0.16
HALLMARK_HYPOXIA 0.28
HALLMARK_INTERFERON_ALPHA_RESPONSE 0.33
HALLMARK_DNA_REPAIR 0.42
HALLMARK_NOTCH_SIGNALING 0.43
HALLMARK_KRAS_SIGNALING_DN 0.60
HALLMARK_PANCREAS_BETA_CELLS 0.59
HALLMARK_MYC_TARGETS_V2 0.77
HALLMARK_MITOTIC_SPINDLE 0.97
HALLMARK_HEDGEHOG_SIGNALING 0.82
HALLMARK_PROTEIN_SECRETION 0.93
HALLMARK_MYC_TARGETS_V1 0.97
HALLMARK_APICAL_SURFACE 0.84
HALLMARK_TGF_BETA_SIGNALING 0.85
HALLMARK_G2M_CHECKPOINT 0.87
HALLMARK_E2F_TARGETS 0.78
HALLMARK_UNFOLDED_PROTEIN_RESPONSE 0.60
HALLMARK_PI3K_AKT_MTOR_SIGNALING 0.57
HALLMARK_MYOGENESIS 0.59
HALLMARK_TNFA_SIGNALING_VIA_NFKB 0.54
HALLMARK_ALLOGRAFT_REJECTION 0.49
HALLMARK_GLYCOLYSIS 0.47
HALLMARK_UV_RESPONSE_DN 0.41
HALLMARK_APOPTOSIS 0.41
HALLMARK_WNT_BETA_CATENIN_SIGNALING 0.40
HALLMARK_COAGULATION 0.38
HALLMARK_HEME_METABOLISM 0.33
HALLMARK_ANDROGEN_RESPONSE 0.32
HALLMARK_COMPLEMENT 0.31
HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION 0.25
HALLMARK_UV_RESPONSE_UP 0.25
HALLMARK_IL2_STAT5_SIGNALING 0.25
HALLMARK_ESTROGEN_RESPONSE_EARLY 0.18
HALLMARK_IL6_JAK_STAT3_SIGNALING 0.20
HALLMARK_MTORC1_SIGNALING 0.10
HALLMARK_ANGIOGENESIS 0.17
HALLMARK_SPERMATOGENESIS 0.07
HALLMARK_ESTROGEN_RESPONSE_LATE 0.03
HALLMARK_KRAS_SIGNALING_UP 0.02
HALLMARK_CHOLESTEROL_HOMEOSTASIS 0.04
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY 0.03
HALLMARK_XENOBIOTIC_METABOLISM 0.00
HALLMARK_BILE_ACID_METABOLISM 0.00
HALLMARK_PEROXISOME 0.00
HALLMARK_OXIDATIVE_PHOSPHORYLATION 0.00
HALLMARK_FATTY_ACID_METABOLISM 0.00
HALLMARK_ADIPOGENESIS 0.00
Pathway
−3
−2
−1
0
1
Normalized Enrichment Score
padj < 0.05
FALSE
TRUE