

GSEA result of RARRES1

Epithelial_cells

+	HALLMARK_ADIPOGENESIS
+	HALLMARK_ALLOGRAFT_REJECTION
+	HALLMARK_ANDROGEN_RESPONSE
+	HALLMARK_ANGIOGENESIS
+	HALLMARK_APICAL_JUNCTION
+	HALLMARK_APICAL_SURFACE
+	HALLMARK_APOPTOSIS
+	HALLMARK_BILE_ACID_METABOLISM
+	HALLMARK_CHOLESTEROL_HOMEOSTASIS
+	HALLMARK_COAGULATION
+	HALLMARK_COMPLEMENT
+	HALLMARK_DNA_REPAIR
+	HALLMARK_E2F_TARGETS
+	HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION
+	HALLMARK_ESTROGEN_RESPONSE_EARLY
+	HALLMARK_ESTROGEN_RESPONSE_LATE
+	HALLMARK_FATTY_ACID_METABOLISM
+	HALLMARK_G2M_CHECKPOINT
+	HALLMARK_GLYCOLYSIS
+	HALLMARK_HEDGEHOG_SIGNALING
+	HALLMARK_HEME_METABOLISM
+	HALLMARK_HYPOXIA
+	HALLMARK_IL2_STAT5_SIGNALING
+	HALLMARK_IL6_JAK_STAT3_SIGNALING
+	HALLMARK_INFLAMMATORY_RESPONSE
+	HALLMARK_INTERFERON_ALPHA_RESPONSE
+	HALLMARK_INTERFERON_GAMMA_RESPONSE
+	HALLMARK_KRAS_SIGNALING_DN
+	HALLMARK_KRAS_SIGNALING_UP
+	HALLMARK_MITOTIC_SPINDLE
+	HALLMARK_MTORC1_SIGNALING
+	HALLMARK_MYC_TARGETS_V1
+	HALLMARK_MYC_TARGETS_V2
+	HALLMARK_MYOGENESIS
+	HALLMARK_NOTCH_SIGNALING
+	HALLMARK_OXIDATIVE_PHOSPHORYLATION
+	HALLMARK_P53_PATHWAY
+	HALLMARK_PANCREAS_BETA_CELLS
+	HALLMARK_PEROXISOME
+	HALLMARK_PI3K_AKT_MTOR_SIGNALING
+	HALLMARK_PROTEIN_SECRETION
+	HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
+	HALLMARK_SPERMATOGENESIS
+	HALLMARK_TGF_BETA_SIGNALING
+	HALLMARK_TNFA_SIGNALING_VIA_NFKB
+	HALLMARK_UNFOLDED_PROTEIN_RESPONSE
+	HALLMARK_UV_RESPONSE_DN
+	HALLMARK_UV_RESPONSE_UP
+	HALLMARK_WNT_BETA_CATENIN_SIGNALING
+	HALLMARK_XENOBIOTIC_METABOLISM