


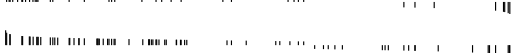




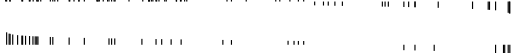








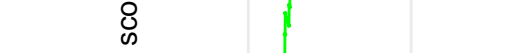


Pathway	Gene ranks	NES	pval	padj
HALLMARK_IL2_STAT5_SIGNALING		-1.29	1.9e-01	5.9e-01
WP_CYTOPLASMIC_RIBOSOMAL_PROTEINS		1.21	2.3e-01	5.9e-01
HALLMARK_INFLAMMATORY_RESPONSE		1.15	3.2e-01	5.9e-01
WP_PI3KAKT_SIGNALING_PATHWAY		-1.15	3.4e-01	5.9e-01
HALLMARK_COAGULATION		1.00	5.2e-01	5.9e-01
HALLMARK_COMPLEMENT		1.00	5.2e-01	5.9e-01
WP_VEGFAVEGFR2_SIGNALING_PATHWAY		0.95	5.2e-01	5.9e-01
HALLMARK_MYC_TARGETS_V1		-0.97	5.3e-01	5.9e-01
TRIAL_RELAXATION_AND_CONTRACTION_PATHWAYS		-0.74	7.7e-01	7.7e-01
TRIAL_RELAXATION_AND_CONTRACTION_PATHWAYS		-0.74	7.7e-01	7.7e-01
HALLMARK_MYC_TARGETS_V1		-0.97	5.3e-01	5.9e-01
WP_VEGFAVEGFR2_SIGNALING_PATHWAY		0.95	5.2e-01	5.9e-01
HALLMARK_COMPLEMENT		1.00	5.2e-01	5.9e-01
HALLMARK_COAGULATION		1.00	5.2e-01	5.9e-01
WP_PI3KAKT_SIGNALING_PATHWAY		-1.15	3.4e-01	5.9e-01
HALLMARK_INFLAMMATORY_RESPONSE		1.15	3.2e-01	5.9e-01
WP_CYTOPLASMIC_RIBOSOMAL_PROTEINS		1.21	2.3e-01	5.9e-01
HALLMARK_IL2_STAT5_SIGNALING		-1.29	1.9e-01	5.9e-01

