

Gene Set

	C8	IC(Δ)	p-value	FDR
VERHAAK_GLIOBLASTOMA_PRONEURAL		0.894(0.005)	2.62e-06	1.82e-05
GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT		0.869(0.005)	2.62e-06	1.82e-05
E2F3_UP.V1		0.847(0.004)	2.62e-06	1.82e-05
REACTOME_VOLTAGE_GATED_POTASSIUM_CHANNELS		0.844(0.007)	2.62e-06	1.82e-05
MICROTUBULE_BASED_MOVEMENT		0.832(0.005)	2.62e-06	1.82e-05
NIELSEN_SYNOVIAL_SARCOMA		0.831(0.005)	2.62e-06	1.82e-05
HUMMEL_BURKITTs_LYMPHOMA		0.831(0.005)	2.62e-06	1.82e-05
GOLUB_ALL_VS_AML		0.828(0.006)	2.62e-06	1.82e-05
NIELSEN_GIST_AND_SYNOVIAL_SARCOMA		0.827(0.005)	2.62e-06	1.82e-05
REACTOME_NEURONAL_SYSTEM		0.826(0.006)	2.62e-06	1.82e-05
REN_ALVEOLAR_RHABDOMYOSARCOMA		0.823(0.006)	2.62e-06	1.82e-05
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_DN		0.819(0.007)	2.62e-06	1.82e-05
REACTOME_POTASSIUM_CHANNELS		0.819(0.007)	2.62e-06	1.82e-05
ASGHARZADEH_NEUROBLASTOMA_POOR_SURVIVAL_DN		0.812(0.006)	2.62e-06	1.82e-05
NEUROTRANSMITTER_SECRETION		0.811(0.006)	2.62e-06	1.82e-05
HALLMARK_SPERMATOGENESIS		0.809(0.006)	2.62e-06	1.82e-05
RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_POORLY		0.809(0.005)	2.62e-06	1.82e-05
REGULATED_SECRETORY_PATHWAY		0.807(0.007)	2.62e-06	1.82e-05
WANG_LSD1_TARGETS_DN		0.805(0.006)	2.62e-06	1.82e-05
BARIS_THYROID_CANCER		0.804(0.007)	2.62e-06	1.82e-05
POSITIVE_REGULATION_OF_PROTEIN_METABOLIC_PROCESS		-0.818(0.005)	2.62e-06	6.70e-06
BLALOCK_ALZHEIMERS_DISEASE		-0.820(0.006)	2.62e-06	6.70e-06
POSITIVE_REGULATION_OF_SIGNAL_TRANSDUCTION		-0.823(0.005)	2.62e-06	6.70e-06
EGFR_UP.V1		-0.823(0.005)	2.62e-06	6.70e-06
XU_HGF_SIGNALING_NOT_VIA_AKT1_6HR		-0.823(0.005)	2.62e-06	6.70e-06
BLALOCK_ALZHEIMERS_DISEASE_INCIPIENT		-0.824(0.006)	2.62e-06	6.70e-06
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION		-0.824(0.007)	2.62e-06	6.70e-06
POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_METABOLIC_PROCESS		-0.826(0.005)	2.62e-06	6.70e-06
JOHNSTONE_PARVB_TARGETS_3_UP		-0.828(0.005)	2.62e-06	6.70e-06
HUMMERICH_SKIN_CANCER_PROGRESSION_UP		-0.829(0.005)	2.62e-06	6.70e-06
NAKAYAMA_SOFT_TISSUE_TUMORS_PCA1		-0.829(0.005)	2.62e-06	6.70e-06
BROWN_MYELOID_CELL_DEVELOPMENT_UP		-0.834(0.006)	2.62e-06	6.70e-06
POSITIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_CASCADE		-0.835(0.005)	2.62e-06	6.70e-06
REACTOME_EXTRINSIC_PATHWAY_FOR_APOPTOSIS		-0.838(0.006)	2.62e-06	6.70e-06
REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_CASCADE		-0.840(0.004)	2.62e-06	6.70e-06
AMIT_EGF_RESPONSE_480_HELA		-0.841(0.005)	2.62e-06	6.70e-06
I_KAPPAB_KINASE_NF_KAPPAB_CASCADE		-0.843(0.005)	2.62e-06	6.70e-06
LI_INDUCED_T_TO_NATURAL_KILLER_UP		-0.850(0.005)	2.62e-06	6.70e-06
VERHAAK_GLIOBLASTOMA_MESENCHYMAL		-0.856(0.005)	2.62e-06	6.70e-06
MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM		-0.867(0.007)	2.62e-06	6.70e-06