

C8 vs Pathway Expression

	C8		IC(Δ)	P-val	FDR
VERHAAK_GLIOBLASTOMA_PRONEURAL			0.894(0.004)	2.62e-06	1.93e-05
GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT			0.869(0.004)	2.62e-06	1.93e-05
E2F3_UP.V1			0.847(0.005)	2.62e-06	1.93e-05
REACTOME_VOLTAGE_GATED_POTASSIUM_CHANNELS			0.844(0.006)	2.62e-06	1.93e-05
MICROTUBULE_BASED_MOVEMENT			0.832(0.006)	2.62e-06	1.93e-05
NIELSEN_SYNOVIAL_SARCOMA			0.831(0.005)	2.62e-06	1.93e-05
HUMMEL_BURKITTS_LYMPHOMA			0.831(0.006)	2.62e-06	1.93e-05
GOLUB_ALL_VS_AML			0.828(0.005)	2.62e-06	1.93e-05
NIELSEN_GIST_AND_SYNOVIAL_SARCOMA			0.827(0.004)	2.62e-06	1.93e-05
REACTOME_NEURONAL_SYSTEM			0.826(0.007)	2.62e-06	1.93e-05
REN_ALVEOLAR_RHABDOMYOSARCOMA			0.823(0.005)	2.62e-06	1.93e-05
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_DN			0.819(0.007)	2.62e-06	1.93e-05
REACTOME_POTASSIUM_CHANNELS			0.819(0.007)	2.62e-06	1.93e-05
ASGHARZADEH_NEUROBLASTOMA_POOR_SURVIVAL_DN			0.812(0.005)	2.62e-06	1.93e-05
NEUROTRANSMITTER_SECRETION			0.811(0.007)	2.62e-06	1.93e-05
HALLMARK_SPERMATOGENESIS			0.809(0.006)	2.62e-06	1.93e-05
RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_POORLY			0.809(0.005)	2.62e-06	1.93e-05
REGULATED_SECRETORY_PATHWAY			0.807(0.007)	2.62e-06	1.93e-05
WANG_LSD1_TARGETS_DN			0.805(0.008)	2.62e-06	1.93e-05
BARIS_THYROID_CANCER			0.804(0.007)	2.62e-06	1.93e-05
POSITIVE_REGULATION_OF_PROTEIN_METABOLIC_PROCESS			-0.818(0.006)	2.62e-06	6.86e-06
BLALOCK_ALZHEIMERS_DISEASE			-0.820(0.007)	2.62e-06	6.86e-06
POSITIVE_REGULATION_OF_SIGNAL_TRANSDUCTION			-0.823(0.005)	2.62e-06	6.86e-06
EGFR_UP.V1			-0.823(0.005)	2.62e-06	6.86e-06
XU_HGF_SIGNALING_NOT_VIA_AKT1_6HR			-0.823(0.006)	2.62e-06	6.86e-06
BLALOCK_ALZHEIMERS_DISEASE_INCIPIENT			-0.824(0.007)	2.62e-06	6.86e-06
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION			-0.824(0.006)	2.62e-06	6.86e-06
POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_METABOLIC_PROCESS			-0.826(0.006)	2.62e-06	6.86e-06
JOHNSTONE_PARVB_TARGETS_3_UP			-0.828(0.005)	2.62e-06	6.86e-06
HUMMERICH_SKIN_CANCER_PROGRESSION_UP			-0.829(0.005)	2.62e-06	6.86e-06
NAKAYAMA_SOFT_TISSUE_TUMORS_PCA1			-0.829(0.005)	2.62e-06	6.86e-06
BROWN_MYELOID_CELL_DEVELOPMENT_UP			-0.834(0.005)	2.62e-06	6.86e-06
POSITIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_CASCADE			-0.835(0.005)	2.62e-06	6.86e-06
REACTOME_EXTRINSIC_PATHWAY_FOR_APOPTOSIS			-0.838(0.005)	2.62e-06	6.86e-06
REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_CASCADE			-0.840(0.005)	2.62e-06	6.86e-06
AMIT_EGF_RESPONSE_480_HELA			-0.841(0.005)	2.62e-06	6.86e-06
I_KAPPAB_KINASE_NF_KAPPAB_CASCADE			-0.843(0.005)	2.62e-06	6.86e-06
LI_INDUCED_T_TO_NATURAL_KILLER_UP			-0.850(0.005)	2.62e-06	6.86e-06
VERHAAK_GLIOBLASTOMA_MESENCHYMAL			-0.856(0.005)	2.62e-06	6.86e-06
MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM			-0.867(0.007)	2.62e-06	6.86e-06