

C1 vs Pathway Expression

Legend: C1

Color scale: -0.908(0.004) to 0.927(0.004)

IC(Δ) P-val FDR

Pathway	IC(Δ)	P-val	FDR
VERHAAK_GLIOBLASTOMA_PRONEURAL	-0.870(0.005)	2.62e-06	8.13e-06
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_DN	-0.859(0.010)	2.62e-06	8.13e-06
ASGHARZADEH_NEUROBLASTOMA_POOR_SURVIVAL_DN	-0.854(0.006)	2.62e-06	8.13e-06
MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_DN	-0.851(0.006)	2.62e-06	8.13e-06
REACTOME_NEURONAL_SYSTEM	-0.842(0.007)	2.62e-06	8.13e-06
REACTOME_VOLTAGE_GATED_POTASSIUM_CHANNELS	-0.840(0.007)	2.62e-06	8.13e-06
MICROTUBULE_BASED_MOVEMENT	-0.840(0.006)	2.62e-06	8.13e-06
REACTOME_POTASSIUM_CHANNELS	-0.838(0.007)	2.62e-06	8.13e-06
HUMMEL_BURKITT_LYMPHOMA	-0.838(0.006)	2.62e-06	8.13e-06
WANG_LSD1_TARGETS_DN	-0.837(0.007)	2.62e-06	8.13e-06
KIM_MYCL1_AMPLIFICATION_TARGETS	-0.837(0.006)	2.62e-06	8.13e-06
CAHOY_NEURONAL	-0.836(0.007)	2.62e-06	8.13e-06
REACTOME_TRANSMISSION_ACROSS_CHEMICAL_SYNAPSES	-0.834(0.006)	2.62e-06	8.13e-06
GOLUB_ALL_VS_AML	-0.834(0.006)	2.62e-06	8.13e-06
LEIN_LOCALIZED_TO_PROXIMAL_DENDRITES	-0.834(0.006)	2.62e-06	8.13e-06
LEIN_NEURON_MARKERS	-0.833(0.006)	2.62e-06	8.13e-06
GLUTAMATE_RECEPTOR_ACTIVITY	-0.833(0.007)	2.62e-06	8.13e-06
REACTOME_GABA_SYNTHESIS_RELEASE_REUPTAKE_AND_DEGRADATION	-0.833(0.006)	2.62e-06	8.13e-06
CHROMATIN_REMODELING_COMPLEX	-0.831(0.007)	2.62e-06	8.13e-06
E2F3_UP.V1	0.846(0.005)	2.62e-06	2.09e-05
VANLOO_SP3_TARGETS_DN	0.847(0.006)	2.62e-06	2.09e-05
DELPUECH_FOXO3_TARGETS	0.847(0.008)	2.62e-06	2.09e-05
I_KAPPAB_KINASE_NF_KAPPAB_CASCADE	0.851(0.006)	2.62e-06	2.09e-05
DELPUECH_FOXO3_TARGETS_UP	0.851(0.006)	2.62e-06	2.09e-05
BIOCARTA_SODD_PATHWAY	0.854(0.005)	2.62e-06	2.09e-05
GRAESSMANN_APOPTOSIS_BY_DOXORUBICIN_UP	0.857(0.006)	2.62e-06	2.09e-05
GOLUB_ALL_VS_AML_DN	0.863(0.006)	2.62e-06	2.09e-05
LI_INDUCED_T_TO_NATURAL_KILLER_UP	0.866(0.006)	2.62e-06	2.09e-05
ENK_UV_RESPONSE_KERATINOCYTE_UP	0.867(0.005)	2.62e-06	2.09e-05
LI_INDUCED_T_TO_NATURAL_KILLER	0.868(0.007)	2.62e-06	2.09e-05
PID_TRAIL_PATHWAY	0.868(0.005)	2.62e-06	2.09e-05
LI_WILMS_TUMOR_VS_FETAL_KIDNEY_1	0.869(0.005)	2.62e-06	2.09e-05
CHUANG_OXIDATIVE_STRESS_RESPONSE_UP	0.872(0.007)	2.62e-06	2.09e-05
VERHAAK_GLIOBLASTOMA_MESENCHYMAL	0.873(0.005)	2.62e-06	2.09e-05
BLALOCK_ALZHEIMERS_DISEASE_INCIPIENT	0.878(0.005)	2.62e-06	2.09e-05
SWEET_KRAS_ONCOGENIC_SIGNATURE	0.882(0.005)	2.62e-06	2.09e-05
BLALOCK_ALZHEIMERS_DISEASE	0.885(0.004)	2.62e-06	2.09e-05
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION	0.889(0.006)	2.62e-06	2.09e-05
REACTOME_EXTRINSIC_PATHWAY_FOR_APOPTOSIS	0.927(0.004)	2.62e-06	2.09e-05
MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM			