

1C_INT_F480_HI_MACROPHAGE_DN, GSE27859_DC_VS_CD11C_INT_F480_HI_MACROPHAGE_DN

GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN
GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_UP, GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_UP
PURBEY_TARGETS_OF_CTBP1_AND_SATB1_DN, PURBEY_TARGETS_OF_CTBP1_AND_SATB1_DN
GO_REGULATION_OF_SMOOTH_MUSCLE_CELL_MIGRATION, GO_REGULATION_OF_SMOOTH_MUSCLE_CELL_MIGRATION
MAHADEVAN_GIST_MORPHOLOGICAL_SWITCH, MAHADEVAN_GIST_MORPHOLOGICAL_SWITCH
GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_DN, GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_DN
GO_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS, GO_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS
GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP
REACTOME_ACTIVATION_OF_NMDA_RECEPTOR_UPON GLUTAMATE_BINDING_AND_POSTSYNAPTIC_EVENTS, REACTOME_ACTIVATION_OF_NMDA_RECEPTOR_UPON GLUTAMATE_BINDING_AND_POSTSYNAPTIC_EVENTS
GO_REGULATION_OF_LIPASE_ACTIVITY, GO_REGULATION_OF_LIPASE_ACTIVITY
AGARWAL_AKT_PATHWAY_TARGETS, AGARWAL_AKT_PATHWAY_TARGETS
GO_EMBRYO_IMPLANTATION, GO_EMBRYO_IMPLANTATION
LUND_SILENCED_BY_METHYLATION, LUND_SILENCED_BY_METHYLATION
GO_CHEMOATTRACTANT_ACTIVITY, GO_CHEMOATTRACTANT_ACTIVITY
GO_PHOTORECEPTOR_OUTER_SEGMENT, GO_PHOTORECEPTOR_OUTER_SEGMENT
EPPERT_CE_HSC_LSC, EPPERT_CE_HSC_LSC