L_MEMBRANES_ACT_MAST_CELL_UP, GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_UP

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GSE25890 CTRL VS IL33 IL7 TREATED NUOCYTES DN, GSE25890 CTRL VS IL33 IL7 TREATED NUOCYTES DN
GSE11924_TFH_VS_TH2_CD4_TCELL_DN, GSE11924_TFH_VS_TH2_CD4_TCELL_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN
GOBP_REGULATION_OF_REGULATED_SECRETORY_PATHWAY, GOBP_REGULATION_OF_REGULATED_SECRETORY_PATHWAY
REACTOME SIGNALING BY VEGF, REACTOME SIGNALING BY VEGF
GOBP_ACTOMYOSIN_STRUCTURE_ORGANIZATION, GOBP_ACTOMYOSIN_STRUCTURE_ORGANIZATION
GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP
GOBP REGULATION OF SYNAPTIC VESICLE EXOCYTOSIS, GOBP REGULATION OF SYNAPTIC VESICLE EXOCYTOSIS
HP_BULBAR_SIGNS, HP_BULBAR_SIGNS
GOBP_POSITIVE_REGULATION_OF_EXOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_EXOCYTOSIS
GOBP_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GOBP_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBL
PTEN_DN.V1_DN, PTEN_DN.V1_DN
MARTIN VIRAL GPCR SIGNALING UP, MARTIN VIRAL GPCR SIGNALING UP
MODULE 201, MODULE 201
GOBP_POSITIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY, GOBP_POSITIVE_REGULATION_OF_REGULATED_SECRET
GOBP_REGULATION_OF_VASCULAR_PERMEABILITY, GOBP_REGULATION_OF_VASCULAR_PERMEABILITY
· HP_FATIGUABLE_WEAKNESS_OF_PROXIMAL_LIMB_MUSCLES, HP_FATIGUABLE_WEAKNESS_OF_PROXIMAL_LIMB_MUSCLES
GOBP_NEGATIVE_REGULATION_OF_VASCULAR_PERMEABILITY, GOBP_NEGATIVE_REGULATION_OF_VASCULAR_PERMEABILITY
GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER
GOBP POSITIVE REGULATION OF ENDOTHELIAL CELL PROLIFERATION, GOBP POSITIVE REGULATION OF ENDOTHELIAL CE
GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION, GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLAT
HP_NEMALINE_BODIES, HP_NEMALINE_BODIES
LE_NEURONAL_DIFFERENTIATION_UP, LE_NEURONAL_DIFFERENTIATION_UP
chrXq12, chrXq12
GOBP REGULATION OF SYNAPTIC TRANSMISSION CHOLINERGIC, GOBP REGULATION OF SYNAPTIC TRANSMISSION CHOLIN
DESCARTES_FETAL_HEART_CARDIOMYOCYTES, DESCARTES_FETAL_HEART_CARDIOMYOCYTES
GOBP POSITIVE REGULATION OF TISSUE REMODELING, GOBP POSITIVE REGULATION OF TISSUE REMODELING
REACTOME_NGF_INDEPENDANT_TRKA_ACTIVATION, REACTOME_NGF_INDEPENDANT_TRKA_ACTIVATION
GOBP_POSITIVE_REGULATION_OF_CIRCADIAN_SLEEP_WAKE_CYCLE_SLEEP, GOBP_POSITIVE_REGULATION_OF_CIRCADIAN_SLE
REACTOME ACTIVATION OF TRKA RECEPTORS, REACTOME ACTIVATION OF TRKA RECEPTORS
chr2q23, chr2q23
MAINA_VHL_TARGETS_UP, MAINA_VHL_TARGETS_UP
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