

TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN

GSE2770_UNTREATED_VS_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_ACT_CD4_TCELL_48H_UP
GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN
GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT, GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT
GSE40274_SATB1_VS_FOXP3_AND_SATB1_TRANSDUCE_ACTIVATED_CD4_TCELL_UP, GSE40274_SATB1_VS_FOXP3_AND_SATB1_TRANSDUCE_ACTIVATED_CD4_TCELL_UP
MIR7156_3P, MIR7156_3P
MIR4645_3P, MIR4645_3P
MIR5572, MIR5572
MIR6881_5P, MIR6881_5P
GOMF_INORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_INORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_INOSITOL_PHOSPHATE_METABOLIC_PROCESS, GOBP_INOSITOL_PHOSPHATE_METABOLIC_PROCESS
GOBP_CHLORIDE_TRANSPORT, GOBP_CHLORIDE_TRANSPORT
GOMF_CHLORIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_CHLORIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
HP_ABNORMALLY_LARGE_GLOBE, HP_ABNORMALLY_LARGE_GLOBE
GOMF_EXTRACELLULAR_LIGAND_GATED_ION_CHANNEL_ACTIVITY, GOMF_EXTRACELLULAR_LIGAND_GATED_ION_CHANNEL_ACTIVITY
MODULE_316, MODULE_316
HAY_BONE_MARROW_CD34_POS_MKP, HAY_BONE_MARROW_CD34_POS_MKP
GOCC_CHLORIDE_CHANNEL_COMPLEX, GOCC_CHLORIDE_CHANNEL_COMPLEX
HP_VERTEBRAL_CLEFTING, HP_VERTEBRAL_CLEFTING
MODULE_214, MODULE_214
HP_CORONAL_CLEFT_VERTEBRAE, HP_CORONAL_CLEFT_VERTEBRAE
GOBP_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS, GOBP_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS
GOMF_LIGAND_GATED_ANION_CHANNEL_ACTIVITY, GOMF_LIGAND_GATED_ANION_CHANNEL_ACTIVITY
GOBP_SUBSTRATE_INDEPENDENT_TELENCEPHALIC_TANGENTIAL_MIGRATION, GOBP_SUBSTRATE_INDEPENDENT_TELENCEPHALIC_TANGENTIAL_MIGRATION
GOMF_GABA_GATED_CHLORIDE_ION_CHANNEL_ACTIVITY, GOMF_GABA_GATED_CHLORIDE_ION_CHANNEL_ACTIVITY
GOCC_GABA_RECEPTOR_COMPLEX, GOCC_GABA_RECEPTOR_COMPLEX
HP_HYPERGRANULOSIS, HP_HYPERGRANULOSIS
GOMF_GABA_RECEPTOR_ACTIVITY, GOMF_GABA_RECEPTOR_ACTIVITY
GOMF_ACETYLCHOLINE_RECEPTOR_INHIBITOR_ACTIVITY, GOMF_ACETYLCHOLINE_RECEPTOR_INHIBITOR_ACTIVITY