
_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP
GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_UP, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_
GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_UP
GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP, GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP
GSE13493_DP_VS_CD4INTCD8POS_THYMOCYTE_UP, GSE13493_DP_VS_CD4INTCD8POS_THYMOCYTE_UP
GSE17721_0.5H_VS_8H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_8H_POLYIC_BMDC_DN
GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_DN
GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP
GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GO_PHOSPHATASE_BINDING, GO_PHOSPHATASE_BINDING
GSE40274_FOXP3_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_FOXP3_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE17721_CTRL_VS_CPG_6H_BMDC_DN, GSE17721_CTRL_VS_CPG_6H_BMDC_DN
GO_PROTEIN_AUTOUBIQUITINATION, GO_PROTEIN_AUTOUBIQUITINATION
WAMUNYOKOLI_OVARIAN_CANCER_LMP_DN, WAMUNYOKOLI_OVARIAN_CANCER_LMP_DN
KEGG_COLORECTAL_CANCER, KEGG_COLORECTAL_CANCER
GSE11961_MEMORY_BCELL_DAY40_VS_GERMINAL_CENTER_BCELL_DAY40_DN, GSE11961_MEMORY_BCELL_DAY40_VS_GERMINAL_CENTER_BCELL_DAY40_DN
GTTATAT_MIR410, GTTATAT_MIR410
GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN
GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE
DARWICHE_PAPILLOMA_RISK_HIGH_DN, DARWICHE_PAPILLOMA_RISK_HIGH_DN
GO_RECEPTOR_SIGNALING_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GO_RECEPTOR_SIGNALING_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY
PID_ECADHERIN_NASCENT_AJ_PATHWAY, PID_ECADHERIN_NASCENT_AJ_PATHWAY
GO_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE, GO_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE
PLASARI_TGFB1_SIGNALING_VIA_NFIC_1HR_UP, PLASARI_TGFB1_SIGNALING_VIA_NFIC_1HR_UP
GO_RESPIRATORY_SYSTEM_DEVELOPMENT, GO_RESPIRATORY_SYSTEM_DEVELOPMENT
GO_ACTOMYOSIN, GO_ACTOMYOSIN
MODULE_486, MODULE_486
GO_SKELETAL_SYSTEM_MORPHOGENESIS, GO_SKELETAL_SYSTEM_MORPHOGENESIS
chr16q21, chr16q21
GO_EMBRYONIC_SKELETAL_SYSTEM_MORPHOGENESIS, GO_EMBRYONIC_SKELETAL_SYSTEM_MORPHOGENESIS
GO_EMBRYONIC_HINDLIMB_MORPHOGENESIS, GO_EMBRYONIC_HINDLIMB_MORPHOGENESIS
GO_DIGESTIVE_TRACT_MORPHOGENESIS, GO_DIGESTIVE_TRACT_MORPHOGENESIS
GO_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE, GO_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE
GO_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING, GO_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING
GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY