

G_VS_CD24HI_TCONV_THYMUS_UP, GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_UP

GSE17721_0.5H_VS_4H_CPG_BMDC_UP, GSE17721_0.5H_VS_4H_CPG_BMDC_UP
GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_45MIN_UP, GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_45MIN_UP
GSE37532_TREG_VS_TCONV_CD4_TCELL_FROM_LN_DN, GSE37532_TREG_VS_TCONV_CD4_TCELL_FROM_LN_DN
GSE34515_CD16_NEG_MONOCYTE_VS_DC_DN, GSE34515_CD16_NEG_MONOCYTE_VS_DC_DN
HALLMARK_ADIPOGENESIS, HALLMARK_ADIPOGENESIS
GSE17721_POLYIC_VS_PAM3CSK4_24H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_24H_BMDC_UP
GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCYTE_DN, GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMO
GSE13411_NAIVE_VS_IGM_MEMORY_BCELL_DN, GSE13411_NAIVE_VS_IGM_MEMORY_BCELL_DN
GSE17721_0.5H_VS_12H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_12H_POLYIC_BMDC_UP
BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS
GSE339_CD4POS_VS_CD4CD8DN_DC_UP, GSE339_CD4POS_VS_CD4CD8DN_DC_UP
GO_REGULATION_OF_PROTEIN_BINDING, GO_REGULATION_OF_PROTEIN_BINDING
GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_DN, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_DN
RAY_TUMORIGENESIS_BY_ERBB2_CDC25A_DN, RAY_TUMORIGENESIS_BY_ERBB2_CDC25A_DN
GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_UP
GO_NEGATIVE_REGULATION_OF_CELL_PROJECTION_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_CELL_PROJECTION_ORGANIZATION
LANDIS_ERBB2_BREAST_TUMORS_324_DN, LANDIS_ERBB2_BREAST_TUMORS_324_DN
GO_NEGATIVE_REGULATION_OF_CELL_GROWTH, GO_NEGATIVE_REGULATION_OF_CELL_GROWTH
ABE_INNER_EAR, ABE_INNER_EAR
GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCECD4_TCELL_UP, GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCECD4_TCELL_UP
GO_CONNECTIVE_TISSUE_DEVELOPMENT, GO_CONNECTIVE_TISSUE_DEVELOPMENT
GO_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE, GO_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE
GSE12505_WT_VS_E2-2_HET_PDC_UP, GSE12505_WT_VS_E2-2_HET_PDC_UP
GSE6875_TCONV_VS_TREG_DN, GSE6875_TCONV_VS_TREG_DN
GO_NEGATIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOL
GSE9650_GP33_VS_GP276_LCMV_SPECIFIC_EXHAUSTED_CD8_TCELL_UP, GSE9650_GP33_VS_GP276_LCMV_SPECIFIC_EXHAUSTED_CD8_TCELL_UP
GO_APPENDAGE_DEVELOPMENT, GO_APPENDAGE_DEVELOPMENT
AR_01, AR_01
GO_CARTILAGE_DEVELOPMENT, GO_CARTILAGE_DEVELOPMENT
KANG_CISPLATIN_RESISTANCE_UP, KANG_CISPLATIN_RESISTANCE_UP
MODULE_343, MODULE_343
GO_MULTICELLULAR_ORGANISM_GROWTH, GO_MULTICELLULAR_ORGANISM_GROWTH
GO_POSITIVE_REGULATION_OF_PHOSPHOLIPID_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_PHOSPHOLIPID_METABOLIC_PROCESS
MODULE_134, MODULE_134
GO_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GO_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS
PR_01, PR_01