RDIQUIMOD\_1H\_BMDC\_DN, GSE17721\_CPG\_VS\_GARDIQUIMOD\_1H\_BMDC\_DN

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GSE23308_CTRL_VS_CORTICOSTERONE_TREATED_MACROPHAGE_MINERALCORTICOID_REC_KO_UP, GSE23308_CTRL_VS_CORTICOSTERONE_TREATED_MACROPHAGE_MINERA
GSE17721_LPS_VS_PAM3CSK4_6H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_6H_BMDC_DN
GSE17721_LPS_VS_CPG_8H_BMDC_DN, GSE17721_LPS_VS_CPG_8H_BMDC_DN
GSE17721_LPS_VS_POLYIC_4H_BMDC_DN, GSE17721_LPS_VS_POLYIC_4H_BMDC_DN
GSE339_CD4POS_VS_CD8POS_DC_DN, GSE339_CD4POS_VS_CD8POS_DC_DN
GSE17721_CTRL_VS_CPG_1H_BMDC_UP, GSE17721_CTRL_VS_CPG_1H_BMDC_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN
GSE17721_LPS_VS_GARDIQUIMOD_8H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_8H_BMDC_DN
GSE17721_CTRL_VS_CPG_2H_BMDC_UP, GSE17721_CTRL_VS_CPG_2H_BMDC_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_4H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_4H_BMDC_DN
GSE39820_CTRL_VS_TGFBETA3_IL6_IL23A_CD4_TCELL_DN, GSE39820_CTRL_VS_TGFBETA3_IL6_IL23A_CD4_TCELL_DN
GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP
GSE8685_IL2_STARVED_VS_IL15_ACT_IL2_STARVED_CD4_TCELL_UP, GSE8685_IL2_STARVED_VS_IL15_ACT_IL2_STARVED_CD4_TCELL_UP
GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_UP, GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_UP
GSE27786_CD8_TCELL_VS_NKTCELL_UP, GSE27786_CD8_TCELL_VS_NKTCELL_UP
GSE14308_TH1_VS_TH17_UP, GSE14308_TH1_VS_TH17_UP
GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_12H_DN, GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_12H_DN
GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN, GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN
IGARASHI_ATF4_TARGETS_DN, IGARASHI_ATF4_TARGETS_DN
GO_MEMBRANE_FUSION, GO_MEMBRANE_FUSION
GO_TRANSLATION_INITIATION_FACTOR_BINDING, GO_TRANSLATION_INITIATION_FACTOR_BINDING
GO_MEMBRANE_DOCKING, GO_MEMBRANE_DOCKING
GO_POSITIVE_REGULATION_OF_PHAGOCYTOSIS, GO_POSITIVE_REGULATION_OF_PHAGOCYTOSIS
LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_UP, LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_UP
GO VESICLE DOCKING, GO VESICLE DOCKING
GO_REGULATION_OF_PHAGOCYTOSIS, GO_REGULATION_OF_PHAGOCYTOSIS
GO_TRANSLATION_REPRESSOR_ACTIVITY, GO_TRANSLATION_REPRESSOR_ACTIVITY
MODULE_229, MODULE_229
GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN, GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN
GO_SNARE_BINDING, GO_SNARE_BINDING
GO_METALLOAMINOPEPTIDASE_ACTIVITY, GO_METALLOAMINOPEPTIDASE_ACTIVITY
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE
GO_POSITIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY
GO_VESICLE_DOCKING_INVOLVED_IN_EXOCYTOSIS, GO_VESICLE_DOCKING_INVOLVED_IN_EXOCYTOSIS
GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT, GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT
GO_PROTEIN_TYROSINE_KINASE_BINDING, GO_PROTEIN_TYROSINE_KINASE_BINDING
DER_IFN_GAMMA_RESPONSE_DN, DER_IFN_GAMMA_RESPONSE_DN
GRATIAS_RETINOBLASTOMA_16Q24, GRATIAS_RETINOBLASTOMA_16Q24
chr7q36, chr7q36
GSE3982_MAST_CELL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_MAST_CELL_VS_EFF_MEMORY_CD4_TCELL_DN
GO_NUCLEOTIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GO_NUCLEOTIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GO_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION, GO_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION
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