

GADIQUIMOD_8H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_8H_BMDC_DN

GSE17721_0.5H_VS_8H_GARDIQUIMOD_BMDC_DN, GSE17721_0.5H_VS_8H_GARDIQUIMOD_BMDC_DN
GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_UP
GSE17721_LPS_VS_PAM3CSK4_8H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_8H_BMDC_UP
KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS, KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS
GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_UP
GSE17721_POLYIC_VS_PAM3CSK4_8H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_8H_BMDC_UP
GSE17721_LPS_VS_CPG_6H_BMDC_UP, GSE17721_LPS_VS_CPG_6H_BMDC_UP
GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_UP, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_UP
GOBP_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY
GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_UP, GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_UP
GSE24210_IL35_TREATED_VS_RESTING_TREG_UP, GSE24210_IL35_TREATED_VS_RESTING_TREG_UP
GSE17721_4_VS_24H_CPG_BMDC_UP, GSE17721_4_VS_24H_CPG_BMDC_UP
WP_APOPTOSIS, WP_APOPTOSIS
GOBP_DNA_MODIFICATION, GOBP_DNA_MODIFICATION
GOBP_NEGATIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION, GOBP_NEGATIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION
MIR4779, MIR4779
KEGG_GLYCEROPHOSPHOLIPID_METABOLISM, KEGG_GLYCEROPHOSPHOLIPID_METABOLISM
GOMF_CCR_CHEMOKINE_RECEPTOR_BINDING, GOMF_CCR_CHEMOKINE_RECEPTOR_BINDING
GOMF_MEMBRANE_INSERTASE_ACTIVITY, GOMF_MEMBRANE_INSERTASE_ACTIVITY
SASAKI_TARGETS_OF_TP73_AND_TP63, SASAKI_TARGETS_OF_TP73_AND_TP63
GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_TCONV_CD4_TCELL_DN, GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_TCONV_CD4_TCELL_DN