

GSE17721_PAM3CSK4_4H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_4H_BMDC_DN

GSE46606_UNSTIM_VS_CD40L_IL2_IL5_3DAY_STIMULATED_IRF4MID_SORTED_BCELL_UP, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_3DAY_STIMULATED_IRF4MID_SORTED_BCELL_UP
GSE17721_4_VS_24H_GARDIQUIMOD_BMDC_DN, GSE17721_4_VS_24H_GARDIQUIMOD_BMDC_DN
GSE17721_PAM3CSK4_VS_CPG_12H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_12H_BMDC_UP
GSE17721_LPS_VS_CPG_8H_BMDC_DN, GSE17721_LPS_VS_CPG_8H_BMDC_DN
GSE17721_PAM3CSK4_VS_CPG_8H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_8H_BMDC_UP
GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_DN
GSE17721_0.5H_VS_24H_GARDIQUIMOD_BMDC_DN, GSE17721_0.5H_VS_24H_GARDIQUIMOD_BMDC_DN
GSE17721_0.5H_VS_24H_PAM3CSK4_BMDC_DN, GSE17721_0.5H_VS_24H_PAM3CSK4_BMDC_DN
GSE17721_0.5H_VS_12H_PAM3CSK4_BMDC_DN, GSE17721_0.5H_VS_12H_PAM3CSK4_BMDC_DN
GSE15624_CTRL_VS_3H_HALOFUGINONE_TREATED_CD4_TCELL_UP, GSE15624_CTRL_VS_3H_HALOFUGINONE_TREATED_CD4_TCELL_UP
GR_01, GR_01
GO_NEGATIVE_REGULATION_OF_HISTONE_ACETYLATION, GO_NEGATIVE_REGULATION_OF_HISTONE_ACETYLATION
REACTOME_SYNTHESIS_OF_VERY_LONG_CHAIN_FATTY_ACYL_COAS, REACTOME_SYNTHESIS_OF_VERY_LONG_CHAIN_FATTY_ACYL_COAS
REACTOME_INWARDLY_RECTIFYING_K_CHANNELS, REACTOME_INWARDLY_RECTIFYING_K_CHANNELS
GO_L_ASCORBIC_ACID_METABOLIC_PROCESS, GO_L_ASCORBIC_ACID_METABOLIC_PROCESS
GO_CYCLOOXYGENASE_PATHWAY, GO_CYCLOOXYGENASE_PATHWAY