

CD11C_INT_F480_HI_MACROPHAGE_DN, GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN

- GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TCONV_DN, GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TCONV_DN
- GSE6674_CPG_VS_PL2_3_STIM_BCELL_UP, GSE6674_CPG_VS_PL2_3_STIM_BCELL_UP
- GSE27859_DC_VS_CD11C_INT_F480_HI_MACROPHAGE_DN, GSE27859_DC_VS_CD11C_INT_F480_HI_MACROPHAGE_DN
- GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM_DN, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM
- GSE9037_CTRL_VS_LPS_4H_STIM_IRAK4_KO_BMDM_UP, GSE9037_CTRL_VS_LPS_4H_STIM_IRAK4_KO_BMDM_UP
- KASLER_HDAC7_TARGETS_1_UP, KASLER_HDAC7_TARGETS_1_UP
- GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_UP, GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_UP
- GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_DN, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_DN
- GSE8685_IL15_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP, GSE8685_IL15_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4
- MIR504_3P, MIR504_3P
- GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_CD8_TCELL_UP, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_CD8_TCELL_UP
- MIR4270, MIR4270
- GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_DN, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TC
- MIR766_3P, MIR766_3P
- GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_DN, GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_I
- GOBP_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACID_BIOSYNTHETIC_PROCESS
- GOCC_FICOLIN_1_RICH_GRANULE_MEMBRANE, GOCC_FICOLIN_1_RICH_GRANULE_MEMBRANE
- GOBP_RUFFLE_ORGANIZATION, GOBP_RUFFLE_ORGANIZATION
- MIR4455, MIR4455
- MIR764, MIR764
- GOBP_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS
- REACTOME_TRANSFERRIN_ENDOCYTOSIS_AND_RECYCLING, REACTOME_TRANSFERRIN_ENDOCYTOSIS_AND_RECYCLING
- HENDRICKS_SMARCA4_TARGETS_UP, HENDRICKS_SMARCA4_TARGETS_UP
- GOBP_UNSATURATED_FATTY_ACID_METABOLIC_PROCESS, GOBP_UNSATURATED_FATTY_ACID_METABOLIC_PROCESS
- REACTOME_ROS_AND_RNS_PRODUCTION_IN_PHAGOCYTES, REACTOME_ROS_AND_RNS_PRODUCTION_IN_PHAGOCYTES
- MIKKELSEN_ES_LCP_WITH_H3K4ME3, MIKKELSEN_ES_LCP_WITH_H3K4ME3
- GOBP_VERY_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GOBP_VERY_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS
- GSE6259_DEC205_POS_DC_VS_BCELL_UP, GSE6259_DEC205_POS_DC_VS_BCELL_UP
- GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS, GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS
- GOBP_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS, GOBP_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS
- REACTOME_ALPHA_LINOLENIC_OMEGA3_AND_LINOLEIC_OMEGA6_ACID_METABOLISM, REACTOME_ALPHA_LINOLENIC_OMEGA3_AND_LINC
- GOBP_STRIATED_MUSCLE_ADAPTATION, GOBP_STRIATED_MUSCLE_ADAPTATION
- HP_HYPERCHLOREMIC_ACIDOSIS, HP_HYPERCHLOREMIC_ACIDOSIS
- GOMF_HORMONE_BINDING, GOMF_HORMONE_BINDING