

L10_STIM_IL6_KO_MACROPHAGE_180MIN_DN, GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_180MIN_DN

- GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN
- GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN
- GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN, GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN
- GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN
- GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP, GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP
- GOBP_CELLULAR_MODIFIED_AMINO_ACID_METABOLIC_PROCESS, GOBP_CELLULAR_MODIFIED_AMINO_ACID_METABOLIC_PROCESS
- MIR548S, MIR548S
- MIR3149, MIR3149
- MIR6510_5P, MIR6510_5P
- MIR4437, MIR4437
- GOBP_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY
- GOCC_INTRINSIC_COMPONENT_OF_PEROXISOMAL_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_PEROXISOMAL_MEMBRANE
- MIR6757_5P, MIR6757_5P
- REACTOME_MET_ACTIVATES_PTK2_SIGNALING, REACTOME_MET_ACTIVATES_PTK2_SIGNALING
- WRN_TARGET_GENES, WRN_TARGET_GENES
- REACTOME_GRB2_SOS_PROVIDES_LINKAGE_TO_MAPK_SIGNALING_FOR_INTEGRINS, REACTOME_GRB2_SOS_PROVIDES_LINKAGE_TO_MAPK_SIGNALING_FOR_INTEGRINS
- GOBP_LUNG_CELL_DIFFERENTIATION, GOBP_LUNG_CELL_DIFFERENTIATION