

_IL6_STIM_SOCS3_KO_MACROPHAGE_UP, GSE411_UNSTIM_VS_100MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_UP

GSE23502_WT_VS_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_BM_UP, GSE23502_WT_VS_HDC_KO_MYELOID_DERIVED_

GSE411_WT_VS_SOCS3_KO_MACROPHAGE_DN, GSE411_WT_VS_SOCS3_KO_MACROPHAGE_DN

GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN

GSE557_WT_VS_I_AB_KO_DC_UP, GSE557_WT_VS_I_AB_KO_DC_UP

GSE411_UNSTIM_VS_400MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_UP, GSE411_UNSTIM_VS_400MIN_IL6_STIM_SOCS3_KO_MAC

GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_TREG_UP, GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_TREG

GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_12H_UP, GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4

GO_PHOSPHATASE_BINDING, GO_PHOSPHATASE_BINDING

GO_PROTEIN_TYROSINE_KINASE_ACTIVITY, GO_PROTEIN_TYROSINE_KINASE_ACTIVITY

GO_AMINOGLYCAN_CATABOLIC_PROCESS, GO_AMINOGLYCAN_CATABOLIC_PROCESS

GSE36078_UNTREATED_VS_AD5_INF_IL1R_KO_MOUSE_LUNG_DC_UP, GSE36078_UNTREATED_VS_AD5_INF_IL1R_KO_MOUSE_L

MODULE_486, MODULE_486

DARWICHE_SQUAMOUS_CELL_CARCINOMA_DN, DARWICHE_SQUAMOUS_CELL_CARCINOMA_DN

PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN, PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN

GO_ENDODERM_DEVELOPMENT, GO_ENDODERM_DEVELOPMENT

GO_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION, GO_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION

chr4q28, chr4q28

MCDOWELL_ACUTE_LUNG_INJURY_DN, MCDOWELL_ACUTE_LUNG_INJURY_DN

GO_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GO_REGULATION_OF_EPITHELIAL_TO_MESENCHYMA

GO_REGULATION_OF_STEM_CELL_DIFFERENTIATION, GO_REGULATION_OF_STEM_CELL_DIFFERENTIATION

GO_AXIS_ELONGATION, GO_AXIS_ELONGATION