


2_TARGETS_DN, WANG_CLIM2_TARGETS_DN



GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_DN
GSE17721_POLYIC_VS_PAM3CSK4_8H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_8H_BMDC_UP
GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_DN, GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_DN
MTOR_UP.N4.V1_UP, MTOR_UP.N4.V1_UP
LTE2_UP.V1_DN, LTE2_UP.V1_DN
GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP, GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP
GSE18893_CTRL_VS_TNF_TREATED_TREG_24H_DN, GSE18893_CTRL_VS_TNF_TREATED_TREG_24H_DN
GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_1H_DN, GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_1H_DN
AIGNER_ZEB1_TARGETS, AIGNER_ZEB1_TARGETS
GO_LIGAND_DEPENDENT_NUCLEAR_RECEPTOR_BINDING, GO_LIGAND_DEPENDENT_NUCLEAR_RECEPTOR_BINDING
GO_BODY_MORPHOGENESIS, GO_BODY_MORPHOGENESIS
TAVOR_CEBPA_TARGETS_DN, TAVOR_CEBPA_TARGETS_DN
GO_CELL_FATE_DETERMINATION, GO_CELL_FATE_DETERMINATION
GO_REGULATION_BY_VIRUS_OF_VIRAL_PROTEIN_LEVELS_IN_HOST_CELL, GO_REGULATION_BY_VIRUS_OF_VIRAL_PROTEIN_LEVELS_IN_HOST_CELL
GSE21546_UNSTIM_VS_ANTI_CD3_STIM_ELK1_KO_DP_THYMOCYTES_UP, GSE21546_UNSTIM_VS_ANTI_CD3_STIM_ELK1_KO_DP_THYMOCYTES_UP
AZARE_NEOPLASTIC_TRANSFORMATION_BY_STAT3_DN, AZARE_NEOPLASTIC_TRANSFORMATION_BY_STAT3_DN
HAN_JNK_SINGALING_UP, HAN_JNK_SINGALING_UP
GO_CARDIAC_RIGHT_VENTRICLE_MORPHOGENESIS, GO_CARDIAC_RIGHT_VENTRICLE_MORPHOGENESIS
GO_CLATHRIN_COAT_OF_TRANS_GOLGI_NETWORK_VESICLE, GO_CLATHRIN_COAT_OF_TRANS_GOLGI_NETWORK_VESICLE
GO_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GO_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_