Y30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN, GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN

GSE41867_DAY6_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN, GSE41867_DAY6_EFFECTOR_VS_DAY30_EXHAUSTED_
GSE41867_DAY6_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_TCELL_
GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_UP, GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_
GOBP_NCRNA_TRANSCRIPTION, GOBP_NCRNA_TRANSCRIPTION
GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG_EFFECTOR_C
MIR3679_3P, MIR3679_3P
GSE42021_TREG_PLN_VS_CD24HI_TREG_THYMUS_UP, GSE42021_TREG_PLN_VS_CD24HI_TREG_THYMUS_UP
GOBP_SNRNA_TRANSCRIPTION, GOBP_SNRNA_TRANSCRIPTION
GSE1448_ANTI_VALPHA2_VS_VBETA5_DP_THYMOCYTE_DN, GSE1448_ANTI_VALPHA2_VS_VBETA5_DP_THYMOCYTE_DN
MIR379_3P_MIR411_3P, MIR379_3P_MIR411_3P
GSE27786_ERYTHROBLAST_VS_NEUTROPHIL_UP, GSE27786_ERYTHROBLAST_VS_NEUTROPHIL_UP
TCCAGAG_MIR518C, TCCAGAG_MIR518C
MIR499B_3P, MIR499B_3P
GSE41176_WT_VS_TAK1_KO_ANTI_IGM_STIM_BCELL_1H_UP, GSE41176_WT_VS_TAK1_KO_ANTI_IGM_STIM_BCELL_1H_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_30H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_30H_UP
MIR1264, MIR1264
MIR2054, MIR2054
HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_PYRAMIDAL_TRACT, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_PYRAMIDAL_TRAC
GOMF_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
MIR5001_3P, MIR5001_3P
MIR6837_3P, MIR6837_3P
MIR6507_3P, MIR6507_3P
MIR885_5P, MIR885_5P
MIR217_3P, MIR217_3P
GSE4590_PRE_BCELL_VS_SMALL_PRE_BCELL_DN, GSE4590_PRE_BCELL_VS_SMALL_PRE_BCELL_DN
HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_CORTICOSPINAL_TRACT, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_CORTICOSPINA
MIR8060, MIR8060
REACTOME_RND2_GTPASE_CYCLE, REACTOME_RND2_GTPASE_CYCLE
MIR4714_5P, MIR4714_5P
GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN, GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN
MIR216B_3P, MIR216B_3P
ACCTGTTG_UNKNOWN, ACCTGTTG_UNKNOWN
MIR6806_5P, MIR6806_5P
FIGUEROA_AML_METHYLATION_CLUSTER_1_DN, FIGUEROA_AML_METHYLATION_CLUSTER_1_DN
REACTOME_RND3_GTPASE_CYCLE, REACTOME_RND3_GTPASE_CYCLE
GOBP_MANNOSYLATION, GOBP_MANNOSYLATION
GOBP_PROTEIN_MANNOSYLATION, GOBP_PROTEIN_MANNOSYLATION
HP_ATRESIA_OF_THE_EXTERNAL_AUDITORY_CANAL, HP_ATRESIA_OF_THE_EXTERNAL_AUDITORY_CANAL
GNF2_SPTA1, GNF2_SPTA1
HP_ABNORMALLY_LARGE_GLOBE, HP_ABNORMALLY_LARGE_GLOBE
GOBP_PROTEIN_NEDDYLATION, GOBP_PROTEIN_NEDDYLATION
HP_ABNORMALITY_OF_THE_AUDITORY_CANAL, HP_ABNORMALITY_OF_THE_AUDITORY_CANAL
CRX_NRL_DN.V1_DN, CRX_NRL_DN.V1_DN
GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN, GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN
ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_UP, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WI
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAGENTA_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAG
GOBP_LIPID_DROPLET_ORGANIZATION, GOBP_LIPID_DROPLET_ORGANIZATION
HP_SCROTAL_HYPOPLASIA, HP_SCROTAL_HYPOPLASIA
HP_BRANCHIAL_ANOMALY, HP_BRANCHIAL_ANOMALY
MIR214_5P, MIR214_5P
MIR548AD_3P, MIR548AD_3P
TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_GRANULOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_GRANULOCYTE_UP
KENNY_CTNNB1_TARGETS_UP, KENNY_CTNNB1_TARGETS_UP
GOBP_PROTEIN_INSERTION_INTO_ER_MEMBRANE, GOBP_PROTEIN_INSERTION_INTO_ER_MEMBRANE
HP_APLASIA_HYPOPLASIA_INVOLVING_THE_VERTEBRAL_COLUMN, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_VERTEBRAL_COLUMN
MIR668_3P, MIR668_3P
GOCC_PRE_SNORNP_COMPLEX, GOCC_PRE_SNORNP_COMPLEX
HP_SHORT_THORAX, HP_SHORT_THORAX
WP_ALZHEIMERS_DISEASE, WP_ALZHEIMERS_DISEASE
GOBP_SMALL_NUCLEOLAR_RIBONUCLEOPROTEIN_COMPLEX_ASSEMBLY, GOBP_SMALL_NUCLEOLAR_RIBONUCLEOPROTEIN_COMPLEX_
GOBP_INTRA_S_DNA_DAMAGE_CHECKPOINT, GOBP_INTRA_S_DNA_DAMAGE_CHECKPOINT
HP_HYPOPLASIA_OF_THE_ODONTOID_PROCESS, HP_HYPOPLASIA_OF_THE_ODONTOID_PROCESS
HP_BARREL_SHAPED_CHEST, HP_BARREL_SHAPED_CHEST
HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY, HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY
SKP2_TARGET_GENES, SKP2_TARGET_GENES
HP_ABNORMAL_MUSCLE_FIBER_PROTEIN_EXPRESSION, HP_ABNORMAL_MUSCLE_FIBER_PROTEIN_EXPRESSION
GSE29615_CTRL_VS_DAY7_LAIV_FLU_VACCINE_PBMC_DN, GSE29615_CTRL_VS_DAY7_LAIV_FLU_VACCINE_PBMC_DN
GOBP_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS
MIR4529_3P, MIR4529_3P
MIR1281, MIR1281
MEF2_04, MEF2_04
GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGNALING_PATHWAY, GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGN
GOBP_POSITIVE_REGULATION_OF_MONOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MONOCYTE_DIFFERENTIATION
HP_ULNAR_DEVIATION_OF_FINGER, HP_ULNAR_DEVIATION_OF_FINGER
HP_ABNORMAL_EPIPHYSIS_MORPHOLOGY_OF_THE_PHALANGES_OF_THE_HAND, HP_ABNORMAL_EPIPHYSIS_MORPHOLOGY_OF_THE_PH
REACTOME_N_GLYCAN_ANTENNAE_ELONGATION_IN_THE_MEDIAL_TRANS_GOLGI, REACTOME_N_GLYCAN_ANTENNAE_ELONGATION
HP_ABNORMALITY_OF_THE_ODONTOID_PROCESS, HP_ABNORMALITY_OF_THE_ODONTOID_PROCESS
chr6q25, chr6q25
GOMF_3_5_CYCLIC_AMP_PHOSPHODIESTERASE_ACTIVITY, GOMF_3_5_CYCLIC_AMP_PHOSPHODIESTERASE_ACTIVITY
GOBP_LEUKEMIA_INHIBITORY_FACTOR_SIGNALING_PATHWAY, GOBP_LEUKEMIA_INHIBITORY_FACTOR_SIGNALING_PATHWAY
HP_ABNORMAL_FOOT_BONE_OSSIFICATION, HP_ABNORMAL_FOOT_BONE_OSSIFICATION
HP_ABSENT_STERNAL_OSSIFICATION, HP_ABSENT_STERNAL_OSSIFICATION
GOBP_REGULATION_OF_NADP_METABOLIC_PROCESS, GOBP_REGULATION_OF_NADP_METABOLIC_PROCESS
GOMF_CILIARY_NEUROTROPHIC_FACTOR_RECEPTOR_ACTIVITY, GOMF_CILIARY_NEUROTROPHIC_FACTOR_RECEPTOR_ACTIVITY
GOMF_INTERLEUKIN_17_RECEPTOR_ACTIVITY, GOMF_INTERLEUKIN_17_RECEPTOR_ACTIVITY
REACTOME_SYNTHESIS_OF_PG, REACTOME_SYNTHESIS_OF_PG
HP_MICROANGIOPATHIC_HEMOLYTIC_ANEMIA, HP_MICROANGIOPATHIC_HEMOLYTIC_ANEMIA
GOMF_ONCOSTATIN_M_RECEPTOR_ACTIVITY, GOMF_ONCOSTATIN_M_RECEPTOR_ACTIVITY
HP_CONE_SHAPED_EPIPHYSES_OF_THE_PHALANGES_OF_THE_HAND, HP_CONE_SHAPED_EPIPHYSES_OF_THE_PHALANGES_OF_THE_HA
GOBP_FAT_PAD_DEVELOPMENT, GOBP_FAT_PAD_DEVELOPMENT
HP_ABNORMAL_ANTERIOR_CHAMBER_MORPHOLOGY, HP_ABNORMAL_ANTERIOR_CHAMBER_MORPHOLOGY