GSE41867_DAY8_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_UP, GSE41867_DAY8_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_UP GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_180MIN_DN, GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_180MIN_DN GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_UP, GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_UP MODULE_195, MODULE_195 GSE17721_CTRL_VS_CPG_24H_BMDC_DN, GSE17721_CTRL_VS_CPG_24H_BMDC_DN MODULE_576, MODULE_576 MODULE_503, MODULE_503 GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_UP, GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_UP GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP, GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP MIR8076, MIR8076 HP_POOR_EYE_CONTACT, HP_POOR_EYE_CONTACT MIR297, MIR297 HP_MITOCHONDRIAL_MYOPATHY, HP_MITOCHONDRIAL_MYOPATHY MIR6892_5P, MIR6892_5P RAGHAVACHARI_PLATELET_SPECIFIC_GENES, RAGHAVACHARI_PLATELET_SPECIFIC_GENES HP_FOCAL_T2_HYPERINTENSE_BRAINSTEM_LESION, HP_FOCAL_T2_HYPERINTENSE_BRAINSTEM_LESION GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_DN, GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_DN HP_ABNORMALLY_FOLDED_HELIX, HP_ABNORMALLY_FOLDED_HELIX GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_DN HP_FETAL_DISTRESS, HP_FETAL_DISTRESS HP PROXIMAL TUBULOPATHY, HP PROXIMAL TUBULOPATHY XU_HGF_SIGNALING_NOT_VIA_AKT1_48HR_UP, XU_HGF_SIGNALING_NOT_VIA_AKT1_48HR_UP MIR4693_3P, MIR4693_3P HP_DOWN_SLOPING_SHOULDERS, HP_DOWN_SLOPING_SHOULDERS MIR5681B, MIR5681B HP_APLASIA_HYPOPLASIA_INVOLVING_THE_VERTEBRAL_COLUMN, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_VERTEBRAL_COLUMN MIR512_5P, MIR512_5P LIAO_HAVE_SOX4_BINDING_SITES, LIAO_HAVE_SOX4_BINDING_SITES ZNF560_TARGET_GENES, ZNF560_TARGET_GENES MIR1231, MIR1231 PID_WNT_NONCANONICAL_PATHWAY, PID_WNT_NONCANONICAL_PATHWAY GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION, GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION MIR3605_5P, MIR3605_5P MIR509_3P, MIR509_3P GOBP_REGULATION_OF_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY, GOBP_REGULATION_OF_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY GOBP_MAINTENANCE_OF_SISTER_CHROMATID_COHESION, GOBP_MAINTENANCE_OF_SISTER_CHROMATID_COHESION GOCC_LAMELLIPODIUM_MEMBRANE, GOCC_LAMELLIPODIUM_MEMBRANE REACTOME_OTHER_SEMAPHORIN_INTERACTIONS, REACTOME_OTHER_SEMAPHORIN_INTERACTIONS GOBP_POSITIVE_REGULATION_OF_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY, GOBP_POSITIVE_REGULATION_OF_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY GOBP_REGULATION_OF_CHROMATIN_ASSEMBLY, GOBP_REGULATION_OF_CHROMATIN_ASSEMBLY MORF_WNT1, MORF_WNT1 HP ABNORMAL GREATER SCIATIC NOTCH MORPHOLOGY, HP ABNORMAL GREATER SCIATIC NOTCH MORPHOLOGY REACTOME_TRNA_MODIFICATION_IN_THE_MITOCHONDRION, REACTOME_TRNA_MODIFICATION_IN_THE_MITOCHONDRION HP_FEMORAL_BOWING, HP_FEMORAL_BOWING GOBP_RETINOL_METABOLIC_PROCESS, GOBP_RETINOL_METABOLIC_PROCESS MORF_IL9, MORF_IL9 HP_SUPRAVENTRICULAR_TACHYCARDIA, HP_SUPRAVENTRICULAR_TACHYCARDIA GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEOLUS, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEOLUS HP_SPONDYLOMETAPHYSEAL_DYSPLASIA, HP_SPONDYLOMETAPHYSEAL_DYSPLASIA GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE, GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE MORF_MAP3K14, MORF_MAP3K14 HP STRIDOR, HP STRIDOR GOBP_CARTILAGE_MORPHOGENESIS, GOBP_CARTILAGE_MORPHOGENESIS WP_EICOSANOID_SYNTHESIS, WP_EICOSANOID_SYNTHESIS WRN_TARGET_GENES, WRN_TARGET_GENES GOBP_PERICARDIUM_MORPHOGENESIS, GOBP_PERICARDIUM_MORPHOGENESIS HP_ESOPHAGEAL_ATRESIA, HP_ESOPHAGEAL_ATRESIA GOMF_ADRENERGIC_RECEPTOR_ACTIVITY, GOMF_ADRENERGIC_RECEPTOR_ACTIVITY REACTOME_ADRENOCEPTORS, REACTOME_ADRENOCEPTORS

HP_RHIZOMELIC_ARM_SHORTENING, HP_RHIZOMELIC_ARM_SHORTENING

GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN, GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7DY_TOP_100_DEG_EX_VIVO_DN, HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7D GOBP_POSITIVE_REGULATION_OF_DNA_METHYLATION_DEPENDENT_HETEROCHROMATIN_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_DNA_METHYLATION_DEPENDENT_HETEROCHROMATIN_ASSEMBLY