GSE41867_NAIVE_VS_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN, GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN GSE5589_WT_VS_IL6_KO_LPS_STIM_MACROPHAGE_45MIN_UP, GSE5589_WT_VS_IL6_KO_LPS_STIM_MACROPHAGE_45MIN_UP GSE11864_CSF1_IFNG_VS_CSF1_PAM3CYS_IN_MAC_UP, GSE11864_CSF1_IFNG_VS_CSF1_PAM3CYS_IN_MAC_UP GSE22432_CDC_VS_COMMON_DC_PROGENITOR_UP, GSE22432_CDC_VS_COMMON_DC_PROGENITOR_UP GSE27786_LSK_VS_BCELL_UP, GSE27786_LSK_VS_BCELL_UP GSE17580_TREG_VS_TEFF_DN, GSE17580_TREG_VS_TEFF_DN GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN, GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN GSE7852_TREG_VS_TCONV_DN, GSE7852_TREG_VS_TCONV_DN GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP, GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP HP_NARROW_PALATE, HP_NARROW_PALATE MIR1825, MIR1825 ZHENG_CORD_BLOOD_C10_MULTILYMPHOID_PROGENITOR, ZHENG_CORD_BLOOD_C10_MULTILYMPHOID_PROGENITOR HP_MANDIBULAR_PROGNATHIA, HP_MANDIBULAR_PROGNATHIA HP_HIGH_NARROW_PALATE, HP_HIGH_NARROW_PALATE RF4MID_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP, GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP IWANAGA_CARCINOGENESIS_BY_KRAS_UP, IWANAGA_CARCINOGENESIS_BY_KRAS_UP MIR4635, MIR4635 HAY_BONE_MARROW_CD34_POS_PRE_B, HAY_BONE_MARROW_CD34_POS_PRE_B NOTCH_DN.V1_UP, NOTCH_DN.V1_UP MIR4708_3P, MIR4708_3P HP_ABNORMALITY_OF_THE_ILIUM, HP_ABNORMALITY_OF_THE_ILIUM GSE32901_TH17_EMRICHED_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH17_EMRICHED_VS_TH17_NEG_CD4_TCELL_UP GOBP_CHONDROCYTE_DIFFERENTIATION, GOBP_CHONDROCYTE_DIFFERENTIATION GOMF_POLY_A_SPECIFIC_RIBONUCLEASE_ACTIVITY, GOMF_POLY_A_SPECIFIC_RIBONUCLEASE_ACTIVITY BRIDEAU_IMPRINTED_GENES, BRIDEAU_IMPRINTED_GENES KOBAYASHI_RESPONSE_TO_ROMIDEPSIN, KOBAYASHI_RESPONSE_TO_ROMIDEPSIN HP_FLAT_ACETABULAR_ROOF, HP_FLAT_ACETABULAR_ROOF HP_DEFORMED_SELLA_TURCICA, HP_DEFORMED_SELLA_TURCICA VANDESLUIS_COMMD1_TARGETS_GROUP_4_DN, VANDESLUIS_COMMD1_TARGETS_GROUP_4_DN GSE37605 C57BL6_VS_NOD_FOXP3_IRES_GFP_TREG_UP, GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TREG_UP WP_ID_SIGNALING_PATHWAY, WP_ID_SIGNALING_PATHWAY GOBP_SODIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_SODIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE HP_OSTEOPOROSIS_OF_VERTEBRAE, HP_OSTEOPOROSIS_OF_VERTEBRAE HP_ABNORMAL_OSSIFICATION_OF_THE_PUBIC_BONE, HP_ABNORMAL_OSSIFICATION_OF_THE_PUBIC_BONE

GSE41867_DAY6_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP, GSE41867_DAY6_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4MID_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4MID_BCELL_DN GSE41867_DAY6_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN, GSE41867_DAY6_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN, GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN GSE43863_DAY6_EFF_VS_DAY150_MEM_LY6C_INT_CXCR5POS_CD4_TCELL_UP, GSE43863_DAY6_EFF_VS_DAY150_MEM_LY6C_INT_CXCR5POS_CD4_TCELL_UP GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_TS_TIM_PPARG_KO_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_TS_TIM_PPARG_KO_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_TS_TIM_PPARG_KO_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_TS_TIM_PPARG_KO_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_TS_TIM_PPARG_KO_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_TS_TIM_PPARG_KO_MACROPHAGE_DAY10_DN, GSE25123_TS_TIM_PPARG_KO_MACROPHAGE_TS_TIM_PPARG_KO_TS_TIM_PPARG_KO_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_TS_TIM_PPARG_ GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10POS_BONE_MARROW_UP, GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10POS_BONE_MARROW_UP GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_DN, GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_DN GOMF_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY, HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY HP_ABNORMAL_EPIPHYSIS_MORPHOLOGY_OF_THE_PHALANGES_OF_THE_HAND, HP_ABNORMAL_EPIPHYSIS_MORPHOLOGY_OF_THE_PHALANGES_OF_THE_HAND WP_H19_ACTION_RBE2F1_SIGNALING_AND_CDKBETACATENIN_ACTIVITY, WP_H19_ACTION_RBE2F1_SIGNALING_AND_CDKBETACATENIN_ACTIVITY