

EMORY_CD8_TCELL_LCMV_ARMSTRONG_DN,GSE41867_DAY6_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN

GSE41867_NAIVE_VS_DAY30_LCMV_CLONE13_EXHAUSTED_CD8_TCELL_DN,GSE41867_NAIVE_VS_DAY30_LCMV_CLONE13_EXHAUSTED_CD8_TCELL_DN

GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP,GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP

GSE23984_CTRL_VS_HYPOCALEMIC_VITAMIND_ANALOG_TCELL_DN,GSE23984_CTRL_VS_HYPOCALEMIC_VITAMIND_ANALOG_TCELL_DN

GSE41867_NAIVE_VS_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN,GSE41867_NAIVE_VS_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN

GSE2789_DC_VS_CD14C_INT_F480_INT_DC_UP,GSE2789_DC_VS_CD14C_INT_F480_INT_DC_UP

GSE24142_DN2_VS_DN3_THYMOCTE_DN,GSE24142_DN2_VS_DN3_THYMOCTE_DN

GSE270_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_4H_UP,GSE270_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_4H_UP

GSE22611_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_2H_UP,GSE22611_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_2H_UP

GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_4H_DN,GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_4H_DN

GSE23502_BM_VS_COLON_TUMOR_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_DN,GSE23502_BM_VS_COLON_TUMOR_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_DN

GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCTE_DN,GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCTE_DN

GSE23432_CDC_VS_COMMON_DC_PROGENITOR_UP,GSE23432_CDC_VS_COMMON_DC_PROGENITOR_UP

GSE3021_UNSTIM_OH_VS_TLR2_2_STIM_MONOCYTE_24H_DN,GSE3021_UNSTIM_OH_VS_TLR2_2_STIM_MONOCYTE_24H_DN

GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP,GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP

GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCTYES_UP,GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCTYES_UP

GSE37301_CD4_TCELL_VS_RAG2_KO_NK_CELL_DN,GSE37301_CD4_TCELL_VS_RAG2_KO_NK_CELL_DN

GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_UP,GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_UP

GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP,GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP

GSE21380_NON_TFH_VS_TFH_CD4_TCELL_UP,GSE21380_NON_TFH_VS_TFH_CD4_TCELL_UP

GSE21063_CTRL_VS_ANTI_IJGM_STIM_BCELL_NFATC1_KO_8H_UP,GSE21063_CTRL_VS_ANTI_IJGM_STIM_BCELL_NFATC1_KO_8H_UP

GSE26077_MPL_VS_R848_STIM_BCELL_DN,GSE26077_MPL_VS_R848_STIM_BCELL_DN

GSE26088_UNTREATED_VS_IL2_TREATED_TCELL_2H_UP,GSE26088_UNTREATED_VS_IL2_TREATED_TCELL_2H_UP

GSE14308_TH17_VS_INDUCED_TREG_UP,GSE14308_TH17_VS_INDUCED_TREG_UP

GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_HET_TCONV_DN,GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_HET_TCONV_DN

GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_4D_CULTURE_UP,GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_4D_CULTURE_UP

GSE41176_UNSTIM_VS_ANTI_IJGM_STIM_TAK1_KO_BCELL_24H_UP,GSE41176_UNSTIM_VS_ANTI_IJGM_STIM_TAK1_KO_BCELL_24H_UP

GSE17580_UNINFECTED_VS_S_MANSONI_INF_TREG_UP,GSE17580_UNINFECTED_VS_S_MANSONI_INF_TREG_UP

GSE7852_TREG_VS_TCONV_UP,GSE7852_TREG_VS_TCONV_UP

GSE309_B2_VS_B1_BCELL_UP,GSE309_B2_VS_B1_BCELL_UP

GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP,GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP

GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_DN,GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_DN

GSE3039_ALPHABETA_CD8_TCELL_VS_B2_BCELL_UP,GSE3039_ALPHABETA_CD8_TCELL_VS_B2_BCELL_UP

GSE15330 GRANULOCYTE_MONOCYTE_PROGENITOR_VS_PRO_BCELL_DN,GSE15330 GRANULOCYTE_MONOCYTE_PROGENITOR_VS_PRO_BCELL_DN

GSE23505_UNTREATED_VS_4DAY_IL6_IL1_IL23_TREATED_CD4_TCELL_UP,GSE23505_UNTREATED_VS_4DAY_IL6_IL1_IL23_TREATED_CD4_TCELL_UP

GSE23502_WT_VS_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_BM_UP,GSE23502_WT_VS_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_BM_UP

GSE24671_CTRL_VS_SENDALVIRUS_INFECTED_MOUSE_SPLENOCTYES_DN,GSE24671_CTRL_VS_SENDALVIRUS_INFECTED_MOUSE_SPLENOCTYES_DN

GSE309_SOCS3_KO_VS_WT_LIVER_NKT_IL6_INJECTION_DN,GSE309_SOCS3_KO_VS_WT_LIVER_NKT_IL6_INJECTION_DN

GSE26168_PDC_VS_MDC_DAY7_FLU_VACCINE_UP,GSE26168_PDC_VS_MDC_DAY7_FLU_VACCINE_UP

GSE3039_NKT_CELL_VS_B2_BCELL_UP,GSE3039_NKT_CELL_VS_B2_BCELL_UP

GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_UP,GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_UP

GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCTE_FETAL_DN,GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCTE_FETAL_DN

GSE7460_CTRL_VS_TGFB_TREATED_ACT_TCONV_DN,GSE7460_CTRL_VS_TGFB_TREATED_ACT_TCONV_DN

GSE26168_MONOCYTE_VS_PDC_DN,GSE26168_MONOCYTE_VS_PDC_DN

GSE8685_IL2_STARVED_VS_IL2_ACT_IL2_STARVED_CD4_TCELL_UP,GSE8685_IL2_STARVED_VS_IL2_ACT_IL2_STARVED_CD4_TCELL_UP

GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_NK_NFkB_KO_DC_UP,GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_NK_NFkB_KO_DC_UP

GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_UP,GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_UP

GSE14308_TH2_VS_NAIVE_CD4_TCELL_DN,GSE14308_TH2_VS_NAIVE_CD4_TCELL_DN

GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_UP,GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_UP

GSE17721_LPS_VS_POLYIC_24H_BMDC_DN,GSE17721_LPS_VS_POLYIC_24H_BMDC_DN

GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_UP,GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_UP

TRAVAGLINI_LUNG_B_CELL,TRAVAGLINI_LUNG_B_CELL

GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_DN,GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_DN

GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP,GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP

GSE38304_MYC_NEG_VS_POS_GC_BCELL_UP,GSE38304_MYC_NEG_VS_POS_GC_BCELL_UP

CTACTGT_MIR199A,CTACTGT_MIR199A

GSE27786_CD8_TCELL_VS_NKTCCELL_DN,GSE27786_CD8_TCELL_VS_NKTCCELL_DN

GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN,GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN

GSE17721_POLYIC_VS_CPG_8H_BMDC_UP,GSE17721_POLYIC_VS_CPG_8H_BMDC_UP

GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN,GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN

ATACCTGT_MIR144,ATACCTGT_MIR144

GSE1047_IL3_VS_IL3_AND_CPG_STIM_PDC_DN,GSE1047_IL3_VS_IL3_AND_CPG_STIM_PDC_DN

GOCC_RECYCLING_ENDOSOME,GOCC_RECYCLING_ENDOSOME

GSE10240_IL22_VS_IL22_AND_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN,GSE10240_IL22_VS_IL22_AND_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN

GOBP_REGULATION_OF_RAS_PROTEIN_SIGNAL_TRANSDUCTION,GOBP_REGULATION_OF_RAS_PROTEIN_SIGNAL_TRANSDUCTION

HAY_BONE_MARROW_FOLLICULAR_B_CELL,HAY_BONE_MARROW_FOLLICULAR_B_CELL

GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_DN,GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_DN

GACAATC_MIR219,GACAATC_MIR219

GSE24671_CTRL_VS_SENDALVIRUS_INFECTED_MOUSE_SPLENOCTYES_UP,GSE24671_CTRL_VS_SENDALVIRUS_INFECTED_MOUSE_SPLENOCTYES_UP

GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN,GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN

PID_MET_PATHWAY,PID_MET_PATHWAY

GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP,GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP

GOMF_NUCLEAR_RECEPTOR_BINDING,GOMF_NUCLEAR_RECEPTOR_BINDING

GSE40274_CTRL_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN,GSE40274_CTRL_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN

GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN,GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN

GSE19401_UNSTIM_VS_PAM2CSK4_STIM_FOLLICULAR_DC_UP,GSE19401_UNSTIM_VS_PAM2CSK4_STIM_FOLLICULAR_DC_UP

GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP,GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP

GOCC_RUFFLE_MEMBRANE,GOCC_RUFFLE_MEMBRANE

MIR1179,MIR1179

MIR130B_5P,MIR130B_5P

MIR3688_5P,MIR3688_5P

GOMF_HORMONE_RECEPTOR_BINDING,GOMF_HORMONE_RECEPTOR_BINDING

REN_ALVEOLAR_RHABDOMYOSARCOMA_UP,REN_ALVEOLAR_RHABDOMYOSARCOMA_UP

GOMF_STEROID_HORMONE_RECEPTOR_BINDING,GOMF_STEROID_HORMONE_RECEPTOR_BINDING

MIR190A_5P,MIR190A_5P

GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP,GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP

GOCC_HISTONE_DEACETYLASE_COMPLEX,GOCC_HISTONE_DEACETYLASE_COMPLEX

GSE360_CTRL_VS_B_MALAYILOW_DOSE_DC_UP,GSE360_CTRL_VS_B_MALAYILOW_DOSE_DC_UP

GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_CD8_TCELL_UP,GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_CD8_TCELL_UP

GSE18281_CORTX_VS_MEDULLA_THYMUS_UP,GSE18281_CORTX_VS_MEDULLA_THYMUS_UP

MIR4473,MIR4473

GSE5542_IFNG_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_DN,GSE5542_IFNG_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_DN

GSE21676_UNTREATED_VS_IL6_TREATED_CD4_TCELL_UP,GSE21676_UNTREATED_VS_IL6_TREATED_CD4_TCELL_UP

GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION,GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION

GOBP_ESTABLISHMENT_OF_CELL_POLARITY,GOBP_ESTABLISHMENT_OF_CELL_POLARITY

GSE6269_STAPH_AUREUS_VS_STREP_PNEUMO_INF_PBMC_DN,GSE6269_STAPH_AUREUS_VS_STREP_PNEUMO_INF_PBMC_DN

MIR942_3P,MIR942_3P

GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_DN,GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_DN

GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP,GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP

GSE32333_MIR17_KO_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_DN,GSE32333_MIR17_KO_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_DN

GSE589_LPS_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_UP,GSE589_LPS_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_UP

GSE7831_IL1_VS_4H_INFLUENZA_STIM_PDC_DN,GSE7831_IL1_VS_4H_INFLUENZA_STIM_PDC_DN

MIR885_5P,MIR885_5P

GCCNNNWTAAAR_UNKNOWN,GCCNNNWTAAAR_UNKNOWN

GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_8D_CULTURE_UP,GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_8D_CULTURE_UP

TRAVAGLINI_LUNG_PLASMACYTOID_DENDRITIC_CELL,TRAVAGLINI_LUNG_PLASMACYTOID_DENDRITIC_CELL

GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN,GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN

PLASARI_TGFB1_SIGNALING_VIA_NFIC_3HR_DN,PLASARI_TGFB1_SIGNALING_VIA_NFIC_3HR_DN

TAGCTCA_MIR192,MIR215,TAGGTCA_MIR192,MIR215

MIR873_5P,MIR873_5P

GSE22342_CD11C_HIGH_VS_LOW_DECIDUAL_MACROPHAGES_UP,GSE22342_CD11C_HIGH_VS_LOW_DECIDUAL_MACROPHAGES_UP

GOBP_ESTABLISHMENT_OF_ENDOTHELIAL_BARRIER,GOBP_ESTABLISHMENT_OF_ENDOTHELIAL_BARRIER

MIR203A_5P,MIR203A_5P

MIR3156_5P,MIR3156_5P

GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MACROPHAGE_UP,GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MACROPHAGE_UP

MIR6829_3P,MIR6829_3P

MIR6791_3P,MIR6791_3P

MODULE_245,MODULE_243

GSE16385_ROSLIGLITAZONE_IL4_VS_ROSLIGLITAZONE_ALONE_STIM_MACROPHAGE_UP,GSE16385_ROSLIGLITAZONE_IL4_VS_ROSLIGLITAZONE_ALONE_STIM_MACROPHAGE_UP

MIR4689,MIR4689

MIR6758_5P,MIR6758_5P

GOBP_POSITIVE_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING,GOBP_POSITIVE_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING

MIR3194_5P,MIR3194_5P

GOBP_ENDOTHELIAL_CELL_DEVELOPMENT,GOBP_ENDOTHELIAL_CELL_DEVELOPMENT

GOBP_PROTEIN_LOCALIZATION_TO_ENDOSOME,GOBP_PROTEIN_LOCALIZATION_TO_ENDOSOME

MIR20B_3P,MIR20B_3P

MIR6856_5P,MIR6856_5P

MIR4691_3P,MIR4691_3P

GSE29614_CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMC_UP,GSE29614_CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMC_UP

GOBP_POSITIVE_REGULATION_OF_COAGULATION,GOBP_POSITIVE_REGULATION_OF_COAGULATION

MIR1246,MIR1246

GSE28130_ACTIVATED_VS_INDUCED_TREG_DN,GSE28130_ACTIVATED_VS_INDUCED_TREG_DN

BAELDE_DIABETIC_NEPHROPATHY_UP,BAELDE_DIABETIC_NEPHROPATHY_UP

PID_NECTIN_PATHWAY,PID_NECTIN_PATHWAY

GOCC_PROTEIN_COMPLEX_INVOLVED_IN_CELL_ADHESION,GOCC_PROTEIN_COMPLEX_INVOLVED_IN_CELL_ADHESION

PID_INTEGRIN_CS_PATHWAY,PID_INTEGRIN_CS_PATHWAY

GOCC_9PLUS2_MOTILE_CILIUM,GOCC_9PLUS2_MOTILE_CILIUM

AMIT_EGF_RESPONSE_120_MCF10A,AMIT_EGF_RESPONSE_120_MCF10A

MIR4721,MIR4721

GOBP TRABECULA MORPHOGENESIS,GOBP TRABECULA MORPHOGENESIS

ZNF502_TARGET_GENES,ZNF502_TARGET_GENES

GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION,GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION

GCTGAGT_MIR5125P,GCTGAGT_MIR5125P

GOBP_RAP_PROTEIN_SIGNAL_TRANSDUCTION,GOBP_RAP_PROTEIN_SIGNAL_TRANSDUCTION

GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING,GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING

GOBP_CELLULAR_RESPONSE_TO_CAMP,GOBP_CELLULAR_RESPONSE_TO_CAMP

HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_VESTIBULE_OF_THE_INNER_EAR,HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_VESTIBULE_OF_THE_INNER_EAR

GOBP_CELLULAR_COMPONENT_MAINTENANCE,GOBP_CELLULAR_COMPONENT_MAINTENANCE

GOBP_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY,GOBP_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY

GOBP_RUFFLE_ORGANIZATION,GOBP_RUFFLE_ORGANIZATION

REACTOME_HEME_SIGNALING,REACTOME_HEME_SIGNALING

ABE_VEGFA_TARGETS_2HR,ABE_VEGFA_TARGETS_2HR

GOBP_CELLULAR_RESPONSE_TO_CALCIIUM_ION,GOBP_CELLULAR_RESPONSE_TO_CALCIIUM_ION

GOBP_FILOPODIUM_ASSEMBLY,GOBP_FILOPODIUM_ASSEMBLY

GOBP_REGULATION_OF_ERAD_PATHWAY,GOBP_REGULATION_OF_ERAD_PATHWAY

HULMAPK14_TARGETS_UP,HULMAPK14_TARGETS_UP

HOLLEMAN_DAUORUBICIN_B_ALL_DN,HOLLEMAN_DAUORUBICIN_B_ALL_DN

BOYAUUT_LIVER_CANCER_SUBCLASS_G23_DN,BOYAUUT_LIVER_CANCER_SUBCLASS_G23_DN

VALK_AML_CLUSTER_10,VALK_AML_CLUSTER_10

HP_ECTOPIA_LENTIS,HP_ECTOPIA_LENTIS

MIR4658,MIR4658

STEIN_ESTROGEN_RESPONSE_NOT_VIA_ESRRA,STEIN_ESTROGEN_RESPONSE_NOT_VIA_ESRRA

MIR3150A_3P,MIR3150A_3P

GOBP_NEUROTROPHIN_SIGNALING_PATHWAY,GOBP_NEUROTROPHIN_SIGNALING_PATHWAY

GOCC_PLATELET_DENSE_TUBULAR_NETWORK,GOCC_PLATELET_DENSE_TUBULAR_NETWORK

GOMF_ACTIN_DEPENDENT_ATPASE_ACTIVITY,GOMF_ACTIN_DEPENDENT_ATPASE_ACTIVITY

GOMF_RETINOID_X_RECEPTOR_BINDING,GOMF_RETINOID_X_RECEPTOR_BINDING

MIR3681_5P,MIR3681_5P

MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_23,MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_23

GOBP_NEGATIVE_REGULATION_OF_BMP_SIGNALING_PATHWAY,GOBP_NEGATIVE_REGULATION_OF_BMP_SIGNALING_PATHWAY

SHIN_B_CELL_LYMPHOMA_CLUSTER_9,SHIN_B_CELL_LYMPHOMA_CLUSTER_9

LI_PROSTATE_CANCER_EPIGENETIC,LI_PROSTATE_CANCER_EPIGENETIC

WP_METABOLISM_OF_SPINGOLIPIDS_IN_ER_AND_GOLGIAPPARATUS,WP_METABOLISM_OF_SPINGOLIPIDS_IN_ER_AND_GOLGIAPPARATUS

GOBP_OLFACTORY_LOBE_DEVELOPMENT,GOBP_OLFACTORY_LOBE_DEVELOPMENT

MIR6849_5P,MIR6849_5P

MIR8074,MIR8074

GOBP_NEGATIVE_REGULATION_OF_MYTUBE_DIFFERENTIATION,GOBP_NEGATIVE_REGULATION_OF_MYTUBE_DIFFERENTIATION

BIOCARTA_PRION_PATHWAY,BIOCARTA_PRION_PATHWAY

MIR451A,MIR451A

GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_5_PHOSPHATASE_ACTIVITY,GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_5_PHOSPHATASE_ACTIVITY

GOMF_MICROFILAMENT_MOTOR_ACTIVITY,GOMF_MICROFILAMENT_MOTOR_ACTIVITY

GOBP_POSITIVE_REGULATION_OF_LIPASE_ACTIVITY,GOBP_POSITIVE_REGULATION_OF_LIPASE_ACTIVITY

GOBP_POSITIVE_REGULATION_OF_OSTEOSTBLAST_PROLIFERATION,GOBP_POSITIVE_REGULATION_OF_OSTEOSTBLAST_PROLIFERATION

GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION,GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION

GOBP_POSITIVE_REGULATION_OF_HAIR_FOLLICLE_DEVELOPMENT,GOBP_POSITIVE_REGULATION_OF_HAIR_FOLLICLE_DEVELOPMENT

HP_URETHRAL_VALVE,HP_URETHRAL_VALVE

GOBP_BUD_ELONGATION_INVOLVED_IN_LUNG_BRANCHING,GOBP_BUD_ELONGATION_INVOLVED_IN_LUNG_BRANCHING

GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_GROWTH_FACTOR_STIMULUS,GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_GROWTH_FACTOR_STIMULUS

HP_PHIMOSIS,HP_PHIMOSIS

WP_ETHANOL_METABOLISM_RESULTING_IN_PRODUCTION_OF_ROS_BY_CYP2E1,WP_ETHANOL_METABOLISM_RESULTING_IN_PRODUCTION_OF_ROS_BY_CYP2E1

GOMF_CO_RECEPTOR_BINDING,GOMF_CO_RECEPTOR_BINDING

GOBP_CELL_JUNCTION_MAINTENANCE,GOBP_CELL_JUNCTION_MAINTENANCE

HP_LENS_SUBLUXATION,HP_LENS_SUBLUXATION

GOBP_TANGENTIAL_MIGRATION_FROM_THE_SUBVENTRICULAR_ZONE_TO_THE_OLFACTORY_BULB,GOBP_TANGENTIAL_MIGRATION_FROM_THE_SUBVENTRICULAR_ZONE_TO_THE_OLFACTORY_BULB

HP_ABNORMALITY_OF_THE_MIDDLE_PHALANX_OF_THE_2ND_FINGER,HP_ABNORMALITY_OF_THE_MIDDLE_PHALANX_OF_THE_2ND_FINGER

GOMF_CARD_DOMAIN_BINDING,GOMF_CARD_DOMAIN_BINDING

HP_ABNORMAL_ARM_SPAN,HP_ABNORMAL_ARM_SPAN

GOBP_PSEUDOPODIUM_ORGANIZATION,GOBP_PSEUDOPODIUM_ORGANIZATION

chr5q33,chr5q33

GOMF_CALCIIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY,GOMF_CALCIIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY

HP_GONADOTROPIN_DEFICIENCY,HP_GONADOTROPIN_DEFICIENCY

HP_HYPOPLASTIC_ANEMIA,HP_HYPOPLASTIC_ANEMIA

HP_ABNORMALITY_OF_THE_2ND_METACARPAL,HP_ABNORMALITY_OF_THE_2ND_METACARPAL

GOBP_RENAL_WATER_TRANSPORT,GOBP_RENAL_WATER_TRANSPORT

GOCC_OUTER_DENSE_FIBER,GOCC_OUTER_DENSE_FIBER

GOBP_NOSE_DEVELOPMENT,GOBP_NOSE_DEVELOPMENT

GOBP_RESPONSE_TO_CGMP,GOBP_RESPONSE_TO_CGMP

GOMF_GALACTOSIDE_BINDING,GOMF_GALACTOSIDE_BINDING

TESAR_JAK_TARGETS_MOUSE_ES_D3_DN,TESAR_JAK_TARGETS_MOUSE_ES_D3_DN

GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY,GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY

GOBP_RESPONSE_TO_LITHIUM_ION,GOBP_RESPONSE_TO_LITHIUM_ION

HP_SHORT_MIDDLE_PHALANX_OF_THE_2ND_FINGER,HP_SHORT_MIDDLE_PHALANX_OF_THE_2ND_FINGER

HP_APLASIA_HYPOPLASIA_OF_THE_2ND_METACARPAL,HP_APLASIA_HYPOPLASIA_OF_THE_2ND_METACARPAL

REACTOME_RUNX2_REGULATES_CHONDRROCYTE_MATURATION,REACTOME_RUNX2_REGULATES_CHONDRROCYTE_MATURATION

GOBP_POSITIVE_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY,GOBP_POSITIVE_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY

GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_OXIDATIVE_STRESS,GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_OXIDATIVE_STRESS

GOBP_ESTABLISHMENT_OF_MEIOTIC_SPINDLE_LOCALIZATION,GOBP_ESTABLISHMENT_OF_MEIOTIC_SPINDLE_LOCALIZATION

PBRM1_TARGET_GENES,PBRM1_TARGET_GENES

HP_INCREASED_ADIPOSE_TISSUE,HP_INCREASED_ADIPOSE_TISSUE

GOBP_REGULATION_OF_RENAL_SYSTEM_PROCESS,GOBP_REGULATION_OF_RENAL_SYSTEM_PROCESS

GOBP_NEGATIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION,GOBP_NEGATIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION

GOBP_NEGATIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING,GOBP_NEGATIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING

GOBP_REGULATION_OF_CGMP_MEDIATED_SIGNALING,GOBP_REGULATION_OF_CGMP_MEDIATED_SIGNALING

HP_LOP_EAR,HP_LOP_EAR

HP_ABNORMAL_OSSIFICATION_OF_THE_PUBIC_BONE,HP_ABNORMAL_OSSIFICATION_OF_THE_PUBIC_BONE

GOBP_NEGATIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY_INVOLVED_IN_APOPTOTIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY_INVOLVED_IN_APOPTOTIC_PROCESS

MIR4749_5P,MIR4749_5P

chr2p21,chr2p21