

| | | |
|--|--|--|
| | GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP, GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP GSE41867_NAIVE_VS_DAY6_LCMV_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY6_LCMV_EFFECTOR_CD8_TCELL_DN GSE14308_INDUCED_VS_NATURAL_TREG_DN, GSE14308_INDUCED_VS_NATURAL_TREG_DN GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_DN, GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_DN GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCYTES_UP, GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCYTES_UP GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_DN, GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_DN GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_DN, GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_DN GSE32164_ALTERNATIVELY_ACT_M2_VS_CMYC_INHIBITED_MACROPHAGE_UP, GSE32164_ALTERNATIVELY_ACT_M2_VS_CMYC_INHIBITED_MACROPHAGE_UP GSE14350_TREG_VS_TEFF_IN_IL2RB_KO_UP, GSE14350_TREG_VS_TEFF_IN_IL2RB_KO_UP GSE44649_WT_VS_MIR155_KO_ACTIVATED_CD8_TCELL_UP, GSE44649_WT_VS_MIR155_KO_ACTIVATED_CD8_TCELL_UP GSE43863_TH1_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN, GSE43863_TH1_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN GSE27786_LIN_NEG_VS_NKTCCELL_DN, GSE27786_LIN_NEG_VS_NKTCCELL_DN GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFB_TCONV_DN, GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFB_TCONV_DN GSE22601_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_DN, GSE22601_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_DN GSE43863_TH1_VS_TFH_EFFECTOR_CD4_TCELL_DN, GSE43863_TH1_VS_TFH_EFFECTOR_CD4_TCELL_DN GSE7568_IL4_VS_IL4_AND_TGFB_TREATED_MACROPHAGE_24H_UP, GSE7568_IL4_VS_IL4_AND_TGFB_TREATED_MACROPHAGE_24H_UP GSE41867_NAIVE_VS_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN GSE40068_BCL6_POS_VS_NEG_CXCR5_POS_TFH_DN, GSE40068_BCL6_POS_VS_NEG_CXCR5_POS_TFH_DN GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_12H_DN, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_12H_DN GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_12H_DN, GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_12H_DN GSE23568_ID3_TRANSDUCED_VS_ID3_KO_CD8_TCELL_DN, GSE23568_ID3_TRANSDUCED_VS_ID3_KO_CD8_TCELL_DN GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_UP, GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_UP GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP, GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP GSE3691_IFN_PRODUCING_KILLER_DC_VS_PLASMACYTOID_DC_SPLEEN_UP, GSE3691_IFN_PRODUCING_KILLER_DC_VS_PLASMACYTOID_DC_SPLEEN_UP GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_INF_DN, GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_INF_DN GSE7852_TREG_VS_TCONV_LN_UP, GSE7852_TREG_VS_TCONV_LN_UP GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_ATOPIC_PATIENT_DN, GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_ATOPIC_PATIENT_DN GSE40274_FOXP3_VS_FOXP3_AND_PBX1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_PBX1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN GSE15324_NAIVE_VS_ACTIVATED_ELF4_KO_CD8_TCELL_UP, GSE15324_NAIVE_VS_ACTIVATED_ELF4_KO_CD8_TCELL_UP GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_12H_IFNA_STIM_UP, GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_12H_IFNA_STIM_UP GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN GSE14699_NAIVE_VS_ACT_CD8_TCELL_UP, GSE14699_NAIVE_VS_ACT_CD8_TCELL_UP GSE27786_CD4_VS_CD8_TCELL_UP, GSE27786_CD4_VS_CD8_TCELL_UP GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY6_POST_IMMUNIZATION_DN, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY6_POST_IMMUNIZATION_DN GSE20366_TREG_VS_NAIVE_CD4_TCELL_HOMEOSTATIC_CONVERSION_DN, GSE20366_TREG_VS_NAIVE_CD4_TCELL_HOMEOSTATIC_CONVERSION_DN GSE37533_PPARG1_FOXP3_VS_PPARG2_FOXP3_TRANSDUCED_CD4_TCELL_DN, GSE37533_PPARG1_FOXP3_VS_PPARG2_FOXP3_TRANSDUCED_CD4_TCELL_DN GCATTGT_MIR105, GCATTGT_MIR105 GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP, GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_CTRL_CD4_TCELL_UP, GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_CTRL_CD4_TCELL_UP GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_TRICHOSTATINA_CD8_TCELL_UP, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_TRICHOSTATINA_CD8_TCELL_UP GSE40274_GATA1_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_GATA1_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP GSE4590_PRE_BCELL_VS_LARGE_PRE_BCELL_DN, GSE4590_PRE_BCELL_VS_LARGE_PRE_BCELL_DN GSE17580_UNINFECTED_VS_S_MANSONI_INF_TREG_UP, GSE17580_UNINFECTED_VS_S_MANSONI_INF_TREG_UP GSE21670_IL6_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_IL6_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_DN GSE37301_PRO_BCELL_VS_CD4_TCELL_UP, GSE37301_PRO_BCELL_VS_CD4_TCELL_UP GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC_UP, GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC_UP GSE7460_TREG_VS_TCONV_ACT_UP, GSE7460_TREG_VS_TCONV_ACT_UP GSE24142_DN2_VS_DN3_THYMOCYTE_DN, GSE24142_DN2_VS_DN3_THYMOCYTE_DN GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_FETAL_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_FETAL_DN WP_INSULIN_SIGNALING, WP_INSULIN_SIGNALING GSE14699_NAIVE_VS_ACT_CD8_TCELL_DN, GSE14699_NAIVE_VS_ACT_CD8_TCELL_DN GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN GSE13547_WT_VS_ZFX_KO_BCELL_UP, GSE13547_WT_VS_ZFX_KO_BCELL_UP GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_UP GSE40685_NAIVE_CD4_TCELL_VS_TREG_DN, GSE40685_NAIVE_CD4_TCELL_VS_TREG_DN GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_IL12_CD8_TCELL_UP, GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_IL12_CD8_TCELL_UP HALLMARK_UV_RESPONSE_DN, HALLMARK_UV_RESPONSE_DN GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP GSE13946_CTRL_VS_DSS_COLITIS_GD_TCELL_FROM_COLON_DN, GSE13946_CTRL_VS_DSS_COLITIS_GD_TCELL_FROM_COLON_DN MIR1184, MIR1184 GSE7852_THYMUS_VS_FAT_TREG_DN, GSE7852_THYMUS_VS_FAT_TREG_DN GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP MIR4426, MIR4426 GSE21033_CTRL_VS_POLYIC_STIM_DC_6H_DN, GSE21033_CTRL_VS_POLYIC_STIM_DC_6H_DN GSE14699_NAIVE_VS_DELETIONAL_TOLERANCE_CD8_TCELL_UP, GSE14699_NAIVE_VS_DELETIONAL_TOLERANCE_CD8_TCELL_UP GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_UP, GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_UP GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_DN GSE13547_CTRL_VS_ANTLIGM_STIM_ZFX_KO_BCELL_12H_DN, GSE13547_CTRL_VS_ANTLIGM_STIM_ZFX_KO_BCELL_12H_DN MIR624_5P, MIR624_5P TBX1_TARGET_GENES, TBX1_TARGET_GENES GSE14415_ACT_TCONV_VS_ACT_NATURAL_TREG_UP, GSE14415_ACT_TCONV_VS_ACT_NATURAL_TREG_UP HP_SELF_INJURIOUS_BEHAVIOR, HP_SELF_INJURIOUS_BEHAVIOR THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_DN, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_DN MIR630, MIR630 GOBP_DENDRITE_MORPHOGENESIS, GOBP_DENDRITE_MORPHOGENESIS HP_ROUND_FACE, HP_ROUND_FACE MIR3977, MIR3977 GSE2935_UV_INACTIVATED_VS_LIVE_SENDAL_VIRUS_INF_MACROPHAGE_UP, GSE2935_UV_INACTIVATED_VS_LIVE_SENDAL_VIRUS_INF_MACROPHAGE_UP REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX, REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_UP, GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_UP MIR1244, MIR1244 KEGG_CHRONIC_MYELOID_LEUKEMIA, KEGG_CHRONIC_MYELOID_LEUKEMIA MIR4782_3P, MIR4782_3P MIR219A_5P, MIR219A_5P MIR6766_3P, MIR6766_3P GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_UP GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_KO_UP, GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_KO_UP GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_CD8_TCELL_UP, GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_CD8_TCELL_UP HP_GENERALIZED_HIRSUTISM, HP_GENERALIZED_HIRSUTISM MIR1279, MIR1279 MIR3667_3P, MIR3667_3P MIR193A_3P, MIR193B_3P, MIR193A_3P, MIR193B_3P GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN ACGTCCT_MIR28, ACGTCCT_MIR28 IRF2_01, IRF2_01 MIR548AT_3P, MIR548AT_3P HP_ABNORMAL_CLAVICLE_MORPHOLOGY, HP_ABNORMAL_CLAVICLE_MORPHOLOGY GSE13306_TREG_VS_TCONV_SPLEEN_UP, GSE13306_TREG_VS_TCONV_SPLEEN_UP MIR548AY_3P, MIR548AY_3P MIR4519, MIR4519 MIR3194_3P, MIR3194_3P MIR3678_3P, MIR3678_3P HP_FOCAL_IMPAIRED_AWARENESS_SEIZURE, HP_FOCAL_IMPAIRED_AWARENESS_SEIZURE MIR642B_5P, MIR642B_5P MIR4786_5P, MIR4786_5P MIR145_3P, MIR145_3P HP_NEOPLASM_OF_THE_SKELETAL_SYSTEM, HP_NEOPLASM_OF_THE_SKELETAL_SYSTEM MIR4691_3P, MIR4691_3P DR3_Q4, DR3_Q4 MIR340_3P, MIR340_3P MIR1324, MIR1324 WP_TCELL_ANTIGEN_RECEPTOR_TCR_SIGNALING_PATHWAY, WP_TCELL_ANTIGEN_RECEPTOR_TCR_SIGNALING_PATHWAY HP_SHORT_PALM, HP_SHORT_PALM HP_LYMPHEDEMA, HP_LYMPHEDEMA MIR4726_5P, MIR4726_5P MIR10401_5P, MIR10401_5P MIR1263, MIR1263 MIR3187_3P, MIR3187_3P KEGG_GNRH_SIGNALING_PATHWAY, KEGG_GNRH_SIGNALING_PATHWAY MIR6782_3P, MIR6782_3P HP_THICKENED_NUCHAL_SKIN_FOLD, HP_THICKENED_NUCHAL_SKIN_FOLD MIR6894_3P, MIR6894_3P GOMF_PRIMARY_ACTIVE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PRIMARY_ACTIVE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY MIR4640_5P, MIR4640_5P TCTATGA_MIR376A_MIR376B, TCTATGA_MIR376A_MIR376B GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY GOBP_REGULATION_OF_CALCIIUM_ION_TRANSMEMBRANE_TRANSPORT, GOBP_REGULATION_OF_CALCIIUM_ION_TRANSMEMBRANE_TRANSPORT MIR3654, MIR3654 GOBP_NEGATIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY REACTOME_FLT3_SIGNALING_IN_DISEASE, REACTOME_FLT3_SIGNALING_IN_DISEASE MIR3976, MIR3976 GOBP_POSITIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION, GOBP_POSITIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION GOMF_CHANNEL_REGULATOR_ACTIVITY, GOMF_CHANNEL_REGULATOR_ACTIVITY HP_TONIC_SEIZURE, HP_TONIC_SEIZURE GOBP_FACE_DEVELOPMENT, GOBP_FACE_DEVELOPMENT GOMF_ION_CHANNEL_REGULATOR_ACTIVITY, GOMF_ION_CHANNEL_REGULATOR_ACTIVITY GOBP_LYMPHOCYTE_MIGRATION, GOBP_LYMPHOCYTE_MIGRATION GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ORGANIZATION, GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ORGANIZATION MIR6754_5P, MIR6754_5P GOBP_NEGATIVE_REGULATION_OF_HISTONE_MODIFICATION, GOBP_NEGATIVE_REGULATION_OF_HISTONE_MODIFICATION GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_2_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_2_PRODUCTION HP_FOCAL_NON_MOTOR_SEIZURE, HP_FOCAL_NON_MOTOR_SEIZURE HP_PROLONGED_BLEEDING_TIME, HP_PROLONGED_BLEEDING_TIME GOCC_SWI_SNF_COMPLEX, GOCC_SWI_SNF_COMPLEX GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN GOBP_BODY_MORPHOGENESIS, GOBP_BODY_MORPHOGENESIS HP_ABNORMAL_MORPHOLOGY_OF_THE_LIMBIC_SYSTEM, HP_ABNORMAL_MORPHOLOGY_OF_THE_LIMBIC_SYSTEM KEGG_AMYOTROPHIC_LATERAL_SCLEROSIS_ALS, KEGG_AMYOTROPHIC_LATERAL_SCLEROSIS_ALS MIR4323, MIR4323 WP_HIPPOYAP_SIGNALING_PATHWAY, WP_HIPPOYAP_SIGNALING_PATHWAY TCCCCAC_MIR491, TCCCCAC_MIR491 KEGG_GLYCEROPHOSPHOLIPID_METABOLISM, KEGG_GLYCEROPHOSPHOLIPID_METABOLISM MIR3137, MIR3137 HP_ABNORMALITY_OF_THE_GASTRIC_MUCOSA, HP_ABNORMALITY_OF_THE_GASTRIC_MUCOSA GOBP_INTERLEUKIN_2_PRODUCTION, GOBP_INTERLEUKIN_2_PRODUCTION WP_BDNFTRKB_SIGNALING, WP_BDNFTRKB_SIGNALING MIR2909, MIR2909 GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_3H_IFNG_STIM_DN, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_3H_IFNG_STIM_DN GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN HU_GENOTOXIN_ACTION_DIRECT_VS_INDIRECT_4HR, HU_GENOTOXIN_ACTION_DIRECT_VS_INDIRECT_4HR HP_EYELID_MYOCLOIUS, HP_EYELID_MYOCLOIUS MIR711, MIR711 GOMF_POTASSIUM_CHANNEL_REGULATOR_ACTIVITY, GOMF_POTASSIUM_CHANNEL_REGULATOR_ACTIVITY WANG_RESPONSE_TO_ANDROGEN_UP, WANG_RESPONSE_TO_ANDROGEN_UP HP_TYPICAL_ABSENCE_SEIZURE, HP_TYPICAL_ABSENCE_SEIZURE GOMF_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING, GOMF_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING REACTOME_REGULATION_OF_SIGNALING_BY_CBL, REACTOME_REGULATION_OF_SIGNALING_BY_CBL GOBP_ACTIVATED_T_CELL_PROLIFERATION, GOBP_ACTIVATED_T_CELL_PROLIFERATION REACTOME_SIGNALING_BY_ERBB4, REACTOME_SIGNALING_BY_ERBB4 GOBP_REGULATION_OF_T_CELL_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_T_CELL_RECEPTOR_SIGNALING_PATHWAY GSE37301_PRO_BCELL_VS_CD4_TCELL_DN, GSE37301_PRO_BCELL_VS_CD4_TCELL_DN HP_FOCAL_EMOTIONAL_SEIZURE, HP_FOCAL_EMOTIONAL_SEIZURE MIZUSHIMA_AUTOPHAGOSOME_FORMATION, MIZUSHIMA_AUTOPHAGOSOME_FORMATION GOBP_HEAD_MORPHOGENESIS, GOBP_HEAD_MORPHOGENESIS HP_OCULOGYRIC_CRISIS, HP_OCULOGYRIC_CRISIS HP_TRANSIENT_ISCHEMIC_ATTACK, HP_TRANSIENT_ISCHEMIC_ATTACK FERRARI_RESPONSE_TO_FENRETINIDE_UP, FERRARI_RESPONSE_TO_FENRETINIDE_UP GOCC_NBAF_COMPLEX, GOCC_NBAF_COMPLEX GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY REACTOME_PROLONGED_ERK_ACTIVATION_EVENTS, REACTOME_PROLONGED_ERK_ACTIVATION_EVENTS GOBP_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION GOMF_EPHRIN_RECEPTOR_BINDING, GOMF_EPHRIN_RECEPTOR_BINDING HP_ISCHEMIC_STROKE, HP_ISCHEMIC_STROKE REACTOME_ERYTHROPOIETIN_ACTIVATES_RAS, REACTOME_ERYTHROPOIETIN_ACTIVATES_RAS HP_RIB_FUSION, HP_RIB_FUSION HP_EMPHYSEMA, HP_EMPHYSEMA HP_OVARIAN_CARCINOMA, HP_OVARIAN_CARCINOMA GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING, GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING HP_WIDELY_SPACED_MAXILLARY_CENTRAL_INCISORS, HP_WIDELY_SPACED_MAXILLARY_CENTRAL_INCISORS HP_BROAD_DISTAL_PHALANX_OF_FINGER, HP_BROAD_DISTAL_PHALANX_OF_FINGER PID_ANTHRAX_PATHWAY, PID_ANTHRAX_PATHWAY ZNF157_TARGET_GENES, ZNF157_TARGET_GENES SHIN_B_CELL_LYMPHOMA_CLUSTER_9, SHIN_B_CELL_LYMPHOMA_CLUSTER_9 HP_APLASIA_OF_THE_SEMICIRCULAR_CANAL, HP_APLASIA_OF_THE_SEMICIRCULAR_CANAL GOBP_LONG_TERM_SYNAPTIC_DEPRESSION, GOBP_LONG_TERM_SYNAPTIC_DEPRESSION GOMF GLUTAMATE_RECEPTOR_BINDING, GOMF GLUTAMATE_RECEPTOR_BINDING REACTOME_CRMP5_IN_SEMA3A_SIGNALING, REACTOME_CRMP5_IN_SEMA3A_SIGNALING CAO_BLOOD_FLUZONE_AGE_05_14YO_CORRELATED_WITH_H3N1_HI_TITTER_1DY_POSITIVE, CAO_BLOOD_FLUZONE_AGE_05_14YO_CORRELATED_WITH_H3N1_HI_TITTER_1DY_POSITIVE GOBP_LOCOMOTORY_BEHAVIOR, GOBP_LOCOMOTORY_BEHAVIOR HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_VESTIBULE_OF_THE_INNER_EAR, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_VESTIBULE_OF_THE_INNER_EAR REACTOME_HDL_REMODELING, REACTOME_HDL_REMODELING REACTOME_REGULATION_OF_KIT_SIGNALING, REACTOME_REGULATION_OF_KIT_SIGNALING BIOCARTA_IL3_PATHWAY, BIOCARTA_IL3_PATHWAY HP_ABNORMAL_SPACED_INCISORS, HP_ABNORMAL_SPACED_INCISORS GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEUBIQUITINATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEUBIQUITINATION chr13q22, chr13q22 GOBP_VENTRICULAR_CARDIAC_MUSCLE_CELL_DIFFERENTIATION, GOBP_VENTRICULAR_CARDIAC_MUSCLE_CELL_DIFFERENTIATION HP_OSTIOMALACIA, HP_OSTIOMALACIA GOBP_RESPONSE_TO_ATP, GOBP_RESPONSE_TO_ATP PID_EPHA_FWDPATHWAY, PID_EPHA_FWDPATHWAY GOBP_PERICARDIUM_DEVELOPMENT, GOBP_PERICARDIUM_DEVELOPMENT GOMF_SUMO_TRANSFERASE_ACTIVITY, GOMF_SUMO_TRANSFERASE_ACTIVITY GOBP_REGULATION_OF_POTASSIUM_ION_TRANSPORT, GOBP_REGULATION_OF_POTASSIUM_ION_TRANSPORT GOBP_MEMBRANE_REPOLARIZATION, GOBP_MEMBRANE_REPOLARIZATION GOMF_ABC_TYPE_XENOBIOTIC_TRANSPORTER_ACTIVITY, GOMF_ABC_TYPE_XENOBIOTIC_TRANSPORTER_ACTIVITY HP_THROMBOPHLEBITIS, HP_THROMBOPHLEBITIS GOBP_NEGATIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT MIR7108_3P, MIR7108_3P BIOCARTA_RHO_PATHWAY, BIOCARTA_RHO_PATHWAY GOBP_EXCRETION, GOBP_EXCRETION REACTOME_NCAM_SIGNALING_FOR_NEURITE_OUT_GROWTH, REACTOME_NCAM_SIGNALING_FOR_NEURITE_OUT_GROWTH REACTOME_SHC1_EVENTS_IN_ERBB4_SIGNALING, REACTOME_SHC1_EVENTS_IN_ERBB4_SIGNALING GOBP_PERICARDIUM_MORPHOGENESIS, GOBP_PERICARDIUM_MORPHOGENESIS WP_15Q133_COPY_NUMBER_VARIATION_SYNDROME, WP_15Q133_COPY_NUMBER_VARIATION_SYNDROME GOMF_PHOSPHORUS_OXYGEN_LYASE_ACTIVITY, GOMF_PHOSPHORUS_OXYGEN_LYASE_ACTIVITY REACTOME_GRB2_EVENTS_IN_ERBB2_SIGNALING, REACTOME_GRB2_EVENTS_IN_ERBB2_SIGNALING HP_PREGNANCY_EXPOSURE, HP_PREGNANCY_EXPOSURE HP_RECURRENT_THROMBOPHLEBITIS, HP_RECURRENT_THROMBOPHLEBITIS GOMF_CALCIIUM_CHANNEL_REGULATOR_ACTIVITY, GOMF_CALCIIUM_CHANNEL_REGULATOR_ACTIVITY | |
|--|--|--|