CTRL_CD8_TCELL_LATE_PASSAGE_CLONE_UP, GSE5142_HTERT_TRANSDUCED_VS_CTRL_CD8_TCELL_LATE_PASSAGE_CLONE_UP
GOCC_CHROMOSOME_CENTROMERIC_REGION, GOCC_CHROMOSOME_CENTROMERIC_REGION
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_24H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_24H_CD4_TCELL_DN
GSE25085_FETAL_LIVER_VS_ADULT_BM_SP4_THYMIC_IMPLANT_DN, GSE25085_FETAL_LIVER_VS_ADULT_BM_SP4_THYMIC_IMPLANT_DN
GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP
KAECH_DAY8_EFF_VS_DAY15_EFF_CD8_TCELL_UP, KAECH_DAY8_EFF_VS_DAY15_EFF_CD8_TCELL_UP
GSE17186_BLOOD_VS_CORD_BLOOD_CD21HIGH_TRANSITIONAL_BCELL_UP, GSE17186_BLOOD_VS_CORD_BLOOD_CD21HIGH_TRANSITIONAL_BCELL_UP
GSE20727_CTRL_VS_ROS_INH_AND_DNFB_ALLERGEN_TREATED_DC_DN, GSE20727_CTRL_VS_ROS_INH_AND_DNFB_ALLERGEN_TREATED_DC_DN
GSE7852_LN_VS_THYMUS_TCONV_DN, GSE7852_LN_VS_THYMUS_TCONV_DN
GSE24634_TREG_VS_TCONV_POST_DAY10_IL4_CONVERSION_UP, GSE24634_TREG_VS_TCONV_POST_DAY10_IL4_CONVERSION_UP
GSE10239_NAIVE_VS_KLRG1HIGH_EFF_CD8_TCELL_DN, GSE10239_NAIVE_VS_KLRG1HIGH_EFF_CD8_TCELL_DN
GSE25085_FETAL_LIVER_VS_ADULT_BM_SP4_THYMIC_IMPLANT_UP, GSE25085_FETAL_LIVER_VS_ADULT_BM_SP4_THYMIC_IMPLANT_UP
REACTOME_MITOTIC_SPINDLE_CHECKPOINT, REACTOME_MITOTIC_SPINDLE_CHECKPOINT
GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_UP, GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_UP
GSE21546_UNSTIM_VS_ANTI_CD3_STIM_DP_THYMOCYTES_DN, GSE21546_UNSTIM_VS_ANTI_CD3_STIM_DP_THYMOCYTES_DN
GSE40068_BCL6_POS_VS_NEG_CXCR5_POS_TFH_UP, GSE40068_BCL6_POS_VS_NEG_CXCR5_POS_TFH_UP
GSE19941_UNSTIM_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN, GSE19941_UNSTIM_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN
GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC_DN, GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC_DN
REACTOME_CHROMOSOME_MAINTENANCE, REACTOME_CHROMOSOME_MAINTENANCE
GSE19888_ADENOSINE_A3R_ACT_VS_A3R_ACT_WITH_A3R_INH_PRETREATMENT_IN_MAST_CELL_UP, GSE19888_ADENOSINE_A3R_ACT_VS_A3R_ACT_WITH_A3R_INH_PRETREATMENT_IN_MAST_CELL_UP
GSE9509_10MIN_VS_30MIN_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_UP, GSE9509_10MIN_VS_30MIN_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_UP
GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_UP
GOCC_KINETOCHORE, GOCC_KINETOCHORE
GSE3982_EOSINOPHIL_VS_BCELL_DN, GSE3982_EOSINOPHIL_VS_BCELL_DN
GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
GOCC_CONDENSED_CHROMOSOME_CENTROMERIC_REGION, GOCC_CONDENSED_CHROMOSOME_CENTROMERIC_REGION
GSE40068_CXCR5NEG_BCL6NEG_CD4_TCELL_VS_CXCR5POS_BCL6NEG_TFH_UP, GSE40068_CXCR5NEG_BCL6NEG_CD4_TCELL_VS_CXCR5POS_BCL6NEG_TFH_UP
WHITFIELD_CELL_CYCLE_G2, WHITFIELD_CELL_CYCLE_G2
GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_IMMATURE_NEURON_CELL_LINE_DN, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_IMMATURE_NEURON_CELL_LINE_DN
GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_DN, GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_DN
KAECH_NAIVE_VS_DAY8_EFF_CD8_TCELL_DN, KAECH_NAIVE_VS_DAY8_EFF_CD8_TCELL_DN
GSE17186_NAIVE_VS_CD21LOW_TRANSITIONAL_BCELL_CORD_BLOOD_DN, GSE17186_NAIVE_VS_CD21LOW_TRANSITIONAL_BCELL_CORD_BLOOD_DN
GSE24210_TCONV_VS_TREG_DN, GSE24210_TCONV_VS_TREG_DN
GSE26030_TH1_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_UP, GSE26030_TH1_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_UP
GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN, GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN
REACTOME_RESOLUTION_OF_SISTER_CHROMATID_COHESION, REACTOME_RESOLUTION_OF_SISTER_CHROMATID_COHESION
GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS_RARA_AGONIST_AM580_TREATED_DC_UP, GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS_RARA_AGONIST_AM580_TREATED_DC_UP
GSE40666_WT_VS_STAT4_KO_CD8_TCELL_WITH_IFNA_STIM_90MIN_DN, GSE40666_WT_VS_STAT4_KO_CD8_TCELL_WITH_IFNA_STIM_90MIN_DN
GSE17580_UNINFECTED_VS_S_MANSONI_INF_TREG_DN, GSE17580_UNINFECTED_VS_S_MANSONI_INF_TREG_DN
GSE369_SOCS3_KO_VS_IFNG_KO_LIVER_DN, GSE369_SOCS3_KO_VS_IFNG_KO_LIVER_DN
GSE45365_HEALTHY_VS_MCMV_INFECTION_BCELL_IFNAR_KO_UP, GSE45365_HEALTHY_VS_MCMV_INFECTION_BCELL_IFNAR_KO_UP
GSE339_CD4POS_VS_CD8POS_DC_DN, GSE339_CD4POS_VS_CD8POS_DC_DN
GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_UP, GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_UP
GSE15624_CTRL_VS_3H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_CTRL_VS_3H_HALOFUGINONE_TREATED_CD4_TCELL_DN
GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_UP
GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_DN, GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_DN
GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_CTRL_CD4_TCELL_DN, GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_CTRL_CD4_TCELL_DN
GSE5455_EX_VIVO_VS_POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_UP, GSE5455_EX_VIVO_VS_POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_UP
GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP
GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY8_EFFECTOR_CD8_TCELL_DN, GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY8_EFFECTOR_CD8_TCELL_DN
GSE27786_NEUTROPHIL_VS_MONO_MAC_DN, GSE27786_NEUTROPHIL_VS_MONO_MAC_DN
GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN, GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN
GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP, GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP
GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_UP, GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_UP
GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_DN, GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_DN
GOBP_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY, GOBP_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY
GSE32901_NAIVE_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_NAIVE_VS_TH17_NEG_CD4_TCELL_UP
REACTOME_RHO_GTPASES_ACTIVATE_FORMINS, REACTOME_RHO_GTPASES_ACTIVATE_FORMINS
GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN
GSE20727_CTRL_VS_DNFB_ALLERGEN_TREATED_DC_UP, GSE20727_CTRL_VS_DNFB_ALLERGEN_TREATED_DC_UP
GSE7460_CTRL_VS_TGFB_TREATED_ACT_CD8_TCELL_UP, GSE7460_CTRL_VS_TGFB_TREATED_ACT_CD8_TCELL_UP
GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_DN, GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_DN
GSE38304_MYC_NEG_VS_POS_GC_BCELL_DN, GSE38304_MYC_NEG_VS_POS_GC_BCELL_DN
GSE36078_UNTREATED_VS_AD5_INF_MOUSE_LUNG_DC_DN, GSE36078_UNTREATED_VS_AD5_INF_MOUSE_LUNG_DC_DN
GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGG_BCELL_UP, GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGG_BCELL_UP
GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_DN, GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_DN
GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN, GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN
GSE11961_UNSTIM_VS_ANTI_IGM_AND_CD40_STIM_6H_FOLLICULAR_BCELL_UP, GSE11961_UNSTIM_VS_ANTI_IGM_AND_CD40_STIM_6H_FOLLICULAR_BCELL_UP
GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNAR_KO_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNAR_KO_DN
GSE40655_FOXO1_KO_VS_WT_NTREG_DN, GSE40655_FOXO1_KO_VS_WT_NTREG_DN
GSE17580_TREG_VS_TEFF_S_MANSONI_INF_UP, GSE17580_TREG_VS_TEFF_S_MANSONI_INF_UP
GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_DN, GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_DN
GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_UP, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_UP
GSE10240_CTRL_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN, GSE10240_CTRL_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN
GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY, GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY
GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_UP, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_UP
GOCC_OXIDOREDUCTASE_COMPLEX, GOCC_OXIDOREDUCTASE_COMPLEX
GOCC_RESPIRATORY_CHAIN_COMPLEX, GOCC_RESPIRATORY_CHAIN_COMPLEX
GOBP_ATP_DEPENDENT_CHROMATIN_REMODELING, GOBP_ATP_DEPENDENT_CHROMATIN_REMODELING
GSE7831_1H_VS_4H_CPG_STIM_PDC_DN, GSE7831_1H_VS_4H_CPG_STIM_PDC_DN
GSE23505_IL6_IL1_VS_IL6_IL1_IL23_TREATED_CD4_TCELL_UP, GSE23505_IL6_IL1_VS_IL6_IL1_IL23_TREATED_CD4_TCELL_UP
GSE17721_LPS_VS_CPG_16H_BMDC_DN, GSE17721_LPS_VS_CPG_16H_BMDC_DN
GSE3982_BCELL_VS_BASOPHIL_UP, GSE3982_BCELL_VS_BASOPHIL_UP
GSE8515_IL1_VS_IL6_4H_STIM_MAC_UP, GSE8515_IL1_VS_IL6_4H_STIM_MAC_UP
GOCC_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT, GOCC_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT
GSE27786_CD8_TCELL_VS_MONO_MAC_DN, GSE27786_CD8_TCELL_VS_MONO_MAC_DN
GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN
GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_DN, GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_DN
REACTOME_COMPLEX_I_BIOGENESIS, REACTOME_COMPLEX_I_BIOGENESIS
REACTOME_MITOCHONDRIAL_PROTEIN_IMPORT, REACTOME_MITOCHONDRIAL_PROTEIN_IMPORT
GSE29614_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_DN, GSE29614_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_DN
GOCC_NADH_DEHYDROGENASE_COMPLEX, GOCC_NADH_DEHYDROGENASE_COMPLEX
GOMF_DNA_DEPENDENT_ATPASE_ACTIVITY, GOMF_DNA_DEPENDENT_ATPASE_ACTIVITY
GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_DN, GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_DN
GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE, GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE
REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS, REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS
GSE34205_HEALTHY_VS_RSV_INF_INFANT_PBMC_DN, GSE34205_HEALTHY_VS_RSV_INF_INFANT_PBMC_DN
GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_DN, GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_DN
GSE41978_WT_VS_ID2_KO_AND_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_WT_VS_ID2_KO_AND_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
GOBP_TRNA_TRANSPORT, GOBP_TRNA_TRANSPORT
REACTOME_NUCLEAR_IMPORT_OF_REV_PROTEIN, REACTOME_NUCLEAR_IMPORT_OF_REV_PROTEIN
GSE10325_BCELL_VS_MYELOID_UP, GSE10325_BCELL_VS_MYELOID_UP
GSE360_DC_VS_MAC_L_DONOVANI_UP, GSE360_DC_VS_MAC_L_DONOVANI_UP
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR
GSE4811_CLASSSICALY_ACTIVATED_VS_TYPE_2_ACTIVATED_MACROPHAGE_UP, GSE4811_CLASSSICALY_ACTIVATED_VS_TYPE_2_ACTIVATED_MACROPHAGE_UP
GSE17721_CTRL_VS_LPS_12H_BMDC_DN, GSE17721_CTRL_VS_LPS_12H_BMDC_DN
GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_UP, GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_UP
GOBP_CHROMATIN_REMODELING_AT_CENTROMERE, GOBP_CHROMATIN_REMODELING_AT_CENTROMERE
HP_ABNORMALITY_OF_THE_TESTIS_SIZE, HP_ABNORMALITY_OF_THE_TESTIS_SIZE
REACTOME_POSTMITOTIC_NUCLEAR_PORE_COMPLEX_NPC_REFORMATION, REACTOME_POSTMITOTIC_NUCLEAR_PORE_COMPLEX_NPC_REFORMATION
PID_AURORA_B_PATHWAY, PID_AURORA_B_PATHWAY
GOMF_NAD_P_H_DEHYDROGENASE_QUINONE_ACTIVITY, GOMF_NAD_P_H_DEHYDROGENASE_QUINONE_ACTIVITY
GUO_TARGETS_OF_IRS1_AND_IRS2, GUO_TARGETS_OF_IRS1_AND_IRS2
REACTOME_E2F_MEDIATED_REGULATION_OF_DNA_REPLICATION, REACTOME_E2F_MEDIATED_REGULATION_OF_DNA_REPLICATION
NAKAYA_PBMC_FLUARIX_FLUVIRIN_AGE_18_50YO_CORRELATED_WITH_HAI_28DY_RESPONSE_AT_7DY_POSITIVE, NAKAYA_PBMC_FLUARIX_FLUVIRIN_AGE_18_50YO_CORRELATED_WITH_HAI_28DY_RESPONSE_AT_7DY_POSITIVE
ESC_J1_UP_LATE.V1_DN, ESC_J1_UP_LATE.V1_DN
GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
REACTOME_EXPORT_OF_VIRAL_RIBONUCLEOPROTEINS_FROM_NUCLEUS, REACTOME_EXPORT_OF_VIRAL_RIBONUCLEOPROTEINS_FROM_NUCLEUS
MODULE_308, MODULE_308
GSE17974_0H_VS_0.5H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_0.5H_IN_VITRO_ACT_CD4_TCELL_DN
GOBP_HISTONE_EXCHANGE, GOBP_HISTONE_EXCHANGE
REACTOME_REGULATION_OF_CHOLESTEROL_BIOSYNTHESIS_BY_SREBP_SREBF, REACTOME_REGULATION_OF_CHOLESTEROL_BIOSYNTHESIS_BY_SREBP_SREBF
HP_BONE_MARROW_HYPOCELLULARITY, HP_BONE_MARROW_HYPOCELLULARITY
CAFFAREL_RESPONSE_TO_THC_24HR_5_DN, CAFFAREL_RESPONSE_TO_THC_24HR_5_DN
HALLMARK_ANDROGEN_RESPONSE, HALLMARK_ANDROGEN_RESPONSE
GSE27786_BCELL_VS_MONO_MAC_DN, GSE27786_BCELL_VS_MONO_MAC_DN
MATZUK_SPERMATOCYTE, MATZUK_SPERMATOCYTE
OUELLET_CULTURED_OVARIAN_CANCER_INVASIVE_VS_LMP_UP, OUELLET_CULTURED_OVARIAN_CANCER_INVASIVE_VS_LMP_UP
GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_FETAL_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_FETAL_DN
GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_DN, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_DN
HP_ABNORMALITY_OF_SKIN_ADNEXA_PHYSIOLOGY, HP_ABNORMALITY_OF_SKIN_ADNEXA_PHYSIOLOGY
GENTILE_RESPONSE_CLUSTER_D3, GENTILE_RESPONSE_CLUSTER_D3
GOBP_PROTEIN_TRANSMEMBRANE_TRANSPORT, GOBP_PROTEIN_TRANSMEMBRANE_TRANSPORT
REACTOME_RHOG_GTPASE_CYCLE, REACTOME_RHOG_GTPASE_CYCLE
KEGG_P53_SIGNALING_PATHWAY, KEGG_P53_SIGNALING_PATHWAY
HP_DECREASED_ACTIVITY_OF_THE_PYRUVATE_DEHYDROGENASE_COMPLEX, HP_DECREASED_ACTIVITY_OF_THE_PYRUVATE_DEHYDROGENASE_COMPLEX
GOCC_NUCLEAR_REPLICATION_FORK, GOCC_NUCLEAR_REPLICATION_FORK
HUANG_GATA2_TARGETS_DN, HUANG_GATA2_TARGETS_DN
GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
REACTOME_RHOC_GTPASE_CYCLE, REACTOME_RHOC_GTPASE_CYCLE
REACTOME_CHOLESTEROL_BIOSYNTHESIS, REACTOME_CHOLESTEROL_BIOSYNTHESIS
BOYLAN_MULTIPLE_MYELOMA_PCA3_DN, BOYLAN_MULTIPLE_MYELOMA_PCA3_DN
GOMF_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE, GOMF_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE
HP_KINETIC_TREMOR, HP_KINETIC_TREMOR
GSE36078_WT_VS_IL1R_KO_LUNG_DC_UP, GSE36078_WT_VS_IL1R_KO_LUNG_DC_UP
EGR3_01, EGR3_01
GOMF_CARBON_OXYGEN_LYASE_ACTIVITY, GOMF_CARBON_OXYGEN_LYASE_ACTIVITY
ZHANG_PROLIFERATING_VS_QUIESCENT, ZHANG_PROLIFERATING_VS_QUIESCENT
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_TRASDUCED_CD4_TCELL_DN, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_TRASDUCED_CD4_TCELL_DN
BOSCO_TH1_CYTOTOXIC_MODULE, BOSCO_TH1_CYTOTOXIC_MODULE
BILD_SRC_ONCOGENIC_SIGNATURE, BILD_SRC_ONCOGENIC_SIGNATURE
GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_ALKYL_OR_ARYL_OTHER_THAN_METHYL_GROUPS, GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_ALKYL_OR_ARYL_OTHER_THAN_METHYL_GROUPS
LU_IL4_SIGNALING, LU_IL4_SIGNALING
HP_DECREASED_ADIPOSE_TISSUE, HP_DECREASED_ADIPOSE_TISSUE
REACTOME_INHIBITION_OF_REPLICATION_INITIATION_OF_DAMAGED_DNA_BY_RB1_E2F1, REACTOME_INHIBITION_OF_REPLICATION_INITIATION_OF_DAMAGED_DNA_BY_RB1_E2F1
GOBP_MEIOTIC_CHROMOSOME_SEGREGATION, GOBP_MEIOTIC_CHROMOSOME_SEGREGATION
GOBP_MICROTUBULE_DEPOLYMERIZATION, GOBP_MICROTUBULE_DEPOLYMERIZATION
REACTOME_RHOD_GTPASE_CYCLE, REACTOME_RHOD_GTPASE_CYCLE
MODULE_320, MODULE_320
PRAMOONJAGO_SOX4_TARGETS_DN, PRAMOONJAGO_SOX4_TARGETS_DN
HORTON_SREBF_TARGETS, HORTON_SREBF_TARGETS
HP_AREFLEXIA_OF_LOWER_LIMBS, HP_AREFLEXIA_OF_LOWER_LIMBS
GOLUB_ALL_VS_AML_UP, GOLUB_ALL_VS_AML_UP
TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_UP, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_UP
PACHER_TARGETS_OF_IGF1_AND_IGF2_UP, PACHER_TARGETS_OF_IGF1_AND_IGF2_UP
WP_CHOLESTEROL_METABOLISM_INCLUDES_BOTH_BLOCH_AND_KANDUTSCHRUSSELL_PATHWAYS, WP_CHOLESTEROL_METABOLISM_INCLUDES_BOTH_BLOCH_AND_KANDUTSCHRUSSELL_PATHWAYS
GOBP_STEROL_BIOSYNTHETIC_PROCESS, GOBP_STEROL_BIOSYNTHETIC_PROCESS
GOMF_ATPASE_REGULATOR_ACTIVITY, GOMF_ATPASE_REGULATOR_ACTIVITY
GOCC_CONDENSED_NUCLEAR_CHROMOSOME_KINETOCHORE, GOCC_CONDENSED_NUCLEAR_CHROMOSOME_KINETOCHORE
GSE17721_POLYIC_VS_PAM3CSK4_2H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_2H_BMDC_DN
HP_PARTIAL_AGENESIS_OF_THE_CORPUS_CALLOSUM, HP_PARTIAL_AGENESIS_OF_THE_CORPUS_CALLOSUM
GOBP_CHROMOSOME_ORGANIZATION_INVOLVED_IN_MEIOTIC_CELL_CYCLE, GOBP_CHROMOSOME_ORGANIZATION_INVOLVED_IN_MEIOTIC_CELL_CYCLE
AIZARANI_LIVER_C12_NK_NKT_CELLS_4, AIZARANI_LIVER_C12_NK_NKT_CELLS_4
HP_HYPERPITUITARISM, HP_HYPERPITUITARISM
GOMF_HYDRO_LYASE_ACTIVITY, GOMF_HYDRO_LYASE_ACTIVITY
HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION
GOBP_RESPONSE_TO_X_RAY, GOBP_RESPONSE_TO_X_RAY
MATZUK_MEIOTIC_AND_DNA_REPAIR, MATZUK_MEIOTIC_AND_DNA_REPAIR
GOBP_PROTEIN_LOCALIZATION_TO_CHROMATIN, GOBP_PROTEIN_LOCALIZATION_TO_CHROMATIN
WP_SMALL_CELL_LUNG_CANCER, WP_SMALL_CELL_LUNG_CANCER
GOBP_MALE_MEIOTIC_NUCLEAR_DIVISION, GOBP_MALE_MEIOTIC_NUCLEAR_DIVISION
WP_CHOLESTEROL_BIOSYNTHESIS_PATHWAY, WP_CHOLESTEROL_BIOSYNTHESIS_PATHWAY
CHOW_RASSF1_TARGETS_DN, CHOW_RASSF1_TARGETS_DN
GOBP_NUCLEAR_PORE_ORGANIZATION, GOBP_NUCLEAR_PORE_ORGANIZATION
GOMF_MACROMOLECULE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_MACROMOLECULE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_REGULATOR_ACTIVITY, GOMF_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_REGULATOR_ACTIVITY
MODULE_306, MODULE_306
GOBP_NEGATIVE_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GOBP_NEGATIVE_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION
GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
HP_PERIODONTITIS, HP_PERIODONTITIS
MARIADASON_RESPONSE_TO_BUTYRATE_SULINDAC_4, MARIADASON_RESPONSE_TO_BUTYRATE_SULINDAC_4
SCHEIDEREIT_IKK_INTERACTING_PROTEINS, SCHEIDEREIT_IKK_INTERACTING_PROTEINS
HP_MUSCLE_FIBRILLATION, HP_MUSCLE_FIBRILLATION
HP_ADIPOSE_TISSUE_LOSS, HP_ADIPOSE_TISSUE_LOSS
BIOCARTA_CASPASE_PATHWAY, BIOCARTA_CASPASE_PATHWAY
GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY, GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY
KEGG_CITRATE_CYCLE_TCA_CYCLE, KEGG_CITRATE_CYCLE_TCA_CYCLE
HP_ABNORMAL_BLISTERING_OF_THE_SKIN, HP_ABNORMAL_BLISTERING_OF_THE_SKIN
WP_TCA_CYCLE_AKA_KREBS_OR_CITRIC_ACID_CYCLE, WP_TCA_CYCLE_AKA_KREBS_OR_CITRIC_ACID_CYCLE
REACTOME_DNA_REPLICATION_INITIATION, REACTOME_DNA_REPLICATION_INITIATION
GOMF_PRENYLTRANSFERASE_ACTIVITY, GOMF_PRENYLTRANSFERASE_ACTIVITY
HP_SLOW_SACCADIC_EYE_MOVEMENTS, HP_SLOW_SACCADIC_EYE_MOVEMENTS
GOCC_CYTOPLASMIC_MICROTUBULE, GOCC_CYTOPLASMIC_MICROTUBULE
GOMF_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR, GOMF_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR
GOBP_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS
GOBP_FEMALE_GAMETE_GENERATION, GOBP_FEMALE_GAMETE_GENERATION
GOBP_REGULATION_OF_STEROID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_STEROID_BIOSYNTHETIC_PROCESS
HP_PRURITUS, HP_PRURITUS
REACTOME_CYTOCHROME_C_MEDIATED_APOPTOTIC_RESPONSE, REACTOME_CYTOCHROME_C_MEDIATED_APOPTOTIC_RESPONSE
GOBP_PORE_COMPLEX_ASSEMBLY, GOBP_PORE_COMPLEX_ASSEMBLY
GOBP_CELLULAR_RESPONSE_TO_GAMMA_RADIATION, GOBP_CELLULAR_RESPONSE_TO_GAMMA_RADIATION
GOBP_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS
MIR382_5P, MIR382_5P
HP_IMPAIRED_SMOOTH_PURSUIT, HP_IMPAIRED_SMOOTH_PURSUIT
GOBP_FEMALE_SEX_DIFFERENTIATION, GOBP_FEMALE_SEX_DIFFERENTIATION
HP_HYPERGONADOTROPIC_HYPOGONADISM, HP_HYPERGONADOTROPIC_HYPOGONADISM
KEGG_PROTEIN_EXPORT, KEGG_PROTEIN_EXPORT
SHAFFER_IRF4_MULTIPLE_MYELOMA_PROGRAM, SHAFFER_IRF4_MULTIPLE_MYELOMA_PROGRAM
GOBP_REGULATION_OF_STEROID_METABOLIC_PROCESS, GOBP_REGULATION_OF_STEROID_METABOLIC_PROCESS
HP_DUPLICATION_OF_THUMB_PHALANX, HP_DUPLICATION_OF_THUMB_PHALANX
MODULE_432, MODULE_432
GOCC_MICROTUBULE_END, GOCC_MICROTUBULE_END
REACTOME_ONCOGENE_INDUCED_SENESCENCE, REACTOME_ONCOGENE_INDUCED_SENESCENCE
GOBP_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS
MIR520G_5P, MIR520G_5P
PID_DELTA_NP63_PATHWAY, PID_DELTA_NP63_PATHWAY
HP_TONGUE_FASCICULATIONS, HP_TONGUE_FASCICULATIONS
ANDERSEN_CHOLANGIOCARCINOMA_CLASS1, ANDERSEN_CHOLANGIOCARCINOMA_CLASS1
GOBP_L_SERINE_METABOLIC_PROCESS, GOBP_L_SERINE_METABOLIC_PROCESS
HP_SQUAMOUS_CELL_CARCINOMA, HP_SQUAMOUS_CELL_CARCINOMA
HP_ABNORMALITY_OF_DNA_REPAIR, HP_ABNORMALITY_OF_DNA_REPAIR
GOBP_NEGATIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GOBP_NEGATIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
GOMF_PROTEIN_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PROTEIN_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
HP_DECREASED_ABSENT_ANKLE_REFLEXES, HP_DECREASED_ABSENT_ANKLE_REFLEXES
GOBP_MITOTIC_RECOMBINATION, GOBP_MITOTIC_RECOMBINATION
REACTOME_METABOLISM_OF_COFACTORS, REACTOME_METABOLISM_OF_COFACTORS
MODULE_196, MODULE_196
GOMF_ATPASE_ACTIVATOR_ACTIVITY, GOMF_ATPASE_ACTIVATOR_ACTIVITY
GOBP_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS
KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM, KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM
SCHAEFFER_SOX9_TARGETS_IN_PROSTATE_DEVELOPMENT_DN, SCHAEFFER_SOX9_TARGETS_IN_PROSTATE_DEVELOPMENT_DN
HP_ABNORMAL_CIRCULATING_GONADOTROPIN_LEVEL, HP_ABNORMAL_CIRCULATING_GONADOTROPIN_LEVEL
HP_INCREASED_CIRCULATING_GONADOTROPIN_LEVEL, HP_INCREASED_CIRCULATING_GONADOTROPIN_LEVEL
MIR7160_3P, MIR7160_3P
HP_ABNORMALITY_OF_HAIR_GROWTH, HP_ABNORMALITY_OF_HAIR_GROWTH
GOBP_NUCLEAR_PORE_COMPLEX_ASSEMBLY, GOBP_NUCLEAR_PORE_COMPLEX_ASSEMBLY
PID_REG_GR_PATHWAY, PID_REG_GR_PATHWAY
BIOCARTA_D4GDI_PATHWAY, BIOCARTA_D4GDI_PATHWAY
HP_COMPLETE_DUPLICATION_OF_PHALANX_OF_HAND, HP_COMPLETE_DUPLICATION_OF_PHALANX_OF_HAND
GOBP_DEVELOPMENT_OF_PRIMARY_FEMALE_SEXUAL_CHARACTERISTICS, GOBP_DEVELOPMENT_OF_PRIMARY_FEMALE_SEXUAL_CHARACTERISTICS
ZFP82_TARGET_GENES, ZFP82_TARGET_GENES
GOBP_DNA_CATABOLIC_PROCESS_ENDONUCLEOLYTIC, GOBP_DNA_CATABOLIC_PROCESS_ENDONUCLEOLYTIC
MODULE_253, MODULE_253
WP_CONSTITUTIVE_ANDROSTANE_RECEPTOR_PATHWAY, WP_CONSTITUTIVE_ANDROSTANE_RECEPTOR_PATHWAY
GOBP_REGULATION_OF_CHOLESTEROL_METABOLIC_PROCESS, GOBP_REGULATION_OF_CHOLESTEROL_METABOLIC_PROCESS
WP_HEPATITIS_C_AND_HEPATOCELLULAR_CARCINOMA, WP_HEPATITIS_C_AND_HEPATOCELLULAR_CARCINOMA
GOBP_HOMOLOGOUS_CHROMOSOME_SEGREGATION, GOBP_HOMOLOGOUS_CHROMOSOME_SEGREGATION
CUI_DEVELOPING_HEART_C2_CARDIOMYOCYTE, CUI_DEVELOPING_HEART_C2_CARDIOMYOCYTE
GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_1H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_1H_ACT_CD4_TCELL_UP
HP_ABNORMAL_SERUM_IRON_CONCENTRATION, HP_ABNORMAL_SERUM_IRON_CONCENTRATION
MOOTHA_TCA, MOOTHA_TCA
MODULE_300, MODULE_300
XU_HGF_SIGNALING_NOT_VIA_AKT1_6HR, XU_HGF_SIGNALING_NOT_VIA_AKT1_6HR
GOCC_FANCONI_ANAEMIA_NUCLEAR_COMPLEX, GOCC_FANCONI_ANAEMIA_NUCLEAR_COMPLEX
GOCC_NUCLEAR_PORE_OUTER_RING, GOCC_NUCLEAR_PORE_OUTER_RING
REACTOME_SYNTHESIS_OF_DIPHTHAMIDE_EEF2, REACTOME_SYNTHESIS_OF_DIPHTHAMIDE_EEF2
GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_REGULATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_REGULATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
REACTOME_RHOF_GTPASE_CYCLE, REACTOME_RHOF_GTPASE_CYCLE
REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION, REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION
HTF_01, HTF_01
GOMF_NUCLEOSIDE_MONOPHOSPHATE_KINASE_ACTIVITY, GOMF_NUCLEOSIDE_MONOPHOSPHATE_KINASE_ACTIVITY
GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
DORSAM_HOXA9_TARGETS_DN, DORSAM_HOXA9_TARGETS_DN
HOEK_T_CELL_2011_2012_TIV_ADULT_3DY_UP, HOEK_T_CELL_2011_2012_TIV_ADULT_3DY_UP
GOBP_ALPHA_AMINO_ACID_CATABOLIC_PROCESS, GOBP_ALPHA_AMINO_ACID_CATABOLIC_PROCESS
GOBP_APOPTOTIC_DNA_FRAGMENTATION, GOBP_APOPTOTIC_DNA_FRAGMENTATION
HP_ABNORMAL_HAIR_LABORATORY_EXAMINATION, HP_ABNORMAL_HAIR_LABORATORY_EXAMINATION
GOBP_GLYCINE_METABOLIC_PROCESS, GOBP_GLYCINE_METABOLIC_PROCESS
HP_CORONAL_CRANIOSYNOSTOSIS, HP_CORONAL_CRANIOSYNOSTOSIS
HP_ABNORMAL_CIRCULATING_SERINE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_SERINE_FAMILY_AMINO_ACID_CONCENTRATION
HP_ABNORMAL_BLOOD_TRANSITION_ELEMENT_CATION_CONCENTRATION, HP_ABNORMAL_BLOOD_TRANSITION_ELEMENT_CATION_CONCENTRATION
MIR219B_5P, MIR219B_5P
WP_HOMOLOGOUS_RECOMBINATION, WP_HOMOLOGOUS_RECOMBINATION
GOBP_PEPTIDYL_DIPHTHAMIDE_BIOSYNTHETIC_PROCESS_FROM_PEPTIDYL_HISTIDINE, GOBP_PEPTIDYL_DIPHTHAMIDE_BIOSYNTHETIC_PROCESS_FROM_PEPTIDYL_HISTIDINE
REACTOME_UBIQUINOL_BIOSYNTHESIS, REACTOME_UBIQUINOL_BIOSYNTHESIS
GOBP_ISOPRENOID_METABOLIC_PROCESS, GOBP_ISOPRENOID_METABOLIC_PROCESS
GOMF_PROTEIN_KINASE_C_BINDING, GOMF_PROTEIN_KINASE_C_BINDING
GOBP_ISOPRENOID_BIOSYNTHETIC_PROCESS, GOBP_ISOPRENOID_BIOSYNTHETIC_PROCESS
OVSYANNIKOVA_PBMC_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_3DY_USED_IN_HAI_AND_VNA_RESPONSE_MODELS_UP, OVSYANNIKOVA_PBMC_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_3DY_U
MIR92A_1_5P, MIR92A_1_5P
OVSYANNIKOVA_PBMC_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_0DY_USED_IN_HAI_AND_VNA_RESPONSE_MODELS_DN, OVSYANNIKOVA_PBMC_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_0DY_U
GOCC_CILIARY_TRANSITION_FIBER, GOCC_CILIARY_TRANSITION_FIBER
PETRETTO_HEART_MASS_QTL_CIS_UP, PETRETTO_HEART_MASS_QTL_CIS_UP
GOCC_HIGH_DENSITY_LIPOPROTEIN_PARTICLE, GOCC_HIGH_DENSITY_LIPOPROTEIN_PARTICLE
GOBP_MAINTENANCE_OF_BLOOD_BRAIN_BARRIER, GOBP_MAINTENANCE_OF_BLOOD_BRAIN_BARRIER
REACTOME_STING_MEDIATED_INDUCTION_OF_HOST_IMMUNE_RESPONSES, REACTOME_STING_MEDIATED_INDUCTION_OF_HOST_IMMUNE_RESPONSES
ZHENG_FOXP3_TARGETS_UP, ZHENG_FOXP3_TARGETS_UP
GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_7MO_CORRELATED_WITH_ANTIBODY_RESPONSE_POSITIVE, GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_7MO_CORRELATED_WITH_ANTIBODY_RESPONSE_POSITIVE
HAHTOLA_SEZARY_SYNDROM_DN, HAHTOLA_SEZARY_SYNDROM_DN
GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
HP_BRITTLE_HAIR, HP_BRITTLE_HAIR
REACTOME_SULFUR_AMINO_ACID_METABOLISM, REACTOME_SULFUR_AMINO_ACID_METABOLISM
GOMF_RNA_POLYMERASE_I_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOMF_RNA_POLYMERASE_I_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING
HP_CARCINOMA, HP_CARCINOMA
GOBP_PYROPTOSIS, GOBP_PYROPTOSIS
LI_CYTIDINE_ANALOGS_CYCTOTOXICITY, LI_CYTIDINE_ANALOGS_CYCTOTOXICITY
HP_LEIOMYOSARCOMA, HP_LEIOMYOSARCOMA
MODULE_471, MODULE_471
HP_NUMEROUS_PIGMENTED_FRECKLES, HP_NUMEROUS_PIGMENTED_FRECKLES
GOBP_DNA_TOPOLOGICAL_CHANGE, GOBP_DNA_TOPOLOGICAL_CHANGE
HP_ABNORMALITY_OF_VITAMIN_K_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_K_METABOLISM
REACTOME_INTERLEUKIN_7_SIGNALING, REACTOME_INTERLEUKIN_7_SIGNALING
HP_MICROSCOPIC_HEMATURIA, HP_MICROSCOPIC_HEMATURIA
GOBP_HOMOLOGOUS_CHROMOSOME_PAIRING_AT_MEIOSIS, GOBP_HOMOLOGOUS_CHROMOSOME_PAIRING_AT_MEIOSIS
GOBP_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DEVELOPMENT, GOBP_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DEVELOPMENT
GOBP_HIPPOCAMPUS_DEVELOPMENT, GOBP_HIPPOCAMPUS_DEVELOPMENT
GOBP_TERPENOID_BIOSYNTHETIC_PROCESS, GOBP_TERPENOID_BIOSYNTHETIC_PROCESS
GOBP_DNA_LIGATION, GOBP_DNA_LIGATION
HP_COLOR_VISION_DEFECT, HP_COLOR_VISION_DEFECT
HP_METAPHYSEAL_SCLEROSIS, HP_METAPHYSEAL_SCLEROSIS
GOCC_TRANSLOCON_COMPLEX, GOCC_TRANSLOCON_COMPLEX
MATHEW_FANCONI_ANEMIA_GENES, MATHEW_FANCONI_ANEMIA_GENES
GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTIBODY_RESPONSE_POSITIVE, GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTIBODY_RESPONSE_POSITIVE
HP_LOW_SET_NIPPLES, HP_LOW_SET_NIPPLES
DESCARTES_FETAL_STOMACH_ERYTHROBLASTS, DESCARTES_FETAL_STOMACH_ERYTHROBLASTS
WP_DISORDERS_OF_THE_KREBS_CYCLE, WP_DISORDERS_OF_THE_KREBS_CYCLE
MODULE_305, MODULE_305
GOBP_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_RESULTING_IN_TRANSCRIPTION, GOBP_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_RESULTING_IN_TRANSCRIPTION
GOBP_PROGRAMMED_NECROTIC_CELL_DEATH, GOBP_PROGRAMMED_NECROTIC_CELL_DEATH
LIU_TARGETS_OF_VMYB_VS_CMYB_DN, LIU_TARGETS_OF_VMYB_VS_CMYB_DN
HP_ABNORMAL_CIRCULATING_MONOCARBOXYLIC_ACID_COCENTRATION, HP_ABNORMAL_CIRCULATING_MONOCARBOXYLIC_ACID_COCENTRATION
HP_ABNORMALITY_OF_IRON_HOMEOSTASIS, HP_ABNORMALITY_OF_IRON_HOMEOSTASIS
GOBP_ASTROCYTE_DEVELOPMENT, GOBP_ASTROCYTE_DEVELOPMENT
GOBP_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY
HP_LONG_TRACT_SIGNS, HP_LONG_TRACT_SIGNS
HP_ABNORMALITY_OF_FACIAL_ADIPOSE_TISSUE, HP_ABNORMALITY_OF_FACIAL_ADIPOSE_TISSUE
NEBEN_AML_WITH_FLT3_OR_NRAS_DN, NEBEN_AML_WITH_FLT3_OR_NRAS_DN
HP_PROXIMAL_FEMORAL_EPIPHYSIOLYSIS, HP_PROXIMAL_FEMORAL_EPIPHYSIOLYSIS
KEGG_TERPENOID_BACKBONE_BIOSYNTHESIS, KEGG_TERPENOID_BACKBONE_BIOSYNTHESIS
MODULE_528, MODULE_528
HP_PREMATURE_LOSS_OF_TEETH, HP_PREMATURE_LOSS_OF_TEETH
WP_HEMATOPOIETIC_STEM_CELL_GENE_REGULATION_BY_GABP_ALPHABETA_COMPLEX, WP_HEMATOPOIETIC_STEM_CELL_GENE_REGULATION_BY_GABP_ALPHABETA_COMPLEX
MODULE_333, MODULE_333
GOMF_ORGANIC_ACID_BINDING, GOMF_ORGANIC_ACID_BINDING
REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS, REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS
MODULE_349, MODULE_349
GOBP_OUTER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_OUTER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
WP_PEROXIREDOXIN_2_INDUCED_OVARIAN_FAILURE, WP_PEROXIREDOXIN_2_INDUCED_OVARIAN_FAILURE
HP_DEFORMED_RIB_CAGE, HP_DEFORMED_RIB_CAGE
GOBP_L_SERINE_BIOSYNTHETIC_PROCESS, GOBP_L_SERINE_BIOSYNTHETIC_PROCESS
GOBP_ESTABLISHMENT_OF_GOLGI_LOCALIZATION, GOBP_ESTABLISHMENT_OF_GOLGI_LOCALIZATION
KEGG_NON_HOMOLOGOUS_END_JOINING, KEGG_NON_HOMOLOGOUS_END_JOINING
GOCC_TELOMERE_CAP_COMPLEX, GOCC_TELOMERE_CAP_COMPLEX
YIH_RESPONSE_TO_ARSENITE_C4, YIH_RESPONSE_TO_ARSENITE_C4
HP_PROLONGED_BLEEDING_FOLLOWING_PROCEDURE, HP_PROLONGED_BLEEDING_FOLLOWING_PROCEDURE
GOBP_HEMATOPOIETIC_STEM_CELL_PROLIFERATION, GOBP_HEMATOPOIETIC_STEM_CELL_PROLIFERATION
HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_3DY_UP, HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_3DY_UP
HP_REDUCTION_OF_NEUTROPHIL_MOTILITY, HP_REDUCTION_OF_NEUTROPHIL_MOTILITY
HP_STEROID_RESISTANT_NEPHROTIC_SYNDROME, HP_STEROID_RESISTANT_NEPHROTIC_SYNDROME
LEE_CALORIE_RESTRICTION_MUSCLE_UP, LEE_CALORIE_RESTRICTION_MUSCLE_UP
GOBP_DNA_REPLICATION_SYNTHESIS_OF_RNA_PRIMER, GOBP_DNA_REPLICATION_SYNTHESIS_OF_RNA_PRIMER
GOBP_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION
REACTOME_SMAC_XIAP_REGULATED_APOPTOTIC_RESPONSE, REACTOME_SMAC_XIAP_REGULATED_APOPTOTIC_RESPONSE
GOBP_REGULATION_OF_CELLULAR_RESPIRATION, GOBP_REGULATION_OF_CELLULAR_RESPIRATION
GOCC_MICROTUBULE_PLUS_END, GOCC_MICROTUBULE_PLUS_END
PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN, PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN
REACTOME_IRF3_MEDIATED_INDUCTION_OF_TYPE_I_IFN, REACTOME_IRF3_MEDIATED_INDUCTION_OF_TYPE_I_IFN
BORLAK_LIVER_CANCER_EGF_UP, BORLAK_LIVER_CANCER_EGF_UP
GOBP_CELLULAR_RESPONSE_TO_ALKALOID, GOBP_CELLULAR_RESPONSE_TO_ALKALOID
HP_PROGRESSIVE_SPASTICITY, HP_PROGRESSIVE_SPASTICITY
GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY
HP_RECURRENT_BRONCHOPULMONARY_INFECTIONS, HP_RECURRENT_BRONCHOPULMONARY_INFECTIONS
GOBP_BONE_MARROW_DEVELOPMENT, GOBP_BONE_MARROW_DEVELOPMENT
GOBP_MEIOSIS_II_CELL_CYCLE_PROCESS, GOBP_MEIOSIS_II_CELL_CYCLE_PROCESS
REACTOME_SERINE_BIOSYNTHESIS, REACTOME_SERINE_BIOSYNTHESIS
HP_POLYCYSTIC_LIVER_DISEASE, HP_POLYCYSTIC_LIVER_DISEASE
GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_TRANSLOCATION, GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_TRANSLOCATION
VANHARANTA_UTERINE_FIBROID_UP, VANHARANTA_UTERINE_FIBROID_UP
HP_ABNORMAL_ECHOCARDIOGRAM, HP_ABNORMAL_ECHOCARDIOGRAM
chr1q22, chr1q22
HP_ANEMIC_PALLOR, HP_ANEMIC_PALLOR
GOCC_PLATELET_DENSE_TUBULAR_NETWORK_MEMBRANE, GOCC_PLATELET_DENSE_TUBULAR_NETWORK_MEMBRANE
GOMF_H3_HISTONE_ACETYLTRANSFERASE_ACTIVITY, GOMF_H3_HISTONE_ACETYLTRANSFERASE_ACTIVITY
SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_DN, SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_DN
WP_TYPE_2_PAPILLARY_RENAL_CELL_CARCINOMA, WP_TYPE_2_PAPILLARY_RENAL_CELL_CARCINOMA
HP_INCREASED_SERUM_IRON, HP_INCREASED_SERUM_IRON
GOBP_GLYCOSIDE_CATABOLIC_PROCESS, GOBP_GLYCOSIDE_CATABOLIC_PROCESS
GOBP_REGULATION_OF_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS, GOBP_REGULATION_OF_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS
HP_CHROMOSOMAL_BREAKAGE_INDUCED_BY_CROSSLINKING_AGENTS, HP_CHROMOSOMAL_BREAKAGE_INDUCED_BY_CROSSLINKING_AGENTS
GOBP_SRP_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_SIGNAL_SEQUENCE_RECOGNITION, GOBP_SRP_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_SIGNAL_SEQUENCE_RECOGNITION
HP_MYOCLONIC_ABSENCE_SEIZURE, HP_MYOCLONIC_ABSENCE_SEIZURE
GOBP_THROMBIN_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY, GOBP_THROMBIN_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY
GOMF_UDP_GLUCOSYLTRANSFERASE_ACTIVITY, GOMF_UDP_GLUCOSYLTRANSFERASE_ACTIVITY
GOBP_LIPID_TRANSPORT_ACROSS_BLOOD_BRAIN_BARRIER, GOBP_LIPID_TRANSPORT_ACROSS_BLOOD_BRAIN_BARRIER
HP_INTESTINAL_BLEEDING, HP_INTESTINAL_BLEEDING
HP_ABNORMAL_SUBCLAVIAN_ARTERY_MORPHOLOGY, HP_ABNORMAL_SUBCLAVIAN_ARTERY_MORPHOLOGY
CAO_BLOOD_FLUMIST_AGE_05_14YO_30DY_UP, CAO_BLOOD_FLUMIST_AGE_05_14YO_30DY_UP
GOBP_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS, GOBP_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS
REACTOME_ACTIVATION_OF_CASPASES_THROUGH_APOPTOSOME_MEDIATED_CLEAVAGE, REACTOME_ACTIVATION_OF_CASPASES_THROUGH_APOPTOSOME_MEDIATED_CLEAVAGE
REACTOME_SCAVENGING_BY_CLASS_F_RECEPTORS, REACTOME_SCAVENGING_BY_CLASS_F_RECEPTORS
HP_MICROVESICULAR_HEPATIC_STEATOSIS, HP_MICROVESICULAR_HEPATIC_STEATOSIS
GOMF_LEUCINE_ZIPPER_DOMAIN_BINDING, GOMF_LEUCINE_ZIPPER_DOMAIN_BINDING
HP_DRY_HAIR, HP_DRY_HAIR
GOBP_LIMBIC_SYSTEM_DEVELOPMENT, GOBP_LIMBIC_SYSTEM_DEVELOPMENT
HP_PAPILLARY_RENAL_CELL_CARCINOMA, HP_PAPILLARY_RENAL_CELL_CARCINOMA
GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS, GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS
HALLMARK_HEDGEHOG_SIGNALING, HALLMARK_HEDGEHOG_SIGNALING
WP_ONE_CARBON_METABOLISM_AND_RELATED_PATHWAYS, WP_ONE_CARBON_METABOLISM_AND_RELATED_PATHWAYS
HP_ABNORMAL_MONOCYTE_MORPHOLOGY, HP_ABNORMAL_MONOCYTE_MORPHOLOGY
REACTOME_SIGNALING_BY_HIPPO, REACTOME_SIGNALING_BY_HIPPO
HP_ERYTHRODERMA, HP_ERYTHRODERMA
MODULE_295, MODULE_295
HP_ABNORMALITY_OF_EXOCRINE_PANCREAS_PHYSIOLOGY, HP_ABNORMALITY_OF_EXOCRINE_PANCREAS_PHYSIOLOGY
GOBP_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS
HP_SPASTIC_PARAPARETIC_GAIT, HP_SPASTIC_PARAPARETIC_GAIT
GOBP_LEARNED_VOCALIZATION_BEHAVIOR_OR_VOCAL_LEARNING, GOBP_LEARNED_VOCALIZATION_BEHAVIOR_OR_VOCAL_LEARNING
GOMF_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY, GOMF_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY
GOBP_RESPONSE_TO_COBALT_ION, GOBP_RESPONSE_TO_COBALT_ION
GOBP_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_III_ASSEMBLY, GOBP_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_III_ASSEMBLY
HP_ABNORMAL_TRANSFERRIN_SATURATION, HP_ABNORMAL_TRANSFERRIN_SATURATION
GOBP_LUTEOLYSIS, GOBP_LUTEOLYSIS
HP_RETINAL_EXUDATE, HP_RETINAL_EXUDATE
GOMF_MUTSALPHA_COMPLEX_BINDING, GOMF_MUTSALPHA_COMPLEX_BINDING
HP_TIGER_TAIL_BANDING, HP_TIGER_TAIL_BANDING
GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OSMOTIC_STRESS, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OSMOTIC_STRESS
JOHANSSON_BRAIN_CANCER_EARLY_VS_LATE_UP, JOHANSSON_BRAIN_CANCER_EARLY_VS_LATE_UP
GLINSKY_CANCER_DEATH_UP, GLINSKY_CANCER_DEATH_UP
HP_CHRONIC_NEUTROPENIA, HP_CHRONIC_NEUTROPENIA
GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_BETA_OXIDATION, GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_BETA_OXIDATION
GOBP_COHESIN_LOADING, GOBP_COHESIN_LOADING
DORN_ADENOVIRUS_INFECTION_48HR_UP, DORN_ADENOVIRUS_INFECTION_48HR_UP
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_GLOBOSE_BASAL_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_GLOBOSE_BASAL_CELLS
GOBP_MAINTENANCE_OF_ORGANELLE_LOCATION, GOBP_MAINTENANCE_OF_ORGANELLE_LOCATION
GOMF_ALDEHYDE_LYASE_ACTIVITY, GOMF_ALDEHYDE_LYASE_ACTIVITY
GOBP_VERY_LOW_DENSITY_LIPOPROTEIN_PARTICLE_ASSEMBLY, GOBP_VERY_LOW_DENSITY_LIPOPROTEIN_PARTICLE_ASSEMBLY
GOMF_LONG_CHAIN_FATTY_ACID_TRANSPORTER_ACTIVITY, GOMF_LONG_CHAIN_FATTY_ACID_TRANSPORTER_ACTIVITY
GOMF_HYDROLASE_ACTIVITY_ACTING_ON_ETHER_BONDS, GOMF_HYDROLASE_ACTIVITY_ACTING_ON_ETHER_BONDS
GOBP_HEMOGLOBIN_BIOSYNTHETIC_PROCESS, GOBP_HEMOGLOBIN_BIOSYNTHETIC_PROCESS
GOBP_PANCREAS_DEVELOPMENT, GOBP_PANCREAS_DEVELOPMENT
JEON_SMAD6_TARGETS_DN, JEON_SMAD6_TARGETS_DN
HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANX_OF_THE_THUMB, HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANX_OF_THE_THUMB
BIOCARTA_TSP1_PATHWAY, BIOCARTA_TSP1_PATHWAY
HP_PYODERMA_GANGRENOSUM, HP_PYODERMA_GANGRENOSUM
GOCC_MITOTIC_SPINDLE_ASTRAL_MICROTUBULE, GOCC_MITOTIC_SPINDLE_ASTRAL_MICROTUBULE
HP_TRANSIENT_NEUTROPENIA, HP_TRANSIENT_NEUTROPENIA
HP_SCOTOMA, HP_SCOTOMA
MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_17, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_17
GOBP_MICROTUBULE_SEVERING, GOBP_MICROTUBULE_SEVERING
GOBP_RESPONSE_TO_STAUROSPORINE, GOBP_RESPONSE_TO_STAUROSPORINE
HP_LIMB_FASCICULATIONS, HP_LIMB_FASCICULATIONS
HP_ABNORMAL_CIRCULATING_SERINE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_SERINE_CONCENTRATION
HP_CLINODACTYLY_OF_THE_2ND_FINGER, HP_CLINODACTYLY_OF_THE_2ND_FINGER
GOBP_REGULATION_OF_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS, GOBP_REGULATION_OF_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS
GOBP_L_SERINE_CATABOLIC_PROCESS, GOBP_L_SERINE_CATABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
MIKKELSEN_PLURIPOTENT_STATE_DN, MIKKELSEN_PLURIPOTENT_STATE_DN
MIR602, MIR602
PBRM1_TARGET_GENES, PBRM1_TARGET_GENES
GOBP_ADP_TRANSPORT, GOBP_ADP_TRANSPORT
GOCC_CENTRAL_ELEMENT, GOCC_CENTRAL_ELEMENT
GOMF_ADP_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_ADP_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_CDP_ALCOHOL_PHOSPHATIDYLTRANSFERASE_ACTIVITY, GOMF_CDP_ALCOHOL_PHOSPHATIDYLTRANSFERASE_ACTIVITY
BIOCARTA_KREB_PATHWAY, BIOCARTA_KREB_PATHWAY
GOBP_CDP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_CDP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS
GOMF_PHOSPHATIDATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDATE_PHOSPHATASE_ACTIVITY
INAMURA_LUNG_CANCER_SCC_UP, INAMURA_LUNG_CANCER_SCC_UP
GSE13485_CTRL_VS_DAY21_YF17D_VACCINE_PBMC_DN, GSE13485_CTRL_VS_DAY21_YF17D_VACCINE_PBMC_DN
WEBER_METHYLATED_LCP_IN_FIBROBLAST_DN, WEBER_METHYLATED_LCP_IN_FIBROBLAST_DN
HP_REDUCED_EJECTION_FRACTION, HP_REDUCED_EJECTION_FRACTION
GOCC_ASTER, GOCC_ASTER
REACTOME_TETRAHYDROBIOPTERIN_BH4_SYNTHESIS_RECYCLING_SALVAGE_AND_REGULATION, REACTOME_TETRAHYDROBIOPTERIN_BH4_SYNTHESIS_RECYCLING_SALVAGE_AND_REGULATION
HP_POST_PARTUM_HEMORRHAGE, HP_POST_PARTUM_HEMORRHAGE
HP_SHYNESS, HP_SHYNESS
GOBP_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION, GOBP_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION
GOCC_PLATELET_DENSE_TUBULAR_NETWORK, GOCC_PLATELET_DENSE_TUBULAR_NETWORK
GOMF_SINGLE_STRANDED_DNA_ENDODEOXYRIBONUCLEASE_ACTIVITY, GOMF_SINGLE_STRANDED_DNA_ENDODEOXYRIBONUCLEASE_ACTIVITY
GOMF_FK506_BINDING, GOMF_FK506_BINDING
REACTOME_SIGNALING_BY_ERBB2_ECD_MUTANTS, REACTOME_SIGNALING_BY_ERBB2_ECD_MUTANTS
PID_ERBB_NETWORK_PATHWAY, PID_ERBB_NETWORK_PATHWAY
REACTOME_CONSTITUTIVE_SIGNALING_BY_EGFRVIII, REACTOME_CONSTITUTIVE_SIGNALING_BY_EGFRVIII
HP_CUTANEOUS_SYNDACTYLY_OF_TOES, HP_CUTANEOUS_SYNDACTYLY_OF_TOES
HP_WIDOW_S_PEAK, HP_WIDOW_S_PEAK
GOBP_SIGNAL_PEPTIDE_PROCESSING, GOBP_SIGNAL_PEPTIDE_PROCESSING
GOBP_METHIONINE_BIOSYNTHETIC_PROCESS, GOBP_METHIONINE_BIOSYNTHETIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_OXIDATION, GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_OXIDATION
HP_SUBSARCOLEMMAL_ACCUMULATIONS_OF_ABNORMALLY_SHAPED_MITOCHONDRIA, HP_SUBSARCOLEMMAL_ACCUMULATIONS_OF_ABNORMALLY_SHAPED_MITOCHONDRIA
GOBP_VOCAL_LEARNING, GOBP_VOCAL_LEARNING
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION
GOBP_TRANSCRIPTION_DEPENDENT_TETHERING_OF_RNA_POLYMERASE_II_GENE_DNA_AT_NUCLEAR_PERIPHERY, GOBP_TRANSCRIPTION_DEPENDENT_TETHERING_OF_RNA_POLYMERASE_II_GENE_DNA_AT_NUCLEAR_PERIPHERY
GOBP_PROTECTION_FROM_NATURAL_KILLER_CELL_MEDIATED_CYTOTOXICITY, GOBP_PROTECTION_FROM_NATURAL_KILLER_CELL_MEDIATED_CYTOTOXICITY
HP_RIDGED_NAIL, HP_RIDGED_NAIL
FOURATI_BLOOD_TWINRIX_AGE_65_81YO_RESPONDERS_VS_POOR_RESPONDERS_TRAINING_SET_0DY_NETWORK_INFERENCE_DN, FOURATI_BLOOD_TWINRIX_AGE_65_81YO_RESPONDERS_VS_POOR_RESPONDERS_TRAINING_SET_0DY_NETWORK_INFERENCE_DN
GOBP_EXOCRINE_PANCREAS_DEVELOPMENT, GOBP_EXOCRINE_PANCREAS_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_BLOOD_BRAIN_BARRIER_PERMEABILITY, GOBP_POSITIVE_REGULATION_OF_BLOOD_BRAIN_BARRIER_PERMEABILITY
GOBP_REGULATION_OF_LIPOPHAGY, GOBP_REGULATION_OF_LIPOPHAGY
HP_ABNORMAL_CIRCULATING_AMYLASE_LEVEL, HP_ABNORMAL_CIRCULATING_AMYLASE_LEVEL
GOBP_CITRATE_METABOLIC_PROCESS, GOBP_CITRATE_METABOLIC_PROCESS
GOBP_PYRIMIDINE_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS
GOBP_MODULATION_BY_SYMBIONT_OF_HOST_APOPTOTIC_PROCESS, GOBP_MODULATION_BY_SYMBIONT_OF_HOST_APOPTOTIC_PROCESS
GOBP_REGULATION_OF_COHESIN_LOADING, GOBP_REGULATION_OF_COHESIN_LOADING
HP_LOSS_OF_SUBCUTANEOUS_ADIPOSE_TISSUE_IN_LIMBS, HP_LOSS_OF_SUBCUTANEOUS_ADIPOSE_TISSUE_IN_LIMBS
HP_HYPERMETHIONINEMIA, HP_HYPERMETHIONINEMIA
HP_MONOCYTOSIS, HP_MONOCYTOSIS
HP_FATIGABLE_WEAKNESS_OF_DISTAL_LIMB_MUSCLES, HP_FATIGABLE_WEAKNESS_OF_DISTAL_LIMB_MUSCLES
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMATIN, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMATIN
ABDULRAHMAN_KIDNEY_CANCER_VHL_DN, ABDULRAHMAN_KIDNEY_CANCER_VHL_DN
GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS
GOMF_DNA_BINDING_BENDING, GOMF_DNA_BINDING_BENDING
HP_CHOLANGITIS, HP_CHOLANGITIS