CD4_TCELL_VS_12H_ACT_TH2_DN, GSE22886_NAIVE_CD4_TCELL_VS_12H_ACT_TH2_DN
GSE26727_WT_VS_KLF2_KO_LPS_STIM_MACROPHAGE_DN, GSE26727_WT_VS_KLF2_KO_LPS_STIM_MACROPHAGE_DN
GSE7568_IL4_VS_IL4_AND_TGFB_TREATED_MACROPHAGE_24H_DN, GSE7568_IL4_VS_IL4_AND_TGFB_TREATED_MACROPHAGE_24H_DN
GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_DN, GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_DN
GSE22886_NAIVE_CD4_TCELL_VS_MEMORY_TCELL_DN, GSE22886_NAIVE_CD4_TCELL_VS_MEMORY_TCELL_DN
GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE22886_NAIVE_CD8_TCELL_VS_NEUTROPHIL_UP, GSE22886_NAIVE_CD8_TCELL_VS_NEUTROPHIL_UP
GSE22886_CD8_VS_CD4_NAIVE_TCELL_UP, GSE22886_CD8_VS_CD4_NAIVE_TCELL_UP
GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_UP, GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_UP
GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN, GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN
GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_DN
GSE17974_0H_VS_72H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_72H_IN_VITRO_ACT_CD4_TCELL_DN
GSE24726_WT_VS_E2_2_KO_PDC_DAY4_POST_DELETION_UP, GSE24726_WT_VS_E2_2_KO_PDC_DAY4_POST_DELETION_UP
GSE3982_MAC_VS_NEUTROPHIL_LPS_STIM_UP, GSE3982_MAC_VS_NEUTROPHIL_LPS_STIM_UP
GSE17974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN
GSE10325_BCELL_VS_LUPUS_BCELL_DN, GSE10325_BCELL_VS_LUPUS_BCELL_DN
GSE36476_CTRL_VS_TSST_ACT_16H_MEMORY_CD4_TCELL_OLD_DN, GSE36476_CTRL_VS_TSST_ACT_16H_MEMORY_CD4_TCELL_OLD_DN
GSE22886_NAIVE_CD4_TCELL_VS_NKCELL_DN, GSE22886_NAIVE_CD4_TCELL_VS_NKCELL_DN
GSE17721_POLYIC_VS_PAM3CSK4_6H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_6H_BMDC_DN
GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP
GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN
GSE41867_DAY8_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_UP, GSE41867_DAY8_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_UP
GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_UP, GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_UP
GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP
GSE40273_XBP1_KO_VS_WT_TREG_UP, GSE40273_XBP1_KO_VS_WT_TREG_UP
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE37416_0H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_6H_DN, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_6H_DN
GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_DN
GSE10325_MYELOID_VS_LUPUS_MYELOID_DN, GSE10325_MYELOID_VS_LUPUS_MYELOID_DN
GSE27786_LIN_NEG_VS_CD4_TCELL_UP, GSE27786_LIN_NEG_VS_CD4_TCELL_UP
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELL_DN
GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_UP, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_UP
IVANOVA_HEMATOPOIESIS_INTERMEDIATE_PROGENITOR, IVANOVA_HEMATOPOIESIS_INTERMEDIATE_PROGENITOR
GSE6269_FLU_VS_STAPH_AUREUS_INF_PBMC_UP, GSE6269_FLU_VS_STAPH_AUREUS_INF_PBMC_UP
MTOR_UP.N4.V1_UP, MTOR_UP.N4.V1_UP
GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_DN, GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_DN
GSE1432_1H_VS_6H_IFNG_MICROGLIA_DN, GSE1432_1H_VS_6H_IFNG_MICROGLIA_DN
GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_UP, GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_UP
GSE17721_POLYIC_VS_CPG_24H_BMDC_DN, GSE17721_POLYIC_VS_CPG_24H_BMDC_DN
REACTOME_ANTIGEN_PROCESSING_CROSS_PRESENTATION, REACTOME_ANTIGEN_PROCESSING_CROSS_PRESENTATION
GSE36826_WT_VS_IL1R_KO_SKIN_DN, GSE36826_WT_VS_IL1R_KO_SKIN_DN
JISON_SICKLE_CELL_DISEASE_UP, JISON_SICKLE_CELL_DISEASE_UP
GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP, GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP
GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN, GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN
GSE17812_WT_VS_THPOK_KO_MEMORY_CD8_TCELL_UP, GSE17812_WT_VS_THPOK_KO_MEMORY_CD8_TCELL_UP
GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_UP, GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_UP
GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
GSE22886_DAY0_VS_DAY1_MONOCYTE_IN_CULTURE_DN, GSE22886_DAY0_VS_DAY1_MONOCYTE_IN_CULTURE_DN
REACTOME_REGULATION_OF_MRNA_STABILITY_BY_PROTEINS_THAT_BIND_AU_RICH_ELEMENTS, REACTOME_REGULATION_OF_MRNA_STABILITY_BY_PROTEINS_THAT_BIND_AU_RICH_ELEMENTS
GSE11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN, GSE11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN
GSE15659_RESTING_VS_ACTIVATED_TREG_DN, GSE15659_RESTING_VS_ACTIVATED_TREG_DN
GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_UP
GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_DN
ZHONG_SECRETOME_OF_LUNG_CANCER_AND_FIBROBLAST, ZHONG_SECRETOME_OF_LUNG_CANCER_AND_FIBROBLAST
MODULE_321, MODULE_321
REACTOME_DEGRADATION_OF_BETA_CATENIN_BY_THE_DESTRUCTION_COMPLEX, REACTOME_DEGRADATION_OF_BETA_CATENIN_BY_THE_DESTRUCTION_COMPLEX
GSE6875_WT_VS_FOXP3_KO_TREG_UP, GSE6875_WT_VS_FOXP3_KO_TREG_UP
HALLMARK_IL2_STAT5_SIGNALING, HALLMARK_IL2_STAT5_SIGNALING
GSE20715_WT_VS_TLR4_KO_24H_OZONE_LUNG_DN, GSE20715_WT_VS_TLR4_KO_24H_OZONE_LUNG_DN
GSE7509_DC_VS_MONOCYTE_DN, GSE7509_DC_VS_MONOCYTE_DN
GSE42088_UNINF_VS_LEISHMANIA_INF_DC_2H_UP, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_2H_UP
REACTOME_REGULATION_OF_HMOX1_EXPRESSION_AND_ACTIVITY, REACTOME_REGULATION_OF_HMOX1_EXPRESSION_AND_ACTIVITY
GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_DN
GSE24210_TCONV_VS_TREG_DN, GSE24210_TCONV_VS_TREG_DN
GSE1432_CTRL_VS_IFNG_6H_MICROGLIA_DN, GSE1432_CTRL_VS_IFNG_6H_MICROGLIA_DN
GSE22611_UNSTIM_VS_2H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_UP, GSE22611_UNSTIM_VS_2H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_UP
GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP, GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP
MEK_UP.V1_DN, MEK_UP.V1_DN
REACTOME_SIGNALING_BY_NOTCH4, REACTOME_SIGNALING_BY_NOTCH4
GOCC_ENDOPEPTIDASE_COMPLEX, GOCC_ENDOPEPTIDASE_COMPLEX
REACTOME_SIGNALING_BY_THE_B_CELL_RECEPTOR_BCR, REACTOME_SIGNALING_BY_THE_B_CELL_RECEPTOR_BCR
REACTOME_C_TYPE_LECTIN_RECEPTORS_CLRS, REACTOME_C_TYPE_LECTIN_RECEPTORS_CLRS
GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_DC_UP, GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_DC_UP
GSE26290_CTRL_VS_AKT_INHIBITOR_TREATED_ANTI_CD3_AND_IL2_STIM_CD8_TCELL_DN, GSE26290_CTRL_VS_AKT_INHIBITOR_TREATED_ANTI_CD3_AND_IL2_STIM_CD8_TCELL_DN
REACTOME_BETA_CATENIN_INDEPENDENT_WNT_SIGNALING, REACTOME_BETA_CATENIN_INDEPENDENT_WNT_SIGNALING
REACTOME_MAPK6_MAPK4_SIGNALING, REACTOME_MAPK6_MAPK4_SIGNALING
REACTOME_REGULATION_OF_PTEN_STABILITY_AND_ACTIVITY, REACTOME_REGULATION_OF_PTEN_STABILITY_AND_ACTIVITY
GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP
TOOKER_GEMCITABINE_RESISTANCE_DN, TOOKER_GEMCITABINE_RESISTANCE_DN
GOBP_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS, GOBP_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS
GSE12839_CTRL_VS_IL12_TREATED_PBMC_UP, GSE12839_CTRL_VS_IL12_TREATED_PBMC_UP
GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_UP
GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP
GOBP_NUCLEOTIDE_PHOSPHORYLATION, GOBP_NUCLEOTIDE_PHOSPHORYLATION
REACTOME_NEGATIVE_REGULATION_OF_NOTCH4_SIGNALING, REACTOME_NEGATIVE_REGULATION_OF_NOTCH4_SIGNALING
GSE42088_UNINF_VS_LEISHMANIA_INF_DC_8H_UP, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_8H_UP
GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA
GSE10325_CD4_TCELL_VS_LUPUS_CD4_TCELL_DN, GSE10325_CD4_TCELL_VS_LUPUS_CD4_TCELL_DN
MORF_GSPT1, MORF_GSPT1
REACTOME_DEGRADATION_OF_AXIN, REACTOME_DEGRADATION_OF_AXIN
GSE15930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_DN
GSE7219_WT_VS_NIK_NFKB2_KO_LPS_AND_ANTI_CD40_STIM_DC_DN, GSE7219_WT_VS_NIK_NFKB2_KO_LPS_AND_ANTI_CD40_STIM_DC_DN
GSE41867_NAIVE_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN
WP_PROTEASOME_DEGRADATION, WP_PROTEASOME_DEGRADATION
HALLMARK_INTERFERON_GAMMA_RESPONSE, HALLMARK_INTERFERON_GAMMA_RESPONSE
REACTOME_DECTIN_1_MEDIATED_NONCANONICAL_NF_KB_SIGNALING, REACTOME_DECTIN_1_MEDIATED_NONCANONICAL_NF_KB_SIGNALING
GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_DN, GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_DN
GSE17721_PAM3CSK4_VS_CPG_0.5H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_0.5H_BMDC_DN
GOBP_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY
HALLMARK_PROTEIN_SECRETION, HALLMARK_PROTEIN_SECRETION
GOBP_VESICLE_TARGETING, GOBP_VESICLE_TARGETING
REACTOME_DEGRADATION_OF_DVL, REACTOME_DEGRADATION_OF_DVL
WP_TNF_ALPHA_SIGNALING_PATHWAY, WP_TNF_ALPHA_SIGNALING_PATHWAY
REACTOME_CELLULAR_RESPONSE_TO_HYPOXIA, REACTOME_CELLULAR_RESPONSE_TO_HYPOXIA
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
MODULE_91, MODULE_91
REACTOME_DEFECTIVE_CFTR_CAUSES_CYSTIC_FIBROSIS, REACTOME_DEFECTIVE_CFTR_CAUSES_CYSTIC_FIBROSIS
REACTOME_AUF1_HNRNP_D0_BINDS_AND_DESTABILIZES_MRNA, REACTOME_AUF1_HNRNP_D0_BINDS_AND_DESTABILIZES_MRNA
PELLICCIOTTA_HDAC_IN_ANTIGEN_PRESENTATION_DN, PELLICCIOTTA_HDAC_IN_ANTIGEN_PRESENTATION_DN
REACTOME_PCP_CE_PATHWAY, REACTOME_PCP_CE_PATHWAY
GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN
REACTOME_HEDGEHOG_LIGAND_BIOGENESIS, REACTOME_HEDGEHOG_LIGAND_BIOGENESIS
REACTOME_REGULATION_OF_RUNX3_EXPRESSION_AND_ACTIVITY, REACTOME_REGULATION_OF_RUNX3_EXPRESSION_AND_ACTIVITY
REACTOME_STABILIZATION_OF_P53, REACTOME_STABILIZATION_OF_P53
MORF_BAG5, MORF_BAG5
ZHU_CMV_ALL_UP, ZHU_CMV_ALL_UP
GOBP_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY
GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN
GSE360_DC_VS_MAC_L_MAJOR_DN, GSE360_DC_VS_MAC_L_MAJOR_DN
MIR216B_3P, MIR216B_3P
GOBP_PROTEIN_MONOUBIQUITINATION, GOBP_PROTEIN_MONOUBIQUITINATION
GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP
MIR6757_3P, MIR6757_3P
JIANG_VHL_TARGETS, JIANG_VHL_TARGETS
NGUYEN_NOTCH1_TARGETS_DN, NGUYEN_NOTCH1_TARGETS_DN
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_4MONTH_POST_DX_DN, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_4MONTH_POST_DX_DN
REACTOME_ASYMMETRIC_LOCALIZATION_OF_PCP_PROTEINS, REACTOME_ASYMMETRIC_LOCALIZATION_OF_PCP_PROTEINS
GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS
REACTOME_CROSS_PRESENTATION_OF_SOLUBLE_EXOGENOUS_ANTIGENS_ENDOSOMES, REACTOME_CROSS_PRESENTATION_OF_SOLUBLE_EXOGENOUS_ANTIGENS_ENDOSOMES
GSE19512_NAUTRAL_VS_INDUCED_TREG_DN, GSE19512_NAUTRAL_VS_INDUCED_TREG_DN
MORI_PLASMA_CELL_UP, MORI_PLASMA_CELL_UP
GSE28130_ACTIVATED_VS_INDUCEED_TREG_UP, GSE28130_ACTIVATED_VS_INDUCEED_TREG_UP
GSE9316_IL6_KO_VS_IFNG_KO_INVIVO_EXPANDED_CD4_TCELL_DN, GSE9316_IL6_KO_VS_IFNG_KO_INVIVO_EXPANDED_CD4_TCELL_DN
REACTOME_METABOLISM_OF_POLYAMINES, REACTOME_METABOLISM_OF_POLYAMINES
GOCC_GOLGI_ASSOCIATED_VESICLE, GOCC_GOLGI_ASSOCIATED_VESICLE
REACTOME_TRANSCRIPTIONAL_REGULATION_BY_RUNX2, REACTOME_TRANSCRIPTIONAL_REGULATION_BY_RUNX2
GOCC_COATED_VESICLE_MEMBRANE, GOCC_COATED_VESICLE_MEMBRANE
GOBP_HISTONE_UBIQUITINATION, GOBP_HISTONE_UBIQUITINATION
GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_DN, GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_DN
REACTOME_COPI_MEDIATED_ANTEROGRADE_TRANSPORT, REACTOME_COPI_MEDIATED_ANTEROGRADE_TRANSPORT
GOBP_PROTEIN_HOMOOLIGOMERIZATION, GOBP_PROTEIN_HOMOOLIGOMERIZATION
GOBP_GLYCOSYL_COMPOUND_METABOLIC_PROCESS, GOBP_GLYCOSYL_COMPOUND_METABOLIC_PROCESS
GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT_MEMBRANE, GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT_MEMBRANE
IGARASHI_ATF4_TARGETS_DN, IGARASHI_ATF4_TARGETS_DN
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_1MONTH_POST_DX_DN, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_1MONTH_POST_DX_DN
GSE360_T_GONDII_VS_B_MALAYI_LOW_DOSE_DC_UP, GSE360_T_GONDII_VS_B_MALAYI_LOW_DOSE_DC_UP
REACTOME_DEGRADATION_OF_GLI1_BY_THE_PROTEASOME, REACTOME_DEGRADATION_OF_GLI1_BY_THE_PROTEASOME
GCM_IL6ST, GCM_IL6ST
HP_CUTANEOUS_PHOTOSENSITIVITY, HP_CUTANEOUS_PHOTOSENSITIVITY
GOBP_VESICLE_TARGETING_TO_FROM_OR_WITHIN_GOLGI, GOBP_VESICLE_TARGETING_TO_FROM_OR_WITHIN_GOLGI
WINTER_HYPOXIA_UP, WINTER_HYPOXIA_UP
REACTOME_REGULATION_OF_RUNX2_EXPRESSION_AND_ACTIVITY, REACTOME_REGULATION_OF_RUNX2_EXPRESSION_AND_ACTIVITY
WP_FERROPTOSIS, WP_FERROPTOSIS
MIR5006_5P, MIR5006_5P
REACTOME_ABC_TRANSPORTER_DISORDERS, REACTOME_ABC_TRANSPORTER_DISORDERS
GOBP_REGULATION_OF_CELLULAR_AMINE_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_AMINE_METABOLIC_PROCESS
REACTOME_COPII_MEDIATED_VESICLE_TRANSPORT, REACTOME_COPII_MEDIATED_VESICLE_TRANSPORT
MIR6837_3P, MIR6837_3P
GOBP_ESTABLISHMENT_OF_TISSUE_POLARITY, GOBP_ESTABLISHMENT_OF_TISSUE_POLARITY
GOBP_COPII_COATED_VESICLE_BUDDING, GOBP_COPII_COATED_VESICLE_BUDDING
GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS
GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP, GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP
GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN, GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN
GOBP_CELLULAR_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES, GOBP_CELLULAR_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES
GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE, GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE
GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
BUFFA_HYPOXIA_METAGENE, BUFFA_HYPOXIA_METAGENE
GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN, GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN
REACTOME_GLYCOLYSIS, REACTOME_GLYCOLYSIS
GSE16450_CTRL_VS_IFNA_12H_STIM_IMMATURE_NEURON_CELL_LINE_DN, GSE16450_CTRL_VS_IFNA_12H_STIM_IMMATURE_NEURON_CELL_LINE_DN
GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10NEG_CD62LPOS_BONE_MARROW_DN, GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10NEG_CD62LPOS_BONE_MARROW_DN
BOSCO_ALLERGEN_INDUCED_TH2_ASSOCIATED_MODULE, BOSCO_ALLERGEN_INDUCED_TH2_ASSOCIATED_MODULE
GOCC_VESICLE_COAT, GOCC_VESICLE_COAT
GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_DN, GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_DN
SESTO_RESPONSE_TO_UV_C8, SESTO_RESPONSE_TO_UV_C8
GSE22432_MULTIPOTENT_PROGENITOR_VS_CDC_UP, GSE22432_MULTIPOTENT_PROGENITOR_VS_CDC_UP
KEGG_PROTEASOME, KEGG_PROTEASOME
GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP
CUI_GLUCOSE_DEPRIVATION, CUI_GLUCOSE_DEPRIVATION
GCM_DLG1, GCM_DLG1
GSE2585_CD80_HIGH_VS_LOW_MTEC_DN, GSE2585_CD80_HIGH_VS_LOW_MTEC_DN
GOCC_PROTEASOME_ACCESSORY_COMPLEX, GOCC_PROTEASOME_ACCESSORY_COMPLEX
MARSON_FOXP3_TARGETS_DN, MARSON_FOXP3_TARGETS_DN
HP_COXA_VALGA, HP_COXA_VALGA
HP_METAPHYSEAL_WIDENING, HP_METAPHYSEAL_WIDENING
GOBP_HISTONE_H2A_UBIQUITINATION, GOBP_HISTONE_H2A_UBIQUITINATION
GOBP_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION, GOBP_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION
GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_DN, GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_DN
HALLMARK_ALLOGRAFT_REJECTION, HALLMARK_ALLOGRAFT_REJECTION
HP_SLENDER_LONG_BONE, HP_SLENDER_LONG_BONE
GOBP_MONOSACCHARIDE_CATABOLIC_PROCESS, GOBP_MONOSACCHARIDE_CATABOLIC_PROCESS
HALLMARK_HYPOXIA, HALLMARK_HYPOXIA
HP_FASCICULATIONS, HP_FASCICULATIONS
GOBP_ORGANOPHOSPHATE_CATABOLIC_PROCESS, GOBP_ORGANOPHOSPHATE_CATABOLIC_PROCESS
GOBP_REGULATION_OF_ANIMAL_ORGAN_MORPHOGENESIS, GOBP_REGULATION_OF_ANIMAL_ORGAN_MORPHOGENESIS
GSE13762_CTRL_VS_125_VITAMIND_DAY12_DC_DN, GSE13762_CTRL_VS_125_VITAMIND_DAY12_DC_DN
GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_DN, GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_DN
GOBP_CELLULAR_MODIFIED_AMINO_ACID_METABOLIC_PROCESS, GOBP_CELLULAR_MODIFIED_AMINO_ACID_METABOLIC_PROCESS
RB_DN.V1_DN, RB_DN.V1_DN
YAP1_UP, YAP1_UP
PROVENZANI_METASTASIS_DN, PROVENZANI_METASTASIS_DN
PURBEY_TARGETS_OF_CTBP1_AND_SATB1_UP, PURBEY_TARGETS_OF_CTBP1_AND_SATB1_UP
MIR3612, MIR3612
GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_TRANSFERASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_TRANSFERASE_ACTIVITY
REACTOME_DISEASES_ASSOCIATED_WITH_GLYCOSYLATION_PRECURSOR_BIOSYNTHESIS, REACTOME_DISEASES_ASSOCIATED_WITH_GLYCOSYLATION_PRECURSOR_BIOSYNTHESIS
MIR3154, MIR3154
GOMF_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY, GOMF_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY
GOBP_AMINE_METABOLIC_PROCESS, GOBP_AMINE_METABOLIC_PROCESS
SESTO_RESPONSE_TO_UV_C1, SESTO_RESPONSE_TO_UV_C1
HOFFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_DN, HOFFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_DN
MIR219A_1_3P, MIR219A_1_3P
VANASSE_BCL2_TARGETS_DN, VANASSE_BCL2_TARGETS_DN
HP_ABNORMALITY_OF_THE_GALLBLADDER, HP_ABNORMALITY_OF_THE_GALLBLADDER
GOCC_GOLGI_ASSOCIATED_VESICLE_MEMBRANE, GOCC_GOLGI_ASSOCIATED_VESICLE_MEMBRANE
MIR650, MIR650
GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_DN, GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_DN
MIR136_5P, MIR136_5P
LU_IL4_SIGNALING, LU_IL4_SIGNALING
GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY, GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY
GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP
MIR127_5P, MIR127_5P
PID_P75_NTR_PATHWAY, PID_P75_NTR_PATHWAY
LINDSTEDT_DENDRITIC_CELL_MATURATION_B, LINDSTEDT_DENDRITIC_CELL_MATURATION_B
MIR4686, MIR4686
BRUNO_HEMATOPOIESIS, BRUNO_HEMATOPOIESIS
MODULE_233, MODULE_233
GOCC_CHAPERONE_COMPLEX, GOCC_CHAPERONE_COMPLEX
DARWICHE_SQUAMOUS_CELL_CARCINOMA_UP, DARWICHE_SQUAMOUS_CELL_CARCINOMA_UP
GOMF_THYROID_HORMONE_RECEPTOR_BINDING, GOMF_THYROID_HORMONE_RECEPTOR_BINDING
GOBP_CELLULAR_METABOLIC_COMPOUND_SALVAGE, GOBP_CELLULAR_METABOLIC_COMPOUND_SALVAGE
REACTOME_INSERTION_OF_TAIL_ANCHORED_PROTEINS_INTO_THE_ENDOPLASMIC_RETICULUM_MEMBRANE, REACTOME_INSERTION_OF_TAIL_ANCHORED_PROTEINS_INTO_THE_ENDOPLASMIC_RETICULUM_MEMBRANE
BARRIER_COLON_CANCER_RECURRENCE_UP, BARRIER_COLON_CANCER_RECURRENCE_UP
XBP1_01, XBP1_01
SCHEIDEREIT_IKK_INTERACTING_PROTEINS, SCHEIDEREIT_IKK_INTERACTING_PROTEINS
REACTOME_INTEGRATION_OF_ENERGY_METABOLISM, REACTOME_INTEGRATION_OF_ENERGY_METABOLISM
MARSON_FOXP3_CORE_DIRECT_TARGETS, MARSON_FOXP3_CORE_DIRECT_TARGETS
GOCC_SNARE_COMPLEX, GOCC_SNARE_COMPLEX
GOBP_PURINE_NUCLEOSIDE_METABOLIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_METABOLIC_PROCESS
GSE9946_IMMATURE_VS_LISTERIA_INF_MATURE_DC_DN, GSE9946_IMMATURE_VS_LISTERIA_INF_MATURE_DC_DN
BIOCARTA_PROTEASOME_PATHWAY, BIOCARTA_PROTEASOME_PATHWAY
GNF2_CDC27, GNF2_CDC27
GOBP_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS, GOBP_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS
CCACACA_MIR147, CCACACA_MIR147
CASTELLANO_NRAS_TARGETS_UP, CASTELLANO_NRAS_TARGETS_UP
GSE3720_UNSTIM_VS_LPS_STIM_VD1_GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_LPS_STIM_VD1_GAMMADELTA_TCELL_DN
MODULE_21, MODULE_21
GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE, GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE
MIR3976, MIR3976
IRX3_TARGET_GENES, IRX3_TARGET_GENES
TBPL1_TARGET_GENES, TBPL1_TARGET_GENES
GOBP_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS
WP_PYRIMIDINE_METABOLISM_AND_RELATED_DISEASES, WP_PYRIMIDINE_METABOLISM_AND_RELATED_DISEASES
GOMF_DNA_POLYMERASE_BINDING, GOMF_DNA_POLYMERASE_BINDING
GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE, GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE
HP_CHOLELITHIASIS, HP_CHOLELITHIASIS
GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH, GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH
HAY_BONE_MARROW_CD34_POS_GRAN, HAY_BONE_MARROW_CD34_POS_GRAN
WNT_SIGNALING, WNT_SIGNALING
GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS, GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION
REACTOME_DISEASES_ASSOCIATED_WITH_N_GLYCOSYLATION_OF_PROTEINS, REACTOME_DISEASES_ASSOCIATED_WITH_N_GLYCOSYLATION_OF_PROTEINS
GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN, GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN
GOCC_ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE
MIR144_5P, MIR144_5P
PID_IL3_PATHWAY, PID_IL3_PATHWAY
GSE13411_IGM_VS_SWITCHED_MEMORY_BCELL_UP, GSE13411_IGM_VS_SWITCHED_MEMORY_BCELL_UP
GOBP_RIBONUCLEOTIDE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOTIDE_CATABOLIC_PROCESS
GOBP_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS, GOBP_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS
GOBP_METHIONINE_METABOLIC_PROCESS, GOBP_METHIONINE_METABOLIC_PROCESS
SCHEIDEREIT_IKK_TARGETS, SCHEIDEREIT_IKK_TARGETS
GOBP_PROTEASOME_ASSEMBLY, GOBP_PROTEASOME_ASSEMBLY
MIR621, MIR621
GOBP_MODULATION_BY_VIRUS_OF_HOST_PROCESS, GOBP_MODULATION_BY_VIRUS_OF_HOST_PROCESS
GOBP_HISTONE_H2A_MONOUBIQUITINATION, GOBP_HISTONE_H2A_MONOUBIQUITINATION
MIR4489, MIR4489
MODY_HIPPOCAMPUS_NEONATAL, MODY_HIPPOCAMPUS_NEONATAL
HP_MOTOR_AXONAL_NEUROPATHY, HP_MOTOR_AXONAL_NEUROPATHY
GOBP_POSITIVE_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_POSITIVE_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION
GOCC_COPII_VESICLE_COAT, GOCC_COPII_VESICLE_COAT
GOBP_PROTEASOMAL_UBIQUITIN_INDEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_PROTEASOMAL_UBIQUITIN_INDEPENDENT_PROTEIN_CATABOLIC_PROCESS
BIOCARTA_TPO_PATHWAY, BIOCARTA_TPO_PATHWAY
MIR9898, MIR9898
GEISS_RESPONSE_TO_DSRNA_UP, GEISS_RESPONSE_TO_DSRNA_UP
AIZARANI_LIVER_C8_RESIDENT_B_CELLS_1, AIZARANI_LIVER_C8_RESIDENT_B_CELLS_1
GOBP_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
MIR6813_5P, MIR6813_5P
GOBP_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY, GOBP_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY
GOBP_ALDITOL_METABOLIC_PROCESS, GOBP_ALDITOL_METABOLIC_PROCESS
GOCC_COPI_COATED_VESICLE, GOCC_COPI_COATED_VESICLE
GOBP_COPPER_ION_TRANSPORT, GOBP_COPPER_ION_TRANSPORT
REACTOME_IL_6_TYPE_CYTOKINE_RECEPTOR_LIGAND_INTERACTIONS, REACTOME_IL_6_TYPE_CYTOKINE_RECEPTOR_LIGAND_INTERACTIONS
HINATA_NFKB_TARGETS_KERATINOCYTE_UP, HINATA_NFKB_TARGETS_KERATINOCYTE_UP
MIR101_5P, MIR101_5P
PID_HIF1_TFPATHWAY, PID_HIF1_TFPATHWAY
GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS, GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS
GSE25677_R848_VS_MPL_AND_R848_STIM_BCELL_UP, GSE25677_R848_VS_MPL_AND_R848_STIM_BCELL_UP
MIR6085, MIR6085
GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS, GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS
HP_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET, HP_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET
HP_ACUTE_HEPATIC_FAILURE, HP_ACUTE_HEPATIC_FAILURE
HP_HYPOPLASIA_OF_DENTAL_ENAMEL, HP_HYPOPLASIA_OF_DENTAL_ENAMEL
GOBP_PYRIMIDINE_NUCLEOBASE_METABOLIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOBASE_METABOLIC_PROCESS
MIR6755_5P, MIR6755_5P
GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN
GNF2_TAL1, GNF2_TAL1
GOBP_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GOBP_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
DER_IFN_GAMMA_RESPONSE_DN, DER_IFN_GAMMA_RESPONSE_DN
HP_ABNORMALITY_OF_THE_WING_OF_THE_ILIUM, HP_ABNORMALITY_OF_THE_WING_OF_THE_ILIUM
GAZDA_DIAMOND_BLACKFAN_ANEMIA_PROGENITOR_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_PROGENITOR_UP
HP_MOTOR_POLYNEUROPATHY, HP_MOTOR_POLYNEUROPATHY
GOBP_ONE_CARBON_METABOLIC_PROCESS, GOBP_ONE_CARBON_METABOLIC_PROCESS
KAMMINGA_SENESCENCE, KAMMINGA_SENESCENCE
HP_PROTRUDING_TONGUE, HP_PROTRUDING_TONGUE
VANDESLUIS_COMMD1_TARGETS_GROUP_2_UP, VANDESLUIS_COMMD1_TARGETS_GROUP_2_UP
GOCC_PROTEASOME_REGULATORY_PARTICLE_BASE_SUBCOMPLEX, GOCC_PROTEASOME_REGULATORY_PARTICLE_BASE_SUBCOMPLEX
GOCC_FLEMMING_BODY, GOCC_FLEMMING_BODY
MIR4791, MIR4791
PID_LIS1_PATHWAY, PID_LIS1_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
REACTOME_OPIOID_SIGNALLING, REACTOME_OPIOID_SIGNALLING
DORMOY_ELAVL1_TARGETS, DORMOY_ELAVL1_TARGETS
HP_ABNORMALITY_OF_PRIMARY_TEETH, HP_ABNORMALITY_OF_PRIMARY_TEETH
WP_BIOMARKERS_FOR_PYRIMIDINE_METABOLISM_DISORDERS, WP_BIOMARKERS_FOR_PYRIMIDINE_METABOLISM_DISORDERS
MIR328_3P, MIR328_3P
GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY
GOBP_GUANOSINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_GUANOSINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
SMID_BREAST_CANCER_ERBB2_UP, SMID_BREAST_CANCER_ERBB2_UP
HWANG_PROSTATE_CANCER_MARKERS, HWANG_PROSTATE_CANCER_MARKERS
WP_EPO_RECEPTOR_SIGNALING, WP_EPO_RECEPTOR_SIGNALING
GOBP_UV_DAMAGE_EXCISION_REPAIR, GOBP_UV_DAMAGE_EXCISION_REPAIR
HP_TURRICEPHALY, HP_TURRICEPHALY
GOCC_PROTEASOME_CORE_COMPLEX, GOCC_PROTEASOME_CORE_COMPLEX
GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLVED_IN_
GOBP_SHORT_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GOBP_SHORT_CHAIN_FATTY_ACID_METABOLIC_PROCESS
GOBP_PURINE_NUCLEOSIDE_BIOSYNTHETIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_BIOSYNTHETIC_PROCESS
GOMF_DISORDERED_DOMAIN_SPECIFIC_BINDING, GOMF_DISORDERED_DOMAIN_SPECIFIC_BINDING
REACTOME_LATE_ENDOSOMAL_MICROAUTOPHAGY, REACTOME_LATE_ENDOSOMAL_MICROAUTOPHAGY
HP_BROAD_PALM, HP_BROAD_PALM
chr1q22, chr1q22
HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_HALLUX, HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_HALLUX
GOBP_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION
KEGG_PENTOSE_PHOSPHATE_PATHWAY, KEGG_PENTOSE_PHOSPHATE_PATHWAY
HP_SELF_MUTILATION, HP_SELF_MUTILATION
HP_CONGENITAL_ICHTHYOSIFORM_ERYTHRODERMA, HP_CONGENITAL_ICHTHYOSIFORM_ERYTHRODERMA
REACTOME_CLEC7A_DECTIN_1_INDUCES_NFAT_ACTIVATION, REACTOME_CLEC7A_DECTIN_1_INDUCES_NFAT_ACTIVATION
VANDESLUIS_COMMD1_TARGETS_GROUP_3_UP, VANDESLUIS_COMMD1_TARGETS_GROUP_3_UP
PID_IL5_PATHWAY, PID_IL5_PATHWAY
HP_COMPLETE_DUPLICATION_OF_PHALANX_OF_HAND, HP_COMPLETE_DUPLICATION_OF_PHALANX_OF_HAND
GOBP_RESPONSE_TO_NITROSATIVE_STRESS, GOBP_RESPONSE_TO_NITROSATIVE_STRESS
REACTOME_GROWTH_HORMONE_RECEPTOR_SIGNALING, REACTOME_GROWTH_HORMONE_RECEPTOR_SIGNALING
NICK_RESPONSE_TO_PROC_TREATMENT_DN, NICK_RESPONSE_TO_PROC_TREATMENT_DN
GNF2_PRDX2, GNF2_PRDX2
HP_ECTOPIA_LENTIS, HP_ECTOPIA_LENTIS
HP_NORMOCYTIC_ANEMIA, HP_NORMOCYTIC_ANEMIA
GOBP_POST_EMBRYONIC_DEVELOPMENT, GOBP_POST_EMBRYONIC_DEVELOPMENT
HP_DIMINISHED_MOVEMENT, HP_DIMINISHED_MOVEMENT
MIR3943, MIR3943
GOBP_MODULATION_BY_SYMBIONT_OF_HOST_PROCESS, GOBP_MODULATION_BY_SYMBIONT_OF_HOST_PROCESS
GOBP_PYRIMIDINE_DIMER_REPAIR, GOBP_PYRIMIDINE_DIMER_REPAIR
HP_FOCAL_DYSTONIA, HP_FOCAL_DYSTONIA
REACTOME_PRE_NOTCH_PROCESSING_IN_GOLGI, REACTOME_PRE_NOTCH_PROCESSING_IN_GOLGI
GOBP_REGULATION_OF_EXOSOMAL_SECRETION, GOBP_REGULATION_OF_EXOSOMAL_SECRETION
REACTOME_ASSEMBLY_OF_THE_HIV_VIRION, REACTOME_ASSEMBLY_OF_THE_HIV_VIRION
GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT, GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT
GOBP_RESPONSE_TO_NUTRIENT, GOBP_RESPONSE_TO_NUTRIENT
HP_SMALL_EPIPHYSES, HP_SMALL_EPIPHYSES
GOBP_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GOBP_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_NEURON_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_NEURON_APOPTOTIC_PROCESS
GOBP_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS
HP_UVEITIS, HP_UVEITIS
GOBP_GALACTOSE_CATABOLIC_PROCESS, GOBP_GALACTOSE_CATABOLIC_PROCESS
HP_JOINT_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET, HP_JOINT_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET
GOBP_REGULATION_OF_HISTONE_UBIQUITINATION, GOBP_REGULATION_OF_HISTONE_UBIQUITINATION
GOBP_RIBONUCLEOSIDE_METABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_METABOLIC_PROCESS
GOMF_MHC_CLASS_IB_PROTEIN_BINDING, GOMF_MHC_CLASS_IB_PROTEIN_BINDING
VALK_AML_CLUSTER_11, VALK_AML_CLUSTER_11
MODULE_50, MODULE_50
HP_MUSCLE_FIBER_INCLUSION_BODIES, HP_MUSCLE_FIBER_INCLUSION_BODIES
HASLINGER_B_CLL_WITH_11Q23_DELETION, HASLINGER_B_CLL_WITH_11Q23_DELETION
KEGG_GALACTOSE_METABOLISM, KEGG_GALACTOSE_METABOLISM
GOMF_EUKARYOTIC_INITIATION_FACTOR_4E_BINDING, GOMF_EUKARYOTIC_INITIATION_FACTOR_4E_BINDING
GOBP_GLYCOLYTIC_PROCESS_THROUGH_FRUCTOSE_6_PHOSPHATE, GOBP_GLYCOLYTIC_PROCESS_THROUGH_FRUCTOSE_6_PHOSPHATE
KIM_LRRC3B_TARGETS, KIM_LRRC3B_TARGETS
GOBP_PURINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_PURINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS
REACTOME_REGULATION_OF_INSULIN_SECRETION, REACTOME_REGULATION_OF_INSULIN_SECRETION
GOBP_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_PARTICLE_METABOLIC_PROCESS, GOBP_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_PARTICLE_METABOLIC_PROCESS
GOBP_VESICLE_CARGO_LOADING, GOBP_VESICLE_CARGO_LOADING
GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS
GOBP_PROTEIN_TRIMERIZATION, GOBP_PROTEIN_TRIMERIZATION
BIOCARTA_TNFR2_PATHWAY, BIOCARTA_TNFR2_PATHWAY
HP_ABNORMALITY_OF_THE_ACHILLES_TENDON, HP_ABNORMALITY_OF_THE_ACHILLES_TENDON
HP_PERSISTENCE_OF_PRIMARY_TEETH, HP_PERSISTENCE_OF_PRIMARY_TEETH
GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS
GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS
REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS, REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS
KANG_FLUOROURACIL_RESISTANCE_UP, KANG_FLUOROURACIL_RESISTANCE_UP
GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN
MARSON_FOXP3_TARGETS_STIMULATED_UP, MARSON_FOXP3_TARGETS_STIMULATED_UP
GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GOMF_CARBOHYDRATE_KINASE_ACTIVITY, GOMF_CARBOHYDRATE_KINASE_ACTIVITY
GOMF_PHOSPHATIDYLINOSITOL_MONOPHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_MONOPHOSPHATE_PHOSPHATASE_ACTIVITY
WANG_ADIPOGENIC_GENES_REPRESSED_BY_SIRT1, WANG_ADIPOGENIC_GENES_REPRESSED_BY_SIRT1
ZNF433_TARGET_GENES, ZNF433_TARGET_GENES
KEGG_FRUCTOSE_AND_MANNOSE_METABOLISM, KEGG_FRUCTOSE_AND_MANNOSE_METABOLISM
WP_THE_HUMAN_IMMUNE_RESPONSE_TO_TUBERCULOSIS, WP_THE_HUMAN_IMMUNE_RESPONSE_TO_TUBERCULOSIS
HP_ADULT_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT, HP_ADULT_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT
GOBP_DTMP_METABOLIC_PROCESS, GOBP_DTMP_METABOLIC_PROCESS
DPPA3_TARGET_GENES, DPPA3_TARGET_GENES
YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UP, YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UP
GOBP_PURINE_CONTAINING_COMPOUND_SALVAGE, GOBP_PURINE_CONTAINING_COMPOUND_SALVAGE
REACTOME_PURINE_SALVAGE, REACTOME_PURINE_SALVAGE