

GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_DN
GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_4H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_4H_CD4_TCELL_DN
GSE28726_ACT_CD4_TCELL_VS_ACT_VA24NEG_NK1CELL_UP, GSE28726_ACT_CD4_TCELL_VS_ACT_VA24NEG_NK1CELL_UP
GSE26727_WT_VS_KLRG1_KO_LPS_STIM_MACROPHAGE_DN, GSE26727_WT_VS_KLRG1_KO_LPS_STIM_MACROPHAGE_DN
GSE4666_INSTIM_VS_CD40L_IL12_IL5_1DAY_STIMULATED_IRF4_KO_BCELL_UP, GSE4666_INSTIM_VS_CD40L_IL12_IL5_1DAY_STIMULATED_IRF4_KO_BCELL_UP
GSE22886_CD8_VS_CD4_NAIVE_TCELL_UP, GSE22886_CD8_VS_CD4_NAIVE_TCELL_UP
GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP
GSE22140_HEALTHY_VS_ATHRITIC_GERMFREE_MOUSE_CD4_TCELL_DN, GSE22140_HEALTHY_VS_ATHRITIC_GERMFREE_MOUSE_CD4_TCELL_DN
GSE41176_WT_VS_TAK1_KO_ANTIL_IQM_STIM_BCELL_6H_DN, GSE41176_WT_VS_TAK1_KO_ANTIL_IQM_STIM_BCELL_6H_DN
GSE26618_BCELL_VS_MDC_DAY7_FLU_VACCINE_DN, GSE26618_BCELL_VS_MDC_DAY7_FLU_VACCINE_DN
GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_UP, GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_UP
GSE36826_WT_VS_IL1R_KO_SKIN_DN, GSE36826_WT_VS_IL1R_KO_SKIN_DN
GSE5099_MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_DN, GSE5099_MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_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
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE36826_NORMAL_VS_STAPH_AURUS_INF_SKIN_UP, GSE36826_NORMAL_VS_STAPH_AURUS_INF_SKIN_UP
GSE1925_CTRL_VS_3H_IFNG_STIM_MACROPHAGE_UP, GSE1925_CTRL_VS_3H_IFNG_STIM_MACROPHAGE_UP
GSE45739_NRAS_KO_VS_WT_UNSTIM_CD4_TCELL_UP, GSE45739_NRAS_KO_VS_WT_UNSTIM_CD4_TCELL_UP
GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_DN
GSE3039_ALPHABETA_CD8_TCELL_VS_B1_BCELL_DN, GSE3039_ALPHABETA_CD8_TCELL_VS_B1_BCELL_DN
GSE2949_MICROGLIA_BRAIN_VS_MONOCYTE_BONE_MARROW_DN, GSE2949_MICROGLIA_BRAIN_VS_MONOCYTE_BONE_MARROW_DN
GSE360_DC_VS_MAC_UP, GSE360_DC_VS_MAC_UP
GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_6H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_6H_CD4_TCELL_DN
GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_MAC_UP, GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_MAC_UP
GSE1432_CTRL_VS_IFNG_24H_MICROGLIA_UP, GSE1432_CTRL_VS_IFNG_24H_MICROGLIA_UP
GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY40_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY40_DN
GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS
GSE539_EX_VIVO_VS_IN_CULTURE_CDPOS_DC_DN, GSE539_EX_VIVO_VS_IN_CULTURE_CDPOS_DC_DN
GSE24726_WT_VS_E2_2_KO_PDC_DN, GSE24726_WT_VS_E2_2_KO_PDC_DN
GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_DN, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_DN
GSE42088_UNINF_VS_LEISHMANIA_INF_DC_24H_DN, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_24H_DN
GSE7509_DC_VS_MONOCYTE_DN, GSE7509_DC_VS_MONOCYTE_DN
GSE3692_TYPE_2_MYELOID_VS_FOSINOPHIL_IL25_TREATED_LUNG_DN, GSE3692_TYPE_2_MYELOID_VS_FOSINOPHIL_IL25_TREATED_LUNG_DN
GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN
GSE360_HIGH_VS_LOW_DOSE_B_MALAYI_MAC_UP, GSE360_HIGH_VS_LOW_DOSE_B_MALAYI_MAC_UP
GSE45881_CXCR6H_VS_CXCR1LO_COLONIC_LAMINA_PROPRIA_DN, GSE45881_CXCR6H_VS_CXCR1LO_COLONIC_LAMINA_PROPRIA_DN
GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_IKAROS_KO_DN, GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_IKAROS_KO_DN
HP_ABNORMALITY_OF_THE_LYMPH_NODES, HP_ABNORMALITY_OF_THE_LYMPH_NODES
GSE37801_COMMON_LYMPHOID_PROGENITOR_VS_CD4_TCELL_UP, GSE37801_COMMON_LYMPHOID_PROGENITOR_VS_CD4_TCELL_UP
GSE28618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MDC_DN, GSE28618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MDC_DN
FERNANDEZ_BOUND_BY_MYC, FERNANDEZ_BOUND_BY_MYC
GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_HIGH_MONOCYTE_UP, GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_HIGH_MONOCYTE_UP
REACTOME_TOLL_LIKE_RECEPTOR_CASCADES, REACTOME_TOLL_LIKE_RECEPTOR_CASCADES
HP_HYPOPIGMENTATION_OF_THE_SKIN, HP_HYPOPIGMENTATION_OF_THE_SKIN
GSE17974_IL4_AND_ANTIL_IL12_VS_UNTREATED_48H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTIL_IL12_VS_UNTREATED_48H_ACT_CD4_TCELL_UP
GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_UP, GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_UP
ZHENG_CORD_BLOOD_C2_PUTATIVE_BASOPHIL_EOSINOPHIL_MAST_CELL_PROGENITOR, ZHENG_CORD_BLOOD_C2_PUTATIVE_BASOPHIL_EOSINOPHIL_MAST_CELL_PROGENITOR
QL_HYPOXIA, QL_HYPOXIA
GSE41978_WT_VS_ID2_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_WT_VS_ID2_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
GSE4666_IRF4HIGH_VS_WT_CD40L_IL12_IL5_1DAY_STIMULATED_BCELL_DN, GSE4666_IRF4HIGH_VS_WT_CD40L_IL12_IL5_1DAY_STIMULATED_BCELL_DN
GSE41978_WT_VS_BM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_WT_VS_BM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
GOBP_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY
GSE360_T_GONDIL_VS_B_MALAYI_LOW_DOSE_DC_UP, GSE360_T_GONDIL_VS_B_MALAYI_LOW_DOSE_DC_UP
GOBP_ALPHA_AMINO_ACID_METABOLIC_PROCESS, GOBP_ALPHA_AMINO_ACID_METABOLIC_PROCESS
CYCLIN_D1_KE_V1_UP, CYCLIN_D1_KE_V1_UP
GSE360_DC_VS_MAC_M_TUBERCULOSIS_UP, GSE360_DC_VS_MAC_M_TUBERCULOSIS_UP
GSE28130_ACTIVATED_VS_INDUCED_TREG_UP, GSE28130_ACTIVATED_VS_INDUCED_TREG_UP
GALINDO_IMMUNE_RESPONSE_TO_ENTEROTOXIN, GALINDO_IMMUNE_RESPONSE_TO_ENTEROTOXIN
GSE360_L_DONOVANI_VS_L_MAJOR_DC_UP, GSE360_L_DONOVANI_VS_L_MAJOR_DC_UP
FLETCHER_PBMc_BCG_10W_INFANT_BCG_STIMULATED_VS_UNSTIMULATED_10W_UP, FLETCHER_PBMc_BCG_10W_INFANT_BCG_STIMULATED_VS_UNSTIMULATED_10W_UP
LU_AGINC_BRAIN_DN, LU_AGINC_BRAIN_DN
HP_RECURRENT_FRACTURES, HP_RECURRENT_FRACTURES
GCCNBNWCTTYS_UNKNOWN, GCCNBNWCTTYS_UNKNOWN
MIR6133, MIR6133
GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_BIOSYNTHETIC_PROCESS
MIR6129, MIR6129
GSE4365_WT_VS_IPNAR_KO_BCELL_UP, GSE4365_WT_VS_IPNAR_KO_BCELL_UP
REACTOME_DUAL_INCISION_IN_GG_NER, REACTOME_DUAL_INCISION_IN_GG_NER
HP_INCREASED_BONE_MINERAL_DENSITY, HP_INCREASED_BONE_MINERAL_DENSITY
MIR6130, MIR6130
GSE22501_PERIPHERAL_BLOOD_VS_CORD_BLOOD_TREG_DN, GSE22501_PERIPHERAL_BLOOD_VS_CORD_BLOOD_TREG_DN
MIR4510, MIR4510
MIR600, MIR600
ZHAN_MULTIPLE_MYELOMA_UP, ZHAN_MULTIPLE_MYELOMA_UP
GOBP_GLYCOSYL_COMPOUND_METABOLIC_PROCESS, GOBP_GLYCOSYL_COMPOUND_METABOLIC_PROCESS
GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN
MENON_FETAL_KIDNEY_6_COLLECTING_DUCT_CELLS, MENON_FETAL_KIDNEY_6_COLLECTING_DUCT_CELLS
REACTOME_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_NLR, REACTOME_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_NLR
REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION, REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION
GOBP_ACTIVATION_OF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOBP_ACTIVATION_OF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN
GOBP_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR, GOBP_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR
WP_APOPTOSIS, WP_APOPTOSIS
GOBP_TRANSLESION_SYNTHESIS, GOBP_TRANSLESION_SYNTHESIS
LE_EGR2_TARGETS_DN, LE_EGR2_TARGETS_DN
REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE, REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE
GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
GATA2_I1, GATA2_I1
TSENG_IRS1_TARGETS_DN, TSENG_IRS1_TARGETS_DN
GOBP_NUCLEOSIDE_PHOSPHATE_CATABOLIC_PROCESS, GOBP_NUCLEOSIDE_PHOSPHATE_CATABOLIC_PROCESS
MIR4709_5P, MIR4709_5P
TC7GAC, MIR198, TC7GAC, MIR198
GOBP_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_ALPHA_BETA_T_CELL_ACTIVATION
TRAVAGLINI_LUNG_ALVEOLAR_EPITHELIAL_TYPE_2_CELL, TRAVAGLINI_LUNG_ALVEOLAR_EPITHELIAL_TYPE_2_CELL
REACTOME_GAP_FILLING_DNA_REPAIR_SYNTHESIS_AND_LIGATION_IN_GG_NER, REACTOME_GAP_FILLING_DNA_REPAIR_SYNTHESIS_AND_LIGATION_IN_GG_NER
GSE28783_ANTL_MIR33_VS_CTRL_ATHEROSCLEROSIS_MACROPHAGE_UP, GSE28783_ANTL_MIR33_VS_CTRL_ATHEROSCLEROSIS_MACROPHAGE_UP
MIR6851_5P, MIR6851_5P
WP_TCELL_ANTIGEN_RECEPTOR_TCR_SIGNALING_PATHWAY, WP_TCELL_ANTIGEN_RECEPTOR_TCR_SIGNALING_PATHWAY
HP_OPSITHOTONUS, HP_OPSITHOTONUS
GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING
HP_ABNORMAL_GLYCOSYLATION, HP_ABNORMAL_GLYCOSYLATION
GOBP_RRNA_TRANSCRIPTION, GOBP_RRNA_TRANSCRIPTION
MIR3689D, MIR3689D
KIM_GASTRIC_CANCER_CHEMOSENSITIVITY, KIM_GASTRIC_CANCER_CHEMOSENSITIVITY
PAPASPYRIDONOS_UNSTABLE_ATHEROSCLEROTIC_PLAQUE_UP, PAPASPYRIDONOS_UNSTABLE_ATHEROSCLEROTIC_PLAQUE_UP
HP_DEATH_IN_CHILDHOOD, HP_DEATH_IN_CHILDHOOD
GOBP_PURINE_NUCLEOSIDE_METABOLIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_METABOLIC_PROCESS
BECKER_TAMOXIFEN_RESISTANCE_DN, BECKER_TAMOXIFEN_RESISTANCE_DN
KIM_MYCN_AMPLIFICATION_TARGETS_UP, KIM_MYCN_AMPLIFICATION_TARGETS_UP
GOBP_REGULATION_OF_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_T_CELL_DIFFERENTIATION
HP_ABNORMAL_PROTEIN_GLYCOSYLATION, HP_ABNORMAL_PROTEIN_GLYCOSYLATION
GOBP_ERROR_PRONE_TRANSLATION_SYNTHESIS, GOBP_ERROR_PRONE_TRANSLATION_SYNTHESIS
HP_ABNORMAL_TONGUE_PHYSIOLOGY, HP_ABNORMAL_TONGUE_PHYSIOLOGY
HP_ADDUCTED_THUMB, HP_ADDUCTED_THUMB
GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_DN, GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_DN
CTCANGCTGNY_UNKNOWN, CTCANGCTGNY_UNKNOWN
GOBP_CELLULAR_AMINO_ACID_CATABOLIC_PROCESS, GOBP_CELLULAR_AMINO_ACID_CATABOLIC_PROCESS
WESTON_VEGFA_TARGETS_3HR, WESTON_VEGFA_TARGETS_3HR
GOBP_INTERFERON_GAMMA_PRODUCTION, GOBP_INTERFERON_GAMMA_PRODUCTION
HARRIS_HYPOXIA, HARRIS_HYPOXIA
PID_TNF_PATHWAY, PID_TNF_PATHWAY
HP_PROURITUS, HP_PROURITUS
HP_RELATIVE_MACROCEPHALY, HP_RELATIVE_MACROCEPHALY
GOBP_LIPID_STORAGE, GOBP_LIPID_STORAGE
OUILLETTE_CL1_13Q14_DELETION_DN, OUILLETTE_CL1_13Q14_DELETION_DN
HP_NAIL_DYSTROPHY, HP_NAIL_DYSTROPHY
MIR4646_3P, MIR4646_3P
MIR1236_5P, MIR1236_5P
GOBP_INACTIVATION_OF_MAPK_ACTIVITY, GOBP_INACTIVATION_OF_MAPK_ACTIVITY
REACTOME_PURINERGIC_SIGNALING_IN_LEISHMANIASIS_INFECTION, REACTOME_PURINERGIC_SIGNALING_IN_LEISHMANIASIS_INFECTION
HP_NEPHROPATHY, HP_NEPHROPATHY
GOBP_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION
HAL1MARK_IL6_JAK_STAT3_SIGNALING, HAL1MARK_IL6_JAK_STAT3_SIGNALING
HP_SEVERE_COMBINED_IMMUNODEFICIENCY, HP_SEVERE_COMBINED_IMMUNODEFICIENCY
REACTOME_REGULATION_OF_TNFR1_SIGNALING, REACTOME_REGULATION_OF_TNFR1_SIGNALING
HP_EPSIDIC_HYPERHIDROSIS, HP_EPSIDIC_HYPERHIDROSIS
MIR6888_5P, MIR6888_5P
IWANAGA_CARCIINOGENESIS_BY_KRAS_DN, IWANAGA_CARCIINOGENESIS_BY_KRAS_DN
SHAFFER_IRF4_MULTIPLE_MYELOMA_PROGRAM, SHAFFER_IRF4_MULTIPLE_MYELOMA_PROGRAM
HP_CELLULAR_IMMUNODEFICIENCY, HP_CELLULAR_IMMUNODEFICIENCY
QL_HYPOXIA_TARGETS_OF_HIF1A_AND_FOXA2, QL_HYPOXIA_TARGETS_OF_HIF1A_AND_FOXA2
GOBP_REGULATION_OF_CELL_KILLING, GOBP_REGULATION_OF_CELL_KILLING
GOBP_ALPHA_AMINO_ACID_CATABOLIC_PROCESS, GOBP_ALPHA_AMINO_ACID_CATABOLIC_PROCESS
GOBP_CELLULAR_EXTRAVASATION, GOBP_CELLULAR_EXTRAVASATION
BICARTA_DEATH_PATHWAY, BICARTA_DEATH_PATHWAY
PID_HIV_NEP_PATHWAY, PID_HIV_NEP_PATHWAY
BICARTA_TNFR1_PATHWAY, BICARTA_TNFR1_PATHWAY
HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_CORTICOSPINAL_TRACT, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_CORTICOSPINAL_TRACT
GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
MODULE_265, MODULE_265
GOBP_PLACENTA_DEVELOPMENT, GOBP_PLACENTA_DEVELOPMENT
GOBP_RIBONUCLEOTIDE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOTIDE_CATABOLIC_PROCESS
CYCLIN_D1_KE_V1_DN, CYCLIN_D1_KE_V1_DN
GOBP_REGULATION_OF_OXIDATIVE_PHOSPHORYLATION, GOBP_REGULATION_OF_OXIDATIVE_PHOSPHORYLATION
HP_SUPRANUCLEAR_GAZE_PALSY, HP_SUPRANUCLEAR_GAZE_PALSY
MODULE_138, MODULE_138
HP_IMPAIRED_SMOOTH_PUJSUIT, HP_IMPAIRED_SMOOTH_PUJSUIT
HP_ABNORMAL_METAPHYSICAL TRABECULATION, HP_ABNORMAL_METAPHYSICAL TRABECULATION
GOBP_CELLULAR_METABOLIC_COMPOUND_SALVAGE, GOBP_CELLULAR_METABOLIC_COMPOUND_SALVAGE
GOBP_RESPONSE_TO_TESTOSTERONE, GOBP_RESPONSE_TO_TESTOSTERONE
GOBP_CELL_KILLING, GOBP_CELL_KILLING
WP_UREA_CYCLE_AND_METABOLISM_OF_AMINO_GROUPS, WP_UREA_CYCLE_AND_METABOLISM_OF_AMINO_GROUPS
MIR6065, MIR6065
GOBP_T_CELL_MEDIATED_CYTOTOXICITY, GOBP_T_CELL_MEDIATED_CYTOTOXICITY
KYNG_DNA_DAMAGE_BY_GAMMA_RADIATION, KYNG_DNA_DAMAGE_BY_GAMMA_RADIATION
REACTOME_INFLAMMASOMES, REACTOME_INFLAMMASOMES
ZHONG_PFC_C8_UNKNOWN_NEUROD2_POS_INTERNEURON, ZHONG_PFC_C8_UNKNOWN_NEUROD2_POS_INTERNEURON
GOBP_NUCLEOTIDE_TRANSPORT, GOBP_NUCLEOTIDE_TRANSPORT
GOBP_POSITIVE_REGULATION_OF_PRL_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_POSITIVE_REGULATION_OF_PRL_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
GSE17974_IL4_AND_ANTIL_IL12_VS_UNTREATED_24H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTIL_IL12_VS_UNTREATED_24H_ACT_CD4_TCELL_UP
XIE_LT_HSC_SIPR3_OE_UP, XIE_LT_HSC_SIPR3_OE_UP
GEISS_RESPONSE_TO_DSRNA_UP, GEISS_RESPONSE_TO_DSRNA_UP
GOBP_POLYAMINE_METABOLIC_PROCESS, GOBP_POLYAMINE_METABOLIC_PROCESS
MCBRYAN_PUBERTAL_TGFB_TARGETS_DN, MCBRYAN_PUBERTAL_TGFB_TARGETS_DN
DACOSTA_ERCC3_ALLELE_VS_TTD_DN, DACOSTA_ERCC3_ALLELE_VS_TTD_DN
HP_GENERALIZED_HYPOPIGMENTATION_OF_HAIR, HP_GENERALIZED_HYPOPIGMENTATION_OF_HAIR
GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY, GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY
GOBP_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS, GOBP_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS
REACTOME_OVARIAN_TUMOR_DOMAIN_PROTEASES, REACTOME_OVARIAN_TUMOR_DOMAIN_PROTEASES
GOBP_RESPONSE_TO_INTERLEUKIN_18, GOBP_RESPONSE_TO_INTERLEUKIN_18
WENG_POR_TARGETS_GLOBAL_DN, WENG_POR_TARGETS_GLOBAL_DN
TSENG_ADIPOGENIC_POTENTIAL_DN, TSENG_ADIPOGENIC_POTENTIAL_DN
HP_LIMB_GIRDLE_MUSCLE_ATROPHY, HP_LIMB_GIRDLE_MUSCLE_ATROPHY
HP_CEREBRAL_EDEMA, HP_CEREBRAL_EDEMA
GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORT
HOWARD_MONOCYTE_INACT_MONOV_INFLUENZA_A_INDONESIA_05_2005_H5N1_AGE_18_49YO_IDY_DN, HOWARD_MONOCYTE_INACT_MONOV_INFLUENZA_A_INDONESIA_05_2005_H5N1_AGE_18_49YO_IDY_DN
REACTOME_THE_NLRP3_INFLAMMASOME, REACTOME_THE_NLRP3_INFLAMMASOME
GOBP_POSITIVE_REGULATION_OF_CELL_KILLING, GOBP_POSITIVE_REGULATION_OF_CELL_KILLING
REACTOME_PURINE_SALVAGE, REACTOME_PURINE_SALVAGE
REACTOME_INSULIN_PROCESSING, REACTOME_INSULIN_PROCESSING
WU_HDX_TARGETS_3_UP, WU_HDX_TARGETS_3_UP
GOBP_NEGATIVE_REGULATION_OF_WOUND_HEALING, GOBP_NEGATIVE_REGULATION_OF_WOUND_HEALING
CAFFARELLI_RESPONSE_TO_THC_3HR_3_DN, CAFFARELLI_RESPONSE_TO_THC_3HR_3_DN
GOME_CARBON_NITROGEN_LYASE_ACTIVITY, GOME_CARBON_NITROGEN_LYASE_ACTIVITY
DURANTE_ADULT_OLFATORY_NEUROEPITHELIUM_RESPIRATORY_HORIZONTAAL_BASAL_CELLS, DURANTE_ADULT_OLFATORY_NEUROEPITHELIUM_RESPIRATORY_HORIZONTAAL_BASAL_CELLS
MIR296_3P, MIR296_3P
GOCC_EUKARYOTIC_TRANSLATION_INITIATION_FACTOR_4F_COMPLEX, GOCC_EUKARYOTIC_TRANSLATION_INITIATION_FACTOR_4F_COMPLEX
BICARTA_MHC_PATHWAY, BICARTA_MHC_PATHWAY
LOPES_METHYLATED_IN_COLON_CANCER_DN, LOPES_METHYLATED_IN_COLON_CANCER_DN
HP_PELVIC_GIRDLE_MUSCLE_WEAKNESS, HP_PELVIC_GIRDLE_MUSCLE_WEAKNESS
GOBP_POLYAMINE_BIOSYNTHETIC_PROCESS, GOBP_POLYAMINE_BIOSYNTHETIC_PROCESS
GOBP_INTERLEUKIN_18_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_18_MEDIATED_SIGNALING_PATHWAY
REACTOME_TNFR1_INDUCED_PROAPOPTOTIC_SIGNALING, REACTOME_TNFR1_INDUCED_PROAPOPTOTIC_SIGNALING
HP_ESOPHAGEAL_STENOSIS, HP_ESOPHAGEAL_STENOSIS
GOBP_PHOSPHATIDYLCHOLINE_ACYL_CHAIN_REMODELING, GOBP_PHOSPHATIDYLCHOLINE_ACYL_CHAIN_REMODELING
GOBP_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION
MODULE_73, MODULE_73
MODULE_322, MODULE_322
MODULE_528, MODULE_528
HP_ORAL_LEUKOPLAKIA, HP_ORAL_LEUKOPLAKIA
GOME_MAP_KINASE_PHOSPHATASE_ACTIVITY, GOME_MAP_KINASE_PHOSPHATASE_ACTIVITY
GOBP_METAL_ION_EXPORT, GOBP_METAL_ION_EXPORT
GOBP_RESPONSE_TO_NITRIC_OXIDE, GOBP_RESPONSE_TO_NITRIC_OXIDE
GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS
HP_ABNORMALITY_OF_SMALL_INTESTINAL_VILLUS_MORPHOLOGY, HP_ABNORMALITY_OF_SMALL_INTESTINAL_VILLUS_MORPHOLOGY
MODULE_78, MODULE_78
MODULE_471, MODULE_471
COLLIS_PRKDC_REGULATORS, COLLIS_PRKDC_REGULATORS
REACTOME_REGULATION_BY_C_FLIP, REACTOME_REGULATION_BY_C_FLIP
GOME_THIOESTERASE_BINDING, GOME_THIOESTERASE_BINDING
GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
MAP2K1_TARGET_GENIS, MAP2K1_TARGET_GENIS
HP_ABNORMALITY_OF_HAND_JOINT_MOBILITY, HP_ABNORMALITY_OF_HAND_JOINT_MOBILITY
HP_GENERALIZED_HYPOPIGMENTATION, HP_GENERALIZED_HYPOPIGMENTATION
GOME_NADPLUS_NUCLEOSIDASE_ACTIVITY, GOME_NADPLUS_NUCLEOSIDASE_ACTIVITY
GOBP_CHEMICAL_HOMEOSTASIS_WITHIN_A_TISSUE, GOBP_CHEMICAL_HOMEOSTASIS_WITHIN_A_TISSUE
GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE
GOBP_HEMATOPOIETIC STEM_CELL_PROLIFERATION, GOBP_HEMATOPOIETIC STEM_CELL_PROLIFERATION
GOBP_CELLULAR_BIOGENIC_AMINE_METABOLIC_PROCESS, GOBP_CELLULAR_BIOGENIC_AMINE_METABOLIC_PROCESS
HP_SKIN_VESICLE, HP_SKIN_VESICLE
REACTOME_CASPASE_ACTIVATION_VIA_DEATH_RECEPTORS_IN_THE_PRESENCE_OF_LIGAND, REACTOME_CASPASE_ACTIVATION_VIA_DEATH_RECEPTORS_IN_THE_PRESENCE_OF_LIGAND
GOBP_SPERMIDINE_METABOLIC_PROCESS, GOBP_SPERMIDINE_METABOLIC_PROCESS
GOBP_ACTIVATION_OF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GOBP_ACTIVATION_OF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY
GOBP_REGULATION_OF_PLASMINOGEN_ACTIVATION, GOBP_REGULATION_OF_PLASMINOGEN_ACTIVATION
BUCKANOVICH_T_LYMPHOCYTE_HOMING_ON_TUMOR_UP, BUCKANOVICH_T_LYMPHOCYTE_HOMING_ON_TUMOR_UP
WP_NANOMATERIAL_INDUCED_APOPTOSIS, WP_NANOMATERIAL_INDUCED_APOPTOSIS
SHIN_B_CELL_LYMPHOMA_CLUSTER_3, SHIN_B_CELL_LYMPHOMA_CLUSTER_3
MCMURRAY_TP53_HRAS_COOPERATION_RESPONSE_UP, MCMURRAY_TP53_HRAS_COOPERATION_RESPONSE_UP
GOBP_LACRIMAL_GLAND_DEVELOPMENT, GOBP_LACRIMAL_GLAND_DEVELOPMENT
GOBP_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS, GOBP_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS
GOME_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GOME_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY
GOBP_REGULATION_OF_T_HELPER_1_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_T_HELPER_1_CELL_DIFFERENTIATION
GOBP_XENOPHAGY, GOBP_XENOPHAGY
ERWIN_COHEN_PBMc_TC_83_AGE_18_45YO_NAIVE_NOT_PREVIOUSLY_IMMUNIZED_24HR_DEC, ERWIN_COHEN_PBMc_TC_83_AGE_18_45YO_NAIVE_NOT_PREVIOUSLY_IMMUNIZED_24HR_DEC
REACTOME_TRAIL_SIGNALING, REACTOME_TRAIL_SIGNALING
PID_IL12_STAT4_PATHWAY, PID_IL12_STAT4_PATHWAY
CUL_DEVELOPING_HEART_COMPACT_VENTRICULAR_CARDIOMYOCYTE, CUL_DEVELOPING_HEART_COMPACT_VENTRICULAR_CARDIOMYOCYTE
HP_HIGH_MYOPIA, HP_HIGH_MYOPIA
TSUNODA_CISPLATIN_RESISTANCE_UP, TSUNODA_CISPLATIN_RESISTANCE_UP
HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_UP, HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_UP
GOBP_POSITIVE_REGULATION_OF_T_HELPER_1_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_T_HELPER_1_CELL_DIFFERENTIATION
GOBP_INTERLEUKIN_1_BETA_PRODUCTION, GOBP_INTERLEUKIN_1_BETA_PRODUCTION
GOBP_NUCLEOSIDE_SALVAGE, GOBP_NUCLEOSIDE_SALVAGE
GOBP_POSITIVE_REGULATION_OF_LIPID_STORAGE, GOBP_POSITIVE_REGULATION_OF_LIPID_STORAGE
GOBP_RECURRENT_UPPER_AND_LOWER_RESPIRATORY_TRACT_INFECTIONS, HP_RECURRENT_UPPER_AND_LOWER_RESPIRATORY_TRACT_INFECTIONS
GOBP_REGULATION_OF_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_REGULATION_OF_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT
HP_CEREBRAL_PALSY, HP_CEREBRAL_PALSY
HP_ABNORMAL_DELAYED_HYPERSENSITIVITY_SKIN_TEST, HP_ABNORMAL_DELAYED_HYPERSENSITIVITY_SKIN_TEST
MODULE_545, MODULE_545
GOBP_RESPONSE_TO_THYROID_HORMONE, GOBP_RESPONSE_TO_THYROID_HORMONE
GOBP_RESPONSE_TO_INACTIVITY, GOBP_RESPONSE_TO_INACTIVITY
ZHONG_PFC_C6_PLXNA4_POS_EXCITATORY_NEURON, ZHONG_PFC_C6_PLXNA4_POS_EXCITATORY_NEURON
GOME_LIM_DOMAIN_BINDING, GOME_LIM_DOMAIN_BINDING