GOMF_SISTER_CHROMATID_SEGREGATION, GOMF_SISTER_CHROMATID_SEGREGATION
GSE16385_IMMATURE_VS_MATURE_NEURON_CELL_LINE_WEST_IQGENE_INC_VIRUS_DN, GSE16385_IMMATURE_VS_MATURE_NEURON_CELL_LINE_WEST_IQGENE_INC_VIRUS_DN
GSE17078_HS_VS_STIM_ICN_CD4_TCELL_DN, GSE17078_HS_VS_12H_IN_STIM_ICN_CD4_TCELL_DN
GSE21546_QC_BCELL_VS_PLASMA_CELL_UP, GSE21546_QC_BCELL_VS_PLASMA_CELL_UP
GSE1460_UP_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_UP, GSE1460_UP_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_UP
GOMF_SISTER_CHROMATID_SEGREGATION, GOMF_MITOTIC_SISTER_CHROMATID_SEGREGATION
GSE22321_EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN, GSE22321_EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN
GSE22886_NAIVE_CD4_TCELL_VS_MEM_CD4_TCELL_2H_STIM_PMA_IONO_DN, GSE22886_NAIVE_CD4_TCELL_VS_MEM_CD4_TCELL_FROM_LN_DN
GSE22886_NAIVE_CD4_TCELL_VS_48H_ACT_TH1_DN, GSE22886_NAIVE_CD4_TCELL_VS_48H_ACT_TH1_DN
GSE22327_ANTI_CD3_INVITRO_VS_UNTREATED_MOUSE_THY_DN, GSE22327_ANTI_CD3_INVIVO_VS_UNTREATED_MOUSE_THY_DN
GSE24634_NAIVE_CD4_TCELL_VS_DAY7_ILA_CONV_THY_DN, GSE24634_NAIVE_CD4_TCELL_VS_DAY7_ILA_CONV_THY_DN
GSE9962_NEUTROPHIL_VS_TH2_DN, GSE9962_NEUTROPHIL_VS_TH2_DN
GSE13485_NAIVE_VS_MEMORY_BCELL_UP, GSE13485_NAIVE_VS_MEMORY_BCELL_UP
GSE4370_NEAS_KO_VS_WT_LIVSTIM_CD4_TCELL_UP, GSE4370_NEAS_KO_VS_WT_LIVSTIM_CD4_TCELL_UP
GSE9962_MAC_VS_NEUTROPHIL_UP, GSE9962_MAC_VS_NEUTROPHIL_UP
WP_DNA_REPAIR_PATHWAYS_FULL_NETWORK, WP_DNA_REPAIR_PATHWAYS_FULL_NETWORK
GSE13765_UNSTIM_VS_PMMB_MTP_VS_BACG_KO_NK_CELL_UP, GSE13765_UNSTIM_PMMB_MTP_VS_BACG_KO_NK_CELL_UP
GOCC_PEROXISOMAL_GOCC_PEROXISOME
GSE9962_MAC_VS_NEUTROPHIL_DN, GSE9962_MAC_VS_NEUTROPHIL_DN
GO:0140130, NAIVE_VS_MEMORY_CD8_TCELL_UP, GO:0140130, NAIVE_VS_MEMORY_CD8_TCELL_UP
GSE9962_CTRL_VS_LPS_48_MAC_UP, GSE9962_CTRL_VS_LPS_48_MAC_UP
GSE17073_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_UP, GSE17073_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY4_DN
GSE17073_PAMOKSA_VS_GTC_4H_BMDC_UP, GSE17073_PAMOKSA_VS_GTC_4H_BMDC_UP
GSE15720_RESTING_VS_HYSTANDER_ACTIVATED_CD4_TCELL_DN, GSE15720_RESTING_VS_HYSTANDER_ACTIVATED_CD4_TCELL_DN
GSE6269_1H20_POS_DC_VS_CD4_TCELL_UP, GSE6269_1H20_POS_DC_VS_CD4_TCELL_UP
GSE17073_LPS_VS_PAMOKSA_24H_BMDC_DN, GSE17073_LPS_VS_PAMOKSA_24H_BMDC_DN
GSE22229_RENAL_TRANSPLANT_VS_HEALTHY_PBMC_UP, GSE22229_RENAL_TRANSPLANT_VS_HEALTHY_PBMC_UP
GSE18346_INTHMYC_2_PROGENITOR_VS_CD4_THYMOCYTE_UP, GSE18346_INTHMYC_2_PROGENITOR_VS_CD4_THYMOCYTE_UP
GSE9962_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_DN, GSE9962_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_DN
GSE9962_MAC_VS_MEM_CD4_TCELL_DN, GSE9962_MAC_VS_MEM_CD4_TCELL_DN
GSE1461_NAIVE_VS_MEMORY_BCELL_DN, GSE1461_NAIVE_VS_MEMORY_BCELL_DN
GSE9962_MAST_CELL_VS_TH1_DN, GSE9962_MAST_CELL_VS_TH1_DN
GSE29618_RCELL_VS_PDC_UP, GSE29618_RCELL_VS_PDC_UP
GSE13485_CTRL_VS_WEST_IQGENE_INC_VIRUS_MATURE_NEURON_CELL_LINE_DN, GSE13485_CTRL_VS_WEST_IQGENE_INC_VIRUS_MATURE_NEURON_CELL_LINE_DN
GSE9468_UNSTIM_VS_RETINOIC_ACID_STIM_FOLLICULAR_DC_DN, GSE9468_UNSTIM_VS_RETINOIC_ACID_STIM_FOLLICULAR_DC_DN
HALLMARK_IL2_STAT5_SIGNALING, HALLMARK_IL2_STAT5_SIGNALING
IVANOVA_HEMATOPOIESIS_INTERMEDIATE_PROGENITOR, IVANOVA_HEMATOPOIESIS_INTERMEDIATE_PROGENITOR
GSE17660_TCONV_VS_TREG_THYMUS_DN, GSE17660_TCONV_VS_TREG_THYMUS_DN
GSE9962_CTRL_VS_LPS_48_MAC_DN, GSE9962_CTRL_VS_LPS_48_MAC_DN
GSE9962_TREG_VS_TCONV_THYMUS_UP, GSE9962_TREG_VS_TCONV_THYMUS_UP
GSE14308_CTRL_VS_STAPH_AUREUS_INF_PBMC_UP, GSE14308_CTRL_VS_STAPH_AUREUS_INF_PBMC_UP
GSE24280_UNSTIM_VS_LISMONASA_2H_DC_48_LIP, GSE24280_UNSTIM_VS_LISMONASA_2H_DC_48_LIP
GSE27273_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_48H_DN, GSE27273_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_48H_DN
GSE6943_DAY8_VS_DAY30_MEM_TH1_CD4_TCELL_DN, GSE6943_DAY8_VS_DAY30_MEM_TH1_CD4_TCELL_DN
GSE15745_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_DN, GSE15745_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_DN
GOMF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GOMF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
ACOSTA_PROLIFERATION_INDEPENDENT_MYC_TARGETS_UP, ACOSTA_PROLIFERATION_INDEPENDENT_MYC_TARGETS_UP
HP_HYPOPIGMENTATION_OF_THE_SKIN, HP_HYPOPIGMENTATION_OF_THE_SKIN
GSE4589_UNSTIM_VS_LISMONASA_2H_DC_24H_DN, GSE4589_UNSTIM_VS_LISMONASA_2H_DC_24H_DN
GSE9962_MAC_VS_MEM_CD4_TCELL_DN, GSE9962_MAC_VS_MEM_CD4_TCELL_DN
GSE16385_VS_WEST_IQGENE_INC_VIRUS_MATURE_NEURON_CELL_LINE_DN, GSE16385_CTRL_VS_WEST_IQGENE_INC_VIRUS_MATURE_NEURON_CELL_LINE_DN
KEGG_PATHOGENIC_ESCHERICHIA_COLI_INFECTION, KEGG_PATHOGENIC_ESCHERICHIA_COLI_INFECTION
WP_PATHOGENIC_ESCHERICHIA_COLI_INFECTION, WP_PATHOGENIC_ESCHERICHIA_COLI_INFECTION
GSE14308_CTRL_VS_STAPH_AUREUS_INF_PBMC_DN, GSE14308_CTRL_VS_STAPH_AUREUS_INF_PBMC_DN
MIR446B_MIR446B
GSE5281_HEMATOPOITIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP, GSE5281_HEMATOPOITIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP
HP_MULTIPLE_CAFE_AU_LAIT_SPOTS, HP_MULTIPLE_CAFE_AU_LAIT_SPOTS
REACTOME_NUCLEAR_ENVELOPE_NEBULIN_RESEMBLE, REACTOME_NUCLEAR_ENVELOPE_NE_REASSEMBLY
HP_INTERNAL_HAEMORRHAGE, HP_INTERNAL_HAEMORRHAGE
GSE46606_DAY3_VS_DAY5_CD4T_IL2_IL15_STIMULATED_IRF4HIGH_BCELL_UP, GSE46606_DAY3_VS_DAY5_CD4T_IL2_IL15_STIMULATED_IRF4HIGH_BCELL_UP
GSE9266_NAIVE_VS_MONOCYTE_UP, GSE9266_NAIVE_BCELL_VS_MONOCYTE_UP
LENAOUR_DENDRITIC_CELL_MATURATION_UP, LENAOUR_DENDRITIC_CELL_MATURATION_UP
ROSOLI_ILEITIS_INDUCED_THU_ASSOCIATED_MODULE, ROSOLI_ILEITIS_INDUCED_THU_ASSOCIATED_MODULE
SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELLS, SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELLS
GOMF_CHROMOSOME_SEGREGATION, GOMF_CHROMOSOME_SEGREGATION
REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX
GSE6269_E_COLI_VS_STREP_PNEUMO_INF_PBMC_UP, GSE6269_E_COLI_VS_STREP_PNEUMO_INF_PBMC_UP
WP_APOPTOSIS, WP_APOPTOSIS
GOBP_TERMINATION_OF_RNA_POLYMERASE_I_TRANSCRIPTION, GOBP_TERMINATION_OF_RNA_POLYMERASE_I_TRANSCRIPTION
REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION_TERMINATION, REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION_TERMINATION
GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
GOBP_PEPTIDYL_THREONINE_MODIFICATION, GOBP_PEPTIDYL_THREONINE_MODIFICATION
MTOR_UP.V1_UP, MTOR_UP.V1_UP
GSE1461_NAIVE_BCELL_VS_PLASMA_CELL_DN, GSE1461_NAIVE_BCELL_VS_PLASMA_CELL_DN
HP_HYPOPIGMENTED_SKIN_PATCHES, HP_HYPOPIGMENTED_SKIN_PATCHES
GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_POTENTIAL, GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_POTENTIAL
GOBP_TRANSLATION_SYNTHESIS, GOBP_TRANSLATION_SYNTHESIS
GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_BIOSYNTHETIC_PROCESS
REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_DNA_REPAIR_GENES, REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_DNA_REPAIR_GENES
GSE27044_ILC_VS_NAIVE_CD4_TCELL_DAY7_POST_CORONAVIRUS_INFECTION_UP, GSE27044_ILC_VS_NAIVE_CD4_TCELL_DAY7_POST_CORONAVIRUS_INFECTION_UP
GOBP_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_I_PROMOTER, GOBP_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_I_PROMOTER
LIND31537_DENDRITIC_CELL_MATURATION_B_UP, LIND31537_DENDRITIC_CELL_MATURATION_B_UP
GOBP_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR, GOBP_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR
HP_ABNORMAL_URETER_MORPHOLOGY, HP_ABNORMAL_URETER_MORPHOLOGY
HP_TRACHEOBRONCHIAL_FISTULA, HP_TRACHEOBRONCHIAL_FISTULA
GSE3178_UNSTIM_VS_LPS_STIM_VDU_GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_LPS_STIM_VDU_GAMMADELTA_TCELL_DN
GOBP_CELL_DEVELOPMENT, GOBP_CELL_DEVELOPMENT
GOBP_HEPATOCELLULAR_SYSTEM_DEVELOPMENT, GOBP_HEPATOCELLULAR_SYSTEM_DEVELOPMENT
HP_APLASIA_HYPOPLASIA_OF_THE_RADIUS, HP_APLASIA_HYPOPLASIA_OF_THE_RADIUS
ZHAN_MULTIPLE_MYELOMA_UP, ZHAN_MULTIPLE_MYELOMA_UP
DESCARTES_FETAL_ADRENAL_LYMPHOID_CELLS, DESCARTES_FETAL_ADRENAL_LYMPHOID_CELLS
SENESE_HDAC2_TARGETS_UP, SENESE_HDAC2_TARGETS_UP
GOMF_MOTOR_ACTIVITY, GOMF_MOTOR_ACTIVITY
NELSON_RESPONSE_TO_ANDROGEN_UP, NELSON_RESPONSE_TO_ANDROGEN_UP
HP_PULMONARY_INFILTRATES, HP_PULMONARY_INFILTRATES
HP_MYOPIA, HP_MYOPIA
HP_HYPOTONIA, HP_HYPOTONIA
HP_FOCAL_DYSTONIA, HP_FOCAL_DYSTONIA
REACTOME_APC_C_COCH_MEDIATED_DEGRADATION_OF_CYCLIN_B, REACTOME_APC_C_CDC20_MEDIATED_DEGRADATION_OF_CYCLIN_B
LIU_PROSTATE_CANCER_UP, LIU_PROSTATE_CANCER_UP
GSE17032_WT_VS_SPNAR_KO_SKIN2_7_CIRCU_7_STRAIN_INF_DN, GSE17032_WT_VS_SPNAR_KO_SKIN2_7_CIRCU_9_STRAIN_INF_DN
ZHAN_MULTIPLE_MYELOMA_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD2_DN
RUFFA_HYPOXIA_METAGENE, RUFFA_HYPOXIA_METAGENE
MIR497_5P, MIR497_5P
KEGG_PATHOGENIC_ESCHERICHIA_COLI_INFECTION, KEGG_PATHOGENIC_ESCHERICHIA_COLI_INFECTION
WP_PATHOGENIC_ESCHERICHIA_COLI_INFECTION, WP_PATHOGENIC_ESCHERICHIA_COLI_INFECTION
GOBP_ADRENAL_GLAND_DEVELOPMENT, GOBP_ABNORMAL_ADRENAL_GLAND_MORPHOLOGY
GOCC_ADRENAL_CORTICAL_CARCINOMA, GOCC_ADRENAL_CORTICAL_CARCINOMA
HP_SKIN_TAGS, HP_SKIN_TAGS
REACTOME_PHOSPHORYLATION_OF_THE_APC_C_REACTOME_PHOSPHORYLATION_OF_THE_APC_C
E2F3_UP.V1_UP, E2F3_UP.V1_UP
BIOCARTA_RESPONSE_TO_CAMPTOTHECIN_DN, BIOCARTA_RESPONSE_TO_CAMPTOTHECIN_DN
REACTOME_INTRAFLAGELLAR_TRANSPORT, REACTOME_INTRAFLAGELLAR_TRANSPORT
HP_PREAURICULAR_SKIN_TAG, HP_PREAURICULAR_SKIN_TAG
ZHONG_PIC_CA_UNKNOWN_INF_ZHONG_PIC_CA_UNKNOWN_INF
DESCARTES_FETAL_LUNG_LYMPHOID_CELLS, DESCARTES_FETAL_LUNG_LYMPHOID_CELLS
REACTOME_ABERRANT_REGULATION_OF_MITOTIC_EXIT_IN_CANCER_DUE_TO_RB1_DEFECTS, REACTOME_ABERRANT_REGULATION_OF_MITOTIC_EXIT_IN_CANCER_DUE_TO_RB1_DEFECTS
GOBP_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN
GOBP_ERROR_FREE_TRANSLATION_SYNTHESIS, GOBP_ERROR_FREE_TRANSLATION_SYNTHESIS
TRAVAGLINE_LUNG_CD4_MEMORY_EFFECTOR_T_CELL, TRAVAGLINE_LUNG_CD4_MEMORY_EFFECTOR_T_CELL
GOCC_PERICENTRIC_HETEROCHROMATIN, GOCC_PERICENTRIC_HETEROCHROMATIN
WANG_METASTASIS_OF_BREAST_CANCER_ESR1_UP, WANG_METASTASIS_OF_BREAST_CANCER_ESR1_UP
REACTOME_KINESINS, REACTOME_KINESINS
GOMF_NUCLEOBASE_CONTAINING_COMPOUND_KINASE_ACTIVITY, GOMF_NUCLEOBASE_CONTAINING_COMPOUND_KINASE_ACTIVITY
GOBP_SPINDLE_LOCALIZATION, GOBP_SPINDLE_LOCALIZATION
MIR4521, MIR4521
GOBP_ESTABLISHMENT_OF_MITOTIC_SPINDLE_LOCALIZATION, GOBP_ESTABLISHMENT_OF_MITOTIC_SPINDLE_LOCALIZATION
HP_ABNORMALITY_OF_PERIARTICULAR_REGION, HP_ABNORMALITY_OF_PERIARTICULAR_REGION
GSE3720_UNSTIM_VS_LPS_STIM_VDU_GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_LPS_STIM_VDU_GAMMADELTA_TCELL_DN
MELISSARI_TUT1_TARGETS_UP, MELISSARI_TUT1_TARGETS_UP
GOCC_PERICENTRIC_HETEROCHROMATIN, GOCC_PERICENTRIC_HETEROCHROMATIN
MODULE_204, MODULE_204
MIR676C_5P, MIR676C_5P
HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING, HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING
AKL_HTLV1_INFECTION_DN, AKL_HTLV1_INFECTION_DN
VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN, VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN
HP_COLON_CANCER, HP_COLON_CANCER
GOCC_ANAPHASE_PROMOTING_COMPLEX, GOCC_ANAPHASE_PROMOTING_COMPLEX
HP_DEVIATION_OF_THE_THUMB, HP_DEVIATION_OF_THE_THUMB
GSE29618_MDC_VS_PDC_UP, GSE29618_MDC_VS_PDC_UP
GOMF_CYTOSKELETAL_ADAPTOR_ACTIVITY, GOMF_CYTOSKELETAL_ADAPTOR_ACTIVITY
HP_ACUTE_LYMPHOBLASTIC_LEUKEMIA, HP_ACUTE_LYMPHOBLASTIC_LEUKEMIA
GOMF_POSITIVE_REGULATION_OF_CHROMOSOME_SEPARATION, GOMF_POSITIVE_REGULATION_OF_CHROMOSOME_SEPARATION
GOMF_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GOMF_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
MIR6665, MIR6665
GOBP_PYRIMIDINE_NUCLEOSIDE_METABOLIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOSIDE_METABOLIC_PROCESS
GSE10325_MYELOID_VS_LUPUS_MYELOID_UP, GSE10325_MYELOID_VS_LUPUS_MYELOID_UP
REACTOME_CONDENSATION_OF_PROMETAPHASE_CHROMOSOMES, REACTOME_CONDENSATION_OF_PROMETAPHASE_CHROMOSOMES
VILIMAS_NOTCH1_TARGETS_UP, VILIMAS_NOTCH1_TARGETS_UP
GOBP_REGULATION_OF_CELL_CYCLE_PROCESS, GOBP_REGULATION_OF_CELL_CYCLE_PROCESS
PID_INTERACTOME_OF_POLYCOMB_REPRESSIVE_COMPLEX_2_PRC2, PID_INTERACTOME_OF_POLYCOMB_REPRESSIVE_COMPLEX_2_PRC2
REACTOME_INITIATION_OF_NUCLEAR_ENVELOPE_NE_REFORMATION, REACTOME_INITIATION_OF_NUCLEAR_ENVELOPE_NE_REFORMATION
REACTOME_SYNTHESIS_OF_PIPS_AT_THE_EARLY_ENDOSOME_MEMBRANE, REACTOME_SYNTHESIS_OF_PIPS_AT_THE_EARLY_ENDOSOME_MEMBRANE
HP_NON_HODGKIN_LYMPHOMA, HP_NON_HODGKIN_LYMPHOMA
DAVICIONI_PAX_FOXO1_SIGNATURE_IN_ARMS_UP, DAVICIONI_PAX_FOXO1_SIGNATURE_IN_ARMS_UP
WOESCHECH_TUMOR_REJECTION_UP, WOESCHECH_TUMOR_REJECTION_UP
MIR644A, MIR644A
WP_RAC2_GAMMA_SYNDROME_PATHWAY, WP_RAC2_GAMMA_SYNDROME_PATHWAY
HP_ABNORMAL_VALVULAR_MORPHOLOGY, HP_ABNORMAL_VALVULAR_MORPHOLOGY
GOBP_REGULATION_OF_CELL_CYCLE_PHASE_TRANSITION, GOBP_REGULATION_OF_CELL_CYCLE_PHASE_TRANSITION
LUND_DISTINCT_BY_MUTATION_STATUS_UP, LUND_SILENCED_BY_METHYLATION
HP_ABNORMAL_LYMPHADENOPATHY, HP_ABNORMAL_LYMPHADENOPATHY
MIR675P, MIR675P
HP_ABNORMAL_EYE_BROW_MORPHOLOGY, HP_ABNORMAL_EYE_BROW_MORPHOLOGY
DUTTA_APOPTOSIS_VIA_NFKB_DUTTA_APOPTOSIS_VIA_NFKB
NEWMAN_BRCC3_TARGETS_UP, NEWMAN_BRCC3_TARGETS_UP
GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY
HOFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_UP, HOFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_UP
MANSAHRIFI_RESPONSE_TO_ESTRADIOL, MANSAHRIFI_RESPONSE_TO_ESTRADIOL
GOCC_ESC_E_2_COMPLEX, GOCC_ESC_E_2_COMPLEX
GOBP_REGULATION_OF_CELL_CELL_ADHESION_PROCESS, GOBP_DISOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS
GOBP_ANEUPLOIDY_CAUSING_CHROMOSOME_LOSS, GOBP_ANEUPLOIDY_CAUSING_CHROMOSOME_LOSS
GOMF_T_CELL_HOMEOSTASIS, GOMF_T_CELL_HOMEOSTASIS
GAZDA_ETOPOSIDE_SILENCING_BY_RAS1, GAZDA_ETOPOSIDE_SILENCING_BY_RAS1
REACTOME_CARBOXYTERMINAL_POST_TRANSLATIONAL_MODIFICATIONS_OF_TUBULIN, REACTOME_CARBOXYTERMINAL_POST_TRANSLATIONAL_MODIFICATIONS_OF_TUBULIN
GOBP_POSITIVE_REGULATION_OF_CHROMOSOME_SEPARATION, GOBP_PYRIMIDINE_NUCLEOTIDE_CATABOLIC_PROCESS
PID_IL27_PATHWAY, PID_IL27_PATHWAY
GSE9635_DAY15_VS_DAY7_LATE_FLU_VACCINE_PBMC_UP, GSE9635_DAY15_VS_DAY7_LATE_FLU_VACCINE_PBMC_UP
HP_BASAL_CELL_CARCINOMA, HP_BASAL_CELL_CARCINOMA
MIR147A, MIR147A
HP_IRIS_STOMA, HP_IRIS_STOMA
MIR1_3P, MIR1_3P
LIU_LIVER_CANCER_MYC_E2F1_UP, LIU_LIVER_CANCER_MYC_E2F1_UP
HP_ABNORMAL_2ND_FINGER_MORPHOLOGY, HP_ABNORMAL_2ND_FINGER_MORPHOLOGY
HP_ABSENT_FOREARM_BONE, HP_ABSENT_FOREARM_BONE
GOBP_MEMBRANE_DISASSEMBLY, GOBP_MEMBRANE_DISASSEMBLY
WP_PHOTODYNAMIC_THERAPYINDUCED_AP1_SURVIVAL_SIGNALING, WP_PHOTODYNAMIC_THERAPYINDUCED_AP1_SURVIVAL_SIGNALING
ZHAN_MULTIPLE_MYELOMA_LB_UP, ZHAN_MULTIPLE_MYELOMA_LB_UP
GOCC_CONDENSIN_COMPLEX, GOCC_CONDENSIN_COMPLEX
FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CONFERRED_ANTIBODY_RESPONSE_AGE_18_49VO_IOV_POSITIVE, FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE
HP_COMBINED_IMMUNODEFICIENCY, HP_COMBINED_IMMUNODEFICIENCY
REACTOME_NUCLEOTIDE_SALVAGE, REACTOME_NUCLEOTIDE_SALVAGE
GOMF_PIGMENT_BINDING, GOMF_PIGMENT_BINDING
GOMF_PIGMENT_GRANULE_LOCALIZATION, GOMF_PIGMENT_GRANULE_LOCALIZATION
MIR539_5P, MIR539_5P
GROSS_ELK3_TARGETS_UP, GROSS_ELK3_TARGETS_UP
ONO_AML1_TARGETS_UP, ONO_AML1_TARGETS_UP
HP_MYOTONIA, HP_MYOTONIA
HUMMEL_BURKITTS_LYMPHOMA_DN, HUMMEL_BURKITTS_LYMPHOMA_DN
BIOCARTA_ECM_PATHWAY, BIOCARTA_ECM_PATHWAY
GOBP_NUCLEOBASE_SALVAGE, GOBP_NUCLEOBASE_SALVAGE
HP_ABNORMALITY_OF_THE_PERITONEUM, HP_ABNORMALITY_OF_THE_PERITONEUM
TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_NORMAL_DN, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_NORMAL_DN
GOBP_PYRIMIDINE_CONTAINING_COMPOUND_SALVAGE, GOBP_PYRIMIDINE_CONTAINING_COMPOUND_SALVAGE
REACTOME_PYRIMIDINE_SALVAGE, REACTOME_PYRIMIDINE_SALVAGE
HP_HYPOPLASTIC_NIPPLES, HP_HYPOPLASTIC_NIPPLES
XU_HGF_T_TARGETS_INDUCED_IN_AKT1_48HR_DN, XU_HGF_T_TARGETS_INDUCED_IN_AKT1_48HR_DN
GSE15097_CTRL_LINE_VS_IMMATURE_T_CELL_LINE_DN, GSE15097_PDC_CELL_LINE_VS_IMMATURE_T_CELL_LINE_DN
GOBP_CENTROMERIC_SISTER_CHROMATID_COHESION, GOBP_CENTROMERIC_SISTER_CHROMATID_COHESION
GOMF_HISTONE_H3_K27_METHYLATED_ARGININE_DEMETHYLIMINASE_ACTIVITY, GOMF_HISTONE_H3_K27_METHYLATED_ARGININE_DEMETHYLIMINASE_ACTIVITY
HP_VAGINAL_FISTULA, HP_VAGINAL_FISTULA
GSE9635_DAY7_VS_DAY0_LATE_FLU_VACCINE_PBMC_UP, GSE9635_DAY7_VS_DAY0_LATE_FLU_VACCINE_PBMC_UP
VOICE_PHMC_FU1ARIS_AGE_36_74YO_FEMALES_VS_MALES_HIV_TO_382_UP, VOICE_PHMC_FU1ARIS_AGE_50_74YO_FEMALES_VS_MALES_HIV_TO_285_UP
HP_BRANCHIAL_FISTULA, HP_BRANCHIAL_FISTULA
REACTOME_INTERLEUKIN_35_SIGNALLING, REACTOME_INTERLEUKIN_35_SIGNALLING
GOBP_MITOTIC_NUCLEAR_ENVELOPE_DISASSEMBLY, GOBP_MITOTIC_NUCLEAR_ENVELOPE_DISASSEMBLY
GOCC_TRANSCRIPTION_FACTOR_TFIIH_HOLO_COMPLEX, GOCC_TRANSCRIPTION_FACTOR_TFIIH_HOLO_COMPLEX
GOCC_RIBOPTEROSOME, GOCC_RIBOPTEROSOME
WP_REGULATION_OF_MICROTUBULE_CYTOSKELETON, WP_REGULATION_OF_MICROTUBULE_CYTOSKELETON
HP_DROWSINESS, HP_DROWSINESS
GSE15097_CTRL_LINE_VS_IMMATURE_T_CELL_LINE_UP, GSE15097_CTRL_LINE_VS_IMMATURE_T_CELL_LINE_UP
GOBP_CENTROMERE_CATABOLIC_PROCESS_MCM_MCCLUNG_DELTA_FOR_TARGETS_2HR, GOBP_CENTROMERE_CATABOLIC_PROCESS_MCM_MCCLUNG_DELTA_FOR_TARGETS_2HR
MAN_B_CELL_LYMPHOMA_CLUSTER_2_MAN_B_CELL_LYMPHOMA_CLUSTER_2
PID_NFAT_TFPATHWAY, PID_NFAT_TFPATHWAY
GOBP_PYRIMIDINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOSIDE_BIOSYNTHETIC_PROCESS
GOBP_TELOMERE_MAINTENANCE_VIA_RECOMBINATION, GOBP_TELOMERE_MAINTENANCE_VIA_RECOMBINATION
GOBP_INTERLEUKIN_35_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_35_MEDIATED_SIGNALING_PATHWAY
MARSON_FOXP3_TARGETS_STIMULATED_DN, MARSON_FOXP3_TARGETS_STIMULATED_DN
HP_ABNORMALLY_LOW_PITCHED_VOICE, HP_ABNORMALLY_LOW_PITCHED_VOICE
REACTOME_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAYS, REACTOME_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAY
GOMF_ATP_DEPENDENT_MICROTUBULE_MOTOR_ACTIVITY_PLUS_END_DIRECTED, GOMF_ATP_DEPENDENT_MICROTUBULE_MOTOR_ACTIVITY_PLUS_END_DIRECTED
HP_FALOPIA_EQUINOVALGUS, HP_FALOPIA_EQUINOVALGUS
GOMF_NUCLEOBASE_KINASE_ACTIVITY, GOMF_NUCLEOBASE_KINASE_ACTIVITY
GOBP_CONSTITUTIVE_SECRETION, GOBP_CONSTITUTIVE_SECRETION
GSE17660_TCONV_VS_TREG_THYMUS_UP, GSE17660_TCONV_VS_TREG_THYMUS_UP
MAGNUSSON_MLL_SUPPLY_MYELOMA_KEL_VS_KGA_UP, MAGNUSSON_MLL_SUPPLY_MYELOMA_KGA_VS_KGK_UP
MIR937_5P, MIR937_5P
GOBP_EPITHELIAL_CELL_PROLIFERATION_INVOLVED_IN_PROSTATE_GLAND_DEVELOPMENT, GOBP_EPITHELIAL_CELL_PROLIFERATION_INVOLVED_IN_PROSTATE_GLAND_DEVELOPMENT
GOMF_COPPER_ION_BINDING, GOMF_COPPER_ION_BINDING
PID_TCR_DOWNSTREAM_PATHWAY, PID_TCR_DOWNSTREAM_PATHWAY
HP_APLASTIC_ANEMIA, HP_APLASTIC_ANEMIA
WP_PHOTODYNAMIC_THERAPYINDUCED_NFKB_SURVIVAL_SIGNALING, WP_PHOTODYNAMIC_THERAPYINDUCED_NFKB_SURVIVAL_SIGNALING
WP_REGULATION_OF_APOPTOSIS_BY_PARATHYROID_HORMONE_RELATED_PROTEIN, WP_REGULATION_OF_APOPTOSIS_BY_PARATHYROID_HORMONE_RELATED_PROTEIN
HP_ABSENT_HAIR, HP_ABSENT_HAIR
GOMF_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON, GOMF_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON
GOBP_ABNORMAL_LUNG_MORPHOLOGY, GOBP_ABNORMALITY_OF_LUNAR_METABOLISM
HP_PARTIAL_ALBINISM, HP_PARTIAL_ALBINISM
HP_ABNORMALITY_OF_UPPER_LIMB_METAPHYSIS, HP_ABNORMALITY_OF_UPPER_LIMB_METAPHYSIS
PAGA_TP53_TARGETS_PAGA_TP53_TARGETS
GOMF_GAMMA_CATENIN_BINDING, GOMF_GAMMA_CATENIN_BINDING
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_COLUMNAR_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_COLUMNAR_CELLS
GOBP_GLOMERULAR_MESANGIAL_CELL_PROLIFERATION, GOBP_GLOMERULAR_MESANGIAL_CELL_PROLIFERATION
MIR30C_5P, MIR30C_5P
REACTOME_CLEC7A_INFLAMMASOME_PATHWAY, REACTOME_CLEC7A_INFLAMMASOME_PATHWAY
HP_ULNAR_DEVIATION_OF_THE_2ND_FINGER, HP_ULNAR_DEVIATION_OF_THE_2ND_FINGER
GOBP_PROTEIN_IMPORT_INTO_NUCLEUS_TRANSLOCATION, GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MEMBRANE
GOBP_NEGATIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY
DESCARTES_MAIN_FETAL_SATELLITE_CELLS, DESCARTES_MAIN_FETAL_SATELLITE_CELLS
HP_NARROW_INTERNAL_AUDITORY_CANAL, HP_NARROW_INTERNAL_AUDITORY_CANAL
SU_THYMUS_SG_THYMUS
STAMBUSKY_BOUND_BY_MUTATED_TP53, STAMBUSKY_BOUND_BY_MUTATED_TP53
TAKAOKA_GASTRIC_CANCER_DPY_NUMBER_DN, TAKAOKA_GASTRIC_CANCER_DPY_NUMBER_DN
GOBP_POSITIV