GSE42724_MEMORY_VS_B1_BCELL_UP, GSE42724_MEMORY_VS_B1_BCELL_UP GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4HIGH_SORTED_BCELL_DN, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4HIGH_SORTED_BCELL_DN IKEDA_MIR30_TARGETS_UP, IKEDA_MIR30_TARGETS_UP GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_UP GO_MEMBRANE_DISASSEMBLY, GO_MEMBRANE_DISASSEMBLY GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_DN, GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_DN GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_UP, GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_UP GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_UP, GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_UP KYNG DNA DAMAGE DN, KYNG DNA DAMAGE DN GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_DN, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_DN GSE45365_NK_CELL_VS_CD8_TCELL_UP, GSE45365_NK_CELL_VS_CD8_TCELL_UP GSE3982_CTRL_VS_LPS_48H_DC_DN, GSE3982_CTRL_VS_LPS_48H_DC_DN GO_NEGATIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY GO_PROTEIN_PHOSPHATASE_BINDING, GO_PROTEIN_PHOSPHATASE_BINDING GO_CHAPERONE_BINDING, GO_CHAPERONE_BINDING GGCNRNWCTTYS_UNKNOWN, GGCNRNWCTTYS_UNKNOWN GO_REGULATION_OF_MEMBRANE_PERMEABILITY, GO_REGULATION_OF_MEMBRANE_PERMEABILITY GO RHO GTPASE BINDING, GO RHO GTPASE BINDING FERRANDO_T_ALL_WITH_MLL_ENL_FUSION_UP, FERRANDO_T_ALL_WITH_MLL_ENL_FUSION_UP GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_DN, GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_DN GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY GO_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLVED GSE22432_MULTIPOTENT_PROGENITOR_VS_PDC_UP, GSE22432_MULTIPOTENT_PROGENITOR_VS_PDC_UP GCM_MAPK10, GCM_MAPK10 MODULE_237, MODULE_237 GSE19923_WT_VS_E2A_KO_DP_THYMOCYTE_UP, GSE19923_WT_VS_E2A_KO_DP_THYMOCYTE_UP AMIT_EGF_RESPONSE_120_HELA, AMIT_EGF_RESPONSE_120_HELA BIOCARTA_DEATH_PATHWAY, BIOCARTA_DEATH_PATHWAY TCTGGAC_MIR198, TCTGGAC_MIR198 PID_HDAC_CLASSII_PATHWAY, PID_HDAC_CLASSII_PATHWAY GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS, GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN_ PID_HIV_NEF_PATHWAY, PID_HIV_NEF_PATHWAY GATA2_01, GATA2_01 SA_PROGRAMMED_CELL_DEATH, SA_PROGRAMMED_CELL_DEATH AMIT_EGF_RESPONSE_120_MCF10A, AMIT_EGF_RESPONSE_120_MCF10A GO_POSITIVE_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS, GO_POSITIVE_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS REACTOME_CHOLESTEROL_BIOSYNTHESIS, REACTOME_CHOLESTEROL_BIOSYNTHESIS GO_STEROL_BIOSYNTHETIC_PROCESS, GO_STEROL_BIOSYNTHETIC_PROCESS GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE ATGCACG MIR517B, ATGCACG MIR517B GSE27786_LSK_VS_NEUTROPHIL_DN, GSE27786_LSK_VS_NEUTROPHIL_DN GO_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS, GO_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS GO_DEATH_RECEPTOR_ACTIVITY, GO_DEATH_RECEPTOR_ACTIVITY TCCCCAC_MIR491, TCCCCAC_MIR491 GO_LEUKOCYTE_HOMEOSTASIS, GO_LEUKOCYTE_HOMEOSTASIS GO_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GO_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY GSE22025_UNTREATED_VS_TGFB1_TREATED_CD4_TCELL_UP, GSE22025_UNTREATED_VS_TGFB1_TREATED_CD4_TCELL_UP ACCAATC_MIR509, ACCAATC_MIR509 GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS GO_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR, GO_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR GO_HYDROGEN_PEROXIDE_METABOLIC_PROCESS, GO_HYDROGEN_PEROXIDE_METABOLIC_PROCESS GO_REGULATION_OF_SULFUR_METABOLIC_PROCESS, GO_REGULATION_OF_SULFUR_METABOLIC_PROCESS SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_UP, SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_UP GO NEGATIVE REGULATION OF TRANSCRIPTION FACTOR IMPORT INTO NUCLEUS, GO NEGATIVE REGULATION OF TRANSCRIPTION FACTOR IMPORT INTO NUCLEUS GAUSSMANN_MLL_AF4_FUSION_TARGETS_B_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_B_UP GO_NEUROMUSCULAR_JUNCTION, GO_NEUROMUSCULAR_JUNCTION GO_T_CELL_APOPTOTIC_PROCESS, GO_T_CELL_APOPTOTIC_PROCESS GO_MICROTUBULE_ANCHORING, GO_MICROTUBULE_ANCHORING VERRECCHIA_RESPONSE_TO_TGFB1_C1, VERRECCHIA_RESPONSE_TO_TGFB1_C1 GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_DN, GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_DN GO_SUMO_TRANSFERASE_ACTIVITY, GO_SUMO_TRANSFERASE_ACTIVITY REACTOME_EXTRINSIC_PATHWAY_FOR_APOPTOSIS, REACTOME_EXTRINSIC_PATHWAY_FOR_APOPTOSIS GO_REGULATION_OF_NFAT_PROTEIN_IMPORT_INTO_NUCLEUS, GO_REGULATION_OF_NFAT_PROTEIN_IMPORT_INTO_NUCLEUS

GO_LYMPHOCYTE_APOPTOTIC_PROCESS, GO_LYMPHOCYTE_APOPTOTIC_PROCESS

BIOCARTA_PML_PATHWAY, BIOCARTA_PML_PATHWAY

GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN

GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN

GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAM

REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE, REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE

GSE23695_CD57_POS_VS_NEG_NK_CELL_DN, GSE23695_CD57_POS_VS_NEG_NK_CELL_DN