GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_UP, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_UP GSE9006 TYPE 1_VS_TYPE 2_DIABETES_PBMC_AT_DX_UP, GSE9006_TYPE 1_VS_TYPE 2_DIABETES_PBMC_AT_DX_UP GSE9006 TYPE 1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMC_UP, GSE9006_TYPE_1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMC_UP GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP GSE17974 0.5H VS 72H IL4 AND ANTI IL12 ACT CD4 TCELL UP, GSE17974 0.5H VS 72H IL4 AND ANTI IL12 ACT CD4 TCELL UP GSE28726_ACT_CD4_TCELL_VS_ACT_VA24NEG_NKTCELL_DN, GSE28726_ACT_CD4_TCELL_VS_ACT_VA24NEG_NKTCELL_DN GSE29618_BCELL_VS_PDC_DN, GSE29618_BCELL_VS_PDC_DN GSE24574_BCL6_HIGH_TFH_VS_TFH_CD4_TCELL_DN, GSE24574_BCL6_HIGH_TFH_VS_TFH_CD4_TCELL_DN GSE34515_CD16_POS_MONOCYTE_VS_DC_UP, GSE34515_CD16_POS_MONOCYTE_VS_DC_UP HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP, HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_UP, GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_UP GSE36476_CTRL_VS_TSST_ACT_72H_MEMORY_CD4_TCELL_YOUNG_UP, GSE36476_CTRL_VS_TSST_ACT_72H_MEMORY_CD4_TCELL_YOUNG_UP GSE36476_CTRL_VS_TSST_ACT_40H_MEMORY_CD4_TCELL_YOUNG_UP, GSE36476_CTRL_VS_TSST_ACT_40H_MEMORY_CD4_TCELL_YOUNG_UP GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN, GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN GSE24574_BCL6_HIGH_VS_LOW_TFH_CD4_TCELL_UP, GSE24574_BCL6_HIGH_VS_LOW_TFH_CD4_TCELL_UP MIR302A_5P, MIR302A_5P GSE22886_NAIVE_CD4_TCELL_VS_DC_UP, GSE22886_NAIVE_CD4_TCELL_VS_DC_UP GSE17580_TREG_VS_TEFF_S_MANSONI_INF_DN, GSE17580_TREG_VS_TEFF_S_MANSONI_INF_DN GSE22886_UNSTIM_VS_STIM_MEMORY_TCELL_UP, GSE22886_UNSTIM_VS_STIM_MEMORY_TCELL_UP MODULE 239, MODULE 239 GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MDC_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MDC_UP GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN, GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN MIR548AV_3P, MIR548AV_3P MIR6739_3P, MIR6739_3P GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP, GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP TOOKER GEMCITABINE RESISTANCE DN, TOOKER GEMCITABINE RESISTANCE DN PICCALUGA_ANGIOIMMUNOBLASTIC_LYMPHOMA_DN, PICCALUGA_ANGIOIMMUNOBLASTIC_LYMPHOMA_DN PDGF_UP.V1_UP, PDGF_UP.V1_UP GSE36476_CTRL_VS_TSST_ACT_40H_MEMORY_CD4_TCELL_OLD_UP, GSE36476_CTRL_VS_TSST_ACT_40H_MEMORY_CD4_TCELL_OLD_UP WP_TGFBETA_SIGNALING_PATHWAY, WP_TGFBETA_SIGNALING_PATHWAY GSE22611_UNSTIM_VS_2H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP, GSE22611_UNSTIM_VS_2H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP NAGASHIMA_NRG1_SIGNALING_UP, NAGASHIMA_NRG1_SIGNALING_UP GCAAAAA_MIR129, GCAAAAA_MIR129 PECE_MAMMARY_STEM_CELL_DN, PECE_MAMMARY_STEM_CELL_DN MODULE_261, MODULE_261 GTGACTT_MIR224, GTGACTT_MIR224 GNF2_DDX5, GNF2_DDX5 KIM_WT1_TARGETS_8HR_UP, KIM_WT1_TARGETS_8HR_UP GOBP POSITIVE REGULATION OF HEMOPOIESIS, GOBP POSITIVE REGULATION OF HEMOPOIESIS MIR6828_5P, MIR6828_5P GSE30971_WBP7_HET_VS_KO_MACROPHAGE_UP, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_UP GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP AACATTC_MIR4093P, AACATTC_MIR4093P GACTGTT_MIR212_MIR132, GACTGTT_MIR212_MIR132 BROCKE_APOPTOSIS_REVERSED_BY_IL6, BROCKE_APOPTOSIS_REVERSED_BY_IL6 GNF2_PTPRC, GNF2_PTPRC GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_UP, GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_UP ZHENG_CORD_BLOOD_C2_PUTATIVE_BASOPHIL_EOSINOPHIL_MAST_CELL_PROGENITOR, ZHENG_CORD_BLOOD_C2_PUTATIVE_BASOPHIL_EOSINOPHIL_MAST_CELL_PROGENITOR SEIDEN_ONCOGENESIS_BY_MET, SEIDEN_ONCOGENESIS_BY_MET GSE36476_CTRL_VS_TSST_ACT_72H_MEMORY_CD4_TCELL_OLD_UP, GSE36476_CTRL_VS_TSST_ACT_72H_MEMORY_CD4_TCELL_OLD_UP MIR6516_3P, MIR6516_3P GNF2 CD53, GNF2 CD53 GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_UP, GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_UP REACTOME_SIGNALING_BY_VEGF, REACTOME_SIGNALING_BY_VEGF MODULE_358, MODULE_358 IVANOVSKA_MIR106B_TARGETS, IVANOVSKA_MIR106B_TARGETS GSE3982_MAC_VS_NEUTROPHIL_DN, GSE3982_MAC_VS_NEUTROPHIL_DN CUI_GLUCOSE_DEPRIVATION, CUI_GLUCOSE_DEPRIVATION GOBP_RESPONSE_TO_STARVATION, GOBP_RESPONSE_TO_STARVATION BYSTROEM_CORRELATED_WITH_IL5_DN, BYSTROEM_CORRELATED_WITH_IL5_DN GSE8515_CTRL_VS_IL1_4H_STIM_MAC_UP, GSE8515_CTRL_VS_IL1_4H_STIM_MAC_UP MIR2110, MIR2110 GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_10DY_DN, GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_10DY_DN PELLICCIOTTA_HDAC_IN_ANTIGEN_PRESENTATION_UP, PELLICCIOTTA_HDAC_IN_ANTIGEN_PRESENTATION_UP FIRESTEIN_PROLIFERATION, FIRESTEIN_PROLIFERATION GOBP_POSITIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION MIR873_5P, MIR873_5P GOBP_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION GOBP_CELLULAR_RESPONSE_TO_STARVATION, GOBP_CELLULAR_RESPONSE_TO_STARVATION SHAFFER_IRF4_TARGETS_IN_PLASMA_CELL_VS_MATURE_B_LYMPHOCYTE, SHAFFER_IRF4_TARGETS_IN_PLASMA_CELL_VS_MATURE_B_LYMPHOCYTE KEGG_RENAL_CELL_CARCINOMA, KEGG_RENAL_CELL_CARCINOMA GOBP_MITOTIC_G1_S_TRANSITION_CHECKPOINT, GOBP_MITOTIC_G1_S_TRANSITION_CHECKPOINT HP_WEBBED_NECK, HP_WEBBED_NECK MIR297, MIR297 MIR450A_2_3P, MIR450A_2_3P WP_HEPATITIS_B_INFECTION, WP_HEPATITIS_B_INFECTION CCAWNWWNNNGGC_UNKNOWN, CCAWNWWNNNGGC_UNKNOWN DE YY1 TARGETS DN, DE YY1 TARGETS DN MODULE_147, MODULE_147 GSE22935_WT_VS_MYD88_KO_MACROPHAGE_12H_MBOVIS_BCG_STIM_UP, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_12H_MBOVIS_BCG_STIM_UP GOBP_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS JI_RESPONSE_TO_FSH_DN, JI_RESPONSE_TO_FSH_DN MIR5100, MIR5100 WP_REGULATION_OF_ACTIN_CYTOSKELETON, WP_REGULATION_OF_ACTIN_CYTOSKELETON MODULE_525, MODULE_525 AIZARANI_LIVER_C28_NK_NKT_CELLS_6, AIZARANI_LIVER_C28_NK_NKT_CELLS_6 MIR1266 5P, MIR1266 5P GTATGAT_MIR154_MIR487, GTATGAT_MIR154_MIR487 MIR148A 5P, MIR148A 5P WP_MET_IN_TYPE_1_PAPILLARY_RENAL_CELL_CARCINOMA, WP_MET_IN_TYPE_1_PAPILLARY_RENAL_CELL_CARCINOMA GSE17721_LPS_VS_CPG_0.5H_BMDC_DN, GSE17721_LPS_VS_CPG_0.5H_BMDC_DN GOBP_MYELOID_CELL_DEVELOPMENT, GOBP_MYELOID_CELL_DEVELOPMENT MIR4518, MIR4518 KEGG T CELL RECEPTOR SIGNALING PATHWAY, KEGG T CELL RECEPTOR SIGNALING PATHWAY PID_VEGFR1_2_PATHWAY, PID_VEGFR1_2_PATHWAY GOBP_ENDOPLASMIC_RETICULUM_ORGANIZATION, GOBP_ENDOPLASMIC_RETICULUM_ORGANIZATION GNF2_RAP1B, GNF2_RAP1B MIR1237_3P, MIR1237_3P NUTT_GBM_VS_AO_GLIOMA_UP, NUTT_GBM_VS_AO_GLIOMA_UP KEGG B CELL RECEPTOR SIGNALING PATHWAY, KEGG B CELL RECEPTOR SIGNALING PATHWAY MORI PLASMA CELL UP, MORI PLASMA CELL UP MIR192_3P, MIR192_3P HP_ECZEMA, HP_ECZEMA GOBP_REGULATION_OF_CIRCADIAN_RHYTHM, GOBP_REGULATION_OF_CIRCADIAN_RHYTHM GNF2_TYK2, GNF2_TYK2 GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCYTES_DN, GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCYTES_DN WP_ERBB_SIGNALING_PATHWAY, WP_ERBB_SIGNALING_PATHWAY GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING, GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING GTCNYYATGR_UNKNOWN, GTCNYYATGR_UNKNOWN PHONG_TNF_TARGETS_UP, PHONG_TNF_TARGETS_UP WP_FAS_LIGAND_FASL_PATHWAY_AND_STRESS_INDUCTION_OF_HEAT_SHOCK_PROTEINS_HSP_REGULATION, WP_FAS_LIGAND_FASL_PATHWAY_AND_STRESS_INDUCTION_OF_HEAT_SHOCK_PROTEINS_HSP_REGULATION GOBP_LEUKOCYTE_HOMEOSTASIS, GOBP_LEUKOCYTE_HOMEOSTASIS FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN, FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN GOMF_PROTEIN_TYROSINE_KINASE_BINDING, GOMF_PROTEIN_TYROSINE_KINASE_BINDING GOBP_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION, GOBP_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION TAGAACC_MIR182, TAGAACC_MIR182 KEGG_MTOR_SIGNALING_PATHWAY, KEGG_MTOR_SIGNALING_PATHWAY GTTATAT_MIR410, GTTATAT_MIR410 MIR4501, MIR4501 MIR562, MIR562 MIR26A_1_3P, MIR26A_1_3P KEGG_ERBB_SIGNALING_PATHWAY, KEGG_ERBB_SIGNALING_PATHWAY GSE6269_FLU_VS_STREP_PNEUMO_INF_PBMC_DN, GSE6269_FLU_VS_STREP_PNEUMO_INF_PBMC_DN PDGF_ERK_DN.V1_UP, PDGF_ERK_DN.V1_UP AGGCACT_MIR5153P, AGGCACT_MIR5153P KEGG_PROSTATE_CANCER, KEGG_PROSTATE_CANCER GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP WP_ATM_SIGNALING_NETWORK_IN_DEVELOPMENT_AND_DISEASE, WP_ATM_SIGNALING_NETWORK_IN_DEVELOPMENT_AND_DISEASE GOBP_RESPONSE_TO_GAMMA_RADIATION, GOBP_RESPONSE_TO_GAMMA_RADIATION AZARE_NEOPLASTIC_TRANSFORMATION_BY_STAT3_UP, AZARE_NEOPLASTIC_TRANSFORMATION_BY_STAT3_UP MCCLUNG_COCAINE_REWARD_5D, MCCLUNG_COCAINE_REWARD_5D MIR4425, MIR4425 GTACAGG_MIR486, GTACAGG_MIR486 FALVELLA_SMOKERS_WITH_LUNG_CANCER, FALVELLA_SMOKERS_WITH_LUNG_CANCER MODULE 456, MODULE 456 SIG_CHEMOTAXIS, SIG_CHEMOTAXIS GSE6269_HEALTHY_VS_E_COLI_INF_PBMC_DN, GSE6269_HEALTHY_VS_E_COLI_INF_PBMC_DN WP_KIT_RECEPTOR_SIGNALING_PATHWAY, WP_KIT_RECEPTOR_SIGNALING_PATHWAY MIR26A_2_3P, MIR26A_2_3P KEGG_ENDOMETRIAL_CANCER, KEGG_ENDOMETRIAL_CANCER AGTCTTA_MIR499, AGTCTTA_MIR499 WP_APOPTOSISRELATED_NETWORK_DUE_TO_ALTERED_NOTCH3_IN_OVARIAN_CANCER, WP_APOPTOSISRELATED_NETWORK_DUE_TO_ALTERED_NOTCH3_IN_OVARIAN_CANCER GOBP_LEARNING, GOBP_LEARNING PID_RHOA_PATHWAY, PID_RHOA_PATHWAY MIR4793 5P, MIR4793 5P SIG_BCR_SIGNALING_PATHWAY, SIG_BCR_SIGNALING_PATHWAY CHEN_LVAD_SUPPORT_OF_FAILING_HEART_UP, CHEN_LVAD_SUPPORT_OF_FAILING_HEART_UP WP_AGERAGE_PATHWAY, WP_AGERAGE_PATHWAY GOBP_VESICLE_MEDIATED_TRANSPORT_BETWEEN_ENDOSOMAL_COMPARTMENTS, GOBP_VESICLE_MEDIATED_TRANSPORT_BETWEEN_ENDOSOMAL_COMPARTMENTS WP_3Q29_COPY_NUMBER_VARIATION_SYNDROME, WP_3Q29_COPY_NUMBER_VARIATION_SYNDROME MIR6895_5P, MIR6895_5P WP_IL6_SIGNALING_PATHWAY, WP_IL6_SIGNALING_PATHWAY JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_UP, JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_UP GGTGAAG_MIR412, GGTGAAG_MIR412 DIRMEIER_LMP1_RESPONSE_EARLY, DIRMEIER_LMP1_RESPONSE_EARLY GOBP_POSITIVE_REGULATION_OF_PEPTIDE_SECRETION, GOBP_POSITIVE_REGULATION_OF_PEPTIDE_SECRETION NAKAYAMA_FRA2_TARGETS, NAKAYAMA_FRA2_TARGETS PID_GMCSF_PATHWAY, PID_GMCSF_PATHWAY MIR1227_3P, MIR1227_3P GOMF_KINASE_ACTIVATOR_ACTIVITY, GOMF_KINASE_ACTIVATOR_ACTIVITY WP IL5 SIGNALING PATHWAY, WP IL5 SIGNALING PATHWAY WP_IL1_SIGNALING_PATHWAY, WP_IL1_SIGNALING_PATHWAY REACTOME_SIGNALING_BY_FGFR, REACTOME_SIGNALING_BY_FGFR GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH, GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH GOMF_RECEPTOR_TYROSINE_KINASE_BINDING, GOMF_RECEPTOR_TYROSINE_KINASE_BINDING GAJATE_RESPONSE_TO_TRABECTEDIN_UP, GAJATE_RESPONSE_TO_TRABECTEDIN_UP REACTOME_L1CAM_INTERACTIONS, REACTOME_L1CAM_INTERACTIONS REACTOME INTERLEUKIN 3 INTERLEUKIN 5 AND GM CSF SIGNALING, REACTOME INTERLEUKIN 3 INTERLEUKIN 5 AND GM CSF SIGNALING GNF2_CASP4, GNF2_CASP4 BRACHAT_RESPONSE_TO_METHOTREXATE_DN, BRACHAT_RESPONSE_TO_METHOTREXATE_DN BIOCARTA_FAS_PATHWAY, BIOCARTA_FAS_PATHWAY KEGG_FC_EPSILON_RI_SIGNALING_PATHWAY, KEGG_FC_EPSILON_RI_SIGNALING_PATHWAY WP_MAPK_CASCADE, WP_MAPK_CASCADE GSE43955_1H_VS_42H_ACT_CD4_TCELL_DN, GSE43955_1H_VS_42H_ACT_CD4_TCELL_DN WP_IL3_SIGNALING_PATHWAY, WP_IL3_SIGNALING_PATHWAY NAGASHIMA_EGF_SIGNALING_UP, NAGASHIMA_EGF_SIGNALING_UP REACTOME_ANTIGEN_PRESENTATION_FOLDING_ASSEMBLY_AND_PEPTIDE_LOADING_OF_CLASS_I_MHC, REACTOME_ANTIGEN_PRESENTATION_FOLDING_ASSEMBLY_AND_PEPTIDE_LOADING_OF_CLASS_I_MHC MIR4684_3P, MIR4684_3P KEGG_NON_SMALL_CELL_LUNG_CANCER, KEGG_NON_SMALL_CELL_LUNG_CANCER MIR3945, MIR3945 HP_ABNORMAL_PLATELET_FUNCTION, HP_ABNORMAL_PLATELET_FUNCTION WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS, WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS GOBP_NEGATIVE_REGULATION_OF_TOR_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_TOR_SIGNALING CAGNYGKNAAA_UNKNOWN, CAGNYGKNAAA_UNKNOWN REACTOME REGULATION_OF_TP53_ACTIVITY_THROUGH_METHYLATION, REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_METHYLATION MIR6776 3P, MIR6776 3P KEGG_VEGF_SIGNALING_PATHWAY, KEGG_VEGF_SIGNALING_PATHWAY BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_DN, BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_DN GOBP_CHAPERONE_COFACTOR_DEPENDENT_PROTEIN_REFOLDING, GOBP_CHAPERONE_COFACTOR_DEPENDENT_PROTEIN_REFOLDING REACTOME SIGNALING BY FGFR2, REACTOME SIGNALING BY FGFR2 REACTOME_APOPTOTIC_EXECUTION_PHASE, REACTOME_APOPTOTIC_EXECUTION_PHASE AGGAAGC MIR5163P, AGGAAGC MIR5163P GOBP_RESPONSE_TO_CALCIUM_ION, GOBP_RESPONSE_TO_CALCIUM_ION MIR4330, MIR4330 MIR5089_3P, MIR5089_3P HP_ABNORMAL_UPPER_MOTOR_NEURON_MORPHOLOGY, HP_ABNORMAL_UPPER_MOTOR_NEURON_MORPHOLOGY MIR937_5P, MIR937_5P PID CXCR3 PATHWAY, PID CXCR3 PATHWAY MIR6813 3P, MIR6813 3P RAO_BOUND_BY_SALL4_ISOFORM_A, RAO_BOUND_BY_SALL4_ISOFORM_A GOBP_CARDIAC_SEPTUM_DEVELOPMENT, GOBP_CARDIAC_SEPTUM_DEVELOPMENT KEGG_LONG_TERM_POTENTIATION, KEGG_LONG_TERM_POTENTIATION KEGG_GLIOMA, KEGG_GLIOMA HP_RIMMED_VACUOLES, HP_RIMMED_VACUOLES WP_PDGFRBETA_PATHWAY, WP_PDGFRBETA_PATHWAY MIR6857_5P, MIR6857_5P WP_NETRINUNC5B_SIGNALING_PATHWAY, WP_NETRINUNC5B_SIGNALING_PATHWAY GOBP_CELLULAR_RESPONSE_TO_OSMOTIC_STRESS, GOBP_CELLULAR_RESPONSE_TO_OSMOTIC_STRESS MIR106A_3P, MIR106A_3P MIR5695, MIR5695 AKL_HTLV1_INFECTION_UP, AKL_HTLV1_INFECTION_UP MIR548Q, MIR548Q HP_ASPIRATION, HP_ASPIRATION GSE22025_TGFB1_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN, GSE22025_TGFB1_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN WP_IL2_SIGNALING_PATHWAY, WP_IL2_SIGNALING_PATHWAY MIR1287 3P, MIR1287 3P GOMF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY, GOMF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY GAJATE_RESPONSE_TO_TRABECTEDIN_DN, GAJATE_RESPONSE_TO_TRABECTEDIN_DN PID_EPHB_FWD_PATHWAY, PID_EPHB_FWD_PATHWAY WP_ESC_PLURIPOTENCY_PATHWAYS, WP_ESC_PLURIPOTENCY_PATHWAYS MODULE 213, MODULE 213 TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_MONOCYTE_DN, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_MONOCYTE_DN KHETCHOUMIAN_TRIM24_TARGETS_UP, KHETCHOUMIAN_TRIM24_TARGETS_UP PID_TRKR_PATHWAY, PID_TRKR_PATHWAY WP_ENVELOPE_PROTEINS_AND_THEIR_POTENTIAL_ROLES_IN_EDMD_PHYSIOPATHOLOGY, WP_ENVELOPE_PROTEINS_AND_THEIR_POTENTIAL_ROLES_IN_EDMD_PHYSIOPATHOLOGY HEDENFALK BREAST CANCER BRACX UP, HEDENFALK BREAST CANCER BRACX UP PID_ER_NONGENOMIC_PATHWAY, PID_ER_NONGENOMIC_PATHWAY CORRE MULTIPLE MYELOMA DN, CORRE MULTIPLE MYELOMA DN GOBP_RESPONSE_TO_TUMOR_CELL, GOBP_RESPONSE_TO_TUMOR_CELL AMIT_EGF_RESPONSE_60_MCF10A, AMIT_EGF_RESPONSE_60_MCF10A KEGG_ANTIGEN_PROCESSING_AND_PRESENTATION, KEGG_ANTIGEN_PROCESSING_AND_PRESENTATION MIR7108_5P, MIR7108_5P KEGG_NATURAL_KILLER_CELL_MEDIATED_CYTOTOXICITY, KEGG_NATURAL_KILLER_CELL_MEDIATED_CYTOTOXICITY PID_ERBB2_ERBB3_PATHWAY, PID_ERBB2_ERBB3_PATHWAY PID_VEGFR1_PATHWAY, PID_VEGFR1_PATHWAY MIR409_5P, MIR409_5P MIR7845_5P, MIR7845_5P GOBP_PEPTIDYL_ASPARAGINE_MODIFICATION, GOBP_PEPTIDYL_ASPARAGINE_MODIFICATION GOBP_PLATELET_AGGREGATION, GOBP_PLATELET_AGGREGATION MIR6804_3P, MIR6804_3P GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION REACTOME_RHO_GTPASES_ACTIVATE_PAKS, REACTOME_RHO_GTPASES_ACTIVATE_PAKS REACTOME_RECYCLING_PATHWAY_OF_L1, REACTOME_RECYCLING_PATHWAY_OF_L1 _DIABETES_PBMC_1MONTH_POST_DX_UP, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_1MONTH_POST_DX_UP GOBP_RESPONSE_TO_INTERLEUKIN_7, GOBP_RESPONSE_TO_INTERLEUKIN_7 WP_METABOLISM_OF_SPINGOLIPIDS_IN_ER_AND_GOLGI_APPARATUS, WP_METABOLISM_OF_SPINGOLIPIDS_IN_ER_AND_GOLGI_APPARATUS HP_THICKENED_HELICES, HP_THICKENED_HELICES REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS, REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS GOBP_ERYTHROCYTE_DEVELOPMENT, GOBP_ERYTHROCYTE_DEVELOPMENT MIR6836 3P, MIR6836 3P LEONARD_HYPOXIA, LEONARD_HYPOXIA MIR3677_5P, MIR3677_5P WANG_TARGETS_OF_MLL_CBP_FUSION_UP, WANG_TARGETS_OF_MLL_CBP_FUSION_UP GOBP_METENCEPHALON_DEVELOPMENT, GOBP_METENCEPHALON_DEVELOPMENT DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXO1_FUSION_UP, DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXO1_FUSION_UP GOBP_POSITIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION GOBP_CELLULAR_HYPEROSMOTIC_RESPONSE, GOBP_CELLULAR_HYPEROSMOTIC_RESPONSE AMIT_EGF_RESPONSE_40_HELA, AMIT_EGF_RESPONSE_40_HELA HP_FATIGABLE_WEAKNESS_OF_BULBAR_MUSCLES, HP_FATIGABLE_WEAKNESS_OF_BULBAR_MUSCLES WP_PRION_DISEASE_PATHWAY, WP_PRION_DISEASE_PATHWAY REACTOME_MAP2K_AND_MAPK_ACTIVATION, REACTOME_MAP2K_AND_MAPK_ACTIVATION DELLA_RESPONSE_TO_TSA_AND_BUTYRATE, DELLA_RESPONSE_TO_TSA_AND_BUTYRATE GOBP_L_GLUTAMATE_TRANSMEMBRANE_TRANSPORT, GOBP_L_GLUTAMATE_TRANSMEMBRANE_TRANSPORT GOBP_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION TCTGATC_MIR383, TCTGATC_MIR383 HP_ABNORMAL_LARYNX_PHYSIOLOGY, HP_ABNORMAL_LARYNX_PHYSIOLOGY GOCC_OLIGOSACCHARYLTRANSFERASE_COMPLEX, GOCC_OLIGOSACCHARYLTRANSFERASE_COMPLEX GENTILE_UV_LOW_DOSE_UP, GENTILE_UV_LOW_DOSE_UP PID_AR_TF_PATHWAY, PID_AR_TF_PATHWAY REACTOME APOPTOSIS INDUCED DNA FRAGMENTATION, REACTOME APOPTOSIS INDUCED DNA FRAGMENTATION GOBP_LYMPH_VESSEL_DEVELOPMENT, GOBP_LYMPH_VESSEL_DEVELOPMENT PLASARI_TGFB1_TARGETS_1HR_UP, PLASARI_TGFB1_TARGETS_1HR_UP GOBP_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK_ORGANIZATION, GOBP_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK_ORGANIZATION WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_MAPK_SIGNALING, WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_MAPK_SIGNALING GOBP_LIPID_IMPORT_INTO_CELL, GOBP_LIPID_IMPORT_INTO_CELL PID_IL3_PATHWAY, PID_IL3_PATHWAY HOWLIN_CITED1_TARGETS_1_DN, HOWLIN_CITED1_TARGETS_1_DN MEF2_01, MEF2_01 HP_HYPOPLASTIC_HEART, HP_HYPOPLASTIC_HEART GOBP_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT, GOBP_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT AMIT_EGF_RESPONSE_40_MCF10A, AMIT_EGF_RESPONSE_40_MCF10A GOBP_CARDIAC_CHAMBER_DEVELOPMENT, GOBP_CARDIAC_CHAMBER_DEVELOPMENT GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_STRESS, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_ GOBP_PHOTOPERIODISM, GOBP_PHOTOPERIODISM GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT, GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT LEIN_LOCALIZED_TO_PROXIMAL_DENDRITES, LEIN_LOCALIZED_TO_PROXIMAL_DENDRITES WP_IL7_SIGNALING_PATHWAY, WP_IL7_SIGNALING_PATHWAY GOBP_POSITIVE_REGULATION_OF_HORMONE_SECRETION, GOBP_POSITIVE_REGULATION_OF_HORMONE_SECRETION ARGGGTTAA_UNKNOWN, ARGGGTTAA_UNKNOWN ZHENG_CORD_BLOOD_C5_SIMILAR_TO_HSC_C6_PUTATIVE_ALTERED_METABOLIC_STATE, ZHENG_CORD_BLOOD_C5_SIMILAR_TO_HSC_C6_PUTATIVE_ALTERED_METABOLIC_STATE GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK, GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK HP_ABNORMALITY_OF_NEUTROPHIL_PHYSIOLOGY, HP_ABNORMALITY_OF_NEUTROPHIL_PHYSIOLOGY SARTIPY_BLUNTED_BY_INSULIN_RESISTANCE_UP, SARTIPY_BLUNTED_BY_INSULIN_RESISTANCE_UP GOBP_ASSOCIATIVE_LEARNING, GOBP_ASSOCIATIVE_LEARNING GOBP_HYPEROSMOTIC_RESPONSE, GOBP_HYPEROSMOTIC_RESPONSE GOBP_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY GOBP_HEART_VALVE_DEVELOPMENT, GOBP_HEART_VALVE_DEVELOPMENT AIZARANI_LIVER_C8_RESIDENT_B_CELLS_1, AIZARANI_LIVER_C8_RESIDENT_B_CELLS_1 GOBP_B_CELL_APOPTOTIC_PROCESS, GOBP_B_CELL_APOPTOTIC_PROCESS SCHRAETS_MLL_TARGETS_UP, SCHRAETS_MLL_TARGETS_UP SIG_REGULATION_OF_THE_ACTIN_CYTOSKELETON_BY_RHO_GTPASES, SIG_REGULATION_OF_THE_ACTIN_CYTOSKELETON_BY_RHO_GTPASES GOBP_TOLERANCE_INDUCTION, GOBP_TOLERANCE_INDUCTION REACTOME_SIGNALING_BY_FLT3_ITD_AND_TKD_MUTANTS, REACTOME_SIGNALING_BY_FLT3_ITD_AND_TKD_MUTANTS WP_ASSOCIATION_BETWEEN_PHYSICOCHEMICAL_FEATURES_AND_TOXICITY_ASSOCIATED_PATHWAYS, WP_ASSOCIATION_BETWEEN_PHYSICOCHEMICAL_FEATURES_AND_TOXICITY_ASSOCIATED_PATHWAYS MIR6798_5P, MIR6798_5P GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREY_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREY_UP GOBP_LYMPH_VESSEL_MORPHOGENESIS, GOBP_LYMPH_VESSEL_MORPHOGENESIS GOBP_ASTRAL_MICROTUBULE_ORGANIZATION, GOBP_ASTRAL_MICROTUBULE_ORGANIZATION GOMF_PYRIMIDINE_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PYRIMIDINE_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_DENDRITIC_TRANSPORT, GOBP_DENDRITIC_TRANSPORT HP_IMPAIRED_NEUTROPHIL_BACTERICIDAL_ACTIVITY, HP_IMPAIRED_NEUTROPHIL_BACTERICIDAL_ACTIVITY GOMF_PROTEIN_TYROSINE_SERINE_THREONINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_TYROSINE_SERINE_THREONINE_PHOSPHATASE_ACTIVITY GOBP_BERGMANN_GLIAL_CELL_DIFFERENTIATION, GOBP_BERGMANN_GLIAL_CELL_DIFFERENTIATION DISTECHE_ESCAPED_FROM_X_INACTIVATION, DISTECHE_ESCAPED_FROM_X_INACTIVATION BONOME OVARIAN CANCER POOR SURVIVAL DN, BONOME OVARIAN CANCER POOR SURVIVAL DN GOBP_POLYSACCHARIDE_CATABOLIC_PROCESS, GOBP_POLYSACCHARIDE_CATABOLIC_PROCESS GOCC_PERINUCLEAR_ENDOPLASMIC_RETICULUM, GOCC_PERINUCLEAR_ENDOPLASMIC_RETICULUM CYCLIN_D1_KE_.V1_DN, CYCLIN_D1_KE_.V1_DN GOBP_CELLULAR_RESPONSE_TO_ETHANOL, GOBP_CELLULAR_RESPONSE_TO_ETHANOL GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS HP_SEVERE_COMBINED_IMMUNODEFICIENCY, HP_SEVERE_COMBINED_IMMUNODEFICIENCY HP_NEOPLASM_OF_THE_MALE_EXTERNAL_GENITALIA, HP_NEOPLASM_OF_THE_MALE_EXTERNAL_GENITALIA NOJIMA_SFRP2_TARGETS_UP, NOJIMA_SFRP2_TARGETS_UP GOBP_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORT, GOBP_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORT GOMF_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORTER_ACTIVITY HOLLEMAN_DAUNORUBICIN_ALL_DN, HOLLEMAN_DAUNORUBICIN_ALL_DN GOCC_MHC_CLASS_I_PEPTIDE_LOADING_COMPLEX, GOCC_MHC_CLASS_I_PEPTIDE_LOADING_COMPLEX SPIELMAN_LYMPHOBLAST_EUROPEAN_VS_ASIAN_2FC_UP, SPIELMAN_LYMPHOBLAST_EUROPEAN_VS_ASIAN_2FC_UP REACTOME_DOWNSTREAM_SIGNAL_TRANSDUCTION, REACTOME_DOWNSTREAM_SIGNAL_TRANSDUCTION GOBP_REGULATION_OF_MONOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_MONOCYTE_DIFFERENTIATION WEINMANN_ADAPTATION_TO_HYPOXIA_DN, WEINMANN_ADAPTATION_TO_HYPOXIA_DN AMIT_DELAYED_EARLY_GENES, AMIT_DELAYED_EARLY_GENES MODULE 131, MODULE 131 HU_ANGIOGENESIS_UP, HU_ANGIOGENESIS_UP GOBP_GLYCOSPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_GLYCOSPHINGOLIPID_BIOSYNTHETIC_PROCESS BIOCARTA_EPO_PATHWAY, BIOCARTA_EPO_PATHWAY WEINMANN_ADAPTATION_TO_HYPOXIA_UP, WEINMANN_ADAPTATION_TO_HYPOXIA_UP GOBP_POSITIVE_REGULATION_OF_PEPTIDE_HORMONE_SECRETION, GOBP_POSITIVE_REGULATION_OF_PEPTIDE_HORMONE_SECRETION GOBP_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE, GOBP_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE GOBP_L_GLUTAMATE_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_L_GLUTAMATE_IMPORT_ACROSS_PLASMA_MEMBRANE GOBP_CARDIAC_VENTRICLE_DEVELOPMENT, GOBP_CARDIAC_VENTRICLE_DEVELOPMENT BIOCARTA_PCAF_PATHWAY, BIOCARTA_PCAF_PATHWAY GOBP_RETINA_VASCULATURE_DEVELOPMENT_IN_CAMERA_TYPE_EYE, GOBP_RETINA_VASCULATURE_DEVELOPMENT_IN_CAMERA_TYPE_EYE PLASARI_TGFB1_SIGNALING_VIA_NFIC_1HR_UP, PLASARI_TGFB1_SIGNALING_VIA_NFIC_1HR_UP REACTOME SIGNALING BY PDGFR IN DISEASE, REACTOME SIGNALING BY PDGFR IN DISEASE WAGNER_APO2_SENSITIVITY, WAGNER_APO2_SENSITIVITY MIR371B 3P, MIR371B 3P ABBUD_LIF_SIGNALING_2_UP, ABBUD_LIF_SIGNALING_2_UP GOBP_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION, GOBP_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION GOBP_POSITIVE_REGULATION_OF_NON_MOTILE_CILIUM_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_NON_MOTILE_CILIUM_ASSEMBLY BUSSLINGER_GASTRIC_PREZYMOGENIC_CELLS, BUSSLINGER_GASTRIC_PREZYMOGENIC_CELLS GOBP_POSITIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT STAT3_01, STAT3_01 GOBP_V_D_J_RECOMBINATION, GOBP_V_D_J_RECOMBINATION GOBP_POSITIVE_REGULATION_OF_ATP_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_ATP_METABOLIC_PROCESS GOBP_AXO_DENDRITIC_PROTEIN_TRANSPORT, GOBP_AXO_DENDRITIC_PROTEIN_TRANSPORT GOBP_SOMITE_DEVELOPMENT, GOBP_SOMITE_DEVELOPMENT HP SEMILOBAR HOLOPROSENCEPHALY, HP SEMILOBAR HOLOPROSENCEPHALY chr1q41, chr1q41 WP_IL9_SIGNALING_PATHWAY, WP_IL9_SIGNALING_PATHWAY WP_SEROTONIN_RECEPTOR_2_AND_ELKSRFGATA4_SIGNALING, WP_SEROTONIN_RECEPTOR_2_AND_ELKSRFGATA4_SIGNALING HP_ABNORMAL_PLATELET_AGGREGATION, HP_ABNORMAL_PLATELET_AGGREGATION PID_DNA_PK_PATHWAY, PID_DNA_PK_PATHWAY GOBP_PRO_B_CELL_DIFFERENTIATION, GOBP_PRO_B_CELL_DIFFERENTIATION GOBP_LYMPHANGIOGENESIS, GOBP_LYMPHANGIOGENESIS KEGG_MELANOMA, KEGG_MELANOMA GOBP_CELLULAR_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE, GOBP_CELLULAR_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE GOBP_REGULATION_OF_POLYSACCHARIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_POLYSACCHARIDE_METABOLIC_PROCESS AIZARANI_LIVER_C38_RESIDENT_B_CELLS_3, AIZARANI_LIVER_C38_RESIDENT_B_CELLS_3 HERNANDEZ ABERRANT MITOSIS BY DOCETACEL 2NM DN, HERNANDEZ ABERRANT MITOSIS BY DOCETACEL 2NM DN REACTOME_SIGNALING_BY_KIT_IN_DISEASE, REACTOME_SIGNALING_BY_KIT_IN_DISEASE GOMF POLYPEPTIDE N ACETYLGALACTOSAMINYLTRANSFERASE ACTIVITY, GOMF POLYPEPTIDE N ACETYLGALACTOSAMINYLTRANSFERASE ACTIVITY GOBP_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME, GOBP_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME HP_ASYMMETRY_OF_THE_THORAX, HP_ASYMMETRY_OF_THE_THORAX HP_CHRONIC_LYMPHATIC_LEUKEMIA, HP_CHRONIC_LYMPHATIC_LEUKEMIA SCHMAHL_PDGF_SIGNALING, SCHMAHL_PDGF_SIGNALING GOBP LABYRINTHINE LAYER DEVELOPMENT, GOBP LABYRINTHINE LAYER DEVELOPMENT GOBP_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_GAMMA_DELTA_T_CELL_DIFFERENTIATION KAPOSI_LIVER_CANCER_MET_UP, KAPOSI_LIVER_CANCER_MET_UP HP_MENORRHAGIA, HP_MENORRHAGIA GOBP CARDIAC CHAMBER MORPHOGENESIS, GOBP CARDIAC CHAMBER MORPHOGENESIS BIOCARTA_UCALPAIN_PATHWAY, BIOCARTA_UCALPAIN_PATHWAY REACTOME CONSTITUTIVE SIGNALING BY LIGAND RESPONSIVE EGFR CANCER VARIANTS, REACTOME CONSTITUTIVE SIGNALING BY LIGAND RESPONSIVE EGFR CANCER VARIANTS BIOCARTA_NKCELLS_PATHWAY, BIOCARTA_NKCELLS_PATHWAY HP_NEOPLASM_OF_THE_LARYNX, HP_NEOPLASM_OF_THE_LARYNX REACTOME_ATF6_ATF6_ALPHA_ACTIVATES_CHAPERONE_GENES, REACTOME_ATF6_ATF6_ALPHA_ACTIVATES_CHAPERONE_GENES GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_SYNAPSE, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_SYNAPSE PID_NECTIN_PATHWAY, PID_NECTIN_PATHWAY REACTOME_SIGNALING_BY_NTRK3_TRKC, REACTOME_SIGNALING_BY_NTRK3_TRKC GOCC_FILOPODIUM_MEMBRANE, GOCC_FILOPODIUM_MEMBRANE GCGCCTT MIR525 MIR524, GCGCCTT MIR525 MIR524 GOMF_CELL_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_ADHESION_MEDIATOR_ACTIVITY GOBP_REGULATION_OF_AMINO_ACID_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_REGULATION_OF_AMINO_ACID_IMPORT_ACROSS_PLASMA_MEMBRANE GOBP_PARENTAL_BEHAVIOR, GOBP_PARENTAL_BEHAVIOR KEGG_NITROGEN_METABOLISM, KEGG_NITROGEN_METABOLISM HP_KERATOSIS_PILARIS, HP_KERATOSIS_PILARIS SCIAN_CELL_CYCLE_TARGETS_OF_TP53_AND_TP73_UP, SCIAN_CELL_CYCLE_TARGETS_OF_TP53_AND_TP73_UP GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_THREONINE, GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_THREONINE REACTOME_INTERLEUKIN_RECEPTOR_SHC_SIGNALING, REACTOME_INTERLEUKIN_RECEPTOR_SHC_SIGNALING GOBP_CELLULAR_RESPONSE_TO_ANTIBIOTIC, GOBP_CELLULAR_RESPONSE_TO_ANTIBIOTIC GOBP_PROTEIN_SULFATION, GOBP_PROTEIN_SULFATION GOBP_POSITIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS WP_MFAP5_EFFECT_ON_PERMEABILITY_AND_MOTILITY_OF_ENDOTHELIAL_CELLS_VIA_CYTOSKELETON_REARRANGEMENT, WP_MFAP5_EFFECT_ON_PERMEABILITY_AND_MOTILITY_OF_ENDOTHELIAL_CELLS_VIA_CYTOSKELETON_RE IVANOVA_HEMATOPOIESIS_STEM_CELL_SHORT_TERM, IVANOVA_HEMATOPOIESIS_STEM_CELL_SHORT_TERM GOBP_NOTCH_RECEPTOR_PROCESSING, GOBP_NOTCH_RECEPTOR_PROCESSING GOBP_REGULATION_OF_CELL_PROJECTION_SIZE, GOBP_REGULATION_OF_CELL_PROJECTION_SIZE HP_SPINAL_CORD_TUMOR, HP_SPINAL_CORD_TUMOR GOBP_POSITIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS ZNF529_TARGET_GENES, ZNF529_TARGET_GENES GOBP_NEGATIVE_REGULATION_OF_STEM_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_STEM_CELL_DIFFERENTIATION WP_BONE_MORPHOGENIC_PROTEIN_BMP_SIGNALLING_AND_REGULATION, WP_BONE_MORPHOGENIC_PROTEIN_BMP_SIGNALLING_AND_REGULATION REACTOME_SIGNALING_BY_PDGFRA_TRANSMEMBRANE_JUXTAMEMBRANE_AND_KINASE_DOMAIN_MUTANTS, REACTOME_SIGNALING_BY_PDGFRA_TRANSMEMBRANE_JUXTAMEMBRANE_AND_KINASE_DOMAIN_MUTANTS HOLLERN_SOLID_NODULAR_BREAST_TUMOR_UP, HOLLERN_SOLID_NODULAR_BREAST_TUMOR_UP PID ALK2 PATHWAY, PID ALK2 PATHWAY MIR101_2_5P, MIR101_2_5P MUELLER_COMMON_TARGETS_OF_AML_FUSIONS_UP, MUELLER_COMMON_TARGETS_OF_AML_FUSIONS_UP HP_CHRONIC_FATIGUE, HP_CHRONIC_FATIGUE WP_OXIDATIVE_STRESS, WP_OXIDATIVE_STRESS GOBP_POSITIVE_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT WANG_ADIPOGENIC_GENES_REPRESSED_BY_SIRT1, WANG_ADIPOGENIC_GENES_REPRESSED_BY_SIRT1 GOBP_LEUKOCYTE_AGGREGATION, GOBP_LEUKOCYTE_AGGREGATION GOBP_PROTEIN_LOCALIZATION_TO_MICROTUBULE, GOBP_PROTEIN_LOCALIZATION_TO_MICROTUBULE MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_VACCINATED_VS_CANDIN_PLACEBO_BCG_PRIMED_1DY_DN, MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_VACCINATED_V GOBP_RESPONSE_TO_GROWTH_HORMONE, GOBP_RESPONSE_TO_GROWTH_HORMONE GSE3720_UNSTIM_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_DN GOBP_NEGATIVE_REGULATION_OF_GLIOGENESIS, GOBP_NEGATIVE_REGULATION_OF_GLIOGENESIS MIR4281, MIR4281 GOBP_TRANS_SYNAPTIC_SIGNALING_MODULATING_SYNAPTIC_TRANSMISSION, GOBP_TRANS_SYNAPTIC_SIGNALING_MODULATING_SYNAPTIC_TRANSMISSION HP_ABNORMALITY_OF_VON_WILLEBRAND_FACTOR, HP_ABNORMALITY_OF_VON_WILLEBRAND_FACTOR DORSEY_GAB2_TARGETS, DORSEY_GAB2_TARGETS VALK AML CLUSTER 2, VALK AML CLUSTER 2 GOBP_REGULATION_OF_MICROVILLUS_LENGTH, GOBP_REGULATION_OF_MICROVILLUS_LENGTH AMIT_EGF_RESPONSE_20_HELA, AMIT_EGF_RESPONSE_20_HELA CLAUS PGR POSITIVE MENINGIOMA DN, CLAUS PGR POSITIVE MENINGIOMA DN GOBP_CEREBELLAR_CORTEX_DEVELOPMENT, GOBP_CEREBELLAR_CORTEX_DEVELOPMENT GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP, GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP GREENBAUM E2A TARGETS DN, GREENBAUM E2A TARGETS DN GOBP_NEGATIVE_REGULATION_OF_T_CELL_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_T_CELL_RECEPTOR_SIGNALING_PATHWAY GOBP_ATRIOVENTRICULAR_VALVE_DEVELOPMENT, GOBP_ATRIOVENTRICULAR_VALVE_DEVELOPMENT GOBP_SPLEEN_DEVELOPMENT, GOBP_SPLEEN_DEVELOPMENT REACTOME_SIGNALING_BY_PDGF, REACTOME_SIGNALING_BY_PDGF GOBP_GLYCOSYLCERAMIDE_METABOLIC_PROCESS, GOBP_GLYCOSYLCERAMIDE_METABOLIC_PROCESS HP_INSULIN_RESISTANT_DIABETES_MELLITUS, HP_INSULIN_RESISTANT_DIABETES_MELLITUS IIZUKA_LIVER_CANCER_EARLY_RECURRENCE, IIZUKA_LIVER_CANCER_EARLY_RECURRENCE HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_UPPER_ARM, HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_UPPER_ARM GOBP_ERBB2_SIGNALING_PATHWAY, GOBP_ERBB2_SIGNALING_PATHWAY GOBP_POSITIVE_REGULATION_OF_LONG_TERM_SYNAPTIC_POTENTIATION, GOBP_POSITIVE_REGULATION_OF_LONG_TERM_SYNAPTIC_POTENTIATION GOBP_VENTRICULAR_SEPTUM_MORPHOGENESIS, GOBP_VENTRICULAR_SEPTUM_MORPHOGENESIS GOMF_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATE_5_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATE_5_PHOSPHATASE_ACTIVITY GOBP_NOTCH_RECEPTOR_PROCESSING_LIGAND_DEPENDENT, GOBP_NOTCH_RECEPTOR_PROCESSING_LIGAND_DEPENDENT GOBP_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBP_REGULATION_OF_ORGANIC_ACID_TRANSPORT REACTOME_SIGNALING_BY_ERBB2_IN_CANCER, REACTOME_SIGNALING_BY_ERBB2_IN_CANCER GOBP_NEGATIVE_REGULATION_OF_TUBULIN_DEACETYLATION, GOBP_NEGATIVE_REGULATION_OF_TUBULIN_DEACETYLATION MIR4737, MIR4737 GOMF_NEUREGULIN_BINDING, GOMF_NEUREGULIN_BINDING GOMF_INSULIN_BINDING, GOMF_INSULIN_BINDING HP_CEPHALOHEMATOMA, HP_CEPHALOHEMATOMA HP_KNEE_DISLOCATION, HP_KNEE_DISLOCATION WP_GLIAL_CELL_DIFFERENTIATION, WP_GLIAL_CELL_DIFFERENTIATION GOBP_REGULATION_OF_GERMINAL_CENTER_FORMATION, GOBP_REGULATION_OF_GERMINAL_CENTER_FORMATION GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS, GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS GOBP_NEUROTRANSMITTER_UPTAKE, GOBP_NEUROTRANSMITTER_UPTAKE GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT HP_ANNULAR_PANCREAS, HP_ANNULAR_PANCREAS REACTOME_SIGNALING_BY_EGFR_IN_CANCER, REACTOME_SIGNALING_BY_EGFR_IN_CANCER BIOCARTA_IL6_PATHWAY, BIOCARTA_IL6_PATHWAY PID_PDGFRA_PATHWAY, PID_PDGFRA_PATHWAY GOBP_ENDOCARDIAL_CUSHION_DEVELOPMENT, GOBP_ENDOCARDIAL_CUSHION_DEVELOPMENT GOBP_REGULATION_OF_MICROTUBULE_BINDING, GOBP_REGULATION_OF_MICROTUBULE_BINDING GOMF_CHLORIDE_CHANNEL_INHIBITOR_ACTIVITY, GOMF_CHLORIDE_CHANNEL_INHIBITOR_ACTIVITY GOBP_NEGATIVE_REGULATION_OF_CELL_MATRIX_ADHESION, GOBP_NEGATIVE_REGULATION_OF_CELL_MATRIX_ADHESION GOBP_REGULATION_OF_GLYCOGEN_METABOLIC_PROCESS, GOBP_REGULATION_OF_GLYCOGEN_METABOLIC_PROCESS HP_SPONTANEOUS_RECURRENT_EPISTAXIS, HP_SPONTANEOUS_RECURRENT_EPISTAXIS REACTOME_ATF6_ATF6_ALPHA_ACTIVATES_CHAPERONES, REACTOME_ATF6_ATF6_ALPHA_ACTIVATES_CHAPERONES GOMF_BMP_RECEPTOR_ACTIVITY, GOMF_BMP_RECEPTOR_ACTIVITY REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_BCR_SIGNALING, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_BCR_SIGNALING GOBP_CARDIAC_ATRIUM_MORPHOGENESIS, GOBP_CARDIAC_ATRIUM_MORPHOGENESIS HP_PULMONARY_INTERSTITIAL_HIGH_RESOLUTION_COMPUTED_TOMOGRAPHY_ABNORMALITY, HP_PULMONARY_INTERSTITIAL_HIGH_RESOLUTION_COMPUTED_TOMOGRAPHY_ABNORMALITY MIR1914_5P, MIR1914_5P REACTOME PI3K EVENTS IN ERBB2 SIGNALING, REACTOME PI3K EVENTS IN ERBB2 SIGNALING HP_ANKLE_WEAKNESS, HP_ANKLE_WEAKNESS PASTURAL_RIZ1_TARGETS_DN, PASTURAL_RIZ1_TARGETS_DN GOCC_CALCINEURIN_COMPLEX, GOCC_CALCINEURIN_COMPLEX GOMF_CYTOSKELETON_NUCLEAR_MEMBRANE_ANCHOR_ACTIVITY, GOMF_CYTOSKELETON_NUCLEAR_MEMBRANE_ANCHOR_ACTIVITY MIR4485_5P, MIR4485_5P GOBP_LYMPHATIC_ENDOTHELIAL_CELL_DIFFERENTIATION, GOBP_LYMPHATIC_ENDOTHELIAL_CELL_DIFFERENTIATION GOBP_DNA_DAMAGE_INDUCED_PROTEIN_PHOSPHORYLATION, GOBP_DNA_DAMAGE_INDUCED_PROTEIN_PHOSPHORYLATION GOBP_ATRIAL_SEPTUM_MORPHOGENESIS, GOBP_ATRIAL_SEPTUM_MORPHOGENESIS