	GSE19825_NAIVE_VS_IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN GSE17721_CPG_VS_GARDIQUIMOD_1H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_1H_BMDC_DN GSE6674_CPG_VS_P12_3_STIM_BCELL_DN, GSE6674_CPG_VS_P12_3_STIM_BCELL_DN GSE15139_GMCSF_TREATED_VS_UNTREATED_NEUTROPHILS_DN, GSE15139_GMCSF_TREATED_VS_UNTREATED_NEUTROPHILS_DN GSE15139_CSE4649_NAIVE_VS_ACTIVATED_CD8_TCELL_MIR155_KO_UP GSE4649_NAIVE_VS_ACTIVATED_ACT_FOXP3_HET_TCONV_UP, GSE4649_NAIVE_VS_ACTIVATED_ACT_FOXP3_HET_TCONV_UP GSE17721_POLYIC_VS_CPG_12H_BMDC_DN, GSE17721_POLYIC_VS_CPG_12H_BMDC_DN GSE17721_POLYIC_VS_CPG_12H_BMDC_DN, GSE17721_POLYIC_VS_CPG_12H_BMDC_DN GSE27786_CD4_VS_CD8_TCELL_DN, CSE27786_CD4_VS_CD8_TCELL_DN
	GSE17301_CTRL_VS_48H_ACD3_ACD28_IFNA2_STIM_CD8_TCELL_UP, GSE17301_CTRL_VS_48H_ACD3_ACD28_IFNA2_STIM_CD8_TCELL_UP GSE6674_PL2_3_VS_ANTI_IGM_AND_CPC_STIM_BCELL_DN, GSE6674_PL2_3_VS_ANTI_IGM_AND_CPC_STIM_BCELL_DN GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_DN, GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_DN GSE17721_LPS_VS_GARDIQUIMOD_SH_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_8H_BMDC_DN GSE40274_CTRL_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP ELVIDGE_HYPOXIA_DN, ELVIDGE_HYPOXIA_DN GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_1H_DN, GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_1H_DN GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_2H_DN, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_2H_DN
	GSE17186_BLOOD_VS_CORD_BLOOD_NAIVE_BCELL_UP, GSE17186_BLOOD_VS_CORD_BLOOD_NAIVE_BCELL_UP GSE6674_CPG_VS_CPG_AND_ANTIIGM_STIM_BCELL_DN, GSE6674_CPG_VS_CPG_AND_ANTIIGM_STIM_BCELL_DN GSE6092_UNSTIM_VS_IFNG_STIM_ENDOTHELIAL_CELL_UP, GSE6781_UNSTIM_VS_IFNG_STIM_ENDOTHELIAL_CELL_UP GSE781_UNSTIM_VS_CPG_STIM_PDC_1H_UP, GSE781_UNSTIM_VS_CPG_STIM_PDC_1H_UP GSE781_UNSTIM_VS_THI7_DAY5_POST_POLARIZATION_DN, GSE2603_THI_VS_THI7_DAY5_POST_POLARIZATION_DN GSE36888_STA75_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_17H_UP, GSE36888_STA75_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_17H_UP GSE17721_CTRL_VS_POLVIC_24H_BMDC_UP, GSE17721_CTRL_VS_POLVIC_24H_BMDC_UP GSE17721_POLVIC_VS_CPG_24H_BMDC_DN, GSE17721_POLVIC_VS_CPG_24H_BMDC_DN GSE17721_POLVIC_VS_CPG_24H_BMDC_DN GSE37872_TIVAMUS_VS_PAT_TCONV_DN_GSE378721_POLVIC_VS_CPG_24H_BMDD_DN
	GSE7852_THYMUS_VS_FAT_TCONV_DN, GSE7852_THYMUS_VS_FAT_TCONV_DN GSE20715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN, GSE20715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN GSE40715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN, GSE40715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN GSE40716_WT_VS_LPS_TIM_DC_DN, GSE40715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN GSE40716_WT_VS_LPS_TIM_DC_DN, GSE40716_WT_VS_LPS_STIM_DC_DN GSE40716_WT_VS_LPS_TIM_DC_DN, GSE40716_WT_VS_LPS_STIM_DC_DN GOLDRATH_NAIVE_VS_EFF_CD8_TCELL_UP, GOLDRATH_NAIVE_VS_EFF_CD8_TCELL_UP GSE4079_DCC205_POS_DC_VS_BCELL_DN, GSE4079_DCC205_POS_DC_VS_BCELL_DN GSE4079_DCC205_POS_DC_VS_BCELL_DN, GSE4079_DCC205_POS_DC_VS_BCELL_DN GSE4079_DCC205_POS_DC_VS_BCELL_DN, GSE4079_DCC_DN, GSE4079_DCC_DN, GSE4079_DCC_DN GSE4079_DCC205_POS_DC_VS_BCELL_DN, GSE4079_DCC_DN, GSE4079_DCC_DN GSE4079_DCC704_VS_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_VS_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_VS_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC704_UB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC704_UB_UB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UP, GSE4070_DCC_DN GSE4079_DCC704_US_ULB_UB_CGCAN_CCAL_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_UB_CCAC_UB_CCAC_UB_UB_CCAC_UB_CCAC_UB_CCAC_UB_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_UB_CCAC_U
	CSE39820_CTRL_VS_IL1B_IL6_CD4_TCELL_UP, CSE39820_CTRL_VS_IL1B_IL6_CD4_TCELL_UP CSE15331_HSC_VS_GRANULOCYTE_MONOCYTE_MONOCYTE_MONOCYTE_MONOCYTE_PROGENITOR_UP GSE17721_POLVIC_VS_GARDIQUIMOD_4H_BMDC_DN, GSE17721_POLVIC_VS_GARDIQUIMOD_4H_BMDC_DN GSE20366_TREG_VS_NAIVE_CD4_TCELL_UP, GSE20366_TREG_VS_NAIVE_CD4_TCELL_UP GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_UP, GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_UP GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_UP, GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_UP GSE40277_EOS_AND_LEFI_TRANSDUCED_VS_CTRL_CD4_TCELL_DN, GSE40277_EOS_AND_LEFI_TRANSDUCED_VS_CTRL_CD4_TCELL_DN HP_ABNORMALITY_OF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_OF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_DF_BONE_MARROW_CELL_MORPHOLOGY GSE47580_TREG_VS_TEFF_S_MANSONI_INF_UP, GSE47580_TREG_VS_TEFF_S_MANSONI_INF_UP, GSE47580_TREG_VS_TEFF_S_MANSONI_INF_UP, GSE47580_TREG_VS_TEFF_S_MANSONI_INF_UP, GSE47580_TREG_VS_TEFF_S_MANSONI_INF_UP
	GOBP_SULFUR_COMPOUND_BIOSYNTHETIC_PROCESS, GOBP_SULFUR_COMPOUND_BIOSYNTHETIC_PROCESS GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_6H_UP GSE17721_CTRL_VS_POLVIC_0.5H_BMDC_UP, GSE17721_CTRL_VS_POLVIC_0.5H_BMDC_UP, GSE17721_CTRL_VS_POLVIC_0.5H_BMDC_UP, GSE17721_CTRL_VS_POLVIC_USH_BMDC_UP, GSE17721_FOLVIC_USH_BMDC_UP, GSE36527_CD62L_HOR_UP, GSE36527_CD62L_HOR_UP, GSE36527_CD62L_HOR_UP, GSE36527_CD62L_HIGH_VS_CD62_LOW_TREG_CD69_NEG_KLRG1_NEG_UP GSE36527_CD62L_HOW_TREG_CD69_LCG_UP, GSE36527_CD62L_HIGH_VS_CD62_LOW_TREG_CD69_NEG_KLRG1_NEG_UP GSE3602_LDAYIC_VS_GARDIQUIMOD_IH_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_IH_BMDC_DN GSE3601_LMAJOR_VS_M_TUBERCULOSIS_DC_UP, GSE3601_LMAJOR_VS_M_TUBERCULOSIS_DC_UP
	HP_NARROW_PALATE GSE14350_IL2RB_KO_VS_WT_TEFF_DN, GSE14350_IL2RB_KO_VS_WT_TEFF_DN GSE17721_[12H_VS_24H_PAM3CSK4_BMDC_UP, GSE17721_12H_VS_24H_PAM3CSK4_BMDC_UP GOBP_REGULATION_OF_GENE_SILENCING, GOBP_REGULATION_OF_GENE_SILENCING GSE17721_CTRL_VS_GARDIQUIMOD_8H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_8H_BMDC_DN RB_P107_DN.V1_DN, RB_P107_DN.V1_DN HALLMARK_ESTROGEN_RESPONSE_EARLY, HALLMARK_ESTROGEN_RESPONSE_EARLY GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_UP, GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_UP GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_UP, GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_UP GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_UP GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_UP
	GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP, GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP GSE41978_WT_VS_ID2_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_WT_VS_ID2_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN HP_IMMUNOLOGIC_HYPERSENSITIVITY, HP_IMMUNOLOGIC_HYPERSENSITIVITY GSE32901_NAIVE_VS_TH17_ENRICHED_CD4_TCELL_DN, GSE32901_NAIVE_VS_TH17_ENRICHED_CD4_TCELL_DN GOBP_REGULATION_OF_GENE_SILENCING_BY_RNA, GOBP_REGULATION_OF_GENE_SILENCING_BY_RNA HALLMARK_ALLOGRAFT_REJECTION, HALLMARK_ALLOGRAFT_REJECTION HP_SKELETAL_DYSPLASIA, HP_SKELETAL_DYSPLASIA GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_DN, GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_DN
	REACTOME_UNFOLDED_PROTEIN_RESPONSE_UPR, REACTOME_UNFOLDED_PROTEIN_RESPONSE_UPR MIR8087, MIR8087 REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS, REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS HP_CRANIOSYNOSTOSIS, HP_CRANIOSYNOSTOSIS HALLMARK_ESTROGEN_RESPONSE_LATE, HALLMARK_ESTROGEN_RESPONSE_LATE MIR1343_3P, MIR1343_3P HP_ABNORMALITY_OF_ODONTOID_TISSUE, HP_ABNORMALITY_OF_ODONTOID_TISSUE GSE4811_CLASSSICALY_ACTIVATED_VS_TYPE_2_ACTIVATED_MACROPHAGE_UP, GSE4811_CLASSSICALY_ACTIVATED_US_TYPE_2_ACTIVATED_MACROPHAGE_UP
	HP_MACROCYTIC_ANEMIA, HP_MACROCYTIC_ANEMIA REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE, REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP, GSE19401_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP MODULE_169, MODULE_169 HP_LIMB_HYPERTONIA, HP_LIMB_HYPERTONIA YYCATTICAWW_UNKNOWN, YYCATTICAWW_UNKNOWN GSE17721_CTRIL_VS_CPG_4H_BMDC_DN, GSE17721_CTRIL_VS_CPG_4H_BMDC_DN MIR6757_3P, MIR6757_3P
	GSE29618_BCELL_VS_MONOCYTE_UP, GSE29618_BCELL_VS_MONOCYTE_UP GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_DN, GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_DN HP_BONE_MARROW_HYPOCELLULARITY, HYBONE_MARROW_HYPOCELLULARITY GOBP_NUCLEOSIDE_BISPHOSPHATE_BIOSYNTHETIC_PROCESS, GDE_BISPHOSPHATE_BIOSYNTHETIC_PROCESS HP_ABNORMAL_LYMPHOCYTE_MORPHOLOGY, HP_ABNORMAL_LYMPHOCYTE_MORPHOLOGY GSE2937_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_DN, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_DN WOO_LIVER_CANCER_RECURRENCE_UP, WOO_LIVER_CANCER_RECURRENCE_UP GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
	HP_ASTHMA, HP_ASTHMA HP_FIBROUS_TISSUE_NEOPLASM GSE20618_DCTLL_VS_MONOCYTE_DAY7_FLU_VACCINE_UP HP_CARIOUS_TEETH, HP_CARIOUS_TEETH JI_RESPONSE_TO_FSH_DN, JI_RESPONSE_TO_FSH_DN GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_CD8_TCELL_UP, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_CD8_TCELL_UP SHAFFER_IRF4_TARGETS_IN_PLASMA_CELL_VS_MATURE_B_LYMPHOCYTE MIR4699_5P, MIR4699_5P
	HP_INCREASED_MEAN_CORPUSCULAR_VOLUME, HP_INCREASED_MEAN_CORPUSCULAR_VOLUME GOBP_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY GOBP_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY, GOBP_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY GOBP_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GOBP_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY NGUYEN_NOTCHI_TARGETS_DN, NGUYEN_NOTCHI_TARGETS_DN HP_ABNORMAL_CIRCULATING_IGG_LEVEL, HP_ABNORMAL_CIRCULATING_IGG_LEVEL MIR589_5P, MIR589_5P GOCC_INCLUSION_BODY, GOCC_INCLUSION_BODY
	NAGASHIMA_EGF_SIGNALING_UP, NAGASHIMA_EGF_SIGNALING_UP KEGG_TGF_BETA_SIGNALING_PATHWAY, KEGG_TGF_BETA_SIGNALING_PATHWAY GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_CH_OH_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_CH_OH_GROUP_OF_DONORS KEGG_STEROID_BIOSYNTHESIS, KEGG_STEROID_BIOSYNTHESIS, KEGG_STEROID_BIOSYNTHESIS, KEGG_STEROID_BIOSYNTHESIS, KEGG_STEROID_BIOSYNTHESIS, TOTAL BIOSYNTHESIS, TOTAL BIOSYNTH B
	MIR517_5P, MIR517_5P RASHI_RESPONSE_TO_IONIZING_RADIATION_3, RASHI_RESPONSE_TO_IONIZING_RADIATION_3 GOBP_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY, GOBP_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY MIR3116 SMIRNOV_RESPONSE_TO_IR_2HR_UP, SMIRNOV_RESPONSE_TO_IR_2HR_UP MIR3927_5P, MIR3927_5P AGGTGCA_MIR500, AGGTGCA_MIR500 SESTO_RESPONSE_TO_UV_C5, SESTO_RESPONSE_TO_UV_C5
	AIZARANI_LIVER_C8. RESIDENT_B_CELLS_1, AIZARANI_LIVER_C8. RESIDENT_B_CELLS_1 HP_ABNORMALITY_OF_VITAMIN_METABOLISM GS_28237_EARLY_VS_LATE_GC_BCELL_UP, GSE28237_EARLY_VS_LATE_GC_BCELL_UP GOBP_GLAND_MORPHOGENESIS, GOBP_GLAND_MORPHOGENESIS WP_CHOLESTEROL_METABOLISM_INCLUDES_BOTH_BLOCH_AND_KANDUTSCHRUSSELL_PATHWAYS, WP_CHOLESTEROL_METABOLISM_INCLUDES_BOTH_BLOCH_AND_KANDUTSCHRUSSELL_PATHWAYS HP_LYMPHOPENIA, HP_LYMPHOPENIA MIR642A_5P, MIR642A_5P COBP_PHOSPHOLIPID_TRANSPORT, GOBP_PHOSPHOLIPID_TRANSPORT GTITNYNNGGTNA_UNKNOWN, GTITNYNNGGTNA_UNKNOWN PLINGTHANSPORT, UNKNOWN STITNYNNGGTNA_UNKNOWN STITNYNNGGTNA_UNKNOWN STITNYNNGGTNA_UNKNOWN
	HP_ABNORMAL_CIRCULATING_IGA_LEVEL, HP_ABNORMAL_CIRCULATING_IGA_LEVEL MIR874_3P, MIR874_3P GOBP_NEGATIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION, GOBP_NEGATIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION GOCC_NUCLEAR_PORE_NUCLEAR_BASKET, GOCC_NUCLEAR_BASKET HP_ABNORMAL_T_CELL_MORPHOLOGY, HP_ABNORMAL_T_CELL_MORPHOLOGY WP_METASTATIC_BRAIN_TUMOR, WP_METASTATIC_BRAIN_TUMOR HP_ABNORMAL_CIRCULATING_IGM_LEVEL, HP_ABNORMAL_CIRCULATING_IGM_LEVEL HP_INFLAMMATION_OF_THE_LARGE_INTESTINE, HP_INFLAMMATION_OF_THE_LARGE_INTESTINE
	GOBP_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS TCTAGAG_MIR517, TCTAGAG_MIR517 HP_ANTINUCLEAR_ANTIBODY_POSITIVITY, HP_ANTINUCLEAR_ANTIBODY_POSITIVITY MIR6505_3P, MIR6505_3P HP_CUPPED_EAR, HP_CUPPED_EAR REACTOME_ONCOGENE_INDUCED_SENESCENCE, REACTOME_ONCOGENE_INDUCED_SENESCENCE MARCHINI_TRABECTEDIN_RESISTANCE_DN, MARCHINI_TRABECTEDIN_RESISTANCE_DN
	HP_ABNORMALITY_OF_EXOCRINE_PANCREAS_PHYSIOLOGY, HP_ABNORMALITY_OF_EXOCRINE_PANCREAS_PHYSIOLOGY HP_RECURRENT_VIRAL_INFECTIONS, HP_RECURRENT_VIRAL_INFECTIONS HP_ABNORMAL_T_CELL_SUBSET_DISTRIBUTION, HP_ABNORMAL_T_CELL_SUBSET_DISTRIBUTION GOBP_REGULATION_OF_MEMBRANE_PERMEABILITY, GOBP_REGULATION_OF_MEMBRANE_PERMEABILITY MIR4704_5P, MIR4704_5P LEIN_MEDULLA_MARKERS, LEIN_MEDULLA_MARKERS NATSUME_RESPONSE_TO_INTERFERON_BETA_UP, NATSUME_RESPONSE_TO_INTERFERON_BETA_UP HAHTOLA_CTCL_PATHOGENESIS, HAHTOLA_CTCL_PATHOGENESIS
	HP_ABNORMAL_ERYTHROCYTE_SEDIMENTATION_RATE, HP_ABNORMAL_ERYTHROCYTE_SEDIMENTATION_RATE HP_POLYARTICULAR_ARTHROPATHY, HP_POLYARTICULAR_ARTHROPATHY WP_IL2_SIGNALING_PATHWAY, WP_IL2_SIGNALING_PATHWAY GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS, GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS HP_COLITIS, HP_COLITIS ZHONG_PFC_C1_ASTROCYTE, ZHONG_PFC_C1_ASTROCYTE HP_ABNORMAL_BRONCHUS_MORPHOLOGY, HP_ABNORMAL_BRONCHUS_MORPHOLOGY HP_ABNORMAL_INCOMMAL_BRONCHUS_MORPHOLOGY, HP_ABNORMAL_INCOMMAL_IN
	HP_BONE_MARROW_HYPERCELLULARITY, HP_BONE_MARROW_HYPERCELLULARITY BRUNO_HEMATOPOIESIS, BRUNO_HEMATOPOIESIS HP_ABNORMAL_B_CELL_MORPHOLOGY, HP_ABNORMAL_B_CELL_MORPHOLOGY PID_IL12_2PATHWAY, PID_IL12_2PATHWAY ROESSLER_LIVER_CANCER_METASTASIS_DN, ROESSLER_LIVER_CANCER_METASTASIS_DN MODULE_73, MODULE_73 GOBP_POSITIVE_REGULATION_OF_RRNA_PROCESSING, GOBP_POSITIVE_REGULATION_OF_RRNA_PROCESSING GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION
	RUAN_RESPONSE_TO_TNF_TROGLITAZONE_DN, RUAN_RESPONSE_TO_TNF_TROGLITAZONE_DN HP_DECREASED_ADIPOSE_TISSUE, HP_DECREASED_ADIPOSE_TISSUE PID_IL2_PI3K_PATHWAY, PID_IL2_PI3K_PATHWAY MODULE_349, MODULE_349, MODULE_349, BIOCARTA_IL2RB_PATHWAY, BIOCARTA_IL2RB_PATHWAY MIR4536_3P, MIR4536_3P AIZARANI_LIVER_C22_RESIDENT_B_CELLS_2, AIZARANI_LIVER_C22_RESIDENT_B_CELLS_2 HP_ANOTIA, HP_ANOTIA
	HP_ATHETOSIS, HP_ATHETOSIS MIR1178_3P, MIR1178_3P HP_T_LYMPHOCYTOPENIA, HP_T_LYMPHOCYTOPENIA MCCLUNG_CREBI_TARGETS_UP, MCCLUNG_CREBI_TARGETS_UP GCM_DEAFI, GCM_DEAFI REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS, REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS MIR5581_5P, MIR5581_5P,
	GOBP_PROTEIN_HETEROOLIGOMERIZATION, GOBP_PROTEIN_HETEROOLIGOMERIZATION HP_ASCENDING_TUBULAR_AORTA_ANEURYSM, HP_ASCENDING_TUBULAR_AORTA_ANEURYSM MIR1229_5P, MIR1229_5P HP_IMPAIRED_ANTIGEN_SPECIFIC_RESPONSE, HP_IMPAIRED_ANTIGEN_SPECIFIC_RESPONSE GOBP_MONOCYTE_DIFFERENTIATION, GOBP_MONOCYTE_DIFFERENTIATION GOBP_MONOCYTE_DIFFERENTIATION, GOBP_MONOCYTE_DIFFERENTIATION GOBTILLE_UV_RESPONSE_CLUSTER_DQ, GENTILE_UV_RESPONSE_CLUSTER_DQ TRAVAGLINI_LUNG_NATURAL_KILLER_T_CELL, TRAVAGLINI_LUNG_NATURAL_KILLER_T_CELL GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS
	OND_FOXP3_TARGETS_UP, OND_FOXP3_TARGETS_UP MIR1322 MORF_ZNF10, MORF_ZNF10 GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS, GOBP_NEGATIVE_REGULATION_TO_NUCLEUS GOBP_CORTICAL_ACTIN_CYTOSKELETON_ORGANIZATION, GOBP_CORTICAL_ACTIN_CYTOSKELETON_ORGANIZATION MARIADASON_RESPONSE_TO_CUCUMIN_SULINDAC_5, MARIADASON_RESPONSE_TO_CURCUMIN_SULINDAC_5 GOBP_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY, GOBP_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY HP_B_LYMPHOCYTOPENIA, HP_B_LYMPHOCYTOPENIA
	GOBP_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY GOUYER_TATI_TARGETS_UP, GOUYER_TATI_TARGETS_UP HP_EXPRESSIVE_LANGUAGE_DELAY, HP_EXPRESSIVE_LANGUAGE_DELAY HP_BROAD_EYEBROW, HP_BROAD_EYEBROW GOBP_REGULATION_OF_GLIOGENESIS, GOBP_REGULATION_OF_GLIOGENESIS LANDIS_BREAST_CANCER_PROGRESSION_UP, LANDIS_BREAST_CANCER_PROGRESSION_UP HP_DECREASED_CIRCULATING_LEVEL_OF_SPECIFIC_ANTIBODY, HP_DECREASED_CIRCULATING_LEVEL_OF_SPECIFIC_ANTIBODY REACTOME_CDO6_ASSOCIATION_WITH_THE_ORC_ORIGIN_COMPLEX, REACTOME_CDC6_ASSOCIATION_WITH_THE_ORC_ORIGIN_COMPLEX
	GOBP_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION HP_AUTOIMMUNE_HEMOLYTIC_ANEMIA, HP_AUTOIMMUNE_HEMOLYTIC_ANEMIA HP_ABNORMALITY_OF_VITAMIN_B12_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_B12_METABOLISM HP_ABNORMAL_CIRCULATING_IGE_LEVEL, HP_ABNORMAL_CIRCULATING_IGE_LEVEL GOBP_REGULATION_OF_STEM_CELL_POPULATION_MAINTENANCE, GOBP_REGULATION_OF_STEM_CELL_POPULATION_MAINTENANCE GOBP_POSITIVE_REGULATION_OF_GLIOGENESIS, GOBP_POSITIVE_REGULATION_OF_GLIOGENESIS WATANABE_COLON_CANCER_MSI_VS_MSS_UP, WATANABE_COLON_CANCER_MSI_VS_MSS_UP HP_11_PAIRS_OF_RIBS, HP_11_PAIRS_OF_RIBS
	GOBP_FATTY_ACID_DERIVATIVE_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACID_DERIVATIVE_BIOSYNTHETIC_PROCESS REACTOME_INTERLEUKIN_2_SIGNALING, REACTOME_INTERLEUKIN_2_SIGNALING GOBP_REGULATION_OF_ENDOTHELIAL_CELL_DEVELOPMENT, GOBP_REGULATION_OF_ENDOTHELIAL_CELL_DEVELOPMENT HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_MITOGEN, HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_MITOGEN HP_CONTROL CLUNG_DISEASE, HP_CHRONIC_LUNG_DISEASE HP_MALNUTRITION, HP_MALNUTRITION HP_FAT_MALABSORPTION, HP_FAT_MALABSORPTION HP_FAT_MALABSORPTION, HP_FAT_MALABSORPTION HP_DECREASED_CIRCULATING_TOTAL_IGM, HP_DECREASED_CIRCULATING_TOTAL_IGM
	GCM_DDX11, GCM_DDX11 AMIT_EGF_RESPONSE_240_HELA, AMIT_EGF_RESPONSE_240_HELA AMIT_EGF_RESPONSE_240_HELA, AMIT_EGF_RESPONSE_240_HELA SHIN_B_CELL_LYMPHOMA_CLUSTER_3, SHIN_B_CELL_LYMPHOMA_CLUSTER_3 GAUTSCHI_SRC_SIGNALING, GAUTSCHI_SRC_SIGNALING GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS, GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS GOBP_SOSTIIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY, GOBP_POSITIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY HP_SOFT_SKIN, HP_SOFT_SKIN AMIT_EGF_RESPONSE_240_MCF10A, AMIT_EGF_RESPONSE_240_MCF10A
	GOMF_ACYL_COA_BINDING, GOMF_ACYL_COA_BINDING GOMF_ACYL_COA_BINDING, GOMF_ACYL_COA_BINDING GOMF_ACYL_COA_BINDING, GOMF_ACYL_COA_BINDING GOMF_ACYL_COA_BINDING, GOMF_ACYL_COA_BINDING GOMF_ACYL_COA_BINDING, GOMF_ACYL_COA_BINDING HP_SMEDDING_BIOSYNTHETIC_PROCESS, GOBP_METHIONINE_BIOSYNTHETIC_PROCESS HP_OSTEOMYELITIS, HP_OSTEOMYELITIS, POSTEOMYELITIS, POSTEOMYELITIS, POSTEOMYELITIS, POSTEOMYELITIS, POSTEOMYELITIS, POSTEOMYELITIS, POSTEOMYELITIS, GOBP_ANGIOGENESIS_INVOLVED_IN_WOUND_HEALING HP_SLENDER_BUILD, HP_SLENDER_BUILD GOBP_NEGATIVE_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_NEGATIVE_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION WP_INFLAMMATORY_RESPONSE_PATHWAY, WP_INFLAMMATORY_RESPONSE_PATHWAY MIR194_3P, MIR194_3P GOMF_THIOLESTER_HYDROLASE_ACTIVITY, GOMF_THIOLESTER_HYDROLASE_ACTIVITY GOMF_THIOLESTER_HYDROLASE_ACTIVITY, GOMF_THIOLESTER_HYDROLASE_ACTIVITY
	GOBP_FAT_CELL_PROLIFERATION, GOBP_FAT_CELL_PROLIFERATION XU_CREBBP_TARGETS_DN, XU_CREBBP_TARGETS_DN HP_ABNORMAL_CELL_PROLIFERATION, HP_ABNORMAL_CELL_PROLIFERATION GOBP_POSITIVE_REGULATION_OF_NON_MOTILE_CILIUM_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_NON_MOTILE_CILIUM_ASSEMBLY REACTOME_INTERLEUKIN_RECEPTOR_SHC_SIGNALING, REACTOME_INTERLEUKIN_RECEPTOR_SHC_SIGNALING GOBP_NEUTRAL_LIPID_METABOLIC_PROCESS, GOBP_NEUTRAL_LIPID_METABOLIC_PROCESS HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS, HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS, KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS
	GOME_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OF_REDUCTION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONOR_AND_INCORPORATION_OF_OXYGEN, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OF_REDUCTION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONOR_PID_RAS_PATHWAY, PID_RAS_PATHWAY PID_RAS_PATHWAY, PID_RAS_PATHWAY PID_RAS_PATHWAY, PID_RAS_PATHWAY PID_RAS_PATHWAY, PID_RAS_PATHWAY P
	HP_TETRAPLEGIA, HP_TETRAPLEGIA GOBP_COENZYME_A_METABOLIC_PROCESS, GOBP_COENZYME_A_METABOLIC_PROCESS ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_7DY_UP, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_7DY_UP PID_IL12_STAT4_PATHWAY, PID_IL12_STAT4_PATHWAY GOBP_NEGATIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION, GOBP_NEGATIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION GOBP_NEGATIVE_REGULATION_OF_EPITHELIAL_CELL_PROLIFERATION, GOBP_NEGATIVE_REGULATION_OF_EPITHELIAL_CELL_PROLIFERATION WP_INTERACTIONS_BETWEEN_IMMUNE_CELLS_AND_MICRORAS_IN_TUMOR_MICROENVIRONMENT, WP_INTERACTIONS_BETWEEN_IMMUNE_CELLS_AND_MICROENVIRONMENT REACTOME_VITAMIN_BS_PANTOTHENATE_METABOLISM, REROAS_IN_TUMOR_MICROENVIRONMENTE_METABOLISM, REROAS_IN_TUMOR_BS_PANTOTHENATE_METABOLISM, REROAS_IN_TUMOR_BS_PANTOTHENATE_METABOLISM.
	LANDIS_ERBB2_BREAST_PRENEOPLASTIC_UP, LANDIS_ERBB2_BREAST_PRENEOPLASTIC_UP GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION_IN_RESPONSE_TO_MITOCHONDRIAL_DEPOLARIZATION, GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRIAL_DEPOLARIZATION HP_CUPPED_RIBS, HP_CUPPED_RIBS (GOBP_REGULATION_OF_TRANSLATIONAL_ELONGATION, GOBP_REGULATION_OF_TRANSLATIONAL_ELONGATION, GOBP_REGULATION_OF_TRANSLATIONAL_ELONGATION, GOBP_REGULATION_OF_TRANSLATIONAL_ELONGATION HP_METAPHYSEAL_DYSPLASIA, HP_METAPPHYSEAL_DYSPLASIA HP_ABNORMAL_VITAMIN_B12_LEVEL, HP_ABNORMAL_VITAMIN_B12_LEVEL HP_INCREASED_CIRCULATING_IGG_LEVEL, HP_INCREASED_CIRCULATING_IGG_LEVEL TING_SILENCED_BY_DICER, TING_SILENCED_BY_DICER
IGM_VS_PL2_3_STIM_BCELL_DN, GSE6674_ANTI_IGM_VS_PL2_3_STIM_BCELL_DN	HP_PERPHERAL_ARTERIAL_STENOSIS, HP_PERPHERAL_ARTERIAL_STENOSIS HP_ACQUIRED_ABNORMAL_HAIR_PATTERN, HP_ACQUIRED_ABNORMAL_HAIR_PATTERN GOBP_GLANDULAR_EPITHELIAL_CELL_DEVELOPMENT, GOBP_GLANDULAR_EPITHELIAL_CELL_DEVELOPMENT HP_WIDE_CRANIAL_SUTURES, HP_WIDE_CRANIAL_SUTURES HP_LOW_LEVELS_OF_VITAMIN_D, HP_LOW_LEVELS_OF_VITAMIN_D WP_MAMMARY_GLAND_DEVELOPMENT_PATHWAY_EMBRYONIC_DEVELOPMENT_STAGE_1_OF_4, WP_MAMMARY_GLAND_DEVELOPMENT_PATHWAY_EMBRYONIC_DEVELOPMENT_STAGE_1_OF_4 HP_ABNORMAL_CIRCULATING_MONOCARBOXYLIC_ACID_COCENTRATION, HP_ABNORMAL_CIRCULATING_MONOCARBOXYLIC_ACID_COCENTRATION HP_HOMOCYSTINURIA, HP_HOMOCYSTINURIA GOBP_CELLULAR_RESPONSE_TO_LEUCINE_STARVATION, GOBP_CELLULAR_RESPONSE_TO_LEUCINE_STARVATION ### COCHECULAR ARTERIAL_STENOSES, HP_ACQUIRED_ABNORMAL_HAIR_PATTERN ### COCHECT ARTERIAL_STENOSES, HP_ACQUIRED_ABNORMAL_HAIR_PATTERN ### COCHECT ARTERIAL_TESTON SET OF A COCHECT ARTERIAL
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	VALK_AML_WITH_FLT3_ITD, VALK_AML_WITH_FLT3_ITD HP_LOWER_LIMB_METAPHYSEAL_IRREGULARITY, HP_LOWER_LIMB_METAPHYSEAL_IRREGULARITY HP_LOWER_LIMB_METAPHYSEAL_IRREGULARITY, HP_LOWER_LIMB_METAPHYSEAL_IRREGULARITY GOBP_REGULATION_OF_MICROTUBULE_BINDING, GOBP_REGULATION_OF_MICROTUBULE_BINDING GOBP_METHIONINE_METABOLIC_PROCESS, GOBP_METHABOLIC_PROCESS, GOBP_METHONINE_METABOLIC_PROCESS, GOBP_METHIONINE_METABOLIC_PROCESS, GOBP_METHIONINE_METABOLIC_PROCESS, GOBP_METHION_OF_INTERLEUKIN_17_PRODUCTION GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_17_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_17_PRODUCTION WP_COVID19_ADVERSE_OUTCOME_PATHWAY, WP_COVID19_ADVERSE_OUTCOME_PATHWAY WP_STATIN_PATHWAY, WP_STATIN_PATHWAY SHIN_B_CELL_LYMPHOMA_CLUSTER_2, SHIN_B_CELL_LYMPHOMA_CLUSTER_2
	GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_CELL_DIFFERENTIATION HP_ABNORMAL_OSSIFICATION_INVOLVING_THE_FEMORAL_HEAD_AND_NECK. HP_ABNORMAL_BLOOD_CHLORIDE_CONCENTRATION GOBP_OSITIVE_REGULATION_OF_CARDIAC_CONCENTRATION HP_ABNORMAL_BLOOD_CHLORIDE_CONCENTRATION GOBP_COLUMNAR_CUBOIDAL_EPIT_CHELIAL_CELL_DEVELOPMENT, GOBP_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DEVELOPMENT GOBP_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION MARCINIAK_ER_STRESS_RESPONSE_VIA_CHOP, MARCINIAK_ER_STRESS_RESPONSE_VIA_CHOP IIZUKA_LIVER_CANCER_PROGRESSION_C2_G3_DN, IIZUKA_LIVER_CANCER_PROGRESSION_C2_G3_DN GOMP_STEROID_DEHYDROCGENASE_ACTIVITY_CRAMES_CENTRATION_DOSENASE_ACTIVITY_CRAMES_CANTURE_CRAVEN_AND CANCER_CRAVEN_AND CANCER_CRAVEN_CR
	GOBP_POSITIVE_REGULATION_OF_FEVER_GENERATION, GOBP_POSITIVE_REGULATION_OF_FEVER_GENERATION GOBP_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_STEROID_FEROID_STERIBUTION, GOBP_MITOCHONDRION_DISTRIBUTION GOBP_MITOCHONDRION_DISTRIBUTION, GOBP_MITOCHONDRION_DISTRIBUTION GOBP_TRIGLYCERIDE_RICH_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_TRIGLYCERIDE_RICH_LIPOPROTEIN_PARTICLE_CLEARANCE GOBP_REGULATION_OF_MONOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_MONOCYTE_DIFFERENTIATION GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION HP_PORE_UNOTHORAX, HP_PNEUMOTHORAX GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION
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	BIOCARTA_IL2_PATHWAY, BIOCARTA_IL2_PATHWAY GOBP_MAMMARY_GLAND_EPITHELIAL_CELL_DIFFERENTIATION, GOBP_MAMMARY_GLAND_EPITHELIAL_CELL_DIFFERENTIATION GOBP_RESPONSE_TO_INTERLEUKIN_2, GOBP_RESPONSE_TO_INTERLEUKIN_2 GOBP_POSITIVE_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION HP_ABNORMAL_PANCREAS_SIZE, HP_ABNORMAL_PANCREAS_SIZE GOBP_REGULATION_OF_LIPID_CATABOLIC_PROCESS, GOBP_REGULATION_OF_LIPID_CATABOLIC_PROCESS REACTOME_METABOLISM_OF_FAT_SOLUBLE_VITAMINS, REACTOME_METABOLISM_OF_FAT_SOLUBLE_VITAMINS HP_ALOPECIA_UNIVERSALIS, HP_ALOPECIA_UNIVERSALIS, HP_ALOPECIA_UNIVERSALIS, HP_ALOPECIA_UNIVERSALIS BIOCARTA_IL2_PATHWAY, BIOCARTABOLISM_OBP_MAMMARY_GLAND_FIREDALIAL_CELL_DIFFERENTIATION GOBP_REGULATION_OF_LIPID_CATABOLISM_OF_FAT_SOLUBLE_VITAMINS HP_ALOPECIA_UNIVERSALIS, HP_ALOPECIA_UNIVERSALIS
	HP_ABNORMAL_T_CELL_ACTIVATION, HP_ABNORMAL_T_CELL_ACTIVATION PID_RXR_VDR_PATHWAY MORF_BMPR2, MORF_BMPR2 GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS REACTOME_EFFECTS_OF_PIP2_HYDROLYSIS, REACTOME_EFFECTS_OF_PIP2_HYDROLYSIS GOBP_POSITIVE_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION REACTOME_SYNTHESIS_OF_PIPS_AT_THE_LATE_ENDOSOME_MEMBRANE, REACTOME_SYNTHESIS_OF_PIPS_AT_THE_LATE_ENDOSOME_MEMBRANE GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_UP
	GOBP_POSITIVE_REGULATION_OF_HEAT_GENERATION, GOBP_POSITIVE_REGULATION_OF_HEAT_GENERATION MIKKELSEN_DEDIFFERENTIATED_STATE_DN, MIKKELSEN_DEDIFFERENTIATED_STATE_DN HP_CERVICAL_SPINE_INSTABILITY, HP_CERVICAL_SPINE_INSTABILITY HP_FOLLICULITIS GOBP_REGULATION_OF_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA, GOBP_REGULATION_OF_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA GOBP_RESPONSE_TO_STEROL, GOBP_RESPONSE_TO_STEROL HP_HYPERTHYROIDISM, HP_HYPERTHYROIDISM MIRISA_3P, MIRISA_3P GENTILE_UV_RESPONSE_CLUSTER_DI, GENTILE_UV_RESPONSE_CLUSTER_DI GENTILE_UV_RESPONSE_CLUSTER_DI, GENTILE_UV_RESPONSE_CLUSTER_DI
	REACTOME_METHYLATION, REACTOME_METHYLATION HP_URINARY_RETENTION GOBP_REGULATION_OF_TRIGLYCERIDE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_TRIGLYCERIDE_BIOSYNTHETIC_PROCESS GOBP_REGULATION_OF_GLUCONEOGENESIS, GOBP_REGULATION_OF_GLUCONEOGENESIS GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_ACTIVATION GOBP_XENDBIOTIC_TRANSPORT, GOBP_XENDBIOTIC_TRANSPORT GOBP_KENDBIOTIC_TRANSPORT, GOBP_XENDBIOTIC_TRANSPORT FRASOR_RESPONSE_TO_ESTRADIOL_UP, FRASOR_RESPONSE_TO_ESTRADIOL_UP FRASOR_RESPONSE_TO_ESTRADIOL_UP, FRASOR_RESPONSE_TO_ESTRADIOL_UP ### PROCESSOR PROCESSOR PROCESS PROCESS PROCESS ### PROCESSOR PROCESSOR PROCESS PROCESS PROCESS ### PROCESSOR PROCESSOR PROCESSOR PROCESSOR PROCESSOR PROCESSOR ### PROCESSOR PROCES
	GOMF_C_METHYLTRANSFERASE_ACTIVITY, GOMF_C_METHYLTRANSFERASE_ACTIVITY GOMF_RRNA_CYTOSINE_METHYLTRANSFERASE_ACTIVITY, GOMF_RRNA_CYTOSINE_METHYLTRANSFERASE_ACTIVITY NME2_TARGET_GENES, NME2_TARGET_GENES HP_ABNORMALITY_OF_VITAMIN_E_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_E_METABOLISM HP_FILL_TORTI, HP_FILL_TORTI GOBP_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION ONO_AMLL_TARGETS_UP, ONO_AMLL_TARGETS_UP WP_STEROID_BIOSYNTHESIS, WP_STEROID_BIOSYNTHESIS
	REACTOME_RHOBTB3_ATPASE_CYCLE, REACTOME_RHOBTB3_ATPASE_CYCLE GOBP_POSITIVE_REGULATION_OF_GLUCONEOGENESIS, GOBP_POSITIVE_REGULATION_OF_GLUCONEOGENESIS BIOCARTA_CIRCADIAN_PATHWAY, BIOCARTA_CIRCADIAN_PATHWAY MCCLUNG_DELTA_FOSB_TARGETS_2WK MCCLUNG_DELTA_FOSB_TARGETS_2WK GOBP_NEGATIVE_REGULATION_OF_HORMONE_SECRETION, GOBP_NEGATIVE_REGULATION_OF_HORMONE_SECRETION ERWIN_COHEN_PBMC_TC_83_AGE_18_45YO_NON_RESPONDERS_PREVIOUSLY_IMMUNIZED_24HR_DEG_CANONICAL_PATHWAY_MEMBERS_UP GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_PROLIFERATION GOBP_NEGATIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GOBP_NEGATIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
	HP_ABNORMALITY_OF_DENTIN, HP_ABNORMALITY_OF_DENTIN HP_SCLEROSIS_OF_SKUIL_BASE, HP_SCLEROSIS_OF_SKUIL_BASE GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_DIFFERENTIATION GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_DIFFERENTIATION ERWIN_COHEN_PBMC_TC_\$3_AGE_18_45YO_NAIVE_NOT_PREVIOUSLY_IMMUNIZED_24HR_DEG_CANONICAL_PATHWAY_MEMBERS_UP, ERWIN_COHEN_PBMC_TC_83_AGE_18_45YO_NAIVE_NOT_PREVIOUSLY_IMMUNIZED_24HR_DEG_CANONICAL_PATHWAY_MEMBERS_UP GOBP_NEGATIVE_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_CYTOLYSIS, GOBP_CYTOLYSIS LIU_IL13_MEMORY_MODEL_DN, LIU_IL13_MEMORY_MODEL_DN HP_ENLARGED_TONSILS, HP_ENLARGED_TONSILS
	GOMF_HEXOKINASE_ACTIVITY, GOMF_HEXOKINASE_ACTIVITY GOBP_NEGATIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_DISASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_DISASSEMBLY GOBP_PROTEIN_OLYUFMYLATION, GOBP_PROTEIN_POLYUFMYLATION GOMF_PROTEIN_GERANYLGERANYLTRANSFERASE_ACTIVITY, GOMF_PROTEIN_GERANYLTRANSFERASE_ACTIVITY, GOMF_PROTEIN_GERANYLTRANSFERASE_ACTIVITY, GOMF_PROTEIN_GERANYLTRANSFERASE_ACTIVITY, GOMF_PROTEIN_DERVANTURE_ARTERIOSCLEROSIS, HP_PREMATURE_ARTERIOSCLEROSIS, HP_PREMATURE_ARTERIOSCLEROSIS GOBP_DNA_DAMAGE_INDUCED_PROTEIN_PHOSPHORYLATION, GOBP_DNA_DAMAGE_INDUCED_PROTEIN_PHOSPHORYLATION KEGG_STEROID_HORMONE_BIOSYNTHESIS GOBP_COBALAMIN_METABOLIC_PROCESS, GOBP_COBALAMIN_METABOLIC_PROCESS HP_NEUTROPENIA_IN_PRESENCE_OF_ANTI_NEUTROPIL_ANTIBODIES, HP_NEUTROPENIA_IN_PRESENCE_OF_ANTI_NEUTROPIL_ANTIBODIES
	GOBP_REGULATION_OF_GLIAL_CELL_PROLIFERATION, GOBP_REGULATION_OF_GLIAL_CELL_PROLIFERATION GOBP_UREA_CYCLE, GOBP_UREA_CYCLE HP_AUTOIMMUNE_THROMBOCYTOPENIA, HP_AUTOIMMUNE_THROMBOCYTOPENIA GOMF_OLIGOPEPTIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_OLIGOPEPTIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_RESPONSE_TO_FRUCTOSE, GOBP_RESPONSE_TO_FRUCTOSE HP_BIFID_RIBS, HP_BIFID_RIBS HP_AORTIC_ATHEROSCLEROTIC_LESION, HP_AORTIC_ATHEROSCLEROTIC_LESION GOMF_AP_2_ADAPTOR_COMPLEX_BINDING, GOMF_AP_2_ADAPTOR_COMPLEX_BINDING
	ZHAN_EARLY_DIFFERENTIATION_GENES_UP, ZHAN_EARLY_DIFFERENTIATION_GENES_UP GOBP_CELLULAR_TRIGLYCERIDE_HOMEOSTASIS, GOBP_CELLULAR_TRIGLYCERIDE_HOMEOSTASIS HP_CHRONIC_NONINFECTIOUS_LYMPHADENOPATHY, HP_CHRONIC_NONINFECTIOUS_LYMPHADENOPATHY BIOCARTA_LDL_PATHWAY, BIOCARTA_LDL_PATHWAY GOBP_PROTEIN_UFMYLATION, GOBP_PROTEIN_UFMYLATION GOBP_REGULATION_OF_CENTROMERE_COMPLEX_ASSEMBLY, GOBP_REGULATION_OF_CENTROMERE_COMPLEX_ASSEMBLY WP_IMATINIB_AND_CHRONIC_MYELOID_LEUKEMIA, WP_IMATINIB_AND_CHRONIC_MYELOID_LEUKEMIA GOBP_VASCULAR_WOUND_HEALING, GOBP_VASCULAR_WOUND_HEALING
	GOBP_REGULATION_OF_MICROGLIAL_CELL_ACTIVATION, GOBP_REGULATION_OF_MICROGLIAL_CELL_ACTIVATION GOBP_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN, GOBP_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN ABBUD_LIF_SIGNALING_1_DN, ABBUD_LIF_SIGNALING_1_DN GOMF_POTASSIUM_ION_LEAK_CHANNEL_ACTIVITY, GOMF_POTASSIUM_ION_LEAK_CHANNEL_ACTIVITY GOBP_LONG_TERM_MEMORY, GOBP_LONG_TERM_MEMORY HP_ATOPIC_DERMATIIIS, HP_ATOPIC_DERMATIIIS, HP_ATOPIC_DERMATIINS, HP_ATOPIC_DERMATIINS, HP_ATOPIC_DERMATIINS, HP_ATOPIC_DERMATIINS, HP_ATOPIC_TERM_MEMORY, GOBP_LONG_TERM_SEGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION GOBP_NEGATIVE_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
	GOMF_ABC_TYPE_XENOBIOTIC_TRANSPORTER_ACTIVITY, GOMF_ABC_TYPE_XENOBIOTIC_TRANSPORTER_ACTIVITY PETRETTO_LEFT_VENTRICLE_MASS_QTL_CIS_DN, PETRETTO_LEFT_VENTRICLE_MASS_QTL_CIS_DN GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K36_SPECIFIC, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K36_SPECIFIC REACTOME_ANDROGEN_BIOSYNTHESIS, REACTOME_ANDROGEN_BIOSYNTHESIS chr4q33, chr24q33 MAP2K1_TARGET_GENES, MAP2K1_TARGET_GENES GOBP_POSITIVE_REGULATION_OF_HEPATOCYTE_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_HEPATOCYTE_PROLIFERATION GOMF_BETA_1_3_GALACTOSYLTRANSFERASE_ACTIVITY, GOMF_BETA_1_3_GALACTOSYLTRANSFERASE_ACTIVITY
	HP_PRECOCIOUS_ATHEROSCLEROSIS, HP_PRECOCIOUS_ATHEROSCLEROSIS HP_SEVERE_EXPRESSIVE_LANGUAGE_DELAY, HP_SEVERE_EXPRESSIVE_LANGUAGE_DELAY HP_CONCENTIAL_FINGER_FLEXION_CONTRACTURES, HP_CONGENITAL_FINGER_FLEXION_CONTRACTURES HP_IMPAIRED_ADP_INDUCED_PLATELET_AGGREGATION, HP_IMPAIRED_ADP_INDUCED_PLATELET_AGGREGATION HP_COTIC_NERVE_COMPRESSION, HP_OPTIC_NERVE_COMPRESSION HP_CORONARY_ARTERY_ANEURYSM, HP_CORONARY_ARTERY_ANEURYSM HP_GASTROINTESTINAL_EOSINOPHILIA, HP_CASTROINTESTINAL_EOSINOPHILIA GOBP_MHC_PROTEIN_COMPLEX_ASSEMBLY, GOBP_MHC_PROTEIN_COMPLEX_ASSEMBLY
	GOBP_TRIPEPTIDE_TRANSPORT, GOBP_TRIPEPTIDE_TRANSPORT GOMF_URIDYLYLTRANSFERASE_ACTIVITY, GOMF_URIDYLYLTRANSFERASE_ACTIVITY HP_LOSS_OF_CONSCIOUSNESS, HP_LOSS_OF_CONSCIOUSNESS chr6p24, chr6p24 GOBP_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION PHONG_TNF_TARGETS_DN, PHONG_TNF_TARGETS_DN GOBP_REGULATION_OF_LYSOSOMAL_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_LYSOSOMAL_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_LYSOSOMAL_PROTEIN_CATABOLIC_PROCESS, GOMF_TYPE_II_TRANSFORMING_GROWTH_FACTOR_BETA_RECEPTOR_BINDING
	GOBP_NEUROINFLAMMATORY_RESPONSE, GOBP_NEUROINFLAMMATORY_RESPONSE MIR7108_3P, MIR7108_3P HP_HYPERSPLENISM, HP_HYPERSPLENISM GOBP_UREA_METABOLIC_PROCESS, GOBP_UREA_METABOLIC_PROCESS GOBP_LIVER_MORPHOGENESIS, GOBP_LIVER_MORPHOGENESIS CONRAD_STEM_CELL, CONRAD_STEM_CELL GRANDVAUX_IRF3_TARGETS_DN, GRANDVAUX_IRF3_TARGETS_DN GOBP_POSITIVE_REGULATION_OF_MICROGLIAL_CELL_ACTIVATION, GOBP_POSITIVE_REGULATION_OF_MICROGLIAL_CELL_ACTIVATION HP_ABNORMAL_PAROTID_GLAND_MORPHOLOGY, HP_ABNORMAL_PAROTID_GLAND_MORPHOLOGY
	REACTOME_UREA_CYCLE, REACTOME_UREA_CYCLE WP_COMPOSITION_OF_LIPID_PARTICLES, WP_COMPOSITION_OF_LIPID_PARTICLES GOBP_NEGATIVE_REGULATION_OF_ASTROCYTE_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_ASTROCYTE_DIFFERENTIATION GOCC_JUXTAPARANODE_REGION_OF_AXON, GOCC_JUXTAPARANODE_REGION_OF_AXON GOOP_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION, GOBP_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION TAXCREB_02, TAXCREB_02 HP_HYPERKALEMIA, HP_HYPERKALEMIA MIR6741_3P, MIR6741_3P MIR6741_3P, MIR6741_3P
	HP_HYPERTONIC_DEHYDRATION, HP_HYPERTONIC_DEHYDRATION GOBP_OSSIFICATION_INVOLVED_IN_BONE_REMODELING, GOBP_OSSIFICATION_INVOLVED_IN_BONE_REMODELING GOBP_POSITIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS_IN_THE_VACUOLE, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MICROTUBULE_END, GOBP_PROTEIN_LOCALIZATION_TO_MICROTUBULE_END HP_HYPEROXALURIA, HP_HYPEROXALURIA HP_SHORT_ILIAC_BONES, HP_SHORT_ILIAC_BONES GOBP_NEUROBLAST_DIFFERENTIATION, GOBP_NEUROBLAST_DIFFERENTIATION GOBP_POSITIVE_REGULATION_OF_GLUCOKINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_GLUCOKINASE_ACTIVITY GOBP_POSITIVE_REGULATION_OF_GLUCOKINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_GLUCOKINASE_ACTIVITY
	REACTOME_CHYLOMICRON_CLEARANCE, REACTOME_CHYLOMICRON_CLEARANCE HP_PROTEIN_AVOIDANCE, HP_PROTEIN_AVOIDANCE HP_BIRTH_LENGTH_GREATER_THAN_97TH_PERCENTILE, HP_BIRTH_LENGTH_GREATER_THAN_97TH_PERCENTILE HP_RENAL_CORTICAL_MICROCYSTS, HP_RENCICRITICAL_MICROCYSTS HP_CERVICAL_LYMPHADENOPATHY, HP_CERVICAL_LYMPHADENOPATHY, HP_CERVICAL_LYMPHADENOPATHY, HP_CERVICAL_LYMPHADENOPATHY, HP_CERVICAL_SOTYPE_SWITCHING_TO_IGA_ISOTYPE_SWITCHING_TO_IGA_ISOTYPE_SWITCHING_TO_IGA_ISOTYPE_SWITCHING_TO_IGA_ISOTYPE_SWITCHING_TO_IGA_ISOTYPE_SWITCHING_TO_IGA_ISOTYPE_SWITCHING_TO_IGA_ISOTYPES
	GOBP_NEGATIVE_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, COBP_NEGATIVE_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION GOMF_RNA_POLYMERASE_L_CORE_BINDING, GOMF_RNA_POLYMERASE_L_CORE_BINDING GOMF_CRCP3_CHEMOKINE_RECEPTOR_BINDING, GOMF_CRCP3_CHEMOKINE_RECEPTOR_BINDING GOMF_TRIPEPTIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_TRIPEPTIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY HP_PERIPAPILLARY_ATROPHY, HP_PERIPAPILLARY_ATROPHY HP_ABNORMALITY_OF_OLFACTORY_LOBE_MORPHOLOGY, HP_ABNORMALITY_OF_OLFACTORY_LOBE_MORPHOLOGY, HP_ABNORMALITY_OF_OLFACTORY_LOBE_MORPHOLOGY, HP_ABNORMALITY_OF_OLFACTORY_LOBE_MORPHOLOGY, HP_ABNORMALITY_OF_OLFACTORY_LOBE_MORPHOLOGY MIR3177_3P WP_CONTROL_OF_IMMUNE_TOLERANCE_BY_VASOACTIVE_INTESTINAL_PEPTIDE, WP_CONTROL_OF_IMMUNE_TOLERANCE_BY_VASOACTIVE_INTESTINAL_PEPTIDE GOBP_CELLULAR_RESPONSE_TO_LITHIUM_ION, GOBP_CELLULAR_RESPONSE_TO_LITHIUM_ION
	WP_BILE_ACIDS_SYNTHESIS_AND_ENTEROHEPATIC_CIRCULATION, WP_BILE_ACIDS_SYNTHESIS_AND_ENTEROHEPATIC_CIRCULATION FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE_AGE_18_40YO_1DY_NEGATIVE SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOXI_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOXI_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOXI_DN SOBOLEV_PBMC_PANDEMRIX_AGE_18_64YO_HIGH_VS_LOW_RESPONDERS_MEDIUM_HIGH_ADVERSE_EVENTS_SCORE_1DY_TRANSIENT_UP GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_PERMEABILITY, GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_PERMEABILITY HP_DECREASED_SPECIFIC_ANTI_POLYSACCHARIDE_ANTIBODY_LEVEL, HP_DECREASED_SPECIFIC_ANTI_POLYSACCHARIDE_ANTIBODY_LEVEL HP_UPPER_EYELID_COLOBOMA, HP_UPPER_EYELID_COLOBOMA RICHERT_PBMC_HIV_LIPO_5_AGE_37_48YO_STIMULATED_14W_SIGNIFICANT_VARIATION_UP, RICHERT_PBMC_HIV_LIPO_5_AGE_37_48YO_STIMULATED_VS_UNSTIMULATE
	GOMF_UDP_GALACTOSE_BETA_N_ACETYLGLUCOSAMINE_BETA_1_3_GALACTOSYLTRANSFERASE_ACTIVITY, GOMF_UDP_GALACTOSE_BETA_N_ACETYLGLUCOSAMINE_BETA_1_3_GALACTOSYLTRANSFERASE_ACTIVITY HP_RESPIRATORY_ALKALOSIS HP_PERIANAL_ABSCESS, HP_PERIANAL_ABSCESS GOBP_POSITIVE_REGULATION_OF_HEXOKINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_HEXOKINASE_ACTIVITY GOBP_POSITIVE_REGULATION_OF_MICROTUBULE_BINDING, GOBP_POSITIVE_REGULATION_OF_MICROTUBULE_BINDING, GOBP_POSITIVE_REGULATION_OF_MICROTUBULE_BINDING GOMF_SALPHA_ANDROSTANE_3BETA_17BETA_DIOL_DEHYDROGENASE_ACTIVITY, GOMF_SALPHA_ANDROSTANE_3BETA_17BETA_DIOL_DEHYDROGENASE_ACTIVITY, GOMF_SALPHA_ANDROSTANE_3BETA_17BETA_DIOL_DEHYDROGENASE_ACTIVITY MIR6775_5P, MIR6775_5P MIR4537, MIR4537
	HP_ATLANTOAXIAL_DISLOCATION, HP_ATLANTOAXIAL_DISLOCATION HP_IMPAIRED_COLLAGEN_INDUCED_PLATELET_AGGREGATION, HP_IMPAIRED_COLLAGEN_INDUCED_PLATELET_AGGREGATION GOBP_RESPONSE_TO_BLUE_LIGHT, GOBP_RESPONSE_TO_BLUE_LIGHT GOBP_TRIPEPTIDE_TRANSMEMBRANE_TRANSPORT, GOBP_TRIPEPTIDE_TRANSMEMBRANE_TRANSPORT GOMF_ATPASE_COUPLED_LIPID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_ATPASE_COUPLED_LIPID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_ATPASE_COUPLED_LIPID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY HP_EMPHYSEMA, HP_EMPHYSEMA SOBOLEV_T_CELL_PANDEMRIX_AGE_18_64YO_1DY_UP, SOBOLEV_T_CELL_PANDEMRIX_AGE_18_64YO_1DY_UP, SOBOLEV_T_CELL_PANDEMRIX_AGE_18_64YO_1DY_UP GOBP_ESTROCEN_BIOSYNTHETIC_PROCESS, COBP_ESTROCEN_BIOSYNTHETIC_PROCESS
	RYTAAWNNNTGAY_UNKNOWN, RYTAAWNNNTGAY_UNKNOWN GOBP_POSITIVE_REGULATION_OF_MHC_CLASS_IL_BIOSYNTHETIC_PROCESS, COBP_POSITIVE_REGULATION_OF_MHC_CLASS_IL_BIOSYNTHETIC_PROCESS WP_UREA_CYCLE_AND_RELATED_DISEASES, WP_UREA_CYCLE_AND_RELATED_DISEASES GOBP_POSITIVE_REGULATION_OF_B_CELL_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_B_CELL_RECEPTOR_SIGNALING_PATHWAY HP_SHORT_NAIL_HP_SHORT_NAIL HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_3DY_UP, HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_3DY_UP, HOEK_MYELOID_DENDRITIC_CELL_2011_COF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY, GOBP_NEGATIVE_REGULATION_OF_PHOSPHOLIC_PROCESS, GOBP_REGULATION_OF_PHOSPHOLIC_PROCESS, GOBP_REGULATION_OF_PHOSPHOLIC_PROCESS, GOBP_REGULATION_OF_BONE_DEVELOPMENT.
	GOBP_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS_IN_THE_VACUOLE, GOBP_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS_IN_THE_VACUOLE chr2q36, chr2q36 GOCC_PLATELET_DENSE_GRANULE_LUMEN, GOCC_PLATELET_DENSE_GRANULE_LUMEN ELLWOOD_MYC_TARGETS_UP, ELLWOOD_MYC_TARGETS_UP HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_4NM_UP, HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_4NM_UP GOBP_CELLULAR_RESPONSE_TO_PRUCTOSE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_FRUCTOSE_STIMULUS GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_MITOCHONDRION, GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_MITOCHONDRION, GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_OF_CD4_NEGATIVE_CD8_NEGATIVE_ALPHA_BETA_REGULATORY_T_CELLS, HP_ELEVATED_PROPORTION_OF_CD4_NEGATIVE_ALPHA_BETA_REGULATORY_T_CELLS
	HP_LARGE_ILIAC_WINGS, HP_LARGE_ILIAC_WINGS HP_MULTIPLE_JOINT_DISLOCATION, HP_MULTIPLE_JOINT_DISLOCATION HP_CHORIORETINAL_HYPOPIGMENTATION, HP_CHORIORETINAL_HYPOPIGMENTATION GOBP_GUSTINION_ETRANSMEMBRANE_TRANSPORT, GOBP_GUSTINION_ETRANSMEMBRANE_TRANSPORT GOBP_POSITIVE_REGULATION_OF_VASCULAR_WOUND_HEALING, GOBP_POSITIVE_REGULATION_OF_VASCULAR_WOUND_HEALING GOMP_ATPASE_INHIBITOR_ACTIVITY, GOMF_ATPASE_INHIBITOR_ACTIVITY GOBP_NEGATIVE_REGULATION_OF_TROPHOBLAST_CELL_MIGRATION, GOBP_NEGATIVE_REGULATION_OF_TROPHOBLAST_CELL_MIGRATION HP_PSEUDOHYPOALDOSTERONISM, HP_PSEUDOHYPOALDOSTERONISM
	MODULE_514, MODULE_514 chr5p13, chr5p13 GOMF_NARROW_PORE_CHANNEL_ACTIVITY, GOMF_NARROW_PORE_CHANNEL_ACTIVITY GOMF_NARROW_PORE_CHANNEL_ACTIVITY, GOMF_NARROW_PORE_CHANNEL_ACTIVITY GOBP_MICROGLIA_DIFFERENTIATION, GOBP_MICROGLIA_DIFFERENTIATION GOMF_LONG_SILTATHIONE_S_CONJUGATE_TRANSPORTER_ACTIVITY, GOMF_ABC_TYPE_GLUTATHIONE_S_CONJUGATE_TRANSPORTER_ACTIVITY GOMF_LONG_CHAIN_FATROCYTE_ACTIVATION, GOBP_REGULATION_OF_ASTROCYTE_ACTIVATION GOMF_LONG_CHAIN_FATTY_ACYL_COA_BINDING, COMF_LONG_CHAIN_FATTY_ACYL_COA_BINDING WP_TFS_REGULATE_MIRNAS_RELATED_TO_CARDIAC_HYPERTROPHY, WP_TFS_REGULATE_MIRNAS_RELATED_TO_CARDIAC_HYPERTROPHY
	HP_ABNORMALITY_OF_FUNDUS_PIGMENTATION, HP_ABNORMALITY_OF_FUNDUS_PIGMENTATION HP_EXTRAMEDULLARY_HEMATOPOIESIS, HP_EXTRAMEDULLARY_HEMATOPOIESIS HP_HYPOPIGMENTATION_OF_THE_FUNDUS, HP_HYPOPIGMENTATION_OF_THE_FUNDUS