

	GSE7852_THYMUS_VS_FAT_TREG_UP, GSE7852_THYMUS_VS_FAT_TREG_UP GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP, GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP, GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP, GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN, GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN GSE17721_POLYIC_VS_GARDIQUIMOD_4H_BMDC, GSE17721_POLYIC_VS_GARDIQUIMOD_6H_BMDC_DN GSE13484_3H_UNSTIM_VS_YF17D_VACCINE_STIM_PBMCL_UP, GSE13484_3H_UNSTIM_VS_YF17D_VACCINE_STIM_PBMCL_UP GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_DN, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_DN GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN, GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN GSE17721_LPS_VS_GARDIQUIMOD_1H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_1H_BMDC_DN GSE17721_CTRL_VS_LPS_1H_BMDC_UP, GSE17721_CTRL_VS_LPS_1H_BMDC_UP GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP GSE28737_WT_VS_BC1A6_KO_MARGINAL_ZONE_BCELL_DN, GSE28737_WT_VS_BC1A6_KO_MARGINAL_ZONE_BCELL_DN GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP GSE5589_LPS_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN, GSE5589_LPS_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP, GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN GCM_HBP1, GCM_HBP1 GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP GSE15659_RESTING_VS_ACTIVATED_TREG_DN, GSE15659_RESTING_VS_ACTIVATED_TREG_DN GOMF_IRRNA_BINDING, GOMF_IRRNA_BINDING GSE30883_SPL_VS_SPS_THYMOCYTE_UP, GSE30883_SPL_VS_SPS_THYMOCYTE_UP GSE17721_0.5H_VS_12H_PAMCSK4_BMDC_UP, GSE17721_0.5H_VS_12H_PAMCSK4_BMDC_UP GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN MIR6715A_3P, MIR6715A_3P MIR603, MIR603 GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP GSE11924_TFH_VS_TH2_CD4_TCELL_DN, GSE11924_TFH_VS_TH2_CD4_TCELL_DN GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP, GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_UP, GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_UP JAZAG_TGFB1_SIGNALING_UP, JAZAG_TGFB1_SIGNALING_UP GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_DN, GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_DN MIR150_3P, MIR150_3P GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_DN, GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_DN FIGUEROA_AML_METHYLATION_CLUSTER_3_UP, FIGUEROA_AML_METHYLATION_CLUSTER_3_UP MIR3927_3P, MIR3927_3P MODULE_277, MODULE_277 MIR4647, MIR4647B, MIR4647, MIR4647B MIR155_3P, MIR155_3P GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THYMUS_UP, GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THYMUS_UP GSE16385_UNTREATED_VS_12H_IL4_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_IL4_TREATED_MACROPHAGE_DN FIGUEROA_AML_METHYLATION_CLUSTER_6_UP, FIGUEROA_AML_METHYLATION_CLUSTER_6_UP FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN, FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN GOBP_MAINTENANCE_OF_CELL_NUMBER, GOBP_MAINTENANCE_OF_CELL_NUMBER GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMCL_DN, GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMCL_DN HP_CUTIS_MARMORATA, HP_CUTIS_MARMORATA GOCC_90S_PRRIBOSOME, GOCC_90S_PRRIBOSOME HP_ABNORMALITY_OF_THE_COAGULATION_CASCADE, HP_ABNORMALITY_OF_THE_COAGULATION_CASCADE MIR3668, MIR3668 MIR376A_3P, MIR376B_3P, MIR376A_3P, MIR376B_3P GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE, GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN FIGUEROA_AML_METHYLATION_CLUSTER_7_UP, FIGUEROA_AML_METHYLATION_CLUSTER_7_UP GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY MIR34A_3P, MIR34A_3P HP_APLASIA_HYPOPLASIA_OF_TOE, HP_APLASIA_HYPOPLASIA_OF_TOE MIR218_1_3P, MIR218_1_3P GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN GCM_BC12L1, GCM_BC12L1 MIR4673, MIR4673 GOCC_PCG_PROTEIN_COMPLEX, GOCC_PCG_PROTEIN_COMPLEX GOBP_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION, GOBP_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN MIR679_5P, MIR679_5P MIR676_3P, MIR676_3P GOBP_REGULATION_OF_PROTEIN_ACETYLATION, GOBP_REGULATION_OF_PROTEIN_ACETYLATION DELASERNA_MYOD_TARGETS_DN, DELASERNA_MYOD_TARGETS_DN AHRARNT_01, AHRARNT_01 BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP, BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP GENTILE_UV_RESPONSE_CLUSTER_D8, GENTILE_UV_RESPONSE_CLUSTER_D8 HP_TRIGONOCEPHALY, HP_TRIGONOCEPHALY MIR684_3P, MIR684_3P ACTCAGC, MIR345, ACTCAGC, MIR345 FIGUEROA_AML_METHYLATION_CLUSTER_4_UP, FIGUEROA_AML_METHYLATION_CLUSTER_4_UP GSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP, GSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP MIR1263, MIR1263 MIR619_3P, MIR619_3P MIR9718, MIR9718 GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE, GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE GROSS_HYPOXIA_VIA_ELFK3_ONLY_DN, GROSS_HYPOXIA_VIA_ELFK3_ONLY_DN WP_MITOCHONDRIAL_GENE_EXPRESSION, WP_MITOCHONDRIAL_GENE_EXPRESSION GTGTGAG, MIR342, GTGTGAG, MIR342 HP_ABNORMAL_CIRCULATING_JGG_LEVEL, HP_ABNORMAL_CIRCULATING_JGG_LEVEL MIR3614_5P, MIR3614_5P MIR628_5P, MIR628_5P MEIS1BHOXA9_01, MEIS1BHOXA9_01 REACTOME_MICRORNA_MIRNA_BIOGENESIS, REACTOME_MICRORNA_MIRNA_BIOGENESIS GOBP_B_CELL_DIFFERENTIATION, GOBP_B_CELL_DIFFERENTIATION MIR5699_3P, MIR5699_3P GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR GOBP_T_CELL_DIFFERENTIATION_IN_THYMUS, GOBP_T_CELL_DIFFERENTIATION_IN_THYMUS MIR4421, MIR4421 IRX3_TARGET_GENES, IRX3_TARGET_GENES MIR647, MIR647 GOBP_CELLULAR_PROTEIN_CONTAINING_COMPLEX_LOCALIZATION, GOBP_CELLULAR_PROTEIN_CONTAINING_COMPLEX_LOCALIZATION GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT_MEMBRANE, GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT_MEMBRANE MIR4423_5P, MIR4423_5P GOBP_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GOBP_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR HP_RECURRENT_VIRAL_INFECTIONS, HP_RECURRENT_VIRAL_INFECTIONS GSE41176_WT_VS_TAK1_KO_ANTLJGM_STIM_BCELL_1H_DN, GSE41176_WT_VS_TAK1_KO_ANTLJGM_STIM_BCELL_1H_DN GOBP_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE, GOBP_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE HP_IMPAIRED_ANTIGEN_SPECIFIC_RESPONSE, HP_IMPAIRED_ANTIGEN_SPECIFIC_RESPONSE GOMF_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS, GOMF_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS MIR642A_5P, MIR642A_5P HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS, HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS HP_DECREASED_CIRCULATING_JGG_LEVEL, HP_DECREASED_CIRCULATING_JGG_LEVEL chr16q22, chr16q22 TATCTCG, MIR488, TATCTCG, MIR488 GSE9946_MATURE_STIMULATORY_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP, GSE9946_MATURE_STIMULATORY_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP HP_PANCREATIC_ADENOCARCINOMA, HP_PANCREATIC_ADENOCARCINOMA GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR HP_ESOPHAGEAL_VARIX, HP_ESOPHAGEAL_VARIX HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_MITOGEN, HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_MITOGEN MIR6793_5P, MIR6793_5P HP_ABNORMAL_CELL_PROLIFERATION, HP_ABNORMAL_CELL_PROLIFERATION PLASARI_TGFB1_SIGNALING_VIA_NFIC_IHR_UP, PLASARI_TGFB1_SIGNALING_VIA_NFIC_IHR_UP CCCACAT, MIR2993P, CCCACAT, MIR2993P HP_ABNORMAL_CIRCULATING_JGA_LEVEL, HP_ABNORMAL_CIRCULATING_JGA_LEVEL MIR3161, MIR3161 MIR6847_3P, MIR6847_3P GOBP_ENDOCARDIAL_CUSHION_MORPHOGENESIS, GOBP_ENDOCARDIAL_CUSHION_MORPHOGENESIS REACTOME_TRANSLATION_OF_SARS_COV_1_STRUCTURAL_PROTEINS, REACTOME_TRANSLATION_OF_SARS_COV_1_STRUCTURAL_PROTEINS HAY_BONE_MARROW_CD14_POS_HSC, HAY_BONE_MARROW_CD14_POS_HSC WP_VITAMIN_D_IN_INFLAMMATORY_DISEASES, WP_VITAMIN_D_IN_INFLAMMATORY_DISEASES FIGUEROA_AML_METHYLATION_CLUSTER_2_UP, FIGUEROA_AML_METHYLATION_CLUSTER_2_UP MIR4662A_3P, MIR4662A_3P REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOCHROME_C_RELEASE, REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOCHROME_C_RELEASE MIR431_5P, MIR431_5P GOBP_RESPONSE_TO_PLATELET_DERIVED_GROWTH_FACTOR, GOBP_RESPONSE_TO_PLATELET_DERIVED_GROWTH_FACTOR HP_MYOPATHIC_FACIES, HP_MYOPATHIC_FACIES MIR200C_5P, MIR200C_5P MIR6826_5P, MIR6826_5P GOMF_FOUR_WAY_JUNCTION_DNA_BINDING, GOMF_FOUR_WAY_JUNCTION_DNA_BINDING GOBP_DOLICHOL_METABOLIC_PROCESS, GOBP_DOLICHOL_METABOLIC_PROCESS BILANGES_RAPAMYCIN_SENSITIVE_GENES, BILANGES_RAPAMYCIN_SENSITIVE_GENES HP_ABNORMAL_B_CELL_MORPHOLOGY, HP_ABNORMAL_B_CELL_MORPHOLOGY GOCC_MITOTIC_SPINDLE_MICROTUBULE, GOCC_MITOTIC_SPINDLE_MICROTUBULE MIR3529_5P, MIR3529_5P HP_T_LYMPHOCYTOPENIA, HP_T_LYMPHOCYTOPENIA HP_DIFFICULTY_RUNNING, HP_DIFFICULTY_RUNNING GOMF_TRANSCRIPTION_COACTIVATOR_BINDING, GOMF_TRANSCRIPTION_COACTIVATOR_BINDING PID_HES_HEY_PATHWAY, PID_HES_HEY_PATHWAY HP_NASOLACRIMAL_DUCT_OBSTRUCTION, HP_NASOLACRIMAL_DUCT_OBSTRUCTION MIR103B, MIR103B GOBP_AMYLOID_FIBRIL_FORMATION, GOBP_AMYLOID_FIBRIL_FORMATION AMIT_EGF_RESPONSE_480_MCF10A, AMIT_EGF_RESPONSE_480_MCF10A MIR631, MIR631 HP_RECTAL_PROLAPSE, HP_RECTAL_PROLAPSE MIR3661, MIR3661 GOBP_DOLICHYL_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_DOLICHYL_DIPHOSPHATE_BIOSYNTHETIC_PROCESS HP_ABNORMAL_CIRCULATING_JGM_LEVEL, HP_ABNORMAL_CIRCULATING_JGM_LEVEL SEIDEN_MET_SIGNALING, SEIDEN_MET_SIGNALING GOCC_AGGRESOME, GOCC_AGGRESOME WP_NOTCH_SIGNALING, WP_NOTCH_SIGNALING GNF2_ATM, GNF2_ATM GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS, GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS SNIPERS_AMPLIFIED_IN_HEAD_AND_NECK_TUMORS, SNIPERS_AMPLIFIED_IN_HEAD_AND_NECK_TUMORS FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE_AGE_18_40YO_14DY_POSITIVE, FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE_AGE_18_40YO_14DY_POSITIVE GOBP_PRE_MIRNA_PROCESSING, GOBP_PRE_MIRNA_PROCESSING GOBP_POLYPRENOL_METABOLIC_PROCESS, GOBP_POLYPRENOL_METABOLIC_PROCESS FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE_AGE_18_40YO_1DY_NEGATIVE, FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE_AGE_18_40YO_1DY_NEGATIVE JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN, JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION, REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION JIANG_HYPOXIA_VIA_VHL, JIANG_HYPOXIA_VIA_VHL GOBP_ENDOCARDIAL_CUSHION_FORMATION, GOBP_ENDOCARDIAL_CUSHION_FORMATION ELLWOOD_MYC_TARGETS_DN, ELLWOOD_MYC_TARGETS_DN GOMF_HISTONE_ACETYLTRANSFERASE_BINDING, GOMF_HISTONE_ACETYLTRANSFERASE_BINDING GOBP_V_D_J_RECOMBINATION, GOBP_V_D_J_RECOMBINATION GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE HP_DECREASED_CIRCULATING_JGA_LEVEL, HP_DECREASED_CIRCULATING_JGA_LEVEL GOBP_HAIR_CELL_DIFFERENTIATION, GOBP_HAIR_CELL_DIFFERENTIATION GOBP_RNA_INTERFERENCE, GOBP_RNA_INTERFERENCE BRACHAT_RESPONSE_TO_CISPLATIN, BRACHAT_RESPONSE_TO_CISPLATIN HP_PRURITUS, HP_PRURITUS MIR943, MIR943 PID_BMP_PATHWAY, PID_BMP_PATHWAY HP_ABNORMAL_NATURAL_KILLER_CELL_MORPHOLOGY, HP_ABNORMAL_NATURAL_KILLER_CELL_MORPHOLOGY HP_THYROIDITIS, HP_THYROIDITIS GOBP_RESPONSE_TO_NICOTINE, GOBP_RESPONSE_TO_NICOTINE GOBP_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION, GOBP_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION GOBP_INNER_EAR_AUDITORY_RECEPTOR_CELL_DIFFERENTIATION, GOBP_INNER_EAR_AUDITORY_RECEPTOR_CELL_DIFFERENTIATION RUSSLINGER_GASTRIC_REG3A_POSITIVE_CELLS, RUSSLINGER_GASTRIC_REG3A_POSITIVE_CELLS HP_B_LYMPHOCYTOPENIA, HP_B_LYMPHOCYTOPENIA HP_SHOULDER_GIRDLE_MUSCLE_WEAKNESS, HP_SHOULDER_GIRDLE_MUSCLE_WEAKNESS HP_RESPONSE_TO_DRUGS_ACTING_ON_NEUROMUSCULAR_TRANSMISSION, HP_RESPONSE_TO_DRUGS_ACTING_ON_NEUROMUSCULAR_TRANSMISSION MIR4683, MIR4683 GOBP_POSITIVE_REGULATION_OF_MYOBLAST_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MYOBLAST_DIFFERENTIATION PAL_PRMT5_TARGETS_DN, PAL_PRMT5_TARGETS_DN MARCNIAK_ER_STRESS_RESPONSE_VIA_CHOP, MARCNIAK_ER_STRESS_RESPONSE_VIA_CHOP HP_INTERMITTENT_DIARRHEA, HP_INTERMITTENT_DIARRHEA MIR4737, MIR4737 REACTOME_RECEPTOR_MEDIATED_MITOPHAGY, REACTOME_RECEPTOR_MEDIATED_MITOPHAGY REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY, REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY HP_ABNORMAL_LACRIMAL_DUCT_MORPHOLOGY, HP_ABNORMAL_LACRIMAL_DUCT_MORPHOLOGY GOBP_CARDIAC_MUSCLE_CELL_DIFFERENTIATION, GOBP_CARDIAC_MUSCLE_CELL_DIFFERENTIATION GOBP_IMMATURE_B_CELL_DIFFERENTIATION, GOBP_IMMATURE_B_CELL_DIFFERENTIATION GOBP_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE, GOBP_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE MIR6802_3P, MIR6802_3P DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_PERICYTES, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_PERICYTES
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