

	GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP, GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP GSE22886_TH1_VS_TH2_48H_ACT_DN, GSE22886_TH1_VS_TH2_48H_ACT_DN GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP GSE38696_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP, GSE38696_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_UP GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP, STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP, GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP GSE3982_BASOPHIL_VS_TH1_UP, GSE3982_BASOPHIL_VS_TH1_UP GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN, GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN MIR194_5P, MIR194_5P GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN GOBP_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GOBP_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP, GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP MIR6768_3P, MIR6768_3P GSE7852_TREG_VS_TCONV_THYMUS_DN, GSE7852_TREG_VS_TCONV_THYMUS_DN GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP MIR558, MIR558 MIR889_5P, MIR889_5P GOMF_HORMONE_RECEPTOR_BINDING, GOMF_HORMONE_RECEPTOR_BINDING MIR8065, MIR8065 REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENTIATION_AND_PLATELET_FUNCTION, REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENTIATION_AND_PLATELET_FUNCTION MIR12132, MIR12132 MIR642B_5P, MIR642B_5P GOBP_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION MIR891A_3P, MIR891A_3P MIR1208, MIR1208 GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_10DY_DN, GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_10DY_DN GOBP_CHROMATIN_SILENCING, GOBP_CHROMATIN_SILENCING MIR4456, MIR4456 GOBP_INTEGRIN_MEDIATED_SIGNALING_PATHWAY, GOBP_INTEGRIN_MEDIATED_SIGNALING_PATHWAY GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN, GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN HP_STAGE_5_CHRONIC_KIDNEY_DISEASE, HP_STAGE_5_CHRONIC_KIDNEY_DISEASE WP_DNA_DAMAGE_RESPONSE_ONLY_ATM_DEPENDENT, WP_DNA_DAMAGE_RESPONSE_ONLY_ATM_DEPENDENT MIR3682_3P, MIR3682_3P MIR1178_3P, MIR1178_3P MIR331_5P, MIR331_5P GOBP_NEPHRON_DEVELOPMENT, GOBP_NEPHRON_DEVELOPMENT GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN MIR211_3P, MIR211_3P MIR4293, MIR4293 GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP, GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP MIR6866_5P, MIR6866_5P REACTOME_ELASTIC_FIBRE_FORMATION, REACTOME_ELASTIC_FIBRE_FORMATION HP_RECURRENT_OTITIS_MEDIA, HP_RECURRENT_OTITIS_MEDIA MIR548Q, MIR548Q LAKE_ADULT_KIDNEY_C25_ENDOTHELIAL_CELLS_UNASSIGNED, LAKE_ADULT_KIDNEY_C25_ENDOTHELIAL_CELLS_UNASSIGNED MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TACI_UP, MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TACI_UP REACTOME_CHONDROITIN_SULFATE_DERMATAN_SULFATE_METABOLISM, REACTOME_CHONDROITIN_SULFATE_DERMATAN_SULFATE_METABOLISM MIR4797_3P, MIR4797_3P BURTON_ADIPOGENESIS_7, BURTON_ADIPOGENESIS_7 REACTOME_MOLECULES_ASSOCIATED_WITH_ELASTIC_FIBRES, REACTOME_MOLECULES_ASSOCIATED_WITH_ELASTIC_FIBRES WP_G_PROTEIN_SIGNALING_PATHWAYS, WP_G_PROTEIN_SIGNALING_PATHWAYS KEGG_NOD_LIKE_RECEPTOR_SIGNALING_PATHWAY, KEGG_NOD_LIKE_RECEPTOR_SIGNALING_PATHWAY GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN MIR4638_3P, MIR4638_3P GOBP_REGULATION_OF_RESPONSE_TO_WOUNDING, GOBP_REGULATION_OF_RESPONSE_TO_WOUNDING MIR198, MIR198 HALLMARK_COAGULATION, HALLMARK_COAGULATION MIR584_5P, MIR584_5P MIR3120_5P, MIR3120_5P BROWNE_HCMV_INFECTION_16HR_DN, BROWNE_HCMV_INFECTION_16HR_DN MIR597_5P, MIR597_5P MIR6851_3P, MIR6851_3P GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_DN, GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_DN WP_FBXL10_ENHANCEMENT_OF_MAPERK_SIGNALING_IN_DIFFUSE_LARGE_BCELL_LYMPHOMA, WP_FBXL10_ENHANCEMENT_OF_MAPERK_SIGNALING_IN_DIFFUSE_LARGE_BCELL_LYMPHOMA GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_UP, GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_UP REACTOME_CA2_PATHWAY, REACTOME_CA2_PATHWAY GOBP_WNT_SIGNALING_PATHWAY_CALCIIUM_MODULATING_PATHWAY, GOBP_WNT_SIGNALING_PATHWAY_CALCIIUM_MODULATING_PATHWAY GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY, GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY GOBP_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT GOBP_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS, GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS GOMF_COLLAGEN_BINDING, GOMF_COLLAGEN_BINDING PID_SIP_SIP3_PATHWAY, PID_SIP_SIP3_PATHWAY GOMF_EXTRACELLULAR_MATRIX_BINDING, GOMF_EXTRACELLULAR_MATRIX_BINDING HP_HYPOPLASTIC_VERTEBRAL_BODIES, HP_HYPOPLASTIC_VERTEBRAL_BODIES GOBP_REGULATION_OF_PLASMINOGEN_ACTIVATION, GOBP_REGULATION_OF_PLASMINOGEN_ACTIVATION PID_SIP_SIP1_PATHWAY, PID_SIP_SIP1_PATHWAY HARALAMBIEVA_PBMC_M_M_R_IL_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP, HARALAMBIEVA_PBMC_M_M_R_IL_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP GOBP_REGULATION_OF_WOUND_HEALING, GOBP_REGULATION_OF_WOUND_HEALING BCAT_GDS748_DN, BCAT_GDS748_DN GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT SIG_CD40PATHWAYMAP, SIG_CD40PATHWAYMAP HP_DISSEMINATED_INTRAVASCULAR_COAGULATION, HP_DISSEMINATED_INTRAVASCULAR_COAGULATION HP_ABNORMAL_PLATELET_FUNCTION, HP_ABNORMAL_PLATELET_FUNCTION KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE REACTOME_FORMATION_OF_APOPTOSOME, REACTOME_FORMATION_OF_APOPTOSOME GOBP_ZYMOGEN_ACTIVATION, GOBP_ZYMOGEN_ACTIVATION GOBP_POLYSACCHARIDE_CATABOLIC_PROCESS, GOBP_POLYSACCHARIDE_CATABOLIC_PROCESS MIR3160_3P, MIR3160_3P FERRARI_RESPONSE_TO_FENRETINIDE_UP, FERRARI_RESPONSE_TO_FENRETINIDE_UP GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS, GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS GOBP_LABYRINTHINE_LAYER_DEVELOPMENT, GOBP_LABYRINTHINE_LAYER_DEVELOPMENT REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS, REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS WP_MFAP5_MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS, WP_MFAP5_MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS, REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS REACTOME_SIGNAL_TRANSDUCTION_BY_L1, REACTOME_SIGNAL_TRANSDUCTION_BY_L1 HASLINGER_B_CLL_WITH_MUTATED_VH_GENES, HASLINGER_B_CLL_WITH_MUTATED_VH_GENES GOBP_PLASMINOGEN_ACTIVATION, GOBP_PLASMINOGEN_ACTIVATION GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION REACTOME_SYNDECAN_INTERACTIONS, REACTOME_SYNDECAN_INTERACTIONS GOMF_MANNOSIDASE_ACTIVITY, GOMF_MANNOSIDASE_ACTIVITY WP_PREGNANE_X_RECEPTOR_PATHWAY, WP_PREGNANE_X_RECEPTOR_PATHWAY GOBP_REGULATION_OF_MACROPHAGE_MIGRATION, GOBP_REGULATION_OF_MACROPHAGE_MIGRATION MIR3677_5P, MIR3677_5P GOBP_REGULATION_OF_MACROPHAGE_CHEMOTAXIS, GOBP_REGULATION_OF_MACROPHAGE_CHEMOTAXIS HP_REFLEX_SEIZURE, HP_REFLEX_SEIZURE GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION GOBP_STRIATED_MUSCLE_CELL_PROLIFERATION, GOBP_STRIATED_MUSCLE_CELL_PROLIFERATION GOBP_KETONE_CATABOLIC_PROCESS, GOBP_KETONE_CATABOLIC_PROCESS REACTOME_VASOPRESSIN_REGULATES_RENAL_WATER_HOMEOSTASIS_VIA_AQUAPORINS, REACTOME_VASOPRESSIN_REGULATES_RENAL_WATER_HOMEOSTASIS_VIA_AQUAPORINS GOBP_ENDOTHELIAL_CELL_PROLIFERATION, GOBP_ENDOTHELIAL_CELL_PROLIFERATION GOBP_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBP_REGULATION_OF_ORGANIC_ACID_TRANSPORT HP_ABNORMAL_DELIVERY, HP_ABNORMAL_DELIVERY PID_INTEGRIN3_PATHWAY, PID_INTEGRIN3_PATHWAY GOMF_GABA_RECEPTOR_BINDING, GOMF_GABA_RECEPTOR_BINDING YANG_BREAST_CANCER_ESR1_DN, YANG_BREAST_CANCER_ESR1_DN REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL, REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL BIOCARTA_BARRESTIN_SRC_PATHWAY, BIOCARTA_BARRESTIN_SRC_PATHWAY HP_MULTIPLE_PRENATAL_FRACTURES, HP_MULTIPLE_PRENATAL_FRACTURES GOBP_REGULATION_OF_GOLGI_ORGANIZATION, GOBP_REGULATION_OF_GOLGI_ORGANIZATION HINATA_NFKB_TARGETS_KERATINOCYTE_DN, HINATA_NFKB_TARGETS_KERATINOCYTE_DN HP_ABNORMAL_PLACENTA_MORPHOLOGY, HP_ABNORMAL_PLACENTA_MORPHOLOGY MATZUK_EARLY_ANTRAL_FOLLICLE, MATZUK_EARLY_ANTRAL_FOLLICLE BIOCARTA_BARR_MAPK_PATHWAY, BIOCARTA_BARR_MAPK_PATHWAY GOBP_NEGATIVE_REGULATION_OF_COAGULATION, GOBP_NEGATIVE_REGULATION_OF_COAGULATION GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY, GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY REACTOME_PROSTACYCLIN_SIGNALLING_THROUGH_PROSTACYCLIN_RECEPTOR, REACTOME_PROSTACYCLIN_SIGNALLING_THROUGH_PROSTACYCLIN_RECEPTOR SERVITJA_ISLET_HNF1A_TARGETS_DN, SERVITJA_ISLET_HNF1A_TARGETS_DN REACTOME_RUNX2_REGULATES_OSTEOLAST_DIFFERENTIATION, REACTOME_RUNX2_REGULATES_OSTEOLAST_DIFFERENTIATION GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS WP_MONOAMINE_TRANSPORT, WP_MONOAMINE_TRANSPORT HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL HP_ABSENT_TOENAIL, HP_ABSENT_TOENAIL DESCARTES_MAIN_FETAL_INHIBITORY_INTERNEURONS, DESCARTES_MAIN_FETAL_INHIBITORY_INTERNEURONS GOMF_FIBROBLAST_GROWTH_FACTOR_BINDING, GOMF_FIBROBLAST_GROWTH_FACTOR_BINDING GOBP_NEUROTRANSMITTER_UPTAKE, GOBP_NEUROTRANSMITTER_UPTAKE GOCC_GLIAL_CELL_PROJECTION, GOCC_GLIAL_CELL_PROJECTION HP_ABNORMALITY_OF_THE_4TH_METACARPAL, HP_ABNORMALITY_OF_THE_4TH_METACARPAL GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY, GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY WP_SEROTONIN_TRANSPORTER_ACTIVITY, WP_SEROTONIN_TRANSPORTER_ACTIVITY GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT, GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT GOBP_DENDRITE_EXTENSION, GOBP_DENDRITE_EXTENSION GOBP_PHOSPHATIDYLGLYCEROL_ACYL_CHAIN_REMODELING, GOBP_PHOSPHATIDYLGLYCEROL_ACYL_CHAIN_REMODELING GOBP_AMINE_CATABOLIC_PROCESS, GOBP_AMINE_CATABOLIC_PROCESS GOBP_ZYMOGEN_INHIBITION, GOBP_ZYMOGEN_INHIBITION MODULE_462, MODULE_462 GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS, GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS GOBP_MESODERMAL_CELL_DIFFERENTIATION, GOBP_MESODERMAL_CELL_DIFFERENTIATION GOBP_SMOOTH_MUSCLE_CELL_MIGRATION, GOBP_SMOOTH_MUSCLE_CELL_MIGRATION HP_ABNORMALITY_OF_THE_5TH_METACARPAL, HP_ABNORMALITY_OF_THE_5TH_METACARPAL GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP, GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP GOBP_FIBRINOLYSIS, GOBP_FIBRINOLYSIS MIR5705, MIR5705 GOBP_NEGATIVE_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_NEGATIVE_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE MODULE_505, MODULE_505 GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_PROLIFERATION, GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_PROLIFERATION GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_MIGRATION, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_MIGRATION GOBP_REGULATION_OF_ADIPONECTIN_SECRETION, GOBP_REGULATION_OF_ADIPONECTIN_SECRETION GOBP_NEUROTRANSMITTER_REUPTAKE, GOBP_NEUROTRANSMITTER_REUPTAKE GSE29614_CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMC_UP, GSE29614_CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMC_UP GOMF_SOLUTE_CATION_ANTIPORTER_ACTIVITY, GOMF_SOLUTE_CATION_ANTIPORTER_ACTIVITY GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS HP_SECRETORY_DIARRHEA, HP_SECRETORY_DIARRHEA GOMF_COLLAGEN_BINDING_INVOLVED_IN_CELL_MATRIX_ADHESION, GOMF_COLLAGEN_BINDING_INVOLVED_IN_CELL_MATRIX_ADHESION GOBP_POSITIVE_REGULATION_OF_AMINE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_AMINE_TRANSPORT HP_ATYPICAL_ABSENCE_SEIZURE, HP_ATYPICAL_ABSENCE_SEIZURE LIU_VAV3_PROSTATE_CARCINOGENESIS_DN, LIU_VAV3_PROSTATE_CARCINOGENESIS_DN GOBP_IMMATURE_T_CELL_PROLIFERATION, GOBP_IMMATURE_T_CELL_PROLIFERATION MONTERO_THYROID_CANCER_POOR_SURVIVAL_DN, MONTERO_THYROID_CANCER_POOR_SURVIVAL_DN BIOCARTA_IL5_PA_THYROID, BIOCARTA_IL5_PA_THYROID GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_2_BINDING, GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_2_BINDING GOMF_CASPASE_BINDING, GOMF_CASPASE_BINDING DESCARTES_FETAL_INTESTINE_LYMPHATIC_ENDOTHELIAL_CELLS, DESCARTES_FETAL_INTESTINE_LYMPHATIC_ENDOTHELIAL_CELLS HP_SPONTANEOUS_RECURRENT_EPISTAXIS, HP_SPONTANEOUS_RECURRENT_EPISTAXIS GOMF_C_X_C_CHEMOKINE_BINDING, GOMF_C_X_C_CHEMOKINE_BINDING GOBP_TRYPTOPHAN_CATABOLIC_PROCESS_TO_KYNURENINE, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS_TO_KYNURENINE GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_UPTAKE, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_UPTAKE HP_CEPHALOHEMATOMA, HP_CEPHALOHEMATOMA GOBP_CARDIAC_MUSCLE_TISSUE_REGENERATION, GOBP_CARDIAC_MUSCLE_TISSUE_REGENERATION REACTOME_CYTOSOLIC_SULFONATION_OF_SMALL_MOLECULES, REACTOME_CYTOSOLIC_SULFONATION_OF_SMALL_MOLECULES chr11q25, chr11q25 GOBP_IMMATURE_T_CELL_PROLIFERATION_IN_THYMUS, GOBP_IMMATURE_T_CELL_PROLIFERATION_IN_THYMUS REACTOME_GLUCAGON_TYPE_LIGAND_RECEPTORS, REACTOME_GLUCAGON_TYPE_LIGAND_RECEPTORS chr4p15, chr4p15 REACTOME_ADORA2B_MEDIATED_ANTINFLAMMATORY_CYTOKINES_PRODUCTION, REACTOME_ADORA2B_MEDIATED_ANTINFLAMMATORY_CYTOKINES_PRODUCTION GOBP_REGULATION_OF_GAP_JUNCTION_ASSEMBLY, GOBP_REGULATION_OF_GAP_JUNCTION_ASSEMBLY GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MOLECULE_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MOLECULE_PRODUCTION BIOCARTA_EXTRINSIC_PATHWAY, BIOCARTA_EXTRINSIC_PATHWAY GOBP_NEGATIVE_REGULATION_OF_LIPID_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_LIPID_TRANSPORT
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