

IL2RA\_HIGH\_DAY3\_EFF\_CD8\_TCELL\_DN,GSE19825\_CD24LOW\_VS\_IL2RA\_HIGH\_DAY3\_EFF\_CD8\_TCELL\_DN

GSE39110\_UNREATED\_VS\_IL2\_TREATED\_CD8\_TCELL\_DAY3\_POST\_IMMUNIZATION\_DN,GSE39110\_UNREATED\_VS\_IL2\_TREATED\_CD8\_TCELL\_DAY3\_POST\_IMMUNIZATION\_DN,GSE6090\_UNSTIM\_VS\_DC\_SIGN\_STIM\_DC\_UP,GSE6090\_UNSTIM\_VS\_DC\_SIGN\_STIM\_DC\_UP,GSE21670\_TGFB\_VS\_TGFB\_AND\_IL6\_TREATED\_STAT3\_KO\_CD4\_TCELL\_DN,GSE21670\_TGFB\_VS\_TGFB\_AND\_IL6\_TREATED\_STAT3\_KO\_CD4\_TCELL\_DN,GSE14308\_TH2\_VS\_TH17\_UP,GSE14308\_TH2\_VS\_TH17\_UP,GSE5589\_UNSTIM\_VS\_45MIN\_LPS\_AND\_IL6\_STIM\_MACROPHAGE\_DN,GSE5589\_UNSTIM\_VS\_45MIN\_LPS\_AND\_IL6\_STIM\_MACROPHAGE\_DN,GSE4748\_CTRL\_VS\_LPS\_STIM\_DC\_3H\_DN,GSE4748\_CTRL\_VS\_LPS\_STIM\_DC\_3H\_DN,GSE4811\_CLASSICALLY\_ACTIVATED\_VS\_TYPE\_2\_ACTIVATED\_MACROPHAGE\_UP,GSE4811\_CLASSICALLY\_ACTIVATED\_VS\_TYPE\_2\_ACTIVATED\_MACROPHAGE\_UP,GSE3982\_BASOPHIL\_VS\_CENT\_MEMORY\_CD4\_TCELL\_DN,GSE3982\_BASOPHIL\_VS\_CENT\_MEMORY\_CD4\_TCELL\_DN,GSE2405\_HEAT\_KILLED\_VS\_LIVE\_A\_PHAGOCYTOPHILUM\_STIM\_NEUTROPHIL\_9H\_DN,GSE2405\_HEAT\_KILLED\_VS\_LIVE\_A\_PHAGOCYTOPHILUM\_STIM\_NEUTROPHIL\_9H\_DN,GSE17580\_TREG\_VS\_TEFF\_UP,GSE17580\_TREG\_VS\_TEFF\_UP,GSE5503\_LIVER\_DC\_VS\_SPLEEN\_DC\_ACTIVATED\_ALLOGENIC\_TCELL\_UP,GSE5503\_LIVER\_DC\_VS\_SPLEEN\_DC\_ACTIVATED\_ALLOGENIC\_TCELL\_UP,GSE17721\_LPS\_VS\_CPG\_16H\_BMDC\_DN,GSE17721\_LPS\_VS\_CPG\_16H\_BMDC\_DN,GSE360\_T\_GONDII\_VS\_M\_TUBERCULOSIS\_DC\_UP,GSE360\_T\_GONDII\_VS\_M\_TUBERCULOSIS\_DC\_UP,MIR4252,MIR4252,GOCC\_PROTEIN\_KINASE\_COMPLEX,GOCC\_PROTEIN\_KINASE\_COMPLEX,MIR6847\_5P,MIR6847\_5P,MIR18A\_5P,MIR18B\_5P,MIR18A\_5P,MIR18B\_5P,GSE42088\_2H\_VS\_24H\_LEISHMANIA\_INF\_DC\_DN,GSE42088\_2H\_VS\_24H\_LEISHMANIA\_INF\_DC\_DN,MIR4735\_3P,MIR4735\_3P,HALLMARK\_ANDROGEN\_RESPONSE,HALLMARK\_ANDROGEN\_RESPONSE,MIR367\_5P,MIR367\_5P,GSE37301\_CD4\_TCELL\_VS\_RAG2\_KO\_NK\_CELL\_UP,GSE37301\_CD4\_TCELL\_VS\_RAG2\_KO\_NK\_CELL\_UP,MIR6761\_5P,MIR6761\_5P,GSE27786\_NKTCCELL\_VS\_ERYTHROBLAST\_DN,GSE27786\_NKTCCELL\_VS\_ERYTHROBLAST\_DN,TCGA\_GLOBLASTOMA\_COPY\_NUMBER\_UP,TCGA\_GLOBLASTOMA\_COPY\_NUMBER\_UP,GSE32423\_MEMORY\_VS\_NAIVE\_CD8\_TCELL\_UP,GSE32423\_MEMORY\_VS\_NAIVE\_CD8\_TCELL\_UP,GSE15659\_CD45RA\_NEG\_CD4\_TCELL\_VS\_RESTING\_TREG\_UP,GSE15659\_CD45RA\_NEG\_CD4\_TCELL\_VS\_RESTING\_TREG\_UP,MIR4777\_5P,MIR4777\_5P,MIR4714\_3P,MIR4714\_3P,RIZKI\_TUMOR\_INVASIVENESS\_2D\_DN,RIZKI\_TUMOR\_INVASIVENESS\_2D\_DN,WP\_ENDOPLASMIC\_RETICULUM\_STRESS\_RESPONSE\_IN\_CORONAVIRUS\_INFECTON,WP\_ENDOPLASMIC\_RETICULUM\_STRESS\_RESPONSE\_IN\_CORONAVIRUS\_INFECTON,MIR455\_5P,MIR455\_5P,MIR644A,MIR644A,GOBP\_NUCLEOBASE\_CONTAINING\_SMALL\_MOLECULE\_CATABOLIC\_PROCESS,GOBP\_NUCLEOBASE\_CONTAINING\_SMALL\_MOLECULE\_CATABOLIC\_PROCESS,GOBP\_REGULATION\_OF\_UBIQUITIN\_PROTEIN\_LIGASE\_ACTIVITY,GOBP\_REGULATION\_OF\_UBIQUITIN\_PROTEIN\_LIGASE\_ACTIVITY,GOMF\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_ACTIVITY,GOMF\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_ACTIVITY,GOBP\_CELLULAR\_RESPONSE\_TO\_OSMOTIC\_STRESS,GOBP\_CELLULAR\_RESPONSE\_TO\_OSMOTIC\_STRESS,MIR4758\_3P,MIR4758\_3P,MIR6861\_3P,MIR6861\_3P,MIR219B\_5P,MIR219B\_5P,MIR1306\_5P,MIR1306\_5P,TSANG\_PBMIC\_FLUVIRIN\_PANDEMRIX\_ADULT\_CORR\_WITH\_CELL\_FREQ\_CD27HI\_CD38HI\_CD20\_NEG\_PLASMABLASTS\_AND\_CD38PLUS\_OF\_IGD\_CD27PLUS\_MEM\_B\_CELLS\_7DY\_POSITIVE,TSANG\_PBMIC\_FLUVIRIN\_PANDEMRIX\_ADULT\_CORR\_WITH\_CELL\_FREQ\_CD27HI\_CD38HI\_CD20\_NEG\_PLASMABLASTS\_AND\_CD38PLUS\_OF\_IGD\_CD27PLUS\_REACTOME\_DETOXIFICATION\_OF\_REACTIVE\_OXYGEN\_SPECIES,REACTOME\_DETOXIFICATION\_OF\_REACTIVE\_OXYGEN\_SPECIES,MIR1243,MIR1243,BARRIER\_COLON\_CANCER\_RECURRENCE\_DN,BARRIER\_COLON\_CANCER\_RECURRENCE\_DN,KEGG\_TIGHT\_JUNCTION,KEGG\_TIGHT\_JUNCTION,GAUSSMANN\_MLL\_AF4\_FUSION\_TARGETS\_D\_UP,GAUSSMANN\_MLL\_AF4\_FUSION\_TARGETS\_D\_UP,HALMOS\_CEBPA\_TARGETS\_UP,HALMOS\_CEBPA\_TARGETS\_UP,BIOCARTA\_PRC2\_PATHWAY,BIOCARTA\_PRC2\_PATHWAY,IL1\_CYTIDINE\_ANALOGS\_CYCOTOXICITY,IL1\_CYTIDINE\_ANALOGS\_CYCOTOXICITY,GOBP\_MAINTENANCE\_OF\_BLOOD\_BRAIN\_BARRIER,GOBP\_MAINTENANCE\_OF\_BLOOD\_BRAIN\_BARRIER,TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_LOBULAR\_NORMAL\_DN,TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_LOBULAR\_NORMAL\_DN,MODULE\_433,MODULE\_433,GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_TARGETING\_TO\_MEMBRANE,GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_TARGETING\_TO\_MEMBRANE,GOBP\_REGULATION\_OF\_PROTEIN\_TARGETING\_TO\_MEMBRANE,GOBP\_REGULATION\_OF\_PROTEIN\_TARGETING\_TO\_MEMBRANE,WANG\_RESPONSE\_TO\_FORSKOLIN\_UP,WANG\_RESPONSE\_TO\_FORSKOLIN\_UP,GOMF\_GDP DISSOCIATION\_INHIBITOR\_ACTIVITY,GOMF\_GDP DISSOCIATION\_INHIBITOR\_ACTIVITY,GOBP\_STRIATUM\_DEVELOPMENT,GOBP\_STRIATUM\_DEVELOPMENT,GOBP\_NEGATIVE\_REGULATION\_OF\_SECRETION,GOBP\_NEGATIVE\_REGULATION\_OF\_SECRETION,ANDERSON\_BLOOD\_CN54GP140\_ADJUVANTED\_WITH\_GLA\_AF\_AGE\_18\_45YO\_3DY\_UP,ANDERSON\_BLOOD\_CN54GP140\_ADJUVANTED\_WITH\_GLA\_AF\_AGE\_18\_45YO\_3DY\_UP,HOLLERN\_MICROACINAR\_BREAST\_TUMOR\_UP,HOLLERN\_MICROACINAR\_BREAST\_TUMOR\_UP,GOMF\_RNA\_POLYMERASE\_II\_CTD\_HEPTAPEPTIDE\_REPEAT\_KINASE\_ACTIVITY,GOMF\_RNA\_POLYMERASE\_II\_CTD\_HEPTAPEPTIDE\_REPEAT\_KINASE\_ACTIVITY,VALK\_AML\_CLUSTER\_4,VALK\_AML\_CLUSTER\_4,ZHANG\_RESPONSE\_TO\_CANTHARIDIN\_UP,ZHANG\_RESPONSE\_TO\_CANTHARIDIN\_UP,GOCC\_INTEGRAL\_COMPONENT\_OF\_SYNAPTIC\_VESICLE\_MEMBRANE,GOCC\_INTEGRAL\_COMPONENT\_OF\_SYNAPTIC\_VESICLE\_MEMBRANE,GOBP\_NEGATIVE\_REGULATION\_BY\_HOST\_OF\_VIRAL\_TRANSCRIPTION,GOBP\_NEGATIVE\_REGULATION\_BY\_HOST\_OF\_VIRAL\_TRANSCRIPTION,WP\_MIRNAS\_INVOLVED\_IN\_DNA\_DAMAGE\_RESPONSE,WP\_MIRNAS\_INVOLVED\_IN\_DNA\_DAMAGE\_RESPONSE,GOBP\_POSITIVE\_REGULATION\_OF\_LEUKOCYTE\_CHEMOTAXIS,GOBP\_POSITIVE\_REGULATION\_OF\_LEUKOCYTE\_CHEMOTAXIS,GOBP\_REGULATION\_OF\_NECROTIC\_CELL\_DEATH,GOBP\_REGULATION\_OF\_NECROTIC\_CELL\_DEATH,GOBP\_SUBPALLIUM\_DEVELOPMENT,GOBP\_SUBPALLIUM\_DEVELOPMENT,HOEK\_NK\_CELL\_2011\_2012\_TIV\_7D\_VS\_0DY\_ADULT\_7D\_DN,HOEK\_NK\_CELL\_2011\_2012\_TIV\_7D\_VS\_0DY\_ADULT\_7D\_DN,GOBP\_NECROPTOTIC\_PROCESS,GOBP\_NECROPTOTIC\_PROCESS,JEISEN\_SMRT\_TARGETS,JEISEN\_SMRT\_TARGETS,chr6p25,chr6p25,GOBP\_DRUG\_METABOLIC\_PROCESS,GOBP\_DRUG\_METABOLIC\_PROCESS,GOMF\_HYDROPEROXY\_ICOSATETRAENOATE\_DEHYDRATASE\_ACTIVITY,GOMF\_HYDROPEROXY\_ICOSATETRAENOATE\_DEHYDRATASE\_ACTIVITY,HE\_PTEN\_TARGETS\_UP,HE\_PTEN\_TARGETS\_UP,GOBP\_POSITIVE\_REGULATION\_OF\_NITRIC\_OXIDE\_SYNTHASE\_BIOSYNTHETIC\_PROCESS,GOBP\_POSITIVE\_REGULATION\_OF\_NITRIC\_OXIDE\_SYNTHASE\_BIOSYNTHETIC\_PROCESS,NABA\_BASEMENT\_MEMBRANES,NABA\_BASEMENT\_MEMBRANES,GOBP\_NEGATIVE\_REGULATION\_OF\_SODIUM\_ION\_TRANSPORT,GOBP\_NEGATIVE\_REGULATION\_OF\_SODIUM\_ION\_TRANSPORT,GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_ETHER\_BONDS,GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_ETHER\_BONDS,REACTOME\_MISCELLANEOUS\_SUBSTRATES,REACTOME\_MISCELLANEOUS\_SUBSTRATES,HP\_SLOWED\_SLURRED\_SPEECH,HP\_SLOWED\_SLURRED\_SPEECH,GOBP\_NEGATIVE\_REGULATION\_OF\_INTERLEUKIN\_1\_ALPHA\_PRODUCTION,GOBP\_NEGATIVE\_REGULATION\_OF\_INTERLEUKIN\_1\_ALPHA\_PRODUCTION,GOBP\_NEGATIVE\_REGULATION\_OF\_CHOLESTEROL\_METABOLIC\_PROCESS,GOBP\_NEGATIVE\_REGULATION\_OF\_CHOLESTEROL\_METABOLIC\_PROCESS,GOBP\_POSITIVE\_REGULATION\_OF\_EXECUTION\_PHASE\_OF\_APOPTOSIS,GOBP\_POSITIVE\_REGULATION\_OF\_EXECUTION\_PHASE\_OF\_APOPTOSIS,GOBP\_NEGATIVE\_REGULATION\_OF\_TYPE\_2\_IMMUNE\_RESPONSE,GOBP\_NEGATIVE\_REGULATION\_OF\_TYPE\_2\_IMMUNE\_RESPONSE,GOBP\_ENDOCRINE\_PANCREAS\_DEVELOPMENT,GOBP\_ENDOCRINE\_PANCREAS\_DEVELOPMENT,GOBP\_T\_CELL\_CHEMOTAXIS,GOBP\_T\_CELL\_CHEMOTAXIS,GOBP\_INSEMINATION,GOBP\_INSEMINATION,WP\_KYNURENINE\_PATHWAY\_AND\_LINKS\_TO\_CELLULAR\_SENESCENCE,WP\_KYNURENINE\_PATHWAY\_AND\_LINKS\_TO\_CELLULAR\_SENESCENCE,GOMF\_SELENOCYSTEINE\_INSERTION\_SEQUENCE\_BINDING,GOMF\_SELENOCYSTEINE\_INSERTION\_SEQUENCE\_BINDING,HP\_PATELLAR\_HYPOPLASIA,HP\_PATELLAR\_HYPOPLASIA,GOBP\_MHC\_CLASS\_II\_BIOSYNTHETIC\_PROCESS,GOBP\_MHC\_CLASS\_II\_BIOSYNTHETIC\_PROCESS,HP\_CONGENITAL\_MUSCULAR\_TORTICOLLIS,HP\_CONGENITAL\_MUSCULAR\_TORTICOLLIS,GOBP\_REGULATION\_OF\_ISOTYPE\_SWITCHING\_TO\_IGE\_ISOTYPES,GOBP\_REGULATION\_OF\_ISOTYPE\_SWITCHING\_TO\_IGE\_ISOTYPES,GOMF\_GLYCERALDEHYDE\_3\_PHOSPHATE\_DEHYDROGENASE\_NADPLUS\_NON\_PHOSPHORYLATING\_ACTIVITY,GOMF\_GLYCERALDEHYDE\_3\_PHOSPHATE\_DEHYDROGENASE\_NADPLUS\_NON\_PHOSPHORYLATING\_ACTIVITY,HP\_HYPEREXTENSIBILITY\_AT\_WRISTS,HP\_HYPEREXTENSIBILITY\_AT\_WRISTS,GOBP\_ANTERIOR\_COMMISSURE\_MORPHOGENESIS,GOBP\_ANTERIOR\_COMMISSURE\_MORPHOGENESIS,GOBP\_CHEMOREPULSION\_OF\_AXON,GOBP\_CHEMOREPULSION\_OF\_AXON,GOBP\_NITRIC\_OXIDE\_SYNTHASE\_BIOSYNTHETIC\_PROCESS,GOBP\_NITRIC\_OXIDE\_SYNTHASE\_BIOSYNTHETIC\_PROCESS,GOBP\_NEGATIVE\_REGULATION\_OF\_BONE\_RESORPTION,GOBP\_NEGATIVE\_REGULATION\_OF\_BONE\_RESORPTION,GOBP\_PANCREATIC\_JUICE\_SECRETION,GOBP\_PANCREATIC\_JUICE\_SECRETION,PID\_INTEGRIN4\_PATHWAY,PID\_INTEGRIN4\_PATHWAY,WP\_METHIONINE\_METABOLISM\_LEADING\_TO\_SULPHUR\_AMINO\_ACIDS\_AND\_RELATED\_DISORDERS,WP\_METHIONINE\_METABOLISM\_LEADING\_TO\_SULPHUR\_AMINO\_ACIDS\_AND\_RELATED\_DISORDERS,GOBP\_INTERLEUKIN\_1\_ALPHA\_PRODUCTION,GOBP\_INTERLEUKIN\_1\_ALPHA\_PRODUCTION,GOMF\_ENDOPEPTIDASE\_REGULATOR\_ACTIVITY,GOMF\_ENDOPEPTIDASE\_REGULATOR\_ACTIVITY,HARALAMBIEVA\_PBMCM\_MR\_IL2AGE\_11\_22YO\_VACCINATED\_VS\_UNVACCINATED\_LOW\_ANTIBODY\_RESPONDERS\_TO\_TREATMENT\_7YR\_UP,HARALAMBIEVA\_PBMCM\_MR\_IL2AGE\_11\_22YO\_VACCINATED\_VS\_UNVACCINATED\_LOW\_ANTIBODY\_RESPONDERS\_TO\_TREATMENT\_7YR\_UP,KEGG\_NICOTINATE\_AND\_NICOTINAMIDE\_METABOLISM,KEGG\_NICOTINATE\_AND\_NICOTINAMIDE\_METABOLISM,ZHENG\_FOXP3\_TARGETS\_DN,ZHENG\_FOXP3\_TARGETS\_DN,HP\_RETINAL\_FOLD,HP\_RETINAL\_FOLD,HP\_MENOMETRORRHAGIA,HP\_MENOMETRORRHAGIA,GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_OSMOTIC\_STRESS,GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_OSMOTIC\_STRESS,GOBP\_CD4\_POSITIVE\_CD25\_POSITIVE\_ALPHA\_BETA\_REGULATORY\_T\_CELL\_DIFFERENTIATION,GOBP\_CD4\_POSITIVE\_CD25\_POSITIVE\_ALPHA\_BETA\_REGULATORY\_T\_CELL\_DIFFERENTIATION,GOBP\_REGULATION\_OF\_POTASSIUM\_ION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY,GOBP\_REGULATION\_OF\_POTASSIUM\_ION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY,GOBP\_INNERVATION,GOBP\_INNERVATION,WP\_TRYPTOPHAN\_CATABOLISM\_LEADING\_TO\_NAD\_PRODUCTION,WP\_TRYPTOPHAN\_CATABOLISM\_LEADING\_TO\_NAD\_PRODUCTION,chr17q12,chr17q12,LE\_SKI\_TARGETS\_UP,LE\_SKI\_TARGETS\_UP,HP\_INCREASED\_LAXITY\_OF\_FINGERS,HP\_INCREASED\_LAXITY\_OF\_FINGERS,GOBP\_REGULATION\_OF\_CD4\_POSITIVE\_CD25\_POSITIVE\_ALPHA\_BETA\_REGULATORY\_T\_CELL\_DIFFERENTIATION,GOBP\_REGULATION\_OF\_CD4\_POSITIVE\_CD25\_POSITIVE\_ALPHA\_BETA\_REGULATORY\_T\_CELL\_DIFFERENTIATION,GOBP\_NEGATIVE\_REGULATION\_OF\_ISOTYPE\_SWITCHING,GOBP\_NEGATIVE\_REGULATION\_OF\_ISOTYPE\_SWITCHING,GOBP\_REGULATION\_OF\_PANCREATIC\_JUICE\_SECRETION,GOBP\_REGULATION\_OF\_PANCREATIC\_JUICE\_SECRETION,GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_RESPIRATION,GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_RESPIRATION,REACTOME\_N\_GLYCAN\_TRIMMING\_AND\_ELONGATION\_IN\_THE\_CIS\_GOLGI,REACTOME\_N\_GLYCAN\_TRIMMING\_AND\_ELONGATION\_IN\_THE\_CIS\_GOLGI,GOBP\_VITAMIN\_K\_METABOLIC\_PROCESS,GOBP\_VITAMIN\_K\_METABOLIC\_PROCESS,GOCC\_EXTRINSIC\_COMPONENT\_OF\_EXTERNAL\_SIDE\_OF\_PLASMA\_MEMBRANE,GOCC\_EXTRINSIC\_COMPONENT\_OF\_EXTERNAL\_SIDE\_OF\_PLASMA\_MEMBRANE,GOBP\_REGULATION\_OF\_PHOSPHOLIPASE\_A2\_ACTIVITY,GOBP\_REGULATION\_OF\_PHOSPHOLIPASE\_A2\_ACTIVITY,GOCC\_LAMININ\_COMPLEX,GOCC\_LAMININ\_COMPLEX,GOBP\_CEREBRAL\_CORTX\_TANGENTIAL\_MIGRATION,GOBP\_CEREBRAL\_CORTX\_TANGENTIAL\_MIGRATION,REACTOME\_CYP2E1\_REACTIONS,REACTOME\_CYP2E1\_REACTIONS,WP\_MIRNA\_TARGETS\_IN\_FCM\_AND\_MEMBRANE\_RECEPTORS,WP\_MIRNA\_TARGETS\_IN\_FCM\_AND\_MEMBRANE\_RECEPTORS,GOBP\_POSITIVE\_REGULATION\_OF\_PROSTAGLANDIN\_BIOSYNTHETIC\_PROCESS,GOBP\_POSITIVE\_REGULATION\_OF\_PROSTAGLANDIN\_BIOSYNTHETIC\_PROCESS,chr7q21,chr7q21,GOBP\_NADH\_OXIDATION,GOBP\_NADH\_OXIDATION,GIAROLA\_SILVA\_BLOOD\_INFLUENZA\_A\_AGE\_21\_51YO\_3DY\_UP,GIAROLA\_SILVA\_BLOOD\_INFLUENZA\_A\_AGE\_21\_51YO\_3DY\_UP,GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_OSMOTIC\_STRESS,GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_OSMOTIC\_STRESS,GOBP\_CELL\_PROLIFERATION\_IN\_MIDBRAIN,GOBP\_CELL\_PROLIFERATION\_IN\_MIDBRAIN,REACTOME\_TIGHT\_JUNCTION\_INTERACTIONS,REACTOME\_TIGHT\_JUNCTION\_INTERACTIONS,REACTOME\_COMMON\_PATHWAY\_OF\_FIBRIN\_CLOT\_FORMATION,REACTOME\_COMMON\_PATHWAY\_OF\_FIBRIN\_CLOT\_FORMATION,GOBP\_HEMIDESMOSOME\_ASSEMBLY,GOBP\_HEMIDESMOSOME\_ASSEMBLY,GOBP\_REGULATION\_OF\_DEVELOPMENT\_HETEROCHRONIC,GOBP\_REGULATION\_OF\_DEVELOPMENT\_HETEROCHRONIC,GIAROLA\_SILVA\_BLOOD\_PANDEMRIX\_AGE\_21\_51YO\_30DY\_UP,GIAROLA\_SILVA\_BLOOD\_PANDEMRIX\_AGE\_21\_51YO\_30DY\_UP,GOMF\_CORECEPTOR\_ACTIVITY\_INVOLVED\_IN\_WNT\_SIGNALING\_PATHWAY\_PLANAR\_CELL\_POLARITY\_PATHWAY,GOMF\_CORECEPTOR\_ACTIVITY\_INVOLVED\_IN\_WNT\_SIGNALING\_PATHWAY\_PLANAR\_CELL\_POLARITY\_PATHWAY,GNF2\_CDKNIC,GNF2\_CDKNIC,GOBP\_POSITIVE\_REGULATION\_OF\_UNSATURATED\_FATTY\_ACID\_BIOSYNTHETIC\_PROCESS,GOBP\_POSITIVE\_REGULATION\_OF\_UNSATURATED\_FATTY\_ACID\_BIOSYNTHETIC\_PROCESS,GOBP\_NEGATIVE\_REGULATION\_OF\_B\_CELL\_MEDIATED\_IMMUNITY,GOBP\_NEGATIVE\_REGULATION\_OF\_B\_CELL\_MEDIATED\_IMMUNITY,GOBP\_NEGATIVE\_REGULATION\_OF\_POTASSIUM\_ION\_TRANSMEMBRANE\_TRANSPORT,GOBP\_NEGATIVE\_REGULATION\_OF\_POTASSIUM\_ION\_TRANSMEMBRANE\_TRANSPORT