GSE17721\_POLYIC\_VS\_CPG\_1H\_BMDC\_UP, GSE17721\_POLYIC\_VS\_CPG\_1H\_BMDC\_UP GOBP\_VESICLE\_TARGETING, GOBP\_VESICLE\_TARGETING GSE20715\_WT\_VS\_TLR4\_KO\_6H\_OZONE\_LUNG\_UP, GSE20715\_WT\_VS\_TLR4\_KO\_6H\_OZONE\_LUNG\_UP GOCC\_MEDIATOR\_COMPLEX, GOCC\_MEDIATOR\_COMPLEX REACTOME\_GLYCEROPHOSPHOLIPID\_BIOSYNTHESIS, REACTOME\_GLYCEROPHOSPHOLIPID\_BIOSYNTHESIS HP\_ABNORMALITY\_OF\_THE\_PARATHYROID\_GLAND, HP\_ABNORMALITY\_OF\_THE\_PARATHYROID\_GLAND HP\_ABNORMAL\_GLYCOSYLATION, HP\_ABNORMAL\_GLYCOSYLATION GOBP\_ER\_NUCLEUS\_SIGNALING\_PATHWAY, GOBP\_ER\_NUCLEUS\_SIGNALING\_PATHWAY GOBP\_PROTEIN\_LOCALIZATION\_TO\_CELL\_SURFACE, GOBP\_PROTEIN\_LOCALIZATION\_TO\_CELL\_SURFACE HP\_NARROW\_PALPEBRAL\_FISSURE, HP\_NARROW\_PALPEBRAL\_FISSURE MIR4726\_3P, MIR4726\_3P HP\_EXCESSIVE\_WRINKLED\_SKIN, HP\_EXCESSIVE\_WRINKLED\_SKIN MIR6508\_3P, MIR6508\_3P GSE13887\_ACT\_CD4\_VS\_NO\_TREATED\_CD4\_TCELL\_DN, GSE13887\_ACT\_CD4\_VS\_NO\_TREATED\_CD4\_TCELL\_DN MIR6501\_5P, MIR6501\_5P MIR3675\_5P, MIR3675\_5P HP\_EPENDYMOMA, HP\_EPENDYMOMA STEGMEIER\_PREMITOTIC\_CELL\_CYCLE\_REGULATORS, STEGMEIER\_PREMITOTIC\_CELL\_CYCLE\_REGULATORS HP\_SMALL\_EARLOBE, HP\_SMALL\_EARLOBE GOBP\_HORMONE\_BIOSYNTHETIC\_PROCESS, GOBP\_HORMONE\_BIOSYNTHETIC\_PROCESS chr1p35, chr1p35 GOCC\_DERLIN\_1\_RETROTRANSLOCATION\_COMPLEX, GOCC\_DERLIN\_1\_RETROTRANSLOCATION\_COMPLEX HP\_ASTROCYTOSIS, HP\_ASTROCYTOSIS BIOCARTA\_ARAP\_PATHWAY, BIOCARTA\_ARAP\_PATHWAY HP\_NEUROFIBRILLARY\_TANGLES, HP\_NEUROFIBRILLARY\_TANGLES WANG\_RESPONSE\_TO\_FORSKOLIN\_UP, WANG\_RESPONSE\_TO\_FORSKOLIN\_UP G\_TREATED\_ENDOTHELIAL\_CELL\_DN, GSE3920\_UNTREATED\_VS\_IFNG\_TREATED\_ENDOTHELIAL\_CELL\_DN HP\_ABNORMAL\_NUMBER\_OF\_PERMANENT\_TEETH, HP\_ABNORMAL\_NUMBER\_OF\_PERMANENT\_TEETH HP\_AGENESIS\_OF\_PERMANENT\_TEETH, HP\_AGENESIS\_OF\_PERMANENT\_TEETH GOBP\_ALCOHOL\_CATABOLIC\_PROCESS, GOBP\_ALCOHOL\_CATABOLIC\_PROCESS OUYANG PROSTATE CANCER PROGRESSION DN, OUYANG PROSTATE CANCER PROGRESSION DN MIR377\_5P, MIR377\_5P GOBP\_STEROID\_HORMONE\_BIOSYNTHETIC\_PROCESS, GOBP\_STEROID\_HORMONE\_BIOSYNTHETIC\_PROCESS GOBP\_ISOPRENOID\_CATABOLIC\_PROCESS, GOBP\_ISOPRENOID\_CATABOLIC\_PROCESS GOMF\_DOPAMINE\_RECEPTOR\_BINDING, GOMF\_DOPAMINE\_RECEPTOR\_BINDING GOBP\_SECRETION\_OF\_LYSOSOMAL\_ENZYMES, GOBP\_SECRETION\_OF\_LYSOSOMAL\_ENZYMES MIR2392, MIR2392 HP\_ABNORMALITY\_OF\_FUNDUS\_PIGMENTATION, HP\_ABNORMALITY\_OF\_FUNDUS\_PIGMENTATION HP\_PLATYBASIA, HP\_PLATYBASIA HP\_ABNORMAL\_RAPID\_EYE\_MOVEMENT\_SLEEP, HP\_ABNORMAL\_RAPID\_EYE\_MOVEMENT\_SLEEP HP\_EPIDERMOID\_CYST, HP\_EPIDERMOID\_CYST GOMF\_ENONE\_REDUCTASE\_ACTIVITY, GOMF\_ENONE\_REDUCTASE\_ACTIVITY GOBP\_SYNAPTIC\_VESICLE\_PRIMING, GOBP\_SYNAPTIC\_VESICLE\_PRIMING GOBP\_MEGAKARYOCYTE\_DEVELOPMENT, GOBP\_MEGAKARYOCYTE\_DEVELOPMENT REACTOME\_HDL\_CLEARANCE, REACTOME\_HDL\_CLEARANCE GOCC\_ENDOLYSOSOME\_LUMEN, GOCC\_ENDOLYSOSOME\_LUMEN GOBP\_ANDROGEN\_BIOSYNTHETIC\_PROCESS, GOBP\_ANDROGEN\_BIOSYNTHETIC\_PROCESS GOBP\_CHOLESTEROL\_CATABOLIC\_PROCESS, GOBP\_CHOLESTEROL\_CATABOLIC\_PROCESS GOBP\_CORTISOL\_METABOLIC\_PROCESS, GOBP\_CORTISOL\_METABOLIC\_PROCESS HP\_DYSESTHESIA, HP\_DYSESTHESIA GOMF\_ALPHA\_2A\_ADRENERGIC\_RECEPTOR\_BINDING, GOMF\_ALPHA\_2A\_ADRENERGIC\_RECEPTOR\_BINDING DESCARTES\_MAIN\_FETAL\_RETINAL\_PROGENITORS\_AND\_MULLER\_GLIA, DESCARTES\_MAIN\_FETAL\_RETINAL\_PROGENITORS\_AND\_MULLER\_GLIA GOBP\_RESPONSE\_TO\_ACIDIC\_PH, GOBP\_RESPONSE\_TO\_ACIDIC\_PH GOBP\_POSITIVE\_REGULATION\_OF\_DEFENSE\_RESPONSE\_TO\_BACTERIUM, GOBP\_POSITIVE\_REGULATION\_OF\_DEFENSE\_RESPONSE\_TO\_BACTERIUM

GSE3920\_IFNB\_VS\_IFNG\_TREATED\_ENDOTHELIAL\_CELL\_DN, GSE3920\_IFNB\_VS\_IFNG\_TREATED\_ENDOTHELIAL\_CELL\_DN GSE7509\_UNSTIM\_VS\_TNFA\_IL1B\_IL6\_PGE\_STIM\_DC\_UP, GSE7509\_UNSTIM\_VS\_TNFA\_IL1B\_IL6\_PGE\_STIM\_DC\_UP GSE40274\_CTRL\_VS\_IRF4\_TRANSDUCED\_ACTIVATED\_CD4\_TCELL\_UP, GSE40274\_CTRL\_VS\_IRF4\_TRANSDUCED\_ACTIVATED\_CD4\_TCELL\_UP GSE18281\_PERIMEDULLARY\_CORTICAL\_REGION\_VS\_WHOLE\_MEDULLA\_THYMUS\_UP, GSE18281\_PERIMEDULLARY\_CORTICAL\_REGION\_VS\_WHOLE GSE39820\_TGFBETA1\_VS\_TGFBETA3\_IN\_IL6\_IL23A\_TREATED\_CD4\_TCELL\_UP, GSE39820\_TGFBETA1\_VS\_TGFBETA3\_IN\_IL6\_IL23A\_TREATED\_CD4\_TCELI GOBP\_PHOSPHATIDYLINOSITOL\_BIOSYNTHETIC\_PROCESS, GOBP\_PHOSPHATIDYLINOSITOL\_BIOSYNTHETIC\_PROCESS GSE16385\_UNTREATED\_VS\_12H\_ROSIGLITAZONE\_TREATED\_MACROPHAGE\_DN, GSE16385\_UNTREATED\_VS\_12H\_ROSIGLITAZONE\_TREATED\_MACR PAPASPYRIDONOS\_UNSTABLE\_ATEROSCLEROTIC\_PLAQUE\_UP, PAPASPYRIDONOS\_UNSTABLE\_ATEROSCLEROTIC\_PLAQUE\_UP GOBP\_PHOSPHATIDYLCHOLINE\_METABOLIC\_PROCESS, GOBP\_PHOSPHATIDYLCHOLINE\_METABOLIC\_PROCESS GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_YELLOW\_DN, GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_YELLOW\_DN GOBP\_REGULATION\_OF\_PROTEIN\_LOCALIZATION\_TO\_CELL\_SURFACE, GOBP\_REGULATION\_OF\_PROTEIN\_LOCALIZATION\_TO\_CELL\_SURFACE GOBP\_NUCLEOSIDE\_TRIPHOSPHATE\_CATABOLIC\_PROCESS, GOBP\_NUCLEOSIDE\_TRIPHOSPHATE\_CATABOLIC\_PROCESS GOBP\_NEGATIVE\_REGULATION\_OF\_INFLAMMATORY\_RESPONSE, GOBP\_NEGATIVE\_REGULATION\_OF\_INFLAMMATORY\_RESPONSE GOMF\_LIPOPOLYSACCHARIDE\_IMMUNE\_RECEPTOR\_ACTIVITY, GOMF\_LIPOPOLYSACCHARIDE\_IMMUNE\_RECEPTOR\_ACTIVITY GOMF\_PROTEIN\_GLUTAMINE\_GAMMA\_GLUTAMYLTRANSFERASE\_ACTIVITY, GOMF\_PROTEIN\_GLUTAMINE\_GAMMA\_GLUTAMYLTRANSFERASE\_A GOMF\_POLYOL\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_POLYOL\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY GOBP\_REGULATION\_OF\_KETONE\_BIOSYNTHETIC\_PROCESS, GOBP\_REGULATION\_OF\_KETONE\_BIOSYNTHETIC\_PROCESS GOBP\_REGULATION\_OF\_THE\_FORCE\_OF\_HEART\_CONTRACTION\_BY\_CHEMICAL\_SIGNAL, GOBP\_REGULATION\_OF\_THE\_FORCE\_OF\_HEART\_CONTI

HP\_ABNORMALITY\_OF\_THE\_ANTITRAGUS, HP\_ABNORMALITY\_OF\_THE\_ANTITRAGUS

GOBP\_NEGATIVE\_REGULATION\_OF\_NODAL\_SIGNALING\_PATHWAY, GOBP\_NEGATIVE\_REGULATION\_OF\_NODAL\_SIGNALING\_PATHWAY