

LEAR\_SPECK, GO\_NUCLEAR\_SPECK

- GSE37533\_PPARG1\_FOXP3\_VS\_PPARG2\_FOXP3\_TRANSDUCECD4\_TCELL\_DN, GSE37533\_PPARG1\_FOXP3\_VS\_PPARG2\_FOXP3\_TRANSDUCECD4\_TCELL\_DN
- GSE25123\_IL4\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_MACROPHAGE\_DAY10\_DN, GSE25123\_IL4\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_MACROPHAGE\_DAY10\_DN
- GO\_PML\_BODY, GO\_PML\_BODY
- DER\_IFN\_BETA\_RESPONSE\_UP, DER\_IFN\_BETA\_RESPONSE\_UP
- STARK\_PREFRONTAL\_CORTEX\_22Q11\_DELETION\_UP, STARK\_PREFRONTAL\_CORTEX\_22Q11\_DELETION\_UP
- MCBRYAN\_PUBERTAL\_BREAST\_5\_6WK\_DN, MCBRYAN\_PUBERTAL\_BREAST\_5\_6WK\_DN
- GO\_HISTONE\_METHYLATION, GO\_HISTONE\_METHYLATION
- MAZ\_Q6, MAZ\_Q6
- GO\_PEPTIDYL\_LYSINE\_METHYLATION, GO\_PEPTIDYL\_LYSINE\_METHYLATION
- GO\_PROTEIN\_KINASE\_COMPLEX, GO\_PROTEIN\_KINASE\_COMPLEX
- GO\_CORE\_PROMOTER\_SEQUENCE\_SPECIFIC\_DNA\_BINDING, GO\_CORE\_PROMOTER\_SEQUENCE\_SPECIFIC\_DNA\_BINDING
- GGTGTGT\_MIR329, GGTGTGT\_MIR329
- GO\_HORMONE\_RECEPTOR\_BINDING, GO\_HORMONE\_RECEPTOR\_BINDING
- MODULE\_486, MODULE\_486
- PID\_HEDGEHOG\_GLI\_PATHWAY, PID\_HEDGEHOG\_GLI\_PATHWAY
- GO\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY, GO\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY
- GO\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR, GO\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR
- SCHAEFFER\_PROSTATE\_DEVELOPMENT\_AND\_CANCER\_BOX4\_DN, SCHAEFFER\_PROSTATE\_DEVELOPMENT\_AND\_CANCER\_BOX4\_DN
- GSE16385\_MONOCYTE\_VS\_12H\_ROSIGLITAZONE\_TREATED\_MACROPHAGE\_UP, GSE16385\_MONOCYTE\_VS\_12H\_ROSIGLITAZONE\_TREATED\_MACROPHAGE\_UP
- GO\_ANDROGEN\_RECEPTOR\_SIGNALING\_PATHWAY, GO\_ANDROGEN\_RECEPTOR\_SIGNALING\_PATHWAY
- GO\_MRNA\_SPLICE\_SITE\_SELECTION, GO\_MRNA\_SPLICE\_SITE\_SELECTION
- GO\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER\_IN\_RESPONSE\_TO\_HYPOXIA, GO\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER\_IN\_RESPONSE\_TO\_HYPOXIA
- GO\_INTRACELLULAR\_STEROID\_HORMONE\_RECEPTOR\_SIGNALING\_PATHWAY, GO\_INTRACELLULAR\_STEROID\_HORMONE\_RECEPTOR\_SIGNALING\_PATHWAY
- AATGGAG\_MIR136, AATGGAG\_MIR136
- GO\_NEPHRON\_EPITHELIUM\_DEVELOPMENT, GO\_NEPHRON\_EPITHELIUM\_DEVELOPMENT
- TAGGTCA\_MIR192\_MIR215, TAGGTCA\_MIR192\_MIR215
- XU\_AKT1\_TARGETS\_6HR, XU\_AKT1\_TARGETS\_6HR
- MATZUK\_SPERMATOGONIA, MATZUK\_SPERMATOGONIA
- GO\_POSITIVE\_REGULATION\_OF\_MESONEPHROS\_DEVELOPMENT, GO\_POSITIVE\_REGULATION\_OF\_MESONEPHROS\_DEVELOPMENT
- GO\_REGULATION\_OF\_MESONEPHROS\_DEVELOPMENT, GO\_REGULATION\_OF\_MESONEPHROS\_DEVELOPMENT
- GO\_BLASTODERM\_SEGMENTATION, GO\_BLASTODERM\_SEGMENTATION
- GO\_ORGAN\_FORMATION, GO\_ORGAN\_FORMATION
- GO\_CILIARY\_BASE, GO\_CILIARY\_BASE
- IIZUKA\_LIVER\_CANCER\_EARLY\_RECURRENCE, IIZUKA\_LIVER\_CANCER\_EARLY\_RECURRENCE
- GO\_REGULATION\_OF\_SMOOTHENED\_SIGNALING\_PATHWAY, GO\_REGULATION\_OF\_SMOOTHENED\_SIGNALING\_PATHWAY
- GO\_SMOOTHENED\_SIGNALING\_PATHWAY, GO\_SMOOTHENED\_SIGNALING\_PATHWAY
- GO\_NUCLEAR\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_HOLOENZYME\_COMPLEX, GO\_NUCLEAR\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_HOLOENZYME\_COMPLEX
- GO\_REGULATION\_OF\_HEART\_GROWTH, GO\_REGULATION\_OF\_HEART\_GROWTH
- GO\_POSITIVE\_REGULATION\_OF\_REPRODUCTIVE\_PROCESS, GO\_POSITIVE\_REGULATION\_OF\_REPRODUCTIVE\_PROCESS
- PAX6\_01, PAX6\_01
- GO\_REGULATION\_OF\_CHONDROCYTE\_DIFFERENTIATION, GO\_REGULATION\_OF\_CHONDROCYTE\_DIFFERENTIATION
- GO\_REGULATION\_OF\_CARTILAGE\_DEVELOPMENT, GO\_REGULATION\_OF\_CARTILAGE\_DEVELOPMENT
- REACTOME\_RESOLUTION\_OF\_AP\_SITES\_VIA\_THE\_SINGLE\_NUCLEOTIDE\_REPLACEMENT\_PATHWAY, REACTOME\_RESOLUTION\_OF\_AP\_SITES\_VIA\_THE\_SINGLE\_NUCLEOTIDE\_REPLACEMENT\_PATHWAY
- VANASSE\_BCL2\_TARGETS\_UP, VANASSE\_BCL2\_TARGETS\_UP
- GO\_PALATE\_DEVELOPMENT, GO\_PALATE\_DEVELOPMENT
- GO\_CRANIAL\_SKELETAL\_SYSTEM\_DEVELOPMENT, GO\_CRANIAL\_SKELETAL\_SYSTEM\_DEVELOPMENT
- GO\_COLUMNAR\_CUBOIDAL\_EPITHELIAL\_CELL\_DIFFERENTIATION, GO\_COLUMNAR\_CUBOIDAL\_EPITHELIAL\_CELL\_DIFFERENTIATION
- GO\_CELLULAR\_RESPONSE\_TO\_TOXIC\_SUBSTANCE, GO\_CELLULAR\_RESPONSE\_TO\_TOXIC\_SUBSTANCE
- REACTOME\_BASE\_FREE\_SUGAR\_PHOSPHATE\_REMOVAL\_VIA\_THE\_SINGLE\_NUCLEOTIDE\_REPLACEMENT\_PATHWAY, REACTOME\_BASE\_FREE\_SUGAR\_PHOSPHATE\_REMOVAL\_VIA\_THE\_SINGLE\_NUCLEOTIDE\_REPLACEMENT\_PATHWAY
- GO\_TRIPARTITE\_REGIONAL\_SUBDIVISION, GO\_TRIPARTITE\_REGIONAL\_SUBDIVISION
- GO\_REGULATION\_OF\_ORGAN\_GROWTH, GO\_REGULATION\_OF\_ORGAN\_GROWTH
- GO\_CELL\_DIFFERENTIATION\_INVOLVED\_IN\_KIDNEY\_DEVELOPMENT, GO\_CELL\_DIFFERENTIATION\_INVOLVED\_IN\_KIDNEY\_DEVELOPMENT
- GO\_MELANOCYTE\_DIFFERENTIATION, GO\_MELANOCYTE\_DIFFERENTIATION
- GO\_MUSCLE\_CELL\_FATE\_COMMITMENT, GO\_MUSCLE\_CELL\_FATE\_COMMITMENT
- GO\_POSITIVE\_REGULATION\_OF\_ORGAN\_GROWTH, GO\_POSITIVE\_REGULATION\_OF\_ORGAN\_GROWTH
- GO\_MESENCHYMAL\_TO\_EPITHELIAL\_TRANSITION, GO\_MESENCHYMAL\_TO\_EPITHELIAL\_TRANSITION
- GO\_KIDNEY\_MESENCHYME\_DEVELOPMENT, GO\_KIDNEY\_MESENCHYME\_DEVELOPMENT
- GO\_PROLINE\_RICH\_REGION\_BINDING, GO\_PROLINE\_RICH\_REGION\_BINDING
- GO\_SPLICEOSOMAL\_TRI\_SNRNP\_COMPLEX\_ASSEMBLY, GO\_SPLICEOSOMAL\_TRI\_SNRNP\_COMPLEX\_ASSEMBLY
- GO\_REGULATION\_OF\_NEUROBLAST\_PROLIFERATION, GO\_REGULATION\_OF\_NEUROBLAST\_PROLIFERATION
- GO\_POSITIVE\_REGULATION\_OF\_HEART\_GROWTH, GO\_POSITIVE\_REGULATION\_OF\_HEART\_GROWTH