DING\_LUNG\_CANCER\_EXPRESSION\_BY\_COPY\_NUMBER, DING\_LUNG\_CANCER\_EXPRESSION\_BY\_COPY\_NUMBER TAATGTG\_MIR323, TAATGTG\_MIR323 ATGTTTC\_MIR494, ATGTTTC\_MIR494 MIR4511, MIR4511 MODULE\_182, MODULE\_182 MIR22\_3P, MIR22\_3P MIR298, MIR298 MIR20A\_3P, MIR20A\_3P GTGCAAA\_MIR507, GTGCAAA\_MIR507 MIR3145\_5P, MIR3145\_5P PR\_01, PR\_01 MIR181A\_2\_3P, MIR181A\_2\_3P MIR3166, MIR3166 MIR491\_3P, MIR491\_3P MIR3922\_5P, MIR3922\_5P GOBP\_MULTICELLULAR\_ORGANISM\_GROWTH, GOBP\_MULTICELLULAR\_ORGANISM\_GROWTH HP\_ABNORMALITY\_OF\_THE\_TARSAL\_BONES, HP\_ABNORMALITY\_OF\_THE\_TARSAL\_BONES MIR5187\_3P, MIR5187\_3P GOCC\_NEURON\_PROJECTION\_CYTOPLASM, GOCC\_NEURON\_PROJECTION\_CYTOPLASM MIR508\_5P, MIR508\_5P MIR4999\_5P, MIR4999\_5P MIR6781\_3P, MIR6781\_3P MIR2682 3P, MIR2682 3P NZA\_AGE\_21\_30YO\_NONRESPONDER\_28DY\_UP, THAKAR\_PBMC\_INACTIVATED\_INFLUENZA\_AGE\_21\_30YO\_NONRESPONDER\_28DY\_UP MIR6800\_5P, MIR6800\_5P GSE15659\_TREG\_VS\_TCONV\_UP, GSE15659\_TREG\_VS\_TCONV\_UP GOBP\_REGULATION\_OF\_PROTEIN\_TYROSINE\_KINASE\_ACTIVITY, GOBP\_REGULATION\_OF\_PROTEIN\_TYROSINE\_KINASE\_ACTIVITY BURTON\_ADIPOGENESIS\_12, BURTON\_ADIPOGENESIS\_12 GACAGGG\_MIR339, GACAGGG\_MIR339 MIR6821\_3P, MIR6821\_3P MIR135A\_3P, MIR135A\_3P GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_TYROSINE\_KINASE\_ACTIVITY, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_TYROSINE\_REGULATION\_OF\_PROTEIN\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROSINE\_TYROS MIR8056, MIR8056 LEE\_METASTASIS\_AND\_ALTERNATIVE\_SPLICING\_DN, LEE\_METASTASIS\_AND\_ALTERNATIVE\_SPLICING\_DN HP\_SYNOSTOSIS\_INVOLVING\_BONES\_OF\_THE\_HAND, HP\_SYNOSTOSIS\_INVOLVING\_BONES\_OF\_THE\_HAND WP\_ENDOCHONDRAL\_OSSIFICATION, WP\_ENDOCHONDRAL\_OSSIFICATION WP\_ENDOCHONDRAL\_OSSIFICATION\_WITH\_SKELETAL\_DYSPLASIAS, WP\_ENDOCHONDRAL\_OSSIFICATION\_WITH\_SKELETAL\_DYSPLA GOBP\_HISTONE\_MRNA\_CATABOLIC\_PROCESS, GOBP\_HISTONE\_MRNA\_CATABOLIC\_PROCESS MIR920, MIR920 MIR6881\_5P, MIR6881\_5P HP\_UNERUPTED\_TOOTH, HP\_UNERUPTED\_TOOTH HP\_UROGENITAL\_SINUS\_ANOMALY, HP\_UROGENITAL\_SINUS\_ANOMALY HP\_SYNOSTOSIS\_INVOLVING\_BONES\_OF\_THE\_LOWER\_LIMBS, HP\_SYNOSTOSIS\_INVOLVING\_BONES\_OF\_THE\_LOWER\_LIMBS GOBP\_EMBRYONIC\_DIGESTIVE\_TRACT\_MORPHOGENESIS, GOBP\_EMBRYONIC\_DIGESTIVE\_TRACT\_MORPHOGENESIS chr5q21, chr5q21 GOBP\_NOSE\_DEVELOPMENT, GOBP\_NOSE\_DEVELOPMENT GOBP\_OUTFLOW\_TRACT\_MORPHOGENESIS, GOBP\_OUTFLOW\_TRACT\_MORPHOGENESIS chr2p24, chr2p24

THAKAR\_PBMC\_INACTIVATED\_INFLUENZA\_AGE\_21\_30YO\_NONRESPONDER\_7DY\_UP, THAKAR\_PBMC\_INACTIVATED\_INFLUENZA\_AGE\_31\_30YO\_NONRESPONDER\_7DY\_UP, THAKAR\_PBMC\_INACTIVATED\_INFLUENTA\_AGE\_31\_30YO\_NONRESPONDER\_7DY\_UP, THAKAR\_PBMC\_INACTIVATED\_INFLUENTA\_AGE\_31\_30YO\_NONRESPONDER\_7DY\_UP, THAKAR\_PBMC\_INACTIVATED\_INFLUENTA\_AGE\_31\_30YO\_NONRESPONDER\_7DY\_UP, THAKAR\_PBMC\_INACTIVATED\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30YO\_NONRESPOND\_INFLUENTA\_AGE\_31\_30