GTCTTCC_MIR7, GTCTTCC_MIR7 TAATGTG_MIR323, TAATGTG_MIR323 DING_LUNG_CANCER_EXPRESSION_BY_COPY_NUMBER, DING_LUNG_CANCER_EXPRESSION_BY_COPY_NUMBER MIR22_3P, MIR22_3P GSE11961 MEMORY BCELL DAY7 VS GERMINAL CENTER BCELL DAY40 UP, GSE11961 MEMORY BCELL DAY7 VS GERMINAL CENTER BCELL DAY40 UP MEK_UP.V1_DN, MEK_UP.V1_DN GCAAAAA_MIR129, GCAAAAA_MIR129 MIR4511, MIR4511 GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN, GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN GSE14415_INDUCED_VS_NATURAL_TREG_UP, GSE14415_INDUCED_VS_NATURAL_TREG_UP MIR3922_5P, MIR3922_5P MIR154_3P, MIR154_3P MIR487A_3P, MIR487A_3P MIR148A_5P, MIR148A_5P ZHENG_CORD_BLOOD_C3_MEGAKARYOCYTE_ERYTHROID_PROGENITOR, ZHENG_CORD_BLOOD_C3_MEGAKARYOCYTE_ERYTHROID_PROGENITOR HAY_BONE_MARROW_FOLLICULAR_B_CELL, HAY_BONE_MARROW_FOLLICULAR_B_CELL MIR3660, MIR3660 MIR181A_2_3P, MIR181A_2_3P MIR4526, MIR4526 GTGCAAA_MIR507, GTGCAAA_MIR507 ATGTTTC_MIR494, ATGTTTC_MIR494 MIR4769_3P, MIR4769_3P GOBP_MACROMOLECULE_DEACYLATION, GOBP_MACROMOLECULE_DEACYLATION GAGACTG_MIR452, GAGACTG_MIR452 MIR4645 3P, MIR4645 3P MIR4502, MIR4502 MIR3145_5P, MIR3145_5P HP_SHORT_FOOT, HP_SHORT_FOOT ACAGGGT_MIR10A_MIR10B, ACAGGGT_MIR10A_MIR10B PR 01, PR 01 GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN, GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN MIR5187_3P, MIR5187_3P PR 02, PR 02 HP_ABNORMALITY_OF_THE_TARSAL_BONES, HP_ABNORMALITY_OF_THE_TARSAL_BONES MIR491_3P, MIR491_3P MIR508_5P, MIR508_5P PID_RAC1_PATHWAY, PID_RAC1_PATHWAY MIR6733_3P, MIR6733_3P GOBP_HEPATICOBILIARY_SYSTEM_DEVELOPMENT, GOBP_HEPATICOBILIARY_SYSTEM_DEVELOPMENT EGR3_01, EGR3_01 MIR4761_3P, MIR4761_3P MIR1324, MIR1324 MIR10A_5P_MIR10B_5P, MIR10A_5P_MIR10B_5P // MIR326, MIR326 MIR5007_3P, MIR5007_3P MIR6874_3P, MIR6874_3P PID_CASPASE_PATHWAY, PID_CASPASE_PATHWAY MIR7703, MIR7703 MIR197_5P, MIR197_5P HP_SMALL_HAND, HP_SMALL_HAND ✓ TACAATC_MIR508, TACAATC_MIR508 — MIR6740_5P, MIR6740_5P HP PETECHIAE, HP PETECHIAE MIR6800_5P, MIR6800_5P MIR6772_3P, MIR6772_3P MIR6780B_3P, MIR6780B_3P CRX_NRL_DN.V1_DN, CRX_NRL_DN.V1_DN GOBP_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY MIR5591_5P, MIR5591_5P GOBP_NEURON_MIGRATION, GOBP_NEURON_MIGRATION MIR1193, MIR1193 MIR4637, MIR4637 MIR4733_3P, MIR4733_3P MIR4300, MIR4300 GENTILE_UV_RESPONSE_CLUSTER_D6, GENTILE_UV_RESPONSE_CLUSTER_D6 PETRETTO_HEART_MASS_QTL_CIS_UP, PETRETTO_HEART_MASS_QTL_CIS_UP HEDENFALK_BREAST_CANCER_BRACX_DN, HEDENFALK_BREAST_CANCER_BRACX_DN TAGGTCA_MIR192_MIR215, TAGGTCA_MIR192_MIR215 GOMF_POLY_PURINE_TRACT_BINDING, GOMF_POLY_PURINE_TRACT_BINDING ZHAN_MULTIPLE_MYELOMA_PR_DN, ZHAN_MULTIPLE_MYELOMA_PR_DN PSIP1_TARGET_GENES, PSIP1_TARGET_GENES MIR6832_5P, MIR6832_5P GOBP_MYD88_DEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY, GOBP_MYD88_DEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY GOBP_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_PATHWAY. AGTTCTC_MIR146A_MIR146B, AGTTCTC_MIR146A_MIR146B GOBP_ESTABLISHMENT_OF_MITOCHONDRION_LOCALIZATION, GOBP_ESTABLISHMENT_OF_MITOCHONDRION_LOCALIZATION KEGG_ALANINE_ASPARTATE_AND_GLUTAMATE_METABOLISM, KEGG_ALANINE_ASPARTATE_AND_GLUTAMATE_METABOLISM MIR920, MIR920 LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN, LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN HP_NON_MIDLINE_CLEFT_LIP, HP_NON_MIDLINE_CLEFT_LIP WP_PRION_DISEASE_PATHWAY, WP_PRION_DISEASE_PATHWAY MIR4791, MIR4791 MIR1909_5P, MIR1909_5P MIR219B_3P, MIR219B_3P GOBP_XENOBIOTIC_TRANSPORT, GOBP_XENOBIOTIC_TRANSPORT GOBP_POSITIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY MIR8056, MIR8056 BIOCARTA_ALK_PATHWAY, BIOCARTA_ALK_PATHWAY GOBP_POSITIVE_REGULATION_OF_BEHAVIOR, GOBP_POSITIVE_REGULATION_OF_BEHAVIOR MIR3909, MIR3909 GOBP_HISTONE_MRNA_CATABOLIC_PROCESS, GOBP_HISTONE_MRNA_CATABOLIC_PROCESS MIR6080, MIR6080 XPO1_TARGET_GENES, XPO1_TARGET_GENES HP_MOOD_SWINGS, HP_MOOD_SWINGS GOBP_POSITIVE_REGULATION_OF_NEURON_MIGRATION, GOBP_POSITIVE_REGULATION_OF_NEURON_MIGRATION GOBP_AXONAL_TRANSPORT_OF_MITOCHONDRION, GOBP_AXONAL_TRANSPORT_OF_MITOCHONDRION chr11q22, chr11q22 GOBP_RESOLUTION_OF_MEIOTIC_RECOMBINATION_INTERMEDIATES, GOBP_RESOLUTION_OF_MEIOTIC_RECOMBINATION_INTERMEDIATES GOCC CD40 RECEPTOR COMPLEX, GOCC CD40 RECEPTOR COMPLEX GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_14DY_DN, GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_14DY_DN GOBP_HISTONE_H3_K9_TRIMETHYLATION, GOBP_HISTONE_H3_K9_TRIMETHYLATION

GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS

GOMF_PROTEIN_TYROSINE_KINASE_ACTIVATOR_ACTIVITY, GOMF_PROTEIN_TYROSINE_KINASE_ACTIVATOR_ACTIVITY

HAHTOLA_CTCL_PATHOGENESIS, HAHTOLA_CTCL_PATHOGENESIS

THAKAR PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_UP

GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_DN, GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_DN

NZA_AGE_21_30YO_NONRESPONDER_28DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_28DY_UP