

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

DING_LUNG_CANCER_EXPRESSION_BY_COPY_NUMBER, DING_LUNG_CANCER_EXPRESSION_BY_COPY_NUMBER

TAAITGTG_MIR323, TAAITGTG_MIR323

THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_NONRESPONDER_2DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_NONRESPONDER_2DY_UP

CSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_5TIM_DC_3H_UP, CSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_5TIM_DC_3H_UP

GSE11961_MEMORY_BCELL_DAY7_VS_GERMINAL_CENTER_BCELL_DAY40_UP, GSE11961_MEMORY_BCELL_DAY7_VS_GERMINAL_CENTER_BCELL_DAY40_UP

GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_DN

MIR183_3P, MIR183_3P

HAY_BONE_MARROW_FOLLICULAR_B_CELL, HAY_BONE_MARROW_FOLLICULAR_B_CELL

MIR22_3P, MIR22_3P

GSE40274_CTRL_VS_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN

MIR148A_5P, MIR148A_5P

TRAVAGLINI_LUNG_B_CELL, TRAVAGLINI_LUNG_B_CELL

GSE14415_INDUCED_VS_NATURAL_TREG_UP, GSE14415_INDUCED_VS_NATURAL_TREG_UP

GCAAAAA_MIR129, GCAAAAA_MIR129

MIR5009_3P, MIR5009_3P

MIR20A_3P, MIR20A_3P

MIR298, MIR298

MIR421, MIR421

GOBP_MACROMOLECULE_DEACYLATION, GOBP_MACROMOLECULE_DEACYLATION

MIR4511, MIR4511

MIR6764_3P, MIR6764_3P

MIR6824_3P, MIR6824_3P

GSE360_L_DONOVANI_VS_M_TUBERCULOSIS_DC_DN, GSE360_L_DONOVANI_VS_M_TUBERCULOSIS_DC_DN

GSE3920_UNTREATED_VS_IFNA_TREATED_ENDOTHELIAL_CELL_DN, GSE3920_UNTREATED_VS_IFNA_TREATED_ENDOTHELIAL_CELL_DN

MODULE_358, MODULE_358

MIR3156_3P, MIR3156_3P

MIR4426, MIR4426

MIR3145_5P, MIR3145_5P

ACAGGGT_MIR10A_MIR10B, ACAGGGT_MIR10A_MIR10B

GSE22886_CD4_TCELL_VS_BCCELL_NAIVE_DN, GSE22886_CD4_TCELL_VS_BCCELL_NAIVE_DN

MIR8962, MIR8962

MIR3611, MIR3611

MIR154_3P, MIR154_3P

MIR320A_5P, MIR320A_5P

MIR6792_5P, MIR6792_5P

MIR103A_2_5P, MIR103A_2_5P

GSE22886_TCELL_VS_BCCELL_NAIVE_DN, GSE22886_TCELL_VS_BCCELL_NAIVE_DN

MIR186_3P, MIR186_3P

MIR5571_5P, MIR5571_5P

GTGCCAAA_MIR507, GTGCCAAA_MIR507

ATTACAT_MIR3803P, ATTACAT_MIR3803P

MIR3977, MIR3977

PR_02, PR_02

MIR671_5P, MIR671_5P

MIR649, MIR649

MIR5187_3P, MIR5187_3P

AAAGGAT_MIR501, AAAGGAT_MIR501

REACTOME_ACTIVATION_OF_GENE_EXPRESSION_BY_SREBF_SREBP, REACTOME_ACTIVATION_OF_GENE_EXPRESSION_BY_SREBF_SREBP

HP_SHORT FOOT, HP_SHORT FOOT

MIR491_3P, MIR491_3P

MIR3922_5P, MIR3922_5P

PID_RACL_PATHWAY, PID_RACL_PATHWAY

MIR181A_2_3P, MIR181A_2_3P

REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION, REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION

MIR3167, MIR3167

GOBP_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY

MIR6848_3P, MIR6848_3P

AAACCAC_MIR140, AAACCAC_MIR140

GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_DN, GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_DN

MIR6843_3P, MIR6843_3P

MIR4502, MIR4502

MIR6874_3P, MIR6874_3P

MIR3660, MIR3660

MIR6772_3P, MIR6772_3P

MIR4526, MIR4526

MIR197_5P, MIR197_5P

MIR5007_3P, MIR5007_3P

GOBP_MULTICELLULAR_ORGANISM_GROWTH, GOBP_MULTICELLULAR_ORGANISM_GROWTH

WP_APOPTOSISRELATED_NETWORK_DUE_TO_ALTERED_NOTCH3_IN_OVARIAN_CANCER, WP_APOPTOSISRELATED_NETWORK_DUE_TO_ALTERED_NOTCH3_IN_OVARIAN_CANCER

GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS

WP_FAS_LIGAND_FASL_PATHWAY_AND_STRESS_INDUCTION_OF_HEAT_SHOCK_PROTEINS_HSP_REGULATION, WP_FAS_LIGAND_FASL_PATHWAY_AND_STRESS_INDUCTION_OF_HEAT_SHOCK_PROTEINS_HSP_REGULATION

GOBP_POSITIVE_REGULATION_OF_ATPASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_ATPASE_ACTIVITY

HP_ABNORMALITY_OF_THE_TARSAL_BONES, HP_ABNORMALITY_OF_THE_TARSAL_BONES

GOCC_AXON_CYTOPLASM, GOCC_AXON_CYTOPLASM

MIR4633_5P, MIR4633_5P

HOLLEMAN_VINCRISTINE_RESISTANCE_B_ALL_UP, HOLLEMAN_VINCRISTINE_RESISTANCE_B_ALL_UP

MIR4300, MIR4300

MIR6740_5P, MIR6740_5P

TAGGTCTA_MIR192_MIR215, TAGGTCTA_MIR192_MIR215

PR_01, PR_01

MIR7151_5P, MIR7151_5P

GOMF_HYDROLASE_ACTIVITY_ACTING_ON CARBON_NITROGEN BUT NOT PEPTIDE BONDS IN LINEAR AMIDES, GOMF_HYDROLASE_ACTIVITY_ACTING_ON CARBON_NITROGEN BUT NOT PEPTIDE BONDS IN LINEAR AMIDES

PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_UP, PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_UP

MIR7703, MIR7703

GOMF_ATPASE_REGULATOR_ACTIVITY, GOMF_ATPASE_REGULATOR_ACTIVITY

MIR4999_3P, MIR4999_3P

BIOCARTA_FAS_PATHWAY, BIOCARTA_FAS_PATHWAY

MIR1258, MIR1258

MIR4777_3P, MIR4777_3P

GOBP_POSITIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY

MIR4662A_3P, MIR4662A_3P

GOBP_NEURON_MIGRATION, GOBP_NEURON_MIGRATION

PID_TNF_PATHWAY, PID_TNF_PATHWAY

LIE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN, LIE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN

GOBP_PLACENTA_DEVELOPMENT, GOBP_PLACENTA_DEVELOPMENT

WP_ENDOCHONDRA ossification, WP_ENDOCHONDRA ossification

WP_ENDOCHONDRA ossification WITH SKELETAL_DYSPLASIAS, WP_ENDOCHONDRA ossification WITH SKELETAL_DYSPLASIAS

CRX_NRL_DN.V1.DN, CRX_NRL_DN.V1.DN

EGR3_01, EGR3_01

ZHAN_MULTIPLE_MYELOMA_HP_DN, ZHAN_MULTIPLE_MYELOMA_HP_DN

MIR6780B_3P, MIR6780B_3P

MIR920, MIR920

GENTILE_UV_RESPONSE_CLUSTER_D6, GENTILE_UV_RESPONSE_CLUSTER_D6

HP_AMBLYOPIA, HP_AMBLYOPIA

HP_ABNORMAL_MORPHOLOGY_OF_THE_RADIUS, HP_ABNORMAL_MORPHOLOGY_OF_THE_RADIUS

MIR1202, MIR1202

MIR4654, MIR4654

MIR135A_3P, MIR135A_3P

HP_PULMONARY_HEMORRHAGE, HP_PULMONARY_HEMORRHAGE

MIR1193, MIR1193

MIR575, MIR575

XPOL_TARGET_GENES, XPOL_TARGET_GENES

HP_ABNORMALITY_OF_CIRCULATING_FIBRINOGEN, HP_ABNORMALITY_OF_CIRCULATING_FIBRINOGEN

GOMF_BHLH_TRANSCRIPTION_FACTOR_BINDING, GOMF_BHLH_TRANSCRIPTION_FACTOR_BINDING

GOMF_5_3_EXONUCLEASE_ACTIVITY, GOMF_5_3_EXONUCLEASE_ACTIVITY

HP_METRORRHAGIA, HP_METRORRHAGIA

WP_SREBF_AND_MIR33_IN_CHOLESTEROL_AND_LIPID_HOMEOSTASIS, WP_SREBF_AND_MIR33_IN_CHOLESTEROL_AND_LIPID_HOMEOSTASIS

GOMF_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY, GOMF_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY

MIR376B_5P, MIR376C_5P, MIR376B_5P, MIR376C_5P

GOMF_PROTEIN_DEACETYLASE_ACTIVITY, GOMF_PROTEIN_DEACETYLASE_ACTIVITY

PSIP1_TARGET_GENES, PSIP1_TARGET_GENES

MIR8621_3P, MIR8621_3P

BIOCARTA_TOLL_PATHWAY, BIOCARTA_TOLL_PATHWAY

BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP

GOBP_FACE_DEVELOPMENT, GOBP_FACE_DEVELOPMENT

GOBP_HISTONE_MRNA_CATABOLIC_PROCESS, GOBP_HISTONE_MRNA_CATABOLIC_PROCESS

HP_SYNOSTOSIS_OF_CARPALS_TARSALS, HP_SYNOSTOSIS_OF_CARPALS_TARSALS

MIR6514_5P, MIR6514_5P

GOBP_SIGNAL_COMPLEX_ASSEMBLY, GOBP_SIGNAL_COMPLEX_ASSEMBLY

HP_ABNORMAL_SUBCUTANEOUS FAT TISSUE DISTRIBUTION, HP_ABNORMAL_SUBCUTANEOUS FAT TISSUE DISTRIBUTION

HP_CHRONIC_INFECTIO, HP_CHRONIC_INFECTIO

HP_NON MIDLINE_CLEFT_LIP, HP_NON MIDLINE_CLEFT_LIP

GOMF_ATPASE_ACTIVATOR_ACTIVITY, GOMF_ATPASE_ACTIVATOR_ACTIVITY

GOMF_NADPLUS_BINDING, GOMF_NADPLUS_BINDING

GOBP_POSITIVE_REGULATION_OF_DEVELOPMENTAL_GROWTH, GOBP_POSITIVE_REGULATION_OF_DEVELOPMENTAL_GROWTH

MCCLUNG_DELTA_FOSB_TARGETS_2WK, MCCLUNG_DELTA_FOSB_TARGETS_2WK

MIR591, MIR591

GOBP_REGULATION_OF_NEURON MIGRATION, GOBP_REGULATION_OF_NEURON MIGRATION

GOMF_PROTEIN_KINASE_A REGULATORY SUBUNIT_BINDING, GOMF_PROTEIN_KINASE_A REGULATORY SUBUNIT_BINDING

GOBP_LIPOPROTEIN_CATABOLIC_PROCESS, GOBP_LIPOPROTEIN_CATABOLIC_PROCESS

GOBP_HISTONE_H3_DEACETYLATION, GOBP_HISTONE_H3_DEACETYLATION

HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_LOWER LIMBS, HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_LOWER LIMBS

GOBP_REGULATION_OF_MYELOID_CELL_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_MYELOID_CELL_APOPTOTIC_PROCESS

HP_ORAL_CAVITY_BLEEDING, HP_ORAL_CAVITY_BLEEDING

MIR6080, MIR6080

GOBP_ESTABLISHMENT_OF_SISTER_CHROMATID_COHESION, GOBP_ESTABLISHMENT_OF_SISTER_CHROMATID_COHESION

GOBP_COHESIN_LOADING, GOBP_COHESIN_LOADING

GOBP_MACROMOLECULE_DEPALMITOYLATION, GOBP_MACROMOLE