

RNA_AGE_21_30YO_NONRESPONDER_28DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_28DY_UP

THAKAR_PBMCI_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_UP, THAKAR_PBMCI_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDI
TAATGTG_MIR323, TAATGTG_MIR323
DING_LUNG_CANCER_EXPRESSION_BY_COPY_NUMBER, DING_LUNG_CANCER_EXPRESSION_BY_COPY_NUMBER
GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_DN, GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_DN
GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP, GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_
MIR4511, MIR4511
PGF_UP.V1_UP, PGF_UP.V1_UP
MIR298, MIR298
MIR520E_5P, MIR520E_5P
MIR6764_3P, MIR6764_3P
MIR6824_3P, MIR6824_3P
GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN, GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN
HAY_BONE_MARROW_FOLLICULAR_B_CELL, HAY_BONE_MARROW_FOLLICULAR_B_CELL
MIR183_3P, MIR183_3P
GSE20366_CD103_POS_VS_NEG_TREG_KLRG1NEG_DN, GSE20366_CD103_POS_VS_NEG_TREG_KLRG1NEG_DN
MIR491_3P, MIR491_3P
MIR3922_5P, MIR3922_5P
MIR3611, MIR3611
EPPERT_HSC_R, EPPERT_HSC_R
MIR508_5P, MIR508_5P
MIR4452, MIR4452
MIR4426, MIR4426
MIR3145_5P, MIR3145_5P
MIR3977, MIR3977
GSE22886_TCELL_VS_BCELL_NAIVE_DN, GSE22886_TCELL_VS_BCELL_NAIVE_DN
MIR3660, MIR3660
MIR4526, MIR4526
HP_SHORT_FOOT, HP_SHORT_FOOT
ACAGGGT_MIR10A_MIR10B, ACAGGGT_MIR10A_MIR10B
MIR181A_2_3P, MIR181A_2_3P
MIR3166, MIR3166
GTGCAAA_MIR507, GTGCAAA_MIR507
EGR3_0l, EGR3_0l
MIR487A_3P, MIR487A_3P
MIR154_3P, MIR154_3P
MIR5187_3P, MIR5187_3P
MIR326, MIR326
HP_ABNORMALITY_OF_THE_TARSAL_BONES, HP_ABNORMALITY_OF_THE_TARSAL_BONES
MIR1324, MIR1324
GOBP_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY
MIR4676_5P, MIR4676_5P
ZHAN_MULTIPLE_MYELOMA_PR_DN, ZHAN_MULTIPLE_MYELOMA_PR_DN
MIR3167, MIR3167
MIR1912_5P, MIR1912_5P
MIR4761_3P, MIR4761_3P
MIR1258, MIR1258
MIR4790_3P, MIR4790_3P
GOBP_REGULATION_OF_DENDRITIC_SPINE_DEVELOPMENT, GOBP_REGULATION_OF_DENDRITIC_SPINE_DEVELOPMENT
REACTOME_APOPTOTIC_EXECUTION_PHASE, REACTOME_APOPTOTIC_EXECUTION_PHASE
TAGGTCA_MIR192_MIR215, TAGGTCA_MIR192_MIR215
MIR4733_3P, MIR4733_3P
GOBP_NEURON_MIGRATION, GOBP_NEURON_MIGRATION
LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN, LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN
MIR1193, MIR1193
CTGYNNCTYTAA_UNKNOWN, CTGYNNCTYTAA_UNKNOWN
GOBP_PIGMENT_CELL_DIFFERENTIATION, GOBP_PIGMENT_CELL_DIFFERENTIATION
WP_ENDOCHONDRAI_OSSIFICATION, WP_ENDOCHONDRAI_OSSIFICATION
WP_ENDOCHONDRAI_OSSIFICATION_WITH_SKELETAL_DYSPLASIAS, WP_ENDOCHONDRAI_OSSIFICATION_WITH_SKELETAL_DYSPLASIAS
HP_ABNORMAL_PERISTALSIS, HP_ABNORMAL_PERISTALSIS
HP_ORAL_CAVITY_BLEEDING, HP_ORAL_CAVITY_BLEEDING
MIR920, MIR920
GOBP_NEGATIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_PLASMA_MEMBRANE.
MIR4654, MIR4654
GOBP_XENOBIOTIC_TRANSPORT, GOBP_XENOBIOTIC_TRANSPORT
GOBP_SMOOTH_MUSCLE_CELL_DIFFERENTIATION, GOBP_SMOOTH_MUSCLE_CELL_DIFFERENTIATION
GOCC_GOLGI_TRANS_CISTERNA, GOCC_GOLGI_TRANS_CISTERNA
HP_DIFFUSE_ALVEOLAR_HEMORRHAGE, HP_DIFFUSE_ALVEOLAR_HEMORRHAGE
HP_ACHALASIA, HP_ACHALASIA
GAUCHER_PBMCI_YF_VAX_STAMARIL_UNKNOWN_AGE_14DY_DN, GAUCHER_PBMCI_YF_VAX_STAMARIL_UNKNOWN_AGE_14DY_DN
HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_LOWER_LIMBS, HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_LOWER_LIMBS
HP_UROGENITAL_SINUS_ANOMALY, HP_UROGENITAL_SINUS_ANOMALY
GOBP_NEGATIVE_REGULATION_OF_TISSUE_REMODELING, GOBP_NEGATIVE_REGULATION_OF_TISSUE_REMODELING
chr2q24, chr2q24
HP_DIGENIC_INHERITANCE, HP_DIGENIC_INHERITANCE
GOBP_NEGATIVE_REGULATION_OF_PROSTAGLANDIN_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROSTAGLANDIN_BIOSYNTHETIC_PROCES
chr9q31, chr9q31
KEGG_NITROGEN_METABOLISM, KEGG_NITROGEN_METABOLISM