GOM_R4301,GOM_R4310
GOM_M5352,GOM_M5352
GOM_C5371,GOM_C5371
GOM_CHON_GOM_CHON
GOM_KAT2,GOM_KAT2
GOM_CTL_VS_QUATERNARY_MEMORY_CD8_TCELL_DN,GSE23280_PRIMARY_VS_QUATERNARY_MEMORY_CD8_TCELL_DN
GSE4367_DAY8_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_UP,GSE4367_DAY8_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_UP
GOM_DF5A_GOM_DF5A
GSE4634_ANTI_IGM_VS_CPG_STIM_BCELL_UP,GSE4634_ANTI_IGM_VS_CPG_STIM_BCELL_UP
GCM_MAP4K4_GCM_MAP4K4
GSE3720_UNSTIM_VS_HELICOBACTER_PYLORI_IFN_STIM_AGS_CELL_DN,GSE3720_UNSTIM_VS_HELICOBACTER_PYLORI_IFN_STIM_AGS_CELL_DN
GOM_ZNF599_GOM_ZNF599
GSE4634_ANTI_IGM_VS_IFN_CD8_TCELL_UP,KAICH_NAIVE_VS_DAY8_IFF_CD8_TCELL_UP
MH0174_UP,MH0174_UP
GOM_BB9MA_GOM_BB9MA
GOM_HH91_GOM_HH91
MH0150_UP,MH0150_UP
GCM_SUH2_GCM_SUH2
GOM_NCAM2_GOM_NCAM2
GSE4367_DAY8_VS_DAY15_CD8_TCELL_UP,KAICH_NAIVE_VS_MEMORY_CD8_TCELL_UP
CSTCTEC_MH57,CSTCTEC_MH57
CAGCCTC_MH0059F,CAGCCTC_MH0059F
GCM_CHEK2_GCM_CHEK2
GSE4748_CTRL_VS_LPS_STIM_DC_3H_UP,GSE4748_CTRL_VS_LPS_STIM_DC_3H_UP
GCM_CTH3K_GCM_CTH3K
MH0170_UP,MH0170_UP
GOM_CTSH_GOM_CTSH_MH92
GCM_TEC2_GCM_TEC2
GOCC_PROTEIN_ACTIVATEDTRANSFERASE_COMPLEX,GOCC_PROTEIN_ACTIVTRANSFERASE_COMPLEX
MORF_BAX19A_MORF_BAX19A
GAGCCAC_MH0149_GAGCCAC_MH0149
GOM_IN2CI_GOM_IN2CI
ATGTTTC_MH0054_ATGTTTC_MH0054
GOM_CTSH_GOM_CTSH_MH0149
GCM_HMMR_GCM_HMMR
MH0155_UP,MH0155_UP
GH023129_CUTBLA_VS_NOD_THYMOCYTE_DN,GH023129_CUTBLA_VS_NOD_THYMOCYTE_DN
MH022_UP,MH022_UP
GCM_SMG_GCM_SMG
GSE4748_CTRL_VS_LPS_STIM_MF_BRD4_LONG_BONES
GH01403_NATURAL_TREG_VS_FOXP3_KO_NATURAL_TREG_DN,GH01403_NATURAL_TREG_VS_FOXP3_KO_NATURAL_TREG_DN
SALE_AIP_WITH_FLT3_MUTATED_UP,SALE_AIP_WITH_FLT3_MUTATED_UP
MH022_UP,MH022_UP
GAGACTC_MH0183_GAGACTC_MH0183
HP_BROAD_FINGER_HP_BROAD_FINGER
GCM_NUMA1_GCM_NUMA1
GOM_TPMT_GOM_TPMT
GOM_TPMT_GOM_TPMT
MH0491_UP,MH0491_UP
GOM_BAG3_GOM_BAG3
MH0594_UP,MH0594_UP
MORF_CNTN1_MORF_CNTN1
MH13A_UP,MH13A_UP
GH023129_CUTBLA_VS_NOD_THYMOCYTE_DP_OCCIPUT_HP_ABNORMAL_SHAPE_OF_THE_OCCIPUT
CTCTGAA_MIR486_UCTCTGA_MIR486
FIGUEROA_AML_METHYLATION_CLUSTER_7_UP,FIGUEROA_AML_METHYLATION_CLUSTER_7_UP
MORF_PDPK1_MORF_PDPK1
MH0497_UP,MH0497_UP
AYAWAGT_MH83_AYAWAGT_MH83
MH0479_UP,MH0479_UP
MH0496_UP,MH0496_UP
MH0470_UP,MH0470_UP
MH0480_MH0480
GH01403_UNSTIM_VS_4HRS_LPS_IMDM_UP,GH01403_UNSTIM_VS_4HRS_LPS_IMDM_UP
AGCGACT_MH0510F_AGCGACT_MH0510F
MH0475_MH0475
MH0432_UP,MH0432_UP
ATGTAGC_MH0251_MH0251 ATGTAGC_MH0251_MH0252
MH432_UP,MH432_UP
MH0495_MH0495
MH0545_MH0545
HP_HYPOTHYLOTARDISH_HP_HYPOTHYLOTARDISH
MH0144_UP,MH0144_UP
MH0464A3_UP,MH0464A3_UP
MH0479_UP,MH0479_UP
GCM_MAA_GCM_MAA
MH0476_UP,MH0476_UP
MH0354_MH0354_MH0355_MH0355B
HP_BROAD_THUMB_HP_BROAD_THUMB
MH0139_MH0139
MH0559_UP,MH0559_UP
MH0439_UP,MH0439_UP
GCM_SCD3_GCM_SCD3
MH0479_UP,MH0479_UP
GOME_CAST_MORF_CAST5
GH0946_LISTERIA_INF_MATURE_VS_PROTAGLANDIN2_TREATED_MATURE_DC_UP,GH0946_LISTERIA_INF_MATURE_VS_PROTAGLANDIN2_TREATED_MATURE_DC_UP
NAGAWHWA_NRG1_SIGNALING_UP,NAGAWHWA_NRG1_SIGNALING_UP
GCM_CASP3_GCM_CASP3
GOM_FAM62A_GOM_FAM62A
MORF_EAD2_MORF_EAD2
MH0477_UP,MH0477_UP
MH0465A_MH0465A
MH0378G_MH0378G
MH0476_UP,MH0476_UP
CACCTTT_MH0302A_CACCTTT_MH0302A
MH0127_UP,MH0127_UP
MH0491_UP,MH0491_UP
MH0477_UP,MH0477_UP
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPID_MAGENTA_UP,GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPID_MAGENTA_UP
MH0398_MH0398
HP_SHRNA_SPLICE_SITE_SELECTION_GOMP_SHRNA_SPLICE_SITE_SELECTION
HP_PAPILLEDEMA_HP_PAPILLEDEMA
GOCC_MRNA_CLEAVAGE_FACTOR_COMPLEX,GOCC_MRNA_CLEAVAGE_FACTOR_COMPLEX
MH0466_MH0466
MH0498_MH0498
MH378A_UP_MH378C_MH0393_MH0394_UP_MH378C_MH0393_MH0393
MORF_JAG1_MORF_JAG1
CTGTCLA_MIR574_CTGTCLA_MIR574
HP_ABNORMAL_GROWTH_HORMONE_LEVEL_HP_ABNORMAL_GROWTH_HORMONE_LEVEL
MH0393_MH0393
MH0432A_MH0432A
MH0477_MH0477
HP_APLASIA_HYPOPLASIA_OF_THE_LUNGS_HP_APLASIA_HYPOPLASIA_OF_THE_LUNGS
MH0446C_UP_MH0446C_UP
MH0183_MH0183
GOMP_SNRNA_METABOLIC_PROCESS_GOMP_SNRNA_METABOLIC_PROCESS
GH0946_IMMATURE_VS_PROSTAGLANDIN2_TREATED_MATURE_DC_UP,GH0946_IMMATURE_VS_PROSTAGLANDIN2_TREATED_MATURE_DC_UP
CTAGGAA_MIR96_CTAGGAA_MIR96
GSE4748_CTRL_VS_LPS_STIM_MF_DC_1H_DN,GSE4591_CTRL_VS_NFACSATLS_VIRUS_DC_1H_DN
MF_VIRAL_ACUTE_MYOCARDITIS_MF_VIRAL_ACUTE_MYOCARDITIS
GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION,GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
GH03705_UNPHOSPHO_PEMSD_MPP_VS_CDK_TCELL_UP,GH03705_UNPHOSPHO_PEMSD_MPP_VS_CDK_TCELL_UP
HP_RECEPTIVE_LANGUAGE_DELAY_HP_RECEPTIVE_LANGUAGE_DELAY
AACCTCA_MIR435B_MIR435A_AACCTCA_MIR435B_MIR435A
MH0560_UP_MH0560_UP
GOBP_PROTEIN_K48_LINKED_DEUBIQUITINATION_GOBP_PROTEIN_K48_LINKED_DEUBIQUITINATION
MH0556_UP_MH0556_UP
HP_ADRENAL_INSUFFICIENCY_HP_ADRENAL_INSUFFICIENCY
MH96_UP_MH96_UP
GOBP_T_BODY_ASSEMBLY_GOBP_T_BODY_ASSEMBLY
MH0793_MH0793
CCATCCA_MIR432_CCATCCA_MIR432
HP_PRIMARY_ADRENAL_INSUFFICIENCY_HP_PRIMARY_ADRENAL_INSUFFICIENCY
MH0795_MH0795
GOME_LYSINE_ACETYLATED_HISTONE_BINDING_GOME_LYSINE_ACETYLATED_HISTONE_BINDING
HP_NEOPLASM_OF_THE_ORAL_CAVITY_HP_NEOPLASM_OF_THE_ORAL_CAVITY
MH03_UP_MH03_UP
GOMP_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER_GOMP_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
REACTOME_DOWNREGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING_REACTOME_DOWNREGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING
MH0794_MH0794
HP_PERIVITELLINA_HP_PERIVITELLINA
HP_PANCREATIC_CYSTS_HP_PANCREATIC_CYSTS
MH0499_UP_MH0499_UP
HP_PROMINENT_OCCIPUT_HP_PROMINENT_OCCIPUT
GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION_GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
BECKATA_CASPASE_PATHWAY_BECKATA_CASPASE_PATHWAY
GOBP_REGULATION_OF_CELL_CYCLE_ARREST_GOBP_REGULATION_OF_CELL_CYCLE_ARREST
CTCTAGA_MIR330C_MIR330F_MIR330A_CTCTAGA_MIR330C_MIR330F_MIR330A
GOBP_SHRNA_PROCESSING_GOBP_SHRNA_PROCESSING
HP_NEOPLASM_OF_THE_TONGUE_HP_NEOPLASM_OF_THE_TONGUE
MH021_UP_MH021_UP
HP_NEOPLASM_OF_THE_PITUITARY_GLAND_HP_NEOPLASM_OF_THE_PITUITARY_GLAND
MH0497_UP_MH0497_UP
GOBP_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER_GOBP_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
GOBP_REGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING_REACTOME_DOWNREGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING
MH0796_MH0796
HP_PERIVITELLINA_HP_PERIVITELLINA
HP_PANCREATIC_CYSTS_HP_PANCREATIC_CYSTS
MH0499_UP_MH0499_UP
HP_PROMINENT_OCCIPUT_HP_PROMINENT_OCCIPUT
GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION_GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
BECKATA_CASPASE_PATHWAY_BECKATA_CASPASE_PATHWAY
GOBP_REGULATION_OF_CELL_CYCLE_ARREST_GOBP_REGULATION_OF_CELL_CYCLE_ARREST
CTCTAGA_MIR330C_MIR330F_MIR330A_CTCTAGA_MIR330C_MIR330F_MIR330A
GOBP_SHRNA_PROCESSING_GOBP_SHRNA_PROCESSING
HP_NEOPLASM_OF_THE_TONGUE_HP_NEOPLASM_OF_THE_TONGUE
MH021_UP_MH021_UP
HP_NEOPLASM_OF_THE_PITUITARY_GLAND_HP_NEOPLASM_OF_THE_PITUITARY_GLAND
MORF_LTK_MORF_LTK
GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA
GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA
HP_ATROPHY_DEGENERATION_AFFECTING_THE_BRAINSTEM_HP_ATROPHY_DEGENERATION_AFFECTING_THE_BRAINSTEM
GOBP_REGULATION_OF_BMP_SIGNALING_PATHWAY_GOBP_REGULATION_OF_BMP_SIGNALING_PATHWAY
HP_SPINAL_CORD_LESION_HP_SPINAL_CORD_LESION
GOMF_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_PHOSPHATASE_ACTIVITY_GOMF_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_PHOSPHATASE_ACTIVITY
GOBP_PHOSPHOLIPID_DEPHOSPHORYLATION_GOBP_PHOSPHOLIPID_DEPHOSPHORYLATION
FLECHNER_BKPV1_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN_FLECHNER_BKPV1_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN
MH0422_UP_MH0422_UP
MH0498_MH0498
ATGCAGC_MIR507B_ATGCAGC_MIR507B
HP_CAVUM_SEPTUM_PELLUCIDUM_HP_CAVUM_SEPTUM_PELLUCIDUM
GOCC_CATALYTIC_STEP_1_SPLICEOSOME_GOCC_CATALYTIC_STEP_1_SPLICEOSOME
HOXD3_TARGET_GENES_HOXD3_TARGET_GENES
MH0496_MH0496_MH0496_MH0496
GAGCTCC_ELF2_SIGNALING_UP_MAGIC_ELF2_SIGNALING_UP
GOCC_ATAC_TYPE_HISTONE_ACETYLTRANSFERASE_COMPLEX_GOCC_ATAC_TYPE_HISTONE_ACETYLTRANSFERASE_COMPLEX
MH0511_UP_MH0511_UP
MH0971_UP_MH0971_UP
MH0446_MH0446
GOCC_CHROMATIN_SILENCING_COMPLEX_GOCC_CHROMATIN_SILENCING_COMPLEX
MH0498_UP_MH0498_UP
MH0446_UP_MH0446_UP
HP_CUTANEOUS_FINGER_SYNDACTYLY_HP_CUTANEOUS_FINGER_SYNDACTYLY
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPID_GREEN_DN_GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPID_GREEN_DN
GOCC_TRAPP_COMPLEX_GOCC_TRAPP_COMPLEX
GOMP_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS_GOMP_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS
MH0498_UP_MH0498_UP
HP_LARGE_HYPERTROPHY_HP_LARGE_HYPERTROPHY
MH0963_MH0963
GOMP_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS_GOBP_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS
MH0498_UP_MH0498_UP
HP_LARGE_HYPERTROPHY_HP_LARGE_HYPERTROPHY
MH0963_MH0963
GOMF_CALCIUM_ION_CONTACTING_PHOSPHOLIPID_CONCENTRATION_HP_INCREASED_CIRCULATING_PROCALCITN_CONCENTRATION
HP_ABNORMAL_THALAMIC_MRI_SIGNAL_INTENSITY_HP_ABNORMAL_THALAMIC_MRI_SIGNAL_INTENSITY
MH0_BB9MA_MORF_BB9MA
MH0644_UP_MH0644_UP
HOFF_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_RCC_VACCINE_AGE_36_49YO_IND_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION_UP,HOFF_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_RCC_VACCINE_AGE_36_49YO_IND_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION_UP
MYELOMA_MAF_UP_MAF_UP
GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_GOMF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY
GOBP_REGULATION_OF_VESICLE_FUSION_GOBP_REGULATION_OF_VESICLE_FUSION
GOMF_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_PHOSPHATASE_ACTIVITY_GOMF_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_PHOSPHATASE_ACTIVITY
GOBP_PHOSPHOLIPID_DEPHOSPHORYLATION_GOBP_PHOSPHOLIPID_DEPHOSPHORYLATION
FLECHNER_BKPV1_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN_FLECHNER_BKPV1_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN
MH0422_UP_MH0422_UP
MH0498_MH0498
ATGCAGC_MIR507B_ATGCAGC_MIR507B
HP_CAVUM_SEPTUM_PELLUCIDUM_HP_CAVUM_SEPTUM_PELLUCIDUM
GOCC_CATALYTIC_STEP_1_SPLICEOSOME_GOCC_CATALYTIC_STEP_1_SPLICEOSOME
HOXD3_TARGET_GENES_HOXD3_TARGET_GENES
MH0496_MH0496_MH0496_MH0496
GAGCTCC_ELF2_SIGNALING_UP_MAGIC_ELF2_SIGNALING_UP
GOCC_CHROMATIN_SILENCING_COMPLEX_GOCC_CHROMATIN_SILENCING_COMPLEX
MH0498_UP_MH0498_UP
MH0446_UP_MH0446_UP
HP_CUTANEOUS_FINGER_SYNDACTYLY_HP_CUTANEOUS_FINGER_SYNDACTYLY
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPID_GREEN_DN_GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPID_GREEN_DN
GOCC_TRAPP_COMPLEX_GOCC_TRAPP_COMPLEX