MIR934, MIR934 Y7_POST_TIV_FLU_VACCINE_PDC_DN, GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_PDC_DN

MIR522 3P, MIR522 3P

GOMF_CYCLASE_REGULATOR_ACTIVITY, GOMF_CYCLASE_REGULATOR_ACTIVITY

WP_DOPAMINERGIC_NEUROGENESIS, WP_DOPAMINERGIC_NEUROGENESIS

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MIR224_3P, MIR224_3P
 LINDGREN_BLADDER_CANCER_CLUSTER_2A_DN, LINDGREN_BLADDER_CANCER_CLUSTER_2A_DN
 BIDUS_METASTASIS_DN, BIDUS_METASTASIS_DN
 REACTOME_PROTEIN_UBIQUITINATION, REACTOME_PROTEIN_UBIQUITINATION
 ZHANG_BREAST_CANCER_PROGENITORS_DN, ZHANG_BREAST_CANCER_PROGENITORS_DN
 GSE4748_CTRL_VS_LPS_STIM_DC_3H_DN, GSE4748_CTRL_VS_LPS_STIM_DC_3H_DN
 REACTOME_PKMTS_METHYLATE_HISTONE_LYSINES, REACTOME_PKMTS_METHYLATE_HISTONE_LYSINES
 GAGCCTG_MIR484, GAGCCTG_MIR484
  REACTOME_NONHOMOLOGOUS_END_JOINING_NHEJ, REACTOME_NONHOMOLOGOUS_END_JOINING_NHEJ
  MIR374A_3P, MIR374A_3P
  REACTOME_DNA_DOUBLE_STRAND_BREAK_RESPONSE, REACTOME_DNA_DOUBLE_STRAND_BREAK_RESPONSE
  CCAWNWWNNNGGC_UNKNOWN, CCAWNWWNNNGGC_UNKNOWN
  MIR1237_3P, MIR1237_3P
  HP_CARIOUS_TEETH, HP_CARIOUS_TEETH
  GSE3982_MAC_VS_NEUTROPHIL_LPS_STIM_DN, GSE3982_MAC_VS_NEUTROPHIL_LPS_STIM_DN
  GOMF_KINASE_INHIBITOR_ACTIVITY, GOMF_KINASE_INHIBITOR_ACTIVITY
  REACTOME_RHOQ_GTPASE_CYCLE, REACTOME_RHOQ_GTPASE_CYCLE
  MODULE_493, MODULE_493
  AGCYRWTTC_UNKNOWN, AGCYRWTTC_UNKNOWN
  MIR5009_5P, MIR5009_5P
  HP MASK LIKE FACIES, HP MASK LIKE FACIES
  GOMF_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING, GOMF_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING
  REACTOME_RND2_GTPASE_CYCLE, REACTOME_RND2_GTPASE_CYCLE
  HP_UROGENITAL_FISTULA, HP_UROGENITAL_FISTULA
  GOBP_ACTION_POTENTIAL, GOBP_ACTION_POTENTIAL
 MAYBURD_RESPONSE_TO_L663536_UP, MAYBURD_RESPONSE_TO_L663536_UP
 GOBP_NEGATIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GOBP_NEGATIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING
MORF_TNFRSF6, MORF_TNFRSF6
 HP_NARROW_PALPEBRAL_FISSURE, HP_NARROW_PALPEBRAL_FISSURE
 HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY, HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY
 HP_DECREASED_ADIPOSE_TISSUE, HP_DECREASED_ADIPOSE_TISSUE
  MIR3157_3P, MIR3157_3P
  HP_INCREASED_NUMBER_OF_TEETH, HP_INCREASED_NUMBER_OF_TEETH
  HP_POOR_WOUND_HEALING, HP_POOR_WOUND_HEALING
  HSF4_TARGET_GENES, HSF4_TARGET_GENES
  GOCC_TRANSCRIPTION_FACTOR_TFIIH_HOLO_COMPLEX, GOCC_TRANSCRIPTION_FACTOR_TFIIH_HOLO_COMPLEX
  GOBP_REGULATION_OF_RYANODINE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY, GOBP_REGULATION_OF_RYANODINE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY
  GOBP_CELL_COMMUNICATION_BY_ELECTRICAL_COUPLING, GOBP_CELL_COMMUNICATION_BY_ELECTRICAL_COUPLING
  PID_UPA_UPAR_PATHWAY, PID_UPA_UPAR_PATHWAY
  GOCC_CALCIUM_CHANNEL_COMPLEX, GOCC_CALCIUM_CHANNEL_COMPLEX
  BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_24HR_DN, BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_24HR_DN
  MODULE_330, MODULE_330
  MODULE 96, MODULE 96
 GOBP_RENAL_FILTRATION, GOBP_RENAL_FILTRATION
  HP_SHAWL_SCROTUM, HP_SHAWL_SCROTUM
  GOMF_ADENYLATE_CYCLASE_BINDING, GOMF_ADENYLATE_CYCLASE_BINDING
 HP_VERTEBRAL_CLEFTING, HP_VERTEBRAL_CLEFTING
 GOBP_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION_BY_CALCIUM_ION_SIGNALING, GOBP_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION_BY_CALCIUM_ION_SIGNALING
 GOBP_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
 GOBP RENAL SYSTEM PROCESS INVOLVED IN REGULATION OF SYSTEMIC ARTERIAL BLOOD PRESSURE, GOBP RENAL SYSTEM PROCESS INVOLVED IN REGULATION OF SYSTEMIC A
 GOBP_RENAL_SYSTEM_PROCESS_INVOLVED_IN_REGULATION_OF_BLOOD_VOLUME, GOBP_RENAL_SYSTEM_PROCESS_INVOLVED_IN_REGULATION_OF_BLOOD_VOLUME
 GOBP_RELEASE_OF_SEQUESTERED_CALCIUM_ION_INTO_CYTOSOL_BY_ENDOPLASMIC_RETICULUM, GOBP_RELEASE_OF_SEQUESTERED_CALCIUM_ION_INTO_CYTOSOL_BY_ENDOPLASMIC
 HP_ABNORMAL_ALDOLASE_LEVEL, HP_ABNORMAL_ALDOLASE_LEVEL
 GOBP_REGULATION_OF_EXTRACELLULAR_MATRIX_DISASSEMBLY, GOBP_REGULATION_OF_EXTRACELLULAR_MATRIX_DISASSEMBLY
 GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION
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