
ROMEDINU_IN_T_CELL_LINE_6H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_DN
GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP, GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP
GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN
GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN
GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN
GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP, GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP
GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP
GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_DN, GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_DN
GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP, GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP
PECE_MAMMARY_STEM_CELL_DN, PECE_MAMMARY_STEM_CELL_DN
MIR12132, MIR12132
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP
MIR6505_5P, MIR6505_5P
GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN, GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN
MIR379_3P_MIR411_3P, MIR379_3P_MIR411_3P
REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENTIATION_AND_PLATELET_FUNCTION, REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENT
MIR1208, MIR1208
GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP
MIR548AI_MIR570_5P, MIR548AI_MIR570_5P
MIR548AO_5P_MIR548AX, MIR548AO_5P_MIR548AX
GOCC_CILIARY_BASAL_BODY, GOCC_CILIARY_BASAL_BODY
MIR558, MIR558
MIR4456, MIR4456
MIR5001_3P, MIR5001_3P
KEGG_AXON_GUIDANCE, KEGG_AXON_GUIDANCE
MIR3682_3P, MIR3682_3P
HP_RECURRENT_OTITIS_MEDIA, HP_RECURRENT_OTITIS_MEDIA
GOBP_EMBRYONIC_PLACENTA_DEVELOPMENT, GOBP_EMBRYONIC_PLACENTA_DEVELOPMENT
GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN
MIR548Q, MIR548Q
MIR331_5P, MIR331_5P
GSE40685_NAIVE_CD4_TCELL_VS_FOXP3_KO_TREG_PRECURSOR_DN, GSE40685_NAIVE_CD4_TCELL_VS_FOXP3_KO_TREG_PRECURSOR_DN
GOBP_LABYRINTHINE_LAYER_DEVELOPMENT, GOBP_LABYRINTHINE_LAYER_DEVELOPMENT
GOBP_POLYSACCHARIDE_METABOLIC_PROCESS, GOBP_POLYSACCHARIDE_METABOLIC_PROCESS
MIR1912_3P, MIR1912_3P
GOBP_NEPHRON_DEVELOPMENT, GOBP_NEPHRON_DEVELOPMENT
HP_FOCAL_SEGMENTAL_GLOMERULOSCLEROSIS, HP_FOCAL_SEGMENTAL_GLOMERULOSCLEROSIS
GOMF_EXTRACELLULAR_MATRIX_BINDING, GOMF_EXTRACELLULAR_MATRIX_BINDING
GOBP_ZYMOGEN_ACTIVATION, GOBP_ZYMOGEN_ACTIVATION
MIR769_5P, MIR769_5P
GOBP_RESPONSE_TO_OSMOTIC_STRESS, GOBP_RESPONSE_TO_OSMOTIC_STRESS
GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
MIR584_5P, MIR584_5P
HARALAMBIEVA_PBMC_M_M_R_II_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP, HARALAMBIEVA_PBMC_M_M_R_II_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_P
MIR1295B_5P, MIR1295B_5P
LAKE_ADULT_KIDNEY_C25_ENDOTHELIAL_CELLS_UNASSIGNED, LAKE_ADULT_KIDNEY_C25_ENDOTHELIAL_CELLS_UNASSIGNED
MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TACI_UP, MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TACI_UP
HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS, HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS
GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS
GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS, GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS
GOBP_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_CELL_ADHESION_MEDIATED_BY_INTEGRIN
GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS
GOBP_REGULATION_OF_GOLGI_ORGANIZATION, GOBP_REGULATION_OF_GOLGI_ORGANIZATION
GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
HP_ATYPICAL_ABSENCE_SEIZURE, HP_ATYPICAL_ABSENCE_SEIZURE
HP_ANKYLOSIS, HP_ANKYLOSIS
HASLINGER_B_CLL_WITH_MUTATED_VH_GENES, HASLINGER_B_CLL_WITH_MUTATED_VH_GENES
WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS, WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANIS
HP_SHORT_RIBS, HP_SHORT_RIBS
GOBP_CHROMATIN_SILENCING_AT_TELOMERE, GOBP_CHROMATIN_SILENCING_AT_TELOMERE
HP_REDUCED_SPERM_MOTILITY, HP_REDUCED_SPERM_MOTILITY
REACTOME_FORMATION_OF_APOPTOSOME, REACTOME_FORMATION_OF_APOPTOSOME
REACTOME_TRYPTOPHAN_CATABOLISM, REACTOME_TRYPTOPHAN_CATABOLISM
HINATA_NFKB_TARGETS_KERATINOCYTE_DN, HINATA_NFKB_TARGETS_KERATINOCYTE_DN
GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
GOBP_KETONE_CATABOLIC_PROCESS, GOBP_KETONE_CATABOLIC_PROCESS
REACTOME_ACTIVATION_OF_THE_AP_1_FAMILY_OF_TRANSCRIPTION_FACTORS, REACTOME_ACTIVATION_OF_THE_AP_1_FAMILY_OF_TRANSCRIPTION_FACTORS
GOCC_AXONAL_GROWTH_CONE, GOCC_AXONAL_GROWTH_CONE
GOBP_KYNURENINE_METABOLIC_PROCESS, GOBP_KYNURENINE_METABOLIC_PROCESS
WP_OSTEOPONTIN_SIGNALING, WP_OSTEOPONTIN_SIGNALING
GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN
TSENG_ADIPOGENIC_POTENTIAL_DN, TSENG_ADIPOGENIC_POTENTIAL_DN
REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS, REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS
WP_PREGNANE_X_RECEPTOR_PATHWAY, WP_PREGNANE_X_RECEPTOR_PATHWAY
WP_PRADERWILLI_AND_ANGELMAN_SYNDROME, WP_PRADERWILLI_AND_ANGELMAN_SYNDROME
MATZUK_EARLY_ANTRAL_FOLLICLE, MATZUK_EARLY_ANTRAL_FOLLICLE
PID_S1P_S1P1_PATHWAY, PID_S1P_S1P1_PATHWAY