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GSE17721_LPS_VS_GARDIQUIMOD_6H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_6H_BMDC_DN
 GSE17721_0.5H_VS_4H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_4H_POLYIC_BMDC_DN
GSE17721_0.5H_VS_24H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_24H_POLYIC_BMDC_DN
GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN
GSE43955_1H_VS_20H_ACT_CD4_TCELL_UP, GSE43955_1H_VS_20H_ACT_CD4_TCELL_UP
GSE3982 BASOPHIL_VS_TH2_UP, GSE3982 BASOPHIL_VS_TH2_UP
GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN
GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN
GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_UP, GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_UP
GSE17721_0.5H_VS_8H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_8H_POLYIC_BMDC_DN
GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_DN
GSE17721 0.5H VS 8H LPS BMDC DN, GSE17721 0.5H VS 8H LPS BMDC DN
GSE17721_CTRL_VS_LPS_0.5H_BMDC_UP, GSE17721_CTRL_VS_LPS_0.5H_BMDC_UP
SEIDEN_ONCOGENESIS_BY_MET, SEIDEN_ONCOGENESIS_BY_MET
MIR5699_5P, MIR5699_5P
GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_UP
MIR624_3P, MIR624_3P
GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_UP
GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DN, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DN
GSE26488 HDAC7 KO VS VP16 TRANSGENIC HDAC7 KO DOUBLE POSITIVE THYMOCYTE UP, GSE26488 HDAC7 KO VS VP16 TRANSGENIC HDAC7 KO DOUBLE POSITIVE THYMOCYTE UP
 GSE17721_CTRL_VS_LPS_4H_BMDC_DN, GSE17721_CTRL_VS_LPS_4H_BMDC_DN
 MIR8061, MIR8061
 MIR4677_3P, MIR4677_3P
 GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN
 WP_PROLACTIN_SIGNALING_PATHWAY, WP_PROLACTIN_SIGNALING_PATHWAY
 TCTATGA_MIR376A_MIR376B, TCTATGA_MIR376A_MIR376B
 MIR127_5P, MIR127_5P
 GOBP_LIPID_PHOSPHORYLATION, GOBP_LIPID_PHOSPHORYLATION
 MIR106A_3P, MIR106A_3P
 MIR1256, MIR1256
 GOBP_GASTRULATION, GOBP_GASTRULATION
 MIR3085 5P, MIR3085 5P
 HP_OLIGODONTIA, HP_OLIGODONTIA
 MIR6743_5P, MIR6743_5P
GSE360_L_DONOVANI_VS_T_GONDII_DC_UP, GSE360_L_DONOVANI_VS_T_GONDII_DC_UP
 GSE40274_IRF4_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_IRF4_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
SERVITJA_LIVER_HNF1A_TARGETS_DN, SERVITJA_LIVER_HNF1A_TARGETS_DN
MIR675 3P, MIR675 3P
HP_ABNORMALITY_OF_PROTHROMBIN, HP_ABNORMALITY_OF_PROTHROMBIN
REACTOME_MET_PROMOTES_CELL_MOTILITY, REACTOME_MET_PROMOTES_CELL_MOTILITY
GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_GLYCOPROTEIN, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_GLYCOPROTEIN
 TRAYNOR_RETT_SYNDROM_DN, TRAYNOR_RETT_SYNDROM_DN
 REACTOME_MET_ACTIVATES_PTK2_SIGNALING, REACTOME_MET_ACTIVATES_PTK2_SIGNALING
GOBP POSITIVE REGULATION OF CELL MIGRATION INVOLVED IN SPROUTING ANGIOGENESIS, GOBP POSITIVE REGULATION OF CELL MIGRATION INVOLVED IN SPROUTING ANGIOGENESIS
GOBP_SPHINGOSINE_BIOSYNTHETIC_PROCESS, GOBP_SPHINGOSINE_BIOSYNTHETIC_PROCESS
NKX2_8_TARGET_GENES, NKX2_8_TARGET_GENES
 PID_INTEGRIN_A9B1_PATHWAY, PID_INTEGRIN_A9B1_PATHWAY
GOBP_AXONAL_FASCICULATION, GOBP_AXONAL_FASCICULATION
HP_AXILLARY_PTERYGIUM, HP_AXILLARY_PTERYGIUM
 GAO_LARGE_INTESTINE_24W_C10_ENTEROCYTE, GAO_LARGE_INTESTINE_24W_C10_ENTEROCYTE
 RGTTAMWNATT_HNF1_01, RGTTAMWNATT_HNF1_01
 RUAN_RESPONSE_TO_TNF_TROGLITAZONE_UP, RUAN_RESPONSE_TO_TNF_TROGLITAZONE_UP
 GOMF N ACETYLLACTOSAMINIDE BETA 1 3 N ACETYLGLUCOSAMINYLTRANSFERASE ACTIVITY, GOMF N ACETYLLACTOSAMINIDE BETA 1 3 N ACETYLGLUCOSAMINYLTRANSFERASE ACTIVITY
GOCC_POLE_PLASM, GOCC_POLE_PLASM
GOMF_RECEPTOR_INHIBITOR_ACTIVITY, GOMF_RECEPTOR_INHIBITOR_ACTIVITY
GOBP_POLY_N_ACETYLLACTOSAMINE_METABOLIC_PROCESS, GOBP_POLY_N_ACETYLLACTOSAMINE_METABOLIC_PROCESS
REACTOME_COLLAGEN_DEGRADATION, REACTOME_COLLAGEN_DEGRADATION
KIM_BIPOLAR_DISORDER_OLIGODENDROCYTE_DENSITY_CORR_DN, KIM_BIPOLAR_DISORDER_OLIGODENDROCYTE_DENSITY_CORR_DN
GOBP_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS, GOBP_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS
LAKE_ADULT_KIDNEY_C1_EPITHELIAL_CELLS_UNASSIGNED, LAKE_ADULT_KIDNEY_C1_EPITHELIAL_CELLS_UNASSIGNED
GOBP_SYNAPTIC_MEMBRANE_ADHESION, GOBP_SYNAPTIC_MEMBRANE_ADHESION
LU_TUMOR_VASCULATURE_UP, LU_TUMOR_VASCULATURE_UP
HP_RENAL_CORTICOMEDULLARY_CYSTS, HP_RENAL_CORTICOMEDULLARY_CYSTS
GOBP_NEGATIVE_REGULATION_OF_CHEMOKINE_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_CHEMOKINE_PRODUCTION
HP_LIMITED_ELBOW_FLEXION, HP_LIMITED_ELBOW_FLEXION
GOBP_LUNG_GROWTH, GOBP_LUNG_GROWTH
MCCOLLUM GELDANAMYCIN RESISTANCE DN, MCCOLLUM GELDANAMYCIN RESISTANCE DN
MIR662, MIR662
GOBP_NEGATIVE_REGULATION_OF_TUBULIN_DEACETYLATION, GOBP_NEGATIVE_REGULATION_OF_TUBULIN_DEACETYLATION
chr6q26, chr6q26
GOBP_PLASMA_MEMBRANE_FUSION, GOBP_PLASMA_MEMBRANE_FUSION
HP_CERVICAL_C2_C3_VERTEBRAL_FUSION, HP_CERVICAL_C2_C3_VERTEBRAL_FUSION
GOBP_RESPONSE_TO_AUDITORY_STIMULUS, GOBP_RESPONSE_TO_AUDITORY_STIMULUS
QIU_PBMC_HEPTATITIS_B_SURFACE_ANTIGEN_AGE_UNDER50_NON_RESPONDERS_VS_RESPONDERS_0_TO_35DY_JOINTLY_UP, QIU_PBMC_HEPTATITIS_B_SURFACE_ANTIGEN_AGE_UNDER50_NON_RESPONDERS_VS_RESPONDERS_
GOCC_NEUROTRANSMITTER_RECEPTOR_COMPLEX, GOCC_NEUROTRANSMITTER_RECEPTOR_COMPLEX
GOMF_KAINATE_SELECTIVE_GLUTAMATE_RECEPTOR_ACTIVITY, GOMF_KAINATE_SELECTIVE_GLUTAMATE_RECEPTOR_ACTIVITY
GOBP_AXIS_ELONGATION_INVOLVED_IN_SOMITOGENESIS, GOBP_AXIS_ELONGATION_INVOLVED_IN_SOMITOGENESIS
GOBP_PITUITARY_GLAND_DEVELOPMENT, GOBP_PITUITARY_GLAND_DEVELOPMENT
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GARDIQUIMOD_0.5H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_UP