## \_VS\_POLYIC\_STIM\_DC\_3H\_DN, GSE21033\_CTRL\_VS\_POLYIC\_STIM\_DC\_3H\_DN

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GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_UP, GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_U
GSE21033 CTRL VS POLYIC STIM DC 3H UP, GSE21033 CTRL VS POLYIC STIM DC 3H UP
GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_UP, GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_UP
GSE1925 3H VS 24H IFNG STIM MACROPHAGE DN, GSE1925 3H VS 24H IFNG STIM MACROPHAGE DN
GSE29618 PRE VS DAY7 POST LAIV FLU VACCINE BCELL UP, GSE29618 PRE VS DAY7 POST LAIV FLU VACCINE BCELL UP
GSE8515_CTRL_VS_IL6_4H_STIM_MAC_UP, GSE8515_CTRL_VS_IL6_4H_STIM_MAC_UP
GSE20152 HTNFA OVERXPRESS ANKLE VS CTRL SPHK1 KO ANKLE DN, GSE20152 HTNFA OVERXPRESS ANKLE VS CTRL SPI
PRC2 SUZ12 UP.V1 DN, PRC2 SUZ12 UP.V1 DN
GSE13306 TREG RA VS TCONV RA DN, GSE13306 TREG RA VS TCONV RA DN
GO_CORONARY_VASCULATURE_DEVELOPMENT, GO_CORONARY_VASCULATURE_DEVELOPMENT
HADDAD T LYMPHOCYTE AND NK PROGENITOR UP, HADDAD T LYMPHOCYTE AND NK PROGENITOR UP
TCCGTCC_MIR184, TCCGTCC_MIR184
GSE43863_TFH_VS_LY6C_INT_CXCR5POS_MEMORY_CD4_TCELL_DN, GSE43863_TFH_VS_LY6C_INT_CXCR5POS_MEMORY_CD4_TCH
PRC2 EED UP.V1_UP, PRC2_EED_UP.V1_UP
GO_REGULATION_OF_MITOCHONDRIAL_FISSION, GO_REGULATION_OF_MITOCHONDRIAL_FISSION
REACTOME DAG AND IP3 SIGNALING, REACTOME DAG AND IP3 SIGNALING
LOPEZ EPITHELIOID MESOTHELIOMA, LOPEZ EPITHELIOID MESOTHELIOMA
GO POSITIVE REGULATION OF SMOOTH MUSCLE CELL MIGRATION, GO POSITIVE REGULATION OF SMOOTH MUSCLE CEL
GO REGULATION OF OSTEOBLAST DIFFERENTIATION, GO REGULATION OF OSTEOBLAST DIFFERENTIATION
GO CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY, GO_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY
GO CELLULAR RESPONSE TO ALKALOID, GO CELLULAR RESPONSE TO ALKALOID
NAKAMURA METASTASIS MODEL UP, NAKAMURA METASTASIS MODEL UP
GO REGULATION OF OSSIFICATION, GO REGULATION OF OSSIFICATION
REACTOME_CA_DEPENDENT_EVENTS, REACTOME_CA_DEPENDENT_EVENTS
RYAAAKNNNNNTTGW UNKNOWN, RYAAAKNNNNNNTTGW UNKNOWN
GO CARDIAC SEPTUM DEVELOPMENT, GO CARDIAC SEPTUM DEVELOPMENT
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