.ARIZED_CD4_TCELL_TREATED_WITH_DIGOXIN_UP, GSE27241_WT_VS_RORGT_KO_TH17_POLARIZED_CD4_TCELL_TREATED_WITH_DIGOXIN_UP	GSI GSI GSI GO CA GO PIE GSI GSI GSI GSI GSI GSI GSI GSI FRI GO GO GO GO GO BAI	E40655_FOXETTOPE IN SET
	GO BAI MC GO GO	D_PHOSPHO NDRES_RE

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DXO1_KO_VS_WT_NTREG_UP, GSE40655_FOXO1_KO_VS_WT_NTREG_UP
STIM_VS_FCGRIIB_STIM_MONOCYTE_DN, GSE7509_UNSTIM_VS_FCGRIIB_STIM_MONOCYTE_DN
VS VEHICLE CTRL TREATED DC DN, GSE4984 LPS VS VEHICLE CTRL TREATED DC DN
VS IL10 KO LPS AND IL10 STIM MACROPHAGE 45MIN UP, GSE5589 WT VS IL10 KO LPS AND IL10 STIM MACRO
TION OF CELL CYCLE G1 S PHASE TRANSITION, GO REGULATION OF CELL CYCLE G1 S PHASE TRANSITION
MIR370, CAGCAGG_MIR370
TRANSDUCTION IN RESPONSE TO DNA DAMAGE, GO SIGNAL TRANSDUCTION IN RESPONSE TO DNA DAMAGE
WNSTREAM PATHWAY, PID P53 DOWNSTREAM PATHWAY
MCSF_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_HIGHDOS
C NEG VS POS GC BCELL UP, GSE38304 MYC NEG VS POS GC BCELL UP
GCM BCL2L1
IR122A, ACACTCC MIR122A
2, MORF_MYST2
PDGF_PATHWAY, BIOCARTA_PDGF_PATHWAY
CARM ER PATHWAY, BIOCARTA CARM ER PATHWAY
_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_DN, GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_S
RECEPTOR PROXIMAL PATHWAY, PID ERBB1 RECEPTOR PROXIMAL PATHWAY
DRMATION, GO_TUBE_FORMATION
FREAC4 01
TORY_SYSTEM_DEVELOPMENT, GO_RESPIRATORY_SYSTEM_DEVELOPMENT
NIC EYE MORPHOGENESIS, GO EMBRYONIC EYE MORPHOGENESIS
OGENESIS OF EMBRYONIC EPITHELIUM, GO MORPHOGENESIS OF EMBRYONIC EPITHELIUM
C_SEPTUM_DEVELOPMENT, GO_CARDIAC_SEPTUM_DEVELOPMENT
IOTRANSFERASE_ACTIVITY_NITROGENOUS_GROUP_AS_ACCEPTOR, GO_PHOSPHOTRANSFERASE_ACTIVITY_NITROG
ESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48HR_UP, BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48H
ORF IL4
EA_DEVELOPMENT, GO_COCHLEA_DEVELOPMENT
GAMETE_GENERATION, GO_FEMALE_GAMETE_GENERATION
NIC HINDLIMB MORPHOGENESIS, GO EMBRYONIC HINDLIMB MORPHOGENESIS
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