IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_6H\_DN, GSE2770\_TGFB\_AND\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_6H\_DN

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GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL4_TCELL_58H_UP, GSE2770_TGFB_AND_IL4_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_58H_UP, GSE27T0_TCELL_5H_UP, GSE27T0_TCELL_5H_UP
 GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN
 GSE7831_CPG_VS_INFLUENZA_STIM_PDC_4H_UP, GSE7831_CPG_VS_INFLUENZA_STIM_PDC_4H_UP
 GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_1H_DN, GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_1H_DN
 MIR653_3P, MIR653_3P
 MIR629_3P, MIR629_3P
 KEGG_PHOSPHATIDYLINOSITOL_SIGNALING_SYSTEM, KEGG_PHOSPHATIDYLINOSITOL_SIGNALING_SYSTEM
 GTAAACC_MIR2995P, GTAAACC_MIR2995P
 GOBP_PROTEIN_AUTOUBIQUITINATION, GOBP_PROTEIN_AUTOUBIQUITINATION
 DARWICHE_PAPILLOMA_RISK_HIGH_DN, DARWICHE_PAPILLOMA_RISK_HIGH_DN
 GOBP_SYNAPTIC_VESICLE_RECYCLING, GOBP_SYNAPTIC_VESICLE_RECYCLING
 GOBP_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC, GOBP_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC
 TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_UP, TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_UP
 PID_CD40_PATHWAY, PID_CD40_PATHWAY
 HP_SUPRANUCLEAR_GAZE_PALSY, HP_SUPRANUCLEAR_GAZE_PALSY
 MIR3188, MIR3188
 GSE8515_IL1_VS_IL6_4H_STIM_MAC_DN, GSE8515_IL1_VS_IL6_4H_STIM_MAC_DN
 MIR4661_5P, MIR4661_5P
 CHANGOLKAR_H2AFY_TARGETS_DN, CHANGOLKAR_H2AFY_TARGETS_DN
 PID_IL12_2PATHWAY, PID_IL12_2PATHWAY
 GOMF_PROLINE_RICH_REGION_BINDING, GOMF_PROLINE_RICH_REGION_BINDING
 GOBP_SPERMATID_DIFFERENTIATION, GOBP_SPERMATID_DIFFERENTIATION
WP_DISRUPTION_OF_POSTSYNAPTIC_SIGNALLING_BY_CNV, WP_DISRUPTION_OF_POSTSYNAPTIC_SIGNALLING_BY_CNV
HP_BILATERAL_CLEFT_LIP_AND_PALATE, HP_BILATERAL_CLEFT_LIP_AND_PALATE
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GOBP\_CELL\_ADHESION\_MEDIATED\_BY\_INTEGRIN, GOBP\_CELL\_ADHESION\_MEDIATED\_BY\_INTEGRIN

HP\_STARING\_GAZE, HP\_STARING\_GAZE

WP\_MIRNAS\_INVOLVEMENT\_IN\_THE\_IMMUNE\_RESPONSE\_IN\_SEPSIS, WP\_MIRNAS\_INVOLVEMENT\_IN\_THE\_IMMUNE\_RESPONSE\_IN\_SEPSIS ZNF626\_TARGET\_GENES, ZNF626\_TARGET\_GENES

WP\_MIR5093P\_ALTERATION\_OF\_YAP1ECM\_AXIS, WP\_MIR5093P\_ALTERATION\_OF\_YAP1ECM\_AXIS

GOBP\_MEMBRANE\_PROTEIN\_PROTEOLYSIS, GOBP\_MEMBRANE\_PROTEIN\_PROTEOLYSIS

WP\_NEUROINFLAMMATION, WP\_NEUROINFLAMMATION

GOBP\_PRESYNAPSE\_ORGANIZATION, GOBP\_PRESYNAPSE\_ORGANIZATION

GOBP\_POSITIVE\_REGULATION\_OF\_EXCITATORY\_POSTSYNAPTIC\_POTENTIAL, GOBP\_POSITIVE\_REGULATION\_OF\_EXCITATORY\_POSTSYNAPTIC\_F

GOCC\_SPANNING\_COMPONENT\_OF\_MEMBRANE, GOCC\_SPANNING\_COMPONENT\_OF\_MEMBRANE
GOBP\_EPITHELIAL\_TUBE\_BRANCHING\_INVOLVED\_IN\_LUNG\_MORPHOGENESIS, GOBP\_EPITHELIAL\_TUBE\_BRANCHING\_INVOLVED\_IN\_LUNG\_M

GOMF\_COLLAGEN\_BINDING, GOMF\_COLLAGEN\_BINDING
GAUSSMANN\_MLL\_AF4\_FUSION\_TARGETS\_B\_UP, GAUSSMANN\_MLL\_AF4\_FUSION\_TARGETS\_B\_UP

GOCC\_SYMMETRIC\_SYNAPSE, GOCC\_SYMMETRIC\_SYNAPSE

chr11q25, chr11q25

SAKAI\_TUMOR\_INFILTRATING\_MONOCYTES\_UP, SAKAI\_TUMOR\_INFILTRATING\_MONOCYTES\_UP

GOBP\_RESPONSE\_TO\_ACIDIC\_PH, GOBP\_RESPONSE\_TO\_ACIDIC\_PH

MIR5195\_5P, MIR5195\_5P

GOBP\_OMEGA\_HYDROXYLASE\_P450\_PATHWAY, GOBP\_OMEGA\_HYDROXYLASE\_P450\_PATHWAY

REACTOME\_SYNTHESIS\_OF\_16\_20\_HYDROXYEICOSATETRAENOIC\_ACIDS\_HETE, REACTOME\_SYNTHESIS\_OF\_16\_20\_HYDROXYEICOSATETRAENOICGOCC\_MANCHETTE, GOCC\_MANCHETTE

REACTOME\_SYNTHESIS\_OF\_IP3\_AND\_IP4\_IN\_THE\_CYTOSOL, REACTOME\_SYNTHESIS\_OF\_IP3\_AND\_IP4\_IN\_THE\_CYTOSOL

GOBP\_MEMBRANE\_PROTEIN\_INTRACELLULAR\_DOMAIN\_PROTEOLYSIS, GOBP\_MEMBRANE\_PROTEIN\_INTRACELLULAR\_DOMAIN\_PROTEOLYSIS