## FB\_TREATED\_ACT\_TCONV\_DN, GSE7460\_CTRL\_VS\_TGFB\_TREATED\_ACT\_TCONV\_DN

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GSE37532_WT_VS_PPARG_KO_LN_TREG_UP, GSE37532_WT_VS_PPARG_KO_LN_TREG_UP
 GSE4535 BM DERIVED DC VS FOLLICULAR DC DN, GSE4535 BM DERIVED DC VS FOLLICULAR DC DN
 GSE39820 CTRL VS_TGFBETA1_IL6_IL23A_CD4_TCELL_DN, GSE39820_CTRL_VS_TGFBETA1_IL6_IL23A_CD4_TCELL_DN
 GSE21379 TFH VS NON TFH SAP KO CD4 TCELL DN, GSE21379 TFH VS NON TFH SAP KO CD4 TCELL DN
 GSE16385 ROSIGLITAZONE IL4 VS ROSIGLITAZONE ALONE STIM MACROPHAGE UP, GSE16385 ROSIGLITAZONE IL4 VS ROSIGLITAZONE ALONE STIM MACRO
 GO T TUBULE, GO T TUBULE
 GSE3203 HEALTHY VS INFLUENZA INFECTED LN BCELL DN, GSE3203 HEALTHY VS INFLUENZA INFECTED LN BCELL DN
 GSE29615 CTRL VS DAY7 LAIV FLU VACCINE PBMC UP, GSE29615 CTRL VS DAY7 LAIV FLU VACCINE PBMC UP
 GO_SPROUTING_ANGIOGENESIS, GO_SPROUTING ANGIOGENESIS
 GO_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT1_PROTEIN, GO_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT1_PROTEIN
 GO GROWTH FACTOR BINDING, GO GROWTH FACTOR BINDING
 GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_KINASE_ACTIVITY, GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_KINASE_ACTIVITY
GO TUMOR NECROSIS FACTOR RECEPTOR BINDING, GO TUMOR NECROSIS FACTOR RECEPTOR BINDING
ZHENG IL22 SIGNALING UP, ZHENG IL22 SIGNALING UP
 GSE29618 PRE VS_DAY7_POST_LAIV_FLU_VACCINE_PDC_UP, GSE29618 PRE VS_DAY7_POST_LAIV_FLU_VACCINE_PDC_UP
 MODULE_80, MODULE_80
 GO CORECEPTOR ACTIVITY, GO CORECEPTOR ACTIVITY
 GO_REGULATION_OF_CALCIUM_MEDIATED_SIGNALING, GO_REGULATION_OF_CALCIUM_MEDIATED_SIGNALING
 MODULE 51, MODULE 51
 GO_POSITIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT1_PROTEIN, GO_POSITIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT1
 REACTOME_A_TETRASACCHARIDE_LINKER_SEQUENCE_IS_REQUIRED_FOR_GAG_SYNTHESIS, REACTOME_A_TETRASACCHARIDE_LINKER_SEQUENCE_IS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_REQUIRED_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHESIS_FOR_GAG_SYNTHES
 GO POSITIVE REGULATION OF CALCIUM MEDIATED SIGNALING, GO POSITIVE REGULATION OF CALCIUM MEDIATED SIGNALING
GO CELLULAR POTASSIUM ION HOMEOSTASIS, GO CELLULAR POTASSIUM ION HOMEOSTASIS
 GO EPITHELIAL CELL CELL ADHESION, GO EPITHELIAL CELL CELL ADHESION
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