## GFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_2H\_DN, GSE2770\_TGFB\_AND\_IL4\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_2H\_DN

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GSE1460_NAIVE_CD4_TCELL_CORD_BLOOD_VS_THYMIC_STROMAL_CELL_DN, GSE1460_NAIVE_CD4_TCELL_CORD_BLOOD_VS_THYMIC_STROMAL_CELL_DN
  GSE13738 RESTING VS_TCR_ACTIVATED_CD4_TCELL_DN, GSE13738 RESTING_VS_TCR_ACTIVATED_CD4_TCELL_DN
  GOBP_RNA_MODIFICATION, GOBP_RNA_MODIFICATION
  GSE2770_UNTREATED_VS_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_ACT_CD4_TCELL_48H_UP
  GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN, GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN
 REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION, REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION
 GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP, GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP
 MIR7156 3P, MIR7156 3P
  REACTOME ERCC6 CSB AND EHMT2 G9A POSITIVELY REGULATE RRNA EXPRESSION, REACTOME ERCC6 CSB AND EHMT2 G9A POSITIVELY REGULATE RRNA EXPRESSION
  GOBP_AMINO_ACID_TRANSPORT, GOBP_AMINO_ACID_TRANSPORT
  MIR5572, MIR5572
  MIR573, MIR573
   REACTOME_BASE_EXCISION_REPAIR_AP_SITE_FORMATION, REACTOME_BASE_EXCISION_REPAIR_AP_SITE_FORMATION
  MIR3616_5P, MIR3616_5P
  GOMF_INORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_INORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
✓ REACTOME_ACTIVATED_PKN1_STIMULATES_TRANSCRIPTION_OF_AR_ANDROGEN_RECEPTOR_REGULATED_GENES_KLK2_AND_KLK3, REACTOME_ACTIVATED_PKN1_STIMULATES_TRANSCRIPTION_OF_AR_ANDROGEN_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_REGULATED_TRANSCRIPTION_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RECEPTOR_RE
 GOMF_CHLORIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_CHLORIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
  WP_HISTONE_MODIFICATIONS, WP_HISTONE_MODIFICATIONS
  HP_SCAPULAR_WINGING, HP_SCAPULAR_WINGING
  GOBP_AMINO_ACID_TRANSMEMBRANE_TRANSPORT, GOBP_AMINO_ACID_TRANSMEMBRANE_TRANSPORT
  GOMF_ANION_CHANNEL_ACTIVITY, GOMF_ANION_CHANNEL_ACTIVITY
  GOBP_POLYOL_BIOSYNTHETIC_PROCESS, GOBP_POLYOL_BIOSYNTHETIC_PROCESS
  GOBP_DICARBOXYLIC_ACID_TRANSPORT, GOBP_DICARBOXYLIC_ACID_TRANSPORT
  MODULE 316, MODULE 316
  GSE37301_HEMATOPOIETIC_STEM_CELL_VS_CD4_TCELL_DN, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_CD4_TCELL_DN
  HP_FAIR_HAIR, HP_FAIR_HAIR
  GOMF_LIGAND_GATED_ANION_CHANNEL_ACTIVITY, GOMF_LIGAND_GATED_ANION_CHANNEL_ACTIVITY
  GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS, GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS
  GOMF_GABA_RECEPTOR_ACTIVITY, GOMF_GABA_RECEPTOR_ACTIVITY
  GOCC_GABA_RECEPTOR_COMPLEX, GOCC_GABA_RECEPTOR_COMPLEX
  BRF1_TARGET_GENES, BRF1_TARGET_GENES
  GOCC CHLORIDE CHANNEL COMPLEX, GOCC CHLORIDE CHANNEL COMPLEX
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