DLAN\_LOWDOSE\_STIM\_DC\_DN, GSE32986\_UNSTIM\_VS\_CURDLAN\_LOWDOSE\_STIM\_DC\_DN

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GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_DN
GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_UP, GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_UP
GSE26928 NAIVE VS_CXCR5_POS_CD4_TCELL_DN, GSE26928_NAIVE_VS_CXCR5_POS_CD4_TCELL_DN
GSE13411_NAIVE_VS_MEMORY_BCELL_DN, GSE13411_NAIVE_VS_MEMORY_BCELL_DN
GSE26928_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN, GSE26928_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN
GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_UP, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_UP
REACTOME_HATS_ACETYLATE_HISTONES, REACTOME_HATS_ACETYLATE_HISTONES
GSE21927_C26GM_VS_4T1_TUMOR_MONOCYTE_BALBC_UP, GSE21927_C26GM_VS_4T1_TUMOR_MONOCYTE_BALBC_UP
GSE27786_LIN_NEG_VS_MONO_MAC_DN, GSE27786_LIN_NEG_VS_MONO_MAC_DN
GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_DN, GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_DN
REACTOME_PROTEIN_UBIQUITINATION, REACTOME_PROTEIN_UBIQUITINATION
ZHENG CORD BLOOD C2 PUTATIVE BASOPHIL EOSINOPHIL MAST CELL PROGENITOR, ZHENG CORD BLOOD C2 PUTATIVE BASOPHIL EOSINOPHIL MAST CELL PROGENIT
REACTOME_FORMATION_OF_THE_BETA_CATENIN_TCF_TRANSACTIVATING_COMPLEX, REACTOME_FORMATION_OF_THE_BETA_CATENIN_TCF_TRANSACTIVATING_COMPLEX
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_UP
MODULE_206, MODULE_206
GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_DN, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_DN
REACTOME_HDACS_DEACETYLATE_HISTONES, REACTOME_HDACS_DEACETYLATE_HISTONES
REACTOME_SIRT1_NEGATIVELY_REGULATES_RRNA_EXPRESSION, REACTOME_SIRT1_NEGATIVELY_REGULATES_RRNA_EXPRESSION
REACTOME_ERCC6_CSB_AND_EHMT2_G9A_POSITIVELY_REGULATE_RRNA_EXPRESSION, REACTOME_ERCC6_CSB_AND_EHMT2_G9A_POSITIVELY_REGULATE_RRNA_EXPRESSION
GSE13306_TREG_RA_VS_TCONV_RA_UP, GSE13306_TREG_RA_VS_TCONV_RA_UP
GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN
KRCTCNNNNMANAGC_UNKNOWN, KRCTCNNNNMANAGC_UNKNOWN
GSE21033_CTRL_VS_POLYIC_STIM_DC_1H_UP, GSE21033_CTRL_VS_POLYIC_STIM_DC_1H_UP
REACTOME_L1CAM_INTERACTIONS, REACTOME_L1CAM_INTERACTIONS
HP_ABNORMALITY_OF_HEAD_BLOOD_VESSEL, HP_ABNORMALITY_OF_HEAD_BLOOD_VESSEL
WP_GENOTOXICITY_PATHWAY, WP_GENOTOXICITY_PATHWAY
GOBP_CARBOHYDRATE_DERIVATIVE_TRANSPORT, GOBP_CARBOHYDRATE_DERIVATIVE_TRANSPORT
REACTOME_GLYCOSAMINOGLYCAN_METABOLISM, REACTOME_GLYCOSAMINOGLYCAN_METABOLISM
GOBP_POSITIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT
NOTCH_DN.V1_DN, NOTCH_DN.V1_DN
GOBP_RDNA_HETEROCHROMATIN_ASSEMBLY, GOBP_RDNA_HETEROCHROMATIN_ASSEMBLY
SENESE_HDAC2_TARGETS_DN, SENESE_HDAC2_TARGETS_DN
GOBP_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOBP_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GSE45365_BCELL_VS_CD8_TCELL_DN, GSE45365_BCELL_VS_CD8_TCELL_DN
GOBP_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION, GOBP_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION
VALK_AML_CLUSTER_8, VALK_AML_CLUSTER_8
MIR4713_3P, MIR4713_3P
REACTOME_GAP_JUNCTION_ASSEMBLY, REACTOME_GAP_JUNCTION_ASSEMBLY
REACTOME_GAP_JUNCTION_TRAFFICKING_AND_REGULATION, REACTOME_GAP_JUNCTION_TRAFFICKING_AND_REGULATION
GOBP_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORT, GOBP_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORT
HP_POOR_WOUND_HEALING, HP_POOR_WOUND_HEALING
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GOBP\_REGULATION\_OF\_WATER\_LOSS\_VIA\_SKIN, GOBP\_REGULATION\_OF\_WATER\_LOSS\_VIA\_SKIN

GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_MORPHOGENESIS\_INVOLVED\_IN\_DIFFERENTIATION, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_MORPHOGENESIS\_INVOLVED\_IN\_DIFI ABBUD\_LIF\_SIGNALING\_2\_UP, ABBUD\_LIF\_SIGNALING\_2\_UP

GOMF\_NUCLEOTIDE\_SUGAR\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_NUCLEOTIDE\_SUGAR\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY

HP\_ELEVATED\_URINARY\_CARBOXYLIC\_ACID, HP\_ELEVATED\_URINARY\_CARBOXYLIC\_ACID

GOBP\_CALCIUM\_ION\_TRANSMEMBRANE\_TRANSPORT\_VIA\_HIGH\_VOLTAGE\_GATED\_CALCIUM\_CHANNEL, GOBP\_CALCIUM\_ION\_TRANSMEMBRANE\_TRANSPORT\_VIA\_HIGH\_V

PLASARI\_NFIC\_TARGETS\_BASAL\_DN, PLASARI\_NFIC\_TARGETS\_BASAL\_DN