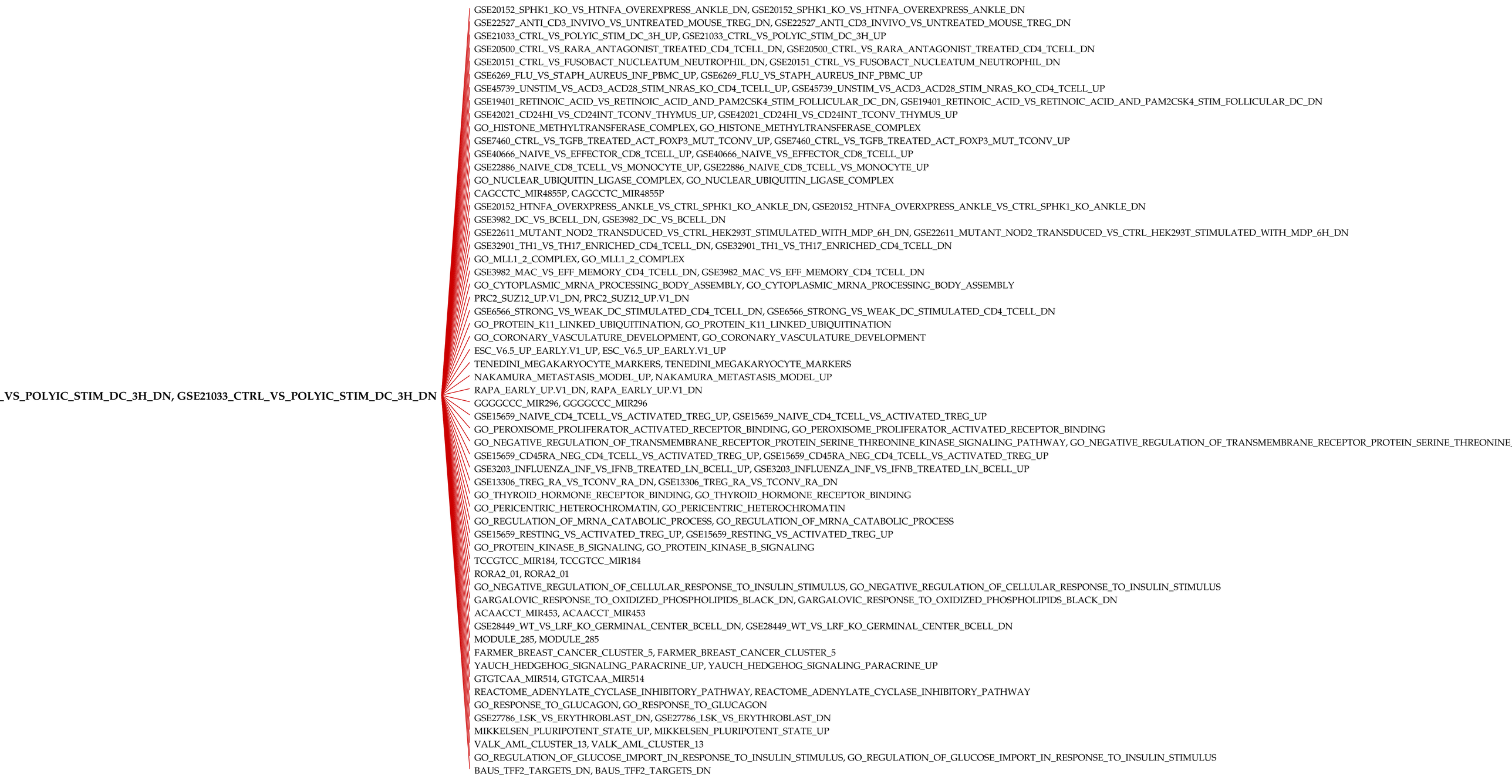
_VS_POLYIC_STIM_DC_3H_DN, GSE21033_CTRL_VS_POLYIC_STIM_DC_3H_DN
GSE20152_SPHK1_KO_VS_HTNFA_OVEREXPRESS_ANKLE_DN, GSE20152_SPHK1_KO_VS_HTNFA_OVEREXPRESS_ANKLE_DN
GSE22527_ANTI_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_DN, GSE22527_ANTI_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_DN
GSE21033_CTRL_VS_POLYIC_STIM_DC_3H_UP, GSE21033_CTRL_VS_POLYIC_STIM_DC_3H_UP
GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_DN, GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_DN
GSE20151_CTRL_VS_FUSOBACT_NUCLEATUM_NEUTROPHIL_DN, GSE20151_CTRL_VS_FUSOBACT_NUCLEATUM_NEUTROPHIL_DN
GSE6269_FLU_VS_STAPH_AUREUS_INF_PBMC_UP, GSE6269_FLU_VS_STAPH_AUREUS_INF_PBMC_UP
GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_NRAS_KO_CD4_TCELL_UP, GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_NRAS_KO_CD4_TCELL_UP
GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN, GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN
GSE42021_CD24HI_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_CD24HI_VS_CD24INT_TCONV_THYMUS_UP
GO_HISTONE_METHYLTRANSFERASE_COMPLEX, GO_HISTONE_METHYLTRANSFERASE_COMPLEX
GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_MUT_TCONV_UP, GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_MUT_TCONV_UP
GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_UP, GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_UP
GSE22886_NAIVE_CD8_TCELL_VS_MONOCYTE_UP, GSE22886_NAIVE_CD8_TCELL_VS_MONOCYTE_UP
GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX, GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX
CAGCCTC_MIR4855P, CAGCCTC_MIR4855P
GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN, GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN
GSE3982_DC_VS_BCELL_DN, GSE3982_DC_VS_BCELL_DN
GSE22611_MUTANT_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_6H_DN, GSE22611_MUTANT_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_6H_DN
GSE32901_TH1_VS_TH17_ENRICHED_CD4_TCELL_DN, GSE32901_TH1_VS_TH17_ENRICHED_CD4_TCELL_DN
GO_MLL1_2_COMPLEX, GO_MLL1_2_COMPLEX
GSE3982_MAC_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_MAC_VS_EFF_MEMORY_CD4_TCELL_DN
GO_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY, GO_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY
PRC2_SUZ12_UP.V1_DN, PRC2_SUZ12_UP.V1_DN
GSE6566_STRONG_VS_WEAK_DC_STIMULATED_CD4_TCELL_DN, GSE6566_STRONG_VS_WEAK_DC_STIMULATED_CD4_TCELL_DN
GO_PROTEIN_K11_LINKED_UBIQUITINATION, GO_PROTEIN_K11_LINKED_UBIQUITINATION
GO_CORONARY_VASCULATURE_DEVELOPMENT, GO_CORONARY_VASCULATURE_DEVELOPMENT
ESC_V6.5_UP_EARLY.V1_UP, ESC_V6.5_UP_EARLY.V1_UP
TENEDINI_MEGAKARYOCYTE_MARKERS, TENEDINI_MEGAKARYOCYTE_MARKERS
NAKAMURA_METASTASIS_MODEL_UP, NAKAMURA_METASTASIS_MODEL_UP
RAPA_EARLY_UP.V1_DN, RAPA_EARLY_UP.V1_DN
GGGGCCC_MIR296, GGGGCCC_MIR296
GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_UP, GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_UP
GO_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_BINDING, GO_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_BINDING
GO_NEGATIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_
GSE15659_CD45RA_NEG_CD4_TCELL_VS_ACTIVATED_TREG_UP, GSE15659_CD45RA_NEG_CD4_TCELL_VS_ACTIVATED_TREG_UP
GSE3203_INFLUENZA_INF_VS_IFNB_TREATED_LN_BCELL_UP, GSE3203_INFLUENZA_INF_VS_IFNB_TREATED_LN_BCELL_UP
GSE13306_TREG_RA_VS_TCONV_RA_DN, GSE13306_TREG_RA_VS_TCONV_RA_DN
GO_THYROID_HORMONE_RECEPTOR_BINDING, GO_THYROID_HORMONE_RECEPTOR_BINDING
GO_PERICENTRIC_HETEROCHROMATIN, GO_PERICENTRIC_HETEROCHROMATIN
GO_REGULATION_OF_MRNA_CATABOLIC_PROCESS, GO_REGULATION_OF_MRNA_CATABOLIC_PROCESS
GSE15659_RESTING_VS_ACTIVATED_TREG_UP, GSE15659_RESTING_VS_ACTIVATED_TREG_UP
GO_PROTEIN_KINASE_B_SIGNALING, GO_PROTEIN_KINASE_B_SIGNALING
TCCGTCC_MIR184, TCCGTCC_MIR184
RORA2_01, RORA2_01
GO_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GO_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLACK_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLACK_DN
ACAACCT_MIR453, ACAACCT_MIR453
GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_DN, GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_DN
MODULE_285, MODULE_285
FARMER_BREAST_CANCER_CLUSTER_5, FARMER_BREAST_CANCER_CLUSTER_5
YAUCH_HEDGEHOG_SIGNALING_PARACRINE_UP, YAUCH_HEDGEHOG_SIGNALING_PARACRINE_UP
GTGTCAA_MIR514, GTGTCAA_MIR514
REACTOME_ADENYLATE_CYCLASE_INHIBITORY_PATHWAY, REACTOME_ADENYLATE_CYCLASE_INHIBITORY_PATHWAY
GO_RESPONSE_TO_GLUCAGON, GO_RESPONSE_TO_GLUCAGON
GSE27786_LSK_VS_ERYTHROBLAST_DN, GSE27786_LSK_VS_ERYTHROBLAST_DN
MIKKELSEN_PLURIPOTENT_STATE_UP, MIKKELSEN_PLURIPOTENT_STATE_UP
VALK_AML_CLUSTER_13, VALK_AML_CLUSTER_13
GO_REGULATION_OF_GLUCOSE_IMPORT_IN_RESPONSE_TO_INSULIN_STIMULUS, GO_REGULATION_OF_GLUCOSE_IMPORT_IN_RESPONSE_TO_INSULIN_STIMULUS
BAUS_TFF2_TARGETS_DN, BAUS_TFF2_TARGETS_DN