OD\_CD21HIGH\_TRANSITIONAL\_BCELL\_UP, GSE17186\_BLOOD\_VS\_CORD\_BLOOD\_CD21HIGH\_TRANSITIONAL\_BCELL\_UP

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GSE17186 NAIVE VS CD21HIGH TRANSITIONAL BCELL DN, GSE17186 NAIVE VS CD21HIGH TRANSITIONAL BCELL DN
 GSE12392_WT_VS_IFNB_KO_CD8A_POS_SPLEEN_DC_UP, GSE12392_WT_VS_IFNB_KO_CD8A_POS_SPLEEN_DC_UP
 GSE17186_MEMORY_VS_CD21LOW_TRANSITIONAL_BCELL_UP, GSE17186_MEMORY_VS_CD21LOW_TRANSITIONAL_BCELL_UP
 GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP, GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP
 GSE30083_SP3_VS_SP4_THYMOCYTE_DN, GSE30083_SP3_VS_SP4_THYMOCYTE_DN
 GSE17186_BLOOD_VS_CORD_BLOOD_NAIVE_BCELL_UP, GSE17186_BLOOD_VS_CORD_BLOOD_NAIVE_BCELL_UP
 GSE17186_MEMORY_VS_NAIVE_BCELL_DN, GSE17186_MEMORY_VS_NAIVE_BCELL_DN
 GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN, GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN
 GO_VACUOLE_ORGANIZATION, GO_VACUOLE_ORGANIZATION
GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_UP, GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_UP
GSE24210_TCONV_VS_TREG_DN, GSE24210_TCONV_VS_TREG_DN
GSE42724_NAIVE_VS_MEMORY_BCELL_DN, GSE42724_NAIVE_VS_MEMORY_BCELL_DN
GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN
GSE5589 IL6 KO_VS IL10 KO_LPS AND IL6 STIM MACROPHAGE 45MIN_DN, GSE5589 IL6 KO_VS IL10 KO_LPS AND IL6 STIM MACROPHAGE 45MIN_DN
 GSE339_CD8POS_VS_CD4CD8DN_DC_IN_CULTURE_UP, GSE339_CD8POS_VS_CD4CD8DN_DC_IN_CULTURE_UP
 GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_UP
GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP
GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_DC_DN, GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_DC_DN
 GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP
 GO_POSITIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
 GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNAR_KO_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNAR_KO_DN
 GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN, GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN
 GSE20366 CD103 KLRG1 DP_VS_DN_TREG_DN, GSE20366 CD103 KLRG1 DP_VS_DN_TREG_DN
 LIU_VMYB_TARGETS_UP, LIU_VMYB_TARGETS_UP
 GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN, GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN
 REACTOME_POST_TRANSLATIONAL_PROTEIN_MODIFICATION, REACTOME_POST_TRANSLATIONAL_PROTEIN_MODIFICATION
 REACTOME_ASPARAGINE_N_LINKED_GLYCOSYLATION, REACTOME_ASPARAGINE_N_LINKED_GLYCOSYLATION
 GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY40_UP, GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY40_UP
 GO_REGULATION_OF_PROTEIN_IMPORT, GO_REGULATION_OF_PROTEIN_IMPORT
 GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP
 GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN, GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN
 GO_MODULATION_BY_VIRUS_OF_HOST_MORPHOLOGY_OR_PHYSIOLOGY, GO_MODULATION_BY_VIRUS_OF_HOST_MORPHOLOGY_OR_PHYSIOLOGY
 KEGG_N_GLYCAN_BIOSYNTHESIS, KEGG_N_GLYCAN_BIOSYNTHESIS
 ST_GAQ_PATHWAY, ST_GAQ_PATHWAY
 GSE37534_GW1929_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP, GSE37534_GW1929_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP
 GO_MODULATION_BY_SYMBIONT_OF_HOST_CELLULAR_PROCESS, GO_MODULATION_BY_SYMBIONT_OF_HOST_CELLULAR_PROCESS
 GO RESPONSE_TO_FLUID_SHEAR_STRESS, GO_RESPONSE_TO_FLUID_SHEAR_STRESS
 GO_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLVED_IN_APOPTOTIC_
 GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_6H_DN, GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_6H_DN
 GO MODIFICATION_BY_SYMBIONT_OF_HOST_MORPHOLOGY_OR_PHYSIOLOGY, GO_MODIFICATION_BY_SYMBIONT_OF_HOST_MORPHOLOGY_OR_PHYSIOLOGY
 GO_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY, GO_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY
 CHEN_HOXA5_TARGETS_9HR_DN, CHEN_HOXA5_TARGETS_9HR_DN
 GO_APOPTOTIC_MITOCHONDRIAL_CHANGES, GO_APOPTOTIC_MITOCHONDRIAL_CHANGES
 GO_KINASE_ACTIVATOR_ACTIVITY, GO_KINASE_ACTIVATOR_ACTIVITY
 GO POSITIVE REGULATION OF MITOCHONDRIAL OUTER MEMBRANE PERMEABILIZATION INVOLVED IN APOPTOTIC SIGNALING PATHWAY, GO POSITIVE REGULATION OF MITOCHONDRIAL OUTER MEMBRANE PERMEABILIZATION INVOLVED.
 LEE_NEURAL_CREST_STEM_CELL_UP, LEE_NEURAL_CREST_STEM_CELL_UP
 GO_NEGATIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY
 BCAT_GDS748_DN, BCAT_GDS748_DN
ST_GA13_PATHWAY, ST_GA13_PATHWAY
 GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH2_GROUP_OF_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH2_GROUP_OF_DONORS
PID_A6B1_A6B4_INTEGRIN_PATHWAY, PID_A6B1_A6B4_INTEGRIN_PATHWAY
GO_RESPONSE_TO_LAMINAR_FLUID_SHEAR_STRESS, GO_RESPONSE_TO_LAMINAR_FLUID_SHEAR_STRESS
GO_SITE_OF_POLARIZED_GROWTH, GO_SITE_OF_POLARIZED_GROWTH
REACTOME_CALNEXIN_CALRETICULIN_CYCLE, REACTOME_CALNEXIN_CALRETICULIN_CYCLE
LUCAS_HNF4A_TARGETS_UP, LUCAS_HNF4A_TARGETS_UP
BIOCARTA_MITOCHONDRIA_PATHWAY, BIOCARTA_MITOCHONDRIA_PATHWAY
GO_PROTEIN_DEMANNOSYLATION, GO_PROTEIN_DEMANNOSYLATION
GO_CELLULAR_CARBOHYDRATE_BIOSYNTHETIC_PROCESS, GO_CELLULAR_CARBOHYDRATE_BIOSYNTHETIC_PROCESS
 REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE, REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE
 GSE21546_UNSTIM_VS_ANTI_CD3_STIM_ELK1_KO_DP_THYMOCYTES_DN, GSE21546_UNSTIM_VS_ANTI_CD3_STIM_ELK1_KO_DP_THYMOCYTES_DN
 TERAMOTO_OPN_TARGETS_CLUSTER_6, TERAMOTO_OPN_TARGETS_CLUSTER_6
 BIOCARTA_BCELLSURVIVAL_PATHWAY, BIOCARTA_BCELLSURVIVAL_PATHWAY
 MEISSNER_ES_ICP_WITH_H3K4ME3, MEISSNER_ES_ICP_WITH_H3K4ME3
 MEISSNER_NPC_ICP_WITH_H3K4ME3, MEISSNER_NPC_ICP_WITH_H3K4ME3
 GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH2_GROUP_OF_DONORS_OXYGEN_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH2_GROUP_OF_DONORS_OXYGEN_AS_ACCEPTOR
 GO_GDP_DISSOCIATION_INHIBITOR_ACTIVITY, GO_GDP_DISSOCIATION_INHIBITOR_ACTIVITY
 GO_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS
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GSE29164 DAY3 VS DAY7 UNTREATED MELANOMA DN, GSE29164 DAY3 VS DAY7 UNTREATED MELANOMA DN