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GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP, GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP
 GSE17721 0.5H VS 24H PAM3CSK4 BMDC DN, GSE17721 0.5H VS 24H PAM3CSK4 BMDC DN
GSE37416_0H_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE37301 MULTIPOTENT PROGENITOR VS RAG2 KO NK CELL DN, GSE37301 MULTIPOTENT PROGENITOR VS RAG2 KO NK CELL DN
UEDA_PERIFERAL_CLOCK, UEDA_PERIFERAL_CLOCK
GSE21546 ELK1 KO_VS SAP1A KO_AND ELK1 KO_ANTI_CD3 STIM_DP_THYMOCYTES_DN, GSE21546 ELK1_KO_VS SAP1A_KO_AND ELK1_KO_ANTI_CD3 STIM_DP_THYMOCYTES_DN
DNMT1_TARGET_GENES, DNMT1_TARGET_GENES
THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_NONRESPONDER_2DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_NONRESPONDER_2DY_UP
GSE21033_CTRL_VS_POLYIC_STIM_DC_6H_UP, GSE21033_CTRL_VS_POLYIC_STIM_DC_6H_UP
MIR21_3P, MIR21_3P
GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_DN, GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_DN
GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN
GOMF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN, GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN
GOBP_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_ALPHA_BETA_T_CELL_ACTIVATION
GSE6566_STRONG_VS_WEAK_DC_STIMULATED_CD4_TCELL_UP, GSE6566_STRONG_VS_WEAK_DC_STIMULATED_CD4_TCELL_UP
GOBP_STEROL_BIOSYNTHETIC_PROCESS, GOBP_STEROL_BIOSYNTHETIC_PROCESS
GCM_BECN1, GCM_BECN1
GSE6269_E_COLI_VS_STREP_PNEUMO_INF_PBMC_UP, GSE6269_E_COLI_VS_STREP_PNEUMO_INF_PBMC_UP
FERRANDO_T_ALL_WITH_MLL_ENL_FUSION_UP, FERRANDO_T_ALL_WITH_MLL_ENL_FUSION_UP
MIR1273H 3P, MIR1273H 3P
GOBP REGULATION OF DENDRITIC SPINE DEVELOPMENT, GOBP REGULATION OF DENDRITIC SPINE DEVELOPMENT
MIR4435, MIR4435
YTCCCRNNAGGY_UNKNOWN, YTCCCRNNAGGY_UNKNOWN
CUI_DEVELOPING_HEART_TRABECULAR_VENTRICULAR_CARDIOMYOCYTE, CUI_DEVELOPING_HEART_TRABECULAR_VENTRICULAR_CARDIOMYOCYTE
MIR4299, MIR4299
GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
MIR6753_3P, MIR6753_3P
MIR7107_3P, MIR7107_3P
GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
HP_DELAYED_PUBERTY, HP_DELAYED_PUBERTY
GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION
MIR7110_5P, MIR7110_5P
MIR10401_5P, MIR10401_5P
GOBP_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS, GOBP_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS
HP_ARTHROPATHY, HP_ARTHROPATHY
GOBP_RESPONSE_TO_OSMOTIC_STRESS, GOBP_RESPONSE_TO_OSMOTIC_STRESS
MIR6842_5P, MIR6842_5P
MIR6752_5P, MIR6752_5P
GOBP_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION
HOEBEKE_LYMPHOID_STEM_CELL_UP, HOEBEKE_LYMPHOID_STEM_CELL_UP
MIR4999 5P, MIR4999 5P
GOMF_CARBOHYDRATE_DERIVATIVE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_CARBOHYDRATE_DERIVATIVE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
LI WILMS TUMOR VS FETAL KIDNEY 2 DN, LI WILMS TUMOR VS FETAL KIDNEY 2 DN
MIR4722 5P, MIR4722 5P
WOOD_EBV_EBNA1_TARGETS_DN, WOOD_EBV_EBNA1_TARGETS_DN
GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_DEVELOPMENT
 MIR4317, MIR4317
 GOBP_PROTEIN_K48_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K48_LINKED_DEUBIQUITINATION
ONO_AML1_TARGETS_DN, ONO_AML1_TARGETS_DN
 MIR6866_5P, MIR6866_5P
 GOCC_NURD_COMPLEX, GOCC_NURD_COMPLEX
 MIR6812_5P, MIR6812_5P
 KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE
 HP_GYNECOMASTIA, HP_GYNECOMASTIA
 MIR323B_3P, MIR323B_3P
CACGTTT_MIR302A, CACGTTT_MIR302A
MIR342_5P, MIR342_5P
GOBP_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION
TAGAACC_MIR182, TAGAACC_MIR182
 MIR4425, MIR4425
GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE
MIR4727_3P, MIR4727_3P
KANG_DOXORUBICIN_RESISTANCE_DN, KANG_DOXORUBICIN_RESISTANCE_DN
GOBP_MEMBRANE_PROTEIN_PROTEOLYSIS, GOBP_MEMBRANE_PROTEIN_PROTEOLYSIS
HP_ABNORMAL_SPERMATOGENESIS, HP_ABNORMAL_SPERMATOGENESIS
MIR4771, MIR4771
MIR4664_5P, MIR4664_5P
MIR6819_5P, MIR6819_5P
 ATGCACG_MIR517B, ATGCACG_MIR517B
KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_LACTO_AND_NEOLACTO_SERIES, KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_LACTO_AND_NEOLACTO_SERIES
HP_IMPOTENCE, HP_IMPOTENCE
GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION
HP_THICKENED_CALVARIA, HP_THICKENED_CALVARIA
GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
HP_EUNUCHOID_HABITUS, HP_EUNUCHOID_HABITUS
MIR6737_5P, MIR6737_5P
GOBP_NITRIC_OXIDE_SYNTHASE_BIOSYNTHETIC_PROCESS, GOBP_NITRIC_OXIDE_SYNTHASE_BIOSYNTHETIC_PROCESS
BIOCARTA_S1P_PATHWAY, BIOCARTA_S1P_PATHWAY
GOBP_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GOBP_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III
WORSCHECH_TUMOR_REJECTION_UP, WORSCHECH_TUMOR_REJECTION_UP
HP_PROMINENT_SUPRAORBITAL_RIDGES, HP_PROMINENT_SUPRAORBITAL_RIDGES
WP_HFE_EFFECT_ON_HEPCIDIN_PRODUCTION, WP_HFE_EFFECT_ON_HEPCIDIN_PRODUCTION
GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION
GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION
GOBP_SPLEEN_DEVELOPMENT, GOBP_SPLEEN_DEVELOPMENT
GOBP_NEGATIVE_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION
GOBP REGULATION OF T HELPER CELL DIFFERENTIATION, GOBP REGULATION OF T HELPER CELL DIFFERENTIATION
GOBP_RESPONSE_TO_GONADOTROPIN, GOBP_RESPONSE_TO_GONADOTROPIN
GOBP_PROTEIN_K11_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K11_LINKED_DEUBIQUITINATION
 GOBP_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE_MEDIATED_BY_CIRCULATING_IMMUNOGLOBULIN, GOBP_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE_MEDIATED_BY_CIRCULATING_IMMUNOGLO
GOBP_GERMINAL_CENTER_FORMATION, GOBP_GERMINAL_CENTER_FORMATION
HP_CAVUM_SEPTUM_PELLUCIDUM, HP_CAVUM_SEPTUM_PELLUCIDUM
GOBP_REGULATION_OF_DENDRITE_MORPHOGENESIS, GOBP_REGULATION_OF_DENDRITE_MORPHOGENESIS
GOBP_RESPONSE_TO_RETINOIC_ACID, GOBP_RESPONSE_TO_RETINOIC_ACID
GOBP_GLUCOSAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_GLUCOSAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
 GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
DESCARTES_MAIN_FETAL_ADRENOCORTICAL_CELLS, DESCARTES_MAIN_FETAL_ADRENOCORTICAL_CELLS
GOBP_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION
GOMF VOLTAGE GATED CALCIUM CHANNEL ACTIVITY, GOMF VOLTAGE GATED CALCIUM CHANNEL ACTIVITY
GOBP_REGULATION_OF_NEURON_MIGRATION, GOBP_REGULATION_OF_NEURON_MIGRATION
 DPPA3_TARGET_GENES, DPPA3_TARGET_GENES
 HP_DECREASED_MUSCLE_MASS, HP_DECREASED_MUSCLE_MASS
HP_SHORTENING_OF_ALL_DISTAL_PHALANGES_OF_THE_FINGERS, HP_SHORTENING_OF_ALL_DISTAL_PHALANGES_OF_THE_FINGERS
chr16q13, chr16q13
HP_HYPOTHALAMIC_GONADOTROPIN_RELEASING_HORMONE_DEFICIENCY, HP_HYPOTHALAMIC_GONADOTROPIN_RELEASING_HORMONE_DEFICIENCY
GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION
HP_ALLERGY, HP_ALLERGY
RICHERT_PBMC_HIV_LIPO_5_AGE_37_48YO_STIMULATED_VS_UNSTIMULATED_0W_14W_METALLOTHIONEIN_SUBSET_UP, RICHERT_PBMC_HIV_LIPO_5_AGE_37_48YO_STIMULATED_VS_UNSTIMULATED_SUBSET_UP, RICHERT_PBMC_HIV_SUBSET_UP, RICHERT_PBMC_HIV_SUBSET_UP, RICHERT_NUP, RICHERT_PBMC_HIV_SUBSET_UP, RICHERT_PBMC_HIV_SUBSET_UP, RICHERT_PBMC_HIV_SUBSET_UP, RICHERT_PBMC_HIV_SUBSET_UP, RICHERT_PBMC_HIV_SUBSET_UP, RICHERT_PBMC_TPAT_UP, RICHERT_PBMC_TPAT_UP, RICHERT_PBMC_TPAT_UP, RICHERT_PBMC_TPAT_UP, RICHERT_PBMC_TPAT_UP, RICH
GOBP_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION, GOBP_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION
GOMF_SULFURTRANSFERASE_ACTIVITY, GOMF_SULFURTRANSFERASE_ACTIVITY
GOBP_REGULATION_OF_BONE_REMODELING, GOBP_REGULATION_OF_BONE_REMODELING
WENG_POR_TARGETS_GLOBAL_UP, WENG_POR_TARGETS_GLOBAL_UP
GOBP POSITIVE REGULATION OF CHOLESTEROL METABOLIC PROCESS, GOBP POSITIVE REGULATION OF CHOLESTEROL METABOLIC PROCESS
 HP_CAMPTODACTYLY_OF_TOE, HP_CAMPTODACTYLY_OF_TOE
 GOMF SELENIUM BINDING, GOMF SELENIUM BINDING
 WP_MEVALONATE_PATHWAY, WP_MEVALONATE_PATHWAY
chr1p22, chr1p22
GOBP_POSITIVE_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS
GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX, GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX
HP_CEREBELLAR_VERMIS_ATROPHY, HP_CEREBELLAR_VERMIS_ATROPHY
HP_PREAXIAL_FOOT_POLYDACTYLY, HP_PREAXIAL_FOOT_POLYDACTYLY
SU_PLACENTA, SU_PLACENTA
 GOBP_N_ACETYLGLUCOSAMINE_METABOLIC_PROCESS, GOBP_N_ACETYLGLUCOSAMINE_METABOLIC_PROCESS
 GOBP_REGULATION_OF_CHRONIC_INFLAMMATORY_RESPONSE, GOBP_REGULATION_OF_CHRONIC_INFLAMMATORY_RESPONSE
 GOBP_REGULATION_OF_T_HELPER_2_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_T_HELPER_2_CELL_DIFFERENTIATION
HP_NUCHAL_RIGIDITY, HP_NUCHAL_RIGIDITY
 WP_SOMITOGENESIS_IN_THE_CONTEXT_OF_SPONDYLOCOSTAL_DYSOSTOSIS, WP_SOMITOGENESIS_IN_THE_CONTEXT_OF_SPONDYLOCOSTAL_DYSOSTOSIS
GOBP_EXCITATORY_CHEMICAL_SYNAPTIC_TRANSMISSION, GOBP_EXCITATORY_CHEMICAL_SYNAPTIC_TRANSMISSION
HP_SHORTENING_OF_ALL_MIDDLE_PHALANGES_OF_THE_FINGERS, HP_SHORTENING_OF_ALL_MIDDLE_PHALANGES_OF_THE_FINGERS
 MIR483_5P, MIR483_5P
GOBP_POSITIVE_REGULATION_OF_T_HELPER_2_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_T_HELPER_2_CELL_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_GROWTH_HORMONE_SECRETION, GOBP_POSITIVE_REGULATION_OF_GROWTH_HORMONE_SECRETION
GOBP_PROTEIN_DEUBIQUITINATION_INVOLVED_IN_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_PROTEIN_DEUBIQUITINATION_INVOLVED_IN_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_
GOBP_CELLULAR_RESPONSE_TO_LOW_DENSITY_LIPOPROTEIN_PARTICLE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_LOW_DENSITY_LIPOPROTEIN_PARTICLE_STIMULUS
 FARMER_BREAST_CANCER_CLUSTER_7, FARMER_BREAST_CANCER_CLUSTER_7
WP_ID_SIGNALING_PATHWAY, WP_ID_SIGNALING_PATHWAY
REACTOME_N_GLYCAN_ANTENNAE_ELONGATION_IN_THE_MEDIAL_TRANS_GOLGI, REACTOME_N_GLYCAN_ANTENNAE_ELONGATION_IN_THE_MEDIAL_TRANS_GOLGI
GOBP_RESPONSE_TO_VITAMIN_E, GOBP_RESPONSE_TO_VITAMIN_E
BAHD1_TARGET_GENES, BAHD1_TARGET_GENES
GOBP_SIALIC_ACID_TRANSPORT, GOBP_SIALIC_ACID_TRANSPORT
GOMF_SIALIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_SIALIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_NMDA_GLUTAMATE_RECEPTOR_ACTIVITY, GOMF_NMDA_GLUTAMATE_RECEPTOR_ACTIVITY
REACTOME_RUNX3_REGULATES_YAP1_MEDIATED_TRANSCRIPTION, REACTOME_RUNX3_REGULATES_YAP1_MEDIATED_TRANSCRIPTION
YAMANAKA_GLIOBLASTOMA_SURVIVAL_DN, YAMANAKA_GLIOBLASTOMA_SURVIVAL_DN
GOBP_GOLGI_RIBBON_FORMATION, GOBP_GOLGI_RIBBON_FORMATION
GOMF_BMP_BINDING, GOMF_BMP_BINDING
GOBP_POSITIVE_REGULATION_OF_ANTIGEN_RECEPTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_ANTIGEN_RECEPTOR_MEDIATED_SIGNALING_PATHWAY
GOCC_ESCRT_I_COMPLEX, GOCC_ESCRT_I_COMPLEX
GOBP_COLLAGEN_BIOSYNTHETIC_PROCESS, GOBP_COLLAGEN_BIOSYNTHETIC_PROCESS
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L\_MEMORY\_VS\_NAIVE\_CD8\_TCELL\_UP, GSE23321\_CD8\_STEM\_CELL\_MEMORY\_VS\_NAIVE\_CD8\_TCELL\_UP

SASAKI\_TARGETS\_OF\_TP73\_AND\_TP63, SASAKI\_TARGETS\_OF\_TP73\_AND\_TP63 GOBP\_POSITIVE\_REGULATION\_OF\_ENDOTHELIAL\_CELL\_DEVELOPMENT, GOBP\_POSITIVE\_REGULATION\_OF\_ENDOTHELIAL\_CELL\_DEVELOPMENT GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT\_PEPTIDE\_BONDS\_IN\_LINEAR\_AMIDINES, GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT\_PEPTIDE\_BONDS\_I REACTOME\_RUNX3\_REGULATES\_IMMUNE\_RESPONSE\_AND\_CELL\_MIGRATION, REACTOME\_RUNX3\_REGULATES\_IMMUNE\_RESPONSE\_AND\_CELL\_MIGRATION

GOMF\_MELANOCORTIN\_RECEPTOR\_BINDING, GOMF\_MELANOCORTIN\_RECEPTOR\_BINDING GOBP\_POSITIVE\_REGULATION\_OF\_CD8\_POSITIVE\_ALPHA\_BETA\_T\_CELL\_DIFFERENTIATION, GOBP\_POSITIVE\_REGULATION\_OF\_CD8\_POSITIVE\_ALPHA\_BETA\_T\_CELL\_DIFFERENTIATION GOBP\_DENDRITE\_ARBORIZATION, GOBP\_DENDRITE\_ARBORIZATION GOBP\_ARGININE\_CATABOLIC\_PROCESS, GOBP\_ARGININE\_CATABOLIC\_PROCESS GOBP\_POSITIVE\_REGULATION\_OF\_ATP\_BIOSYNTHETIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_ATP\_BIOSYNTHETIC\_PROCESS

GOBP\_NEGATIVE\_REGULATION\_OF\_CD4\_POSITIVE\_ALPHA\_BETA\_T\_CELL\_ACTIVATION, GOBP\_NEGATIVE\_REGULATION\_OF\_CD4\_POSITIVE\_ALPHA\_BETA\_T\_CELL\_ACTIVATION GOBP\_TYPE\_2\_IMMUNE\_RESPONSE, GOBP\_TYPE\_2\_IMMUNE\_RESPONSE GOBP\_T\_HELPER\_2\_CELL\_DIFFERENTIATION, GOBP\_T\_HELPER\_2\_CELL\_DIFFERENTIATION WP\_GENE\_REGULATORY\_NETWORK\_MODELLING\_SOMITOGENESIS, WP\_GENE\_REGULATORY\_NETWORK\_MODELLING\_SOMITOGENESIS HOLLERN SOLID NODULAR BREAST TUMOR UP, HOLLERN SOLID NODULAR BREAST TUMOR UP

GOBP\_MYOBLAST\_FUSION\_INVOLVED\_IN\_SKELETAL\_MUSCLE\_REGENERATION, GOBP\_MYOBLAST\_FUSION\_INVOLVED\_IN\_SKELETAL\_MUSCLE\_REGENERATION

GOBP T HELPER 1 CELL DIFFERENTIATION, GOBP T HELPER 1 CELL DIFFERENTIATION GOBP\_MATING\_BEHAVIOR, GOBP\_MATING\_BEHAVIOR GOBP\_REGULATION\_OF\_COLLAGEN\_METABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_COLLAGEN\_METABOLIC\_PROCESS BANDRES\_RESPONSE\_TO\_CARMUSTIN\_WITHOUT\_MGMT\_24HR\_DN, BANDRES\_RESPONSE\_TO\_CARMUSTIN\_WITHOUT\_MGMT\_24HR\_DN GAO\_LARGE\_INTESTINE\_24W\_C4\_PROM1LOW\_PROGENITOR, GAO\_LARGE\_INTESTINE\_24W\_C4\_PROM1LOW\_PROGENITOR

GOMF\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_COACTIVATOR\_BINDING, GOMF\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_COACTIVATOR\_BINDING

GOBP\_POSITIVE\_REGULATION\_OF\_COLLAGEN\_METABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_COLLAGEN\_METABOLIC\_PROCESS

GOMF\_GLUTAMATE\_GATED\_CALCIUM\_ION\_CHANNEL\_ACTIVITY, GOMF\_GLUTAMATE\_GATED\_CALCIUM\_ION\_CHANNEL\_ACTIVITY

HP\_ABNORMAL\_NASAL\_MUCUS\_SECRETION, HP\_ABNORMAL\_NASAL\_MUCUS\_SECRETION

HP\_SPARSE\_BODY\_HAIR, HP\_SPARSE\_BODY\_HAIR

HP\_FOCAL\_SENSORY\_SEIZURE, HP\_FOCAL\_SENSORY\_SEIZURE

GOBP\_NEGATIVE\_REGULATION\_OF\_T\_HELPER\_1\_TYPE\_IMMUNE\_RESPONSE, GOBP\_NEGATIVE\_REGULATION\_OF\_T\_HELPER\_1\_TYPE\_IMMUNE\_RESPONSE GOMF\_CARBOHYDRATE\_PROTON\_SYMPORTER\_ACTIVITY, GOMF\_CARBOHYDRATE\_PROTON\_SYMPORTER\_ACTIVITY GOBP\_REGULATION\_OF\_CD8\_POSITIVE\_ALPHA\_BETA\_T\_CELL\_DIFFERENTIATION, GOBP\_REGULATION\_OF\_CD8\_POSITIVE\_ALPHA\_BETA\_T\_CELL\_DIFFERENTIATION GTF3C1\_TARGET\_GENES, GTF3C1\_TARGET\_GENES GOBP\_INTERLEUKIN\_4\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_INTERLEUKIN\_4\_MEDIATED\_SIGNALING\_PATHWAY chr1q25, chr1q25 MODULE\_560, MODULE\_560

GOBP\_MYOBLAST\_FUSION, GOBP\_MYOBLAST\_FUSION GOMF\_INTERLEUKIN\_17\_RECEPTOR\_ACTIVITY, GOMF\_INTERLEUKIN\_17\_RECEPTOR\_ACTIVITY GOBP REGULATION OF GERMINAL CENTER FORMATION, GOBP REGULATION OF GERMINAL CENTER FORMATION

GOBP\_NEGATIVE\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_4\_SIGNALING\_PATHWAY, GOBP\_NEGATIVE\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_4\_SIGNALING\_PATHWAY BIOCARTA NGF PATHWAY, BIOCARTA NGF PATHWAY GOBP\_REGULATION\_OF\_T\_HELPER\_1\_CELL\_DIFFERENTIATION, GOBP\_REGULATION\_OF\_T\_HELPER\_1\_CELL\_DIFFERENTIATION GOBP\_POSITIVE\_REGULATION\_OF\_TYPE\_2\_IMMUNE\_RESPONSE, GOBP\_POSITIVE\_REGULATION\_OF\_TYPE\_2\_IMMUNE\_RESPONSE GOCC\_CALYX\_OF\_HELD, GOCC\_CALYX\_OF\_HELD

GOBP\_NEGATIVE\_REGULATION\_OF\_ACUTE\_INFLAMMATORY\_RESPONSE, GOBP\_NEGATIVE\_REGULATION\_OF\_ACUTE\_INFLAMMATORY\_RESPONSE

GOBP\_REGULATION\_OF\_TYPE\_2\_IMMUNE\_RESPONSE, GOBP\_REGULATION\_OF\_TYPE\_2\_IMMUNE\_RESPONSE GOBP NEGATIVE REGULATION OF ACUTE INFLAMMATORY RESPONSE TO ANTIGENIC STIMULUS, GOBP NEGATIVE REGULATION OF ACUTE INFLAMMATORY RESPONSE TO ANTIGENIC STIMULUS HP\_SPONTANEOUS\_PAIN\_SENSATION, HP\_SPONTANEOUS\_PAIN\_SENSATION GOBP\_NEGATIVE\_REGULATION\_OF\_NITRIC\_OXIDE\_SYNTHASE\_BIOSYNTHETIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_NITRIC\_OXIDE\_SYNTHASE\_BIOSYNTHETIC\_PROCESS GOBP PROTEIN K29 LINKED DEUBIQUITINATION, GOBP PROTEIN K29 LINKED DEUBIQUITINATION

GOBP\_CELLULAR\_RESPONSE\_TO\_AMMONIUM\_ION, GOBP\_CELLULAR\_RESPONSE\_TO\_AMMONIUM\_ION GOBP\_VACUOLE\_FUSION, GOBP\_VACUOLE\_FUSION GOBP\_RETINAL\_CELL\_APOPTOTIC\_PROCESS, GOBP\_RETINAL\_CELL\_APOPTOTIC\_PROCESS

REACTOME\_ELECTRIC\_TRANSMISSION\_ACROSS\_GAP\_JUNCTIONS, REACTOME\_ELECTRIC\_TRANSMISSION\_ACROSS\_GAP\_JUNCTIONS IYENGAR\_RESPONSE\_TO\_ADIPOCYTE\_FACTORS, IYENGAR\_RESPONSE\_TO\_ADIPOCYTE\_FACTORS MIR6090, MIR6090 SCHERER\_PBMC\_YF\_VAX\_AGE\_18\_40YO\_ANYD\_UP, SCHERER\_PBMC\_YF\_VAX\_AGE\_18\_40YO\_ANYD\_UP

GOBP\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_2\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_2\_SIGNALING\_PATHWAY

GOBP\_HEPARAN\_SULFATE\_PROTEOGLYCAN\_BIOSYNTHETIC\_PROCESS\_ENZYMATIC\_MODIFICATION, GOBP\_HEPARAN\_SULFATE\_PROTEOGLYCAN\_BIOSYNTHETIC\_PROCESS\_ENZYMATIC\_MODIFICATION GOBP\_NEGATIVE\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_2\_SIGNALING\_PATHWAY, GOBP\_NEGATIVE\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_2\_SIGNALING\_PATHWAY MIR1973, MIR1973 HP\_SEESAW\_NYSTAGMUS, HP\_SEESAW\_NYSTAGMUS GOBP\_POSITIVE\_REGULATION\_OF\_ALCOHOL\_BIOSYNTHETIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_ALCOHOL\_BIOSYNTHETIC\_PROCESS

MIR6795 3P, MIR6795 3P GOBP\_NEGATIVE\_REGULATION\_OF\_BONE\_RESORPTION, GOBP\_NEGATIVE\_REGULATION\_OF\_BONE\_RESORPTION KRAS.50 UP.V1 DN, KRAS.50 UP.V1 DN HP\_GAZE\_EVOKED\_HORIZONTAL\_NYSTAGMUS, HP\_GAZE\_EVOKED\_HORIZONTAL\_NYSTAGMUS GOBP\_POSITIVE\_REGULATION\_OF\_T\_HELPER\_2\_CELL\_CYTOKINE\_PRODUCTION, GOBP\_POSITIVE\_REGULATION\_OF\_T\_HELPER\_2\_CELL\_CYTOKINE\_PRODUCTION

REACTOME\_METABOLISM\_OF\_STEROID\_HORMONES, REACTOME\_METABOLISM\_OF\_STEROID\_HORMONES

HP\_ABNORMAL\_VESTIBULO\_OCULAR\_REFLEX, HP\_ABNORMAL\_VESTIBULO\_OCULAR\_REFLEX

GOBP REGULATION OF PRESYNAPTIC MEMBRANE POTENTIAL, GOBP REGULATION OF PRESYNAPTIC MEMBRANE POTENTIAL HP\_LEYDIG\_CELL\_NEOPLASIA, HP\_LEYDIG\_CELL\_NEOPLASIA GOBP OSTEOCLAST PROLIFERATION, GOBP OSTEOCLAST PROLIFERATION HP\_FACIAL\_PARALYSIS, HP\_FACIAL\_PARALYSIS FINAK\_BREAST\_CANCER\_SDPP\_SIGNATURE, FINAK\_BREAST\_CANCER\_SDPP\_SIGNATURE

GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_MATURATION, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_MATURATION

GOBP\_NEGATIVE\_REGULATION\_OF\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION, GOBP\_NEGATIVE\_REGULATION\_OF\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION GOBP NEGATIVE REGULATION OF EXCITATORY POSTSYNAPTIC POTENTIAL, GOBP NEGATIVE REGULATION OF EXCITATORY POSTSYNAPTIC POTENTIAL