

GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE37416_0H_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN, GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN
GSE45365_WT_VS_IFNAR_KO_CD8A_DC_DN, GSE45365_WT_VS_IFNAR_KO_CD8A_DC_DN
GSE22140_HEALTHY_VS_ARTHRITIC_GERMFREE_MOUSE_CD4_TCELL_DN, GSE22140_HEALTHY_VS_ARTHRITIC_GERMFREE_MOUSE_CD4_TCELL_DN
GSE22886_DAY0_VS_DAY1_MONOCYTE_IN_CULTURE_DN, GSE22886_DAY0_VS_DAY1_MONOCYTE_IN_CULTURE_DN
GSE14769_UNSTIM_VS_40MIN_LPS_BMDM_DN, GSE14769_UNSTIM_VS_40MIN_LPS_BMDM_DN
GSE7219_WT_VS_NIK_NFKB2_KO_DC_UP, GSE7219_WT_VS_NIK_NFKB2_KO_DC_UP
GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN, GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN
GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP
GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_24H_DN, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_24H_DN
GSE9960_GRAM_NEG_VS_GRAM_POS_SEPSIS_PPMC_UP, GSE9960_GRAM_NEG_VS_GRAM_POS_SEPSIS_PPMC_UP
GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_2H_UP, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_2H_UP
GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_UP, GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_UP
GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_DN
GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PPMC_DN, GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PPMC_DN
HALLMARK_HYPOXIA, HALLMARK_HYPOXIA
REACTOME_RAB_REGULATION_OF_TRAFFICKING, REACTOME_RAB_REGULATION_OF_TRAFFICKING
MIR4436B_5P, MIR4436B_5P
GSE17721_0.5H_VS_4H_CPG_BMDC_DN, GSE17721_0.5H_VS_4H_CPG_BMDC_DN
OKUMURA_INFLAMMATORY_RESPONSE_LPS, OKUMURA_INFLAMMATORY_RESPONSE_LPS
GSE8515_IL1_VS_IL6_4H_STIM_MAC_UP, GSE8515_IL1_VS_IL6_4H_STIM_MAC_UP
GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_UP, GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_UP
GOMF_ISOMERASE_ACTIVITY, GOMF_ISOMERASE_ACTIVITY
GSE9988_ANTL_TREM1_AND_LPS_VS_CTRL_TREATED_MONOCYTES_UP, GSE9988_ANTL_TREM1_AND_LPS_VS_CTRL_TREATED_MONOCYTES_UP
GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE37605_FOXP3_FUSION_GFP_VS_IRES_GFP_TREG_C57BL6_UP, GSE37605_FOXP3_FUSION_GFP_VS_IRES_GFP_TREG_C57BL6_UP
GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE2706_2H_VS_8H_R848_AND_LPS_STIM_DC_DN, GSE2706_2H_VS_8H_R848_AND_LPS_STIM_DC_DN
GSE21360_SECONDARY_VS_QUATERNARY_MEMORY_CD8_TCELL_DN, GSE21360_SECONDARY_VS_QUATERNARY_MEMORY_CD8_TCELL_DN
GROSS_HYPOXIA_VIA_ELK3_DN, GROSS_HYPOXIA_VIA_ELK3_DN
GSE360_CTRL_VS_T_GONDIL_DC_DN, GSE360_CTRL_VS_T_GONDIL_DC_DN
GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_DC_UP, GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_DC_UP
MODULE_97, MODULE_97
GSE27241_CTRL_VS_DIGOXIN_TREATED_CD4_TCELL_IN_THI17_POLARIZING_CONDITIONS_UP, GSE27241_CTRL_VS_DIGOXIN_TREATED_CD4_TCELL_IN_THI17_POLARIZING_CONDITIONS_UP
TRAVAGLINI_LUNG_LIPOFIBROBLAST_CELL, TRAVAGLINI_LUNG_LIPOFIBROBLAST_CELL
MODULE_182, MODULE_182
ELVIDGE_HYPOXIA_UP, ELVIDGE_HYPOXIA_UP
GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP, GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP
GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_2H_UP, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_2H_UP
GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_UP
GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_DN, GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_DN
ELVIDGE_HYPOXIA_BY_DMOG_UP, ELVIDGE_HYPOXIA_BY_DMOG_UP
GSE13484_12H_UNSTIM_VS_YF17D_VACCINE_STIM_PPMC_DN, GSE13484_12H_UNSTIM_VS_YF17D_VACCINE_STIM_PPMC_DN
GSE2706_UNSTIM_VS_2H_LPS_DC_DN, GSE2706_UNSTIM_VS_2H_LPS_DC_DN
MIR4713_5P, MIR4713_5P
TGFB_UP.V1_UP, TGFB_UP.V1_UP
MIR3944_5P, MIR3944_5P
GSE7218_IGM_VS_JGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_DN, GSE7218_IGM_VS_JGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_DN
GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP, GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP
GSE35825_UNTREATED_VS_IFNA_STIM_MACROPHAGE_DN, GSE35825_UNTREATED_VS_IFNA_STIM_MACROPHAGE_DN
GSE19374_UNINF_VS_LISTERIA_INFECTED_MACROPHAGE_DN, GSE19374_UNINF_VS_LISTERIA_INFECTED_MACROPHAGE_DN
GSE3039_CD4_TCELL_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_CD4_TCELL_VS_ALPHABETA_CD8_TCELL_UP
GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_UP, GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_UP
GSE9946_IMMATURE_VS_MATURE_STIMULATORY_DC_UP, GSE9946_IMMATURE_VS_MATURE_STIMULATORY_DC_UP
MODULE_358, MODULE_358
GSE41176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_3H_UP, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_3H_UP
GSE17721_12H_VS_24H_CPG_BMDC_UP, GSE17721_12H_VS_24H_CPG_BMDC_UP
MIR4712_5P, MIR4712_5P
GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_DN, GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_DN
KEGG_B_CELL_RECEPTOR_SIGNALING_PATHWAY, KEGG_B_CELL_RECEPTOR_SIGNALING_PATHWAY
MIR376C_3P, MIR376C_3P
MIR770_5P, MIR770_5P
GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCYTES_DN, GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCYTES_DN
XBP1_01, XBP1_01
KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY, KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY
KEGG_PROGESTERONE_MEDIATED_OOCYTE_MATURATION, KEGG_PROGESTERONE_MEDIATED_OOCYTE_MATURATION
GOMF_SIGNALING_ADAPTOR_ACTIVITY, GOMF_SIGNALING_ADAPTOR_ACTIVITY
TGCAAAC_MIR452, TGCAAAC_MIR452
GSE6259_33D1_POS_VS_DEC205_POS_FLT3L_INDUCED_SPLENIC_DC_UP, GSE6259_33D1_POS_VS_DEC205_POS_FLT3L_INDUCED_SPLENIC_DC_UP
GSE3920_UNTREATED_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_UNTREATED_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP
SANA_TNF_SIGNALING_UP, SANA_TNF_SIGNALING_UP
PAX3_B, PAX3_B
FREACT_01, FREACT_01
KEGG_SMALL_CELL_LUNG_CANCER, KEGG_SMALL_CELL_LUNG_CANCER
PID_BCR_5PATHWAY, PID_BCR_5PATHWAY
GTTNYNNGGTNA_UNKNOWN, GTTNYNNGGTNA_UNKNOWN
CHANDRAN_METASTASIS_TOP50_DN, CHANDRAN_METASTASIS_TOP50_DN
CATRRAGC_UNKNOWN, CATRRAGC_UNKNOWN
MODULE_456, MODULE_456
GSE15324_ELF4_KO_VS_WT_NAIVE_CD8_TCELL_DN, GSE15324_ELF4_KO_VS_WT_NAIVE_CD8_TCELL_DN
GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_RAG2_KO_NK_CELL_UP, GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_RAG2_KO_NK_CELL_UP
AMIT_SERUM_RESPONSE_60_MCF10A, AMIT_SERUM_RESPONSE_60_MCF10A
WP_GLUCOCORTICOID_RECEPTOR_PATHWAY, WP_GLUCOCORTICOID_RECEPTOR_PATHWAY
PRAMOONJAGO_SOX4_TARGETS_UP, PRAMOONJAGO_SOX4_TARGETS_UP
ONDER_CDH1_SIGNALING_VIA_CTNNB1, ONDER_CDH1_SIGNALING_VIA_CTNNB1
KEGG_T_CELL_RECEPTOR_SIGNALING_PATHWAY, KEGG_T_CELL_RECEPTOR_SIGNALING_PATHWAY
MIR6836_3P, MIR6836_3P
BERENJENO_TRANSFORMED_BY_RHOA_REVERSIBLY_DN, BERENJENO_TRANSFORMED_BY_RHOA_REVERSIBLY_DN
FRIDMAN_IMMORTALIZATION_DN, FRIDMAN_IMMORTALIZATION_DN
LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_DN, LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_DN
TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_DN
AMIT_EGF_RESPONSE_60_MCF10A, AMIT_EGF_RESPONSE_60_MCF10A
KASLER_HDAC7_TARGETS_2_DN, KASLER_HDAC7_TARGETS_2_DN
MIR3659, MIR3659
GOBP_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT, GOBP_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT
MIR7161_3P, MIR7161_3P
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_YELLOW_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_YELLOW_UP
GOBP_AMINO_ACID_IMPORT, GOBP_AMINO_ACID_IMPORT
BROWNE_HCMV_INFECTION_2HR_UP, BROWNE_HCMV_INFECTION_2HR_UP
GOMF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY, GOMF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY
AMIT_SERUM_RESPONSE_40_MCF10A, AMIT_SERUM_RESPONSE_40_MCF10A
YEGNASUBRAMANIAN_PROSTATE_CANCER, YEGNASUBRAMANIAN_PROSTATE_CANCER
LINDSTEDT_DENDRITIC_CELL_MATURATION_A, LINDSTEDT_DENDRITIC_CELL_MATURATION_A
GOMF_DEMETHYLASE_ACTIVITY, GOMF_DEMETHYLASE_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_NF_KAPPA_B_TRANSCRIPTION_FACTOR_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_NF_KAPPA_B_TRANSCRIPTION_FACTOR_ACTIVITY
GHANDHL_BYSTANDER_IRRADIATION_UP, GHANDHL_BYSTANDER_IRRADIATION_UP
GOMF_NEUTRAL_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_NEUTRAL_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
WP_VITAMIN_DSENSITIVE_CALCIIUM_SIGNALING_IN_DEPRESSION, WP_VITAMIN_DSENSITIVE_CALCIIUM_SIGNALING_IN_DEPRESSION
CROONQUIST_STROMAL_STIMULATION_UP, CROONQUIST_STROMAL_STIMULATION_UP
ACAWYAAAG_UNKNOWN, ACAWYAAAG_UNKNOWN
REACTOME_TRANSLATION_OF_REPLICASE_AND_ASSEMBLY_OF_THE_REPLICATION_TRANSCRIPTION_COMPLEX, REACTOME_TRANSLATION_OF_REPLICASE_AND_ASSEMBLY_OF_THE_REPLICATION_TRANSCRIPTION_COMPLEX
RNTCANNRRNNYNATTW_UNKNOWN, RNTCANNRRNNYNATTW_UNKNOWN
MULLIGAN_NTF3_SIGNALING_VIA_INSR_AND_IGF1R_DN, MULLIGAN_NTF3_SIGNALING_VIA_INSR_AND_IGF1R_DN
GOBP_REGULATION_OF_LEUKOCYTE_CHEMOTAXIS, GOBP_REGULATION_OF_LEUKOCYTE_CHEMOTAXIS
IL15_UP.V1_DN, IL15_UP.V1_DN
MIR490_5P, MIR490_5P
GOCC_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK, GOCC_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK
KRAS.BREAST_UP.V1_UP, KRAS.BREAST_UP.V1_UP
HP_ABNORMALITY_OF_THE_COMMON_COAGULATION_PATHWAY, HP_ABNORMALITY_OF_THE_COMMON_COAGULATION_PATHWAY
SHIPP_DLCL_CURED_VS_FATAL_UP, SHIPP_DLCL_CURED_VS_FATAL_UP
MIR4323, MIR4323
GOBP_REGULATION_OF_NERVOUS_SYSTEM_PROCESS, GOBP_REGULATION_OF_NERVOUS_SYSTEM_PROCESS
GOCC_INTEGRATOR_COMPLEX, GOCC_INTEGRATOR_COMPLEX
GOMF_AROMATIC_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AROMATIC_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_NUCLEOSIDE_MONOPHOSPHATE_KINASE_ACTIVITY, GOMF_NUCLEOSIDE_MONOPHOSPHATE_KINASE_ACTIVITY
GOBP_AROMATIC_AMINO_ACID_TRANSPORT, GOBP_AROMATIC_AMINO_ACID_TRANSPORT
PID_TCR_RAS_PATHWAY, PID_TCR_RAS_PATHWAY
GOBP_POSITIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT
GENTLES_LEUKEMIC_STEM_CELL_UP, GENTLES_LEUKEMIC_STEM_CELL_UP
GOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE, GOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE
GOBP_ALANINE_TRANSPORT, GOBP_ALANINE_TRANSPORT
HP_DIABETES_INSPIDUS, HP_DIABETES_INSPIDUS
WANG_TNF_TARGETS, WANG_TNF_TARGETS
GOBP_AMELOGENESIS, GOBP_AMELOGENESIS
SCHWAB_TARGETS_OF_BMYB_POLYMORPHIC_VARIANTS_UP, SCHWAB_TARGETS_OF_BMYB_POLYMORPHIC_VARIANTS_UP
DAZARD_UV_RESPONSE_CLUSTER_G28, DAZARD_UV_RESPONSE_CLUSTER_G28
HP_EXCESSIVE_DAYTIME_SOMNOLENCE, HP_EXCESSIVE_DAYTIME_SOMNOLENCE
CHO_NR4A1_TARGETS, CHO_NR4A1_TARGETS
KRAS.LUNG.BREAST_UP.V1_UP, KRAS.LUNG.BREAST_UP.V1_UP
HINATA_NFKB_MATRIX, HINATA_NFKB_MATRIX
GOBP_NUCLEOSIDE_MONOPHOSPHATE_PHOSPHORYLATION, GOBP_NUCLEOSIDE_MONOPHOSPHATE_PHOSPHORYLATION
DAZARD_UV_RESPONSE_CLUSTER_G4, DAZARD_UV_RESPONSE_CLUSTER_G4
MOREIRA_RESPONSE_TO_TSA_DN, MOREIRA_RESPONSE_TO_TSA_DN
GOBP_AUTOPHAGY_OF_NUCLEUS, GOBP_AUTOPHAGY_OF_NUCLEUS
GOBP_ARTERY_SMOOTH_MUSCLE_CONTRACTION, GOBP_ARTERY_SMOOTH_MUSCLE_CONTRACTION
HP_HIP_SUBLUXATION, HP_HIP_SUBLUXATION
GOBP_POSITIVE_REGULATION_OF_COAGULATION, GOBP_POSITIVE_REGULATION_OF_COAGULATION
HP_CONCAVE_NASAL_RIDGE, HP_CONCAVE_NASAL_RIDGE
REACTOME_PROLONGED_ERK_ACTIVATION_EVENTS, REACTOME_PROLONGED_ERK_ACTIVATION_EVENTS
GOBP_POSITIVE_REGULATION_OF_INTEGRIN_MEDIATED_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_INTEGRIN_MEDIATED_SIGNALING_PATHWAY
AMIT_EGF_RESPONSE_20_HELA, AMIT_EGF_RESPONSE_20_HELA
GOBP_NEGATIVE_REGULATION_OF_TRANSMISSION_OF_NERVE_IMPULSE, GOBP_NEGATIVE_REGULATION_OF_TRANSMISSION_OF_NERVE_IMPULSE
GARCIA_PINERES_PPMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTIBODY_RESPONSE_NEGATIVE, GARCIA_PINERES_PPMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTIBODY_RESPONSE_NEGATIVE
HP_MENORRHAGIA, HP_MENORRHAGIA
GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS, GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS
GOBP_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE, GOBP_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE
YIH_RESPONSE_TO_ARSENITE_C5, YIH_RESPONSE_TO_ARSENITE_C5
GOBP_D_AMINO_ACID_TRANSPORT, GOBP_D_AMINO_ACID_TRANSPORT
GOBP_MESENCHYMAL_TO_EPITHELIAL_TRANSITION_INVOLVED_IN_METANEPHROS_MORPHOGENESIS, GOBP_MESENCHYMAL_TO_EPITHELIAL_TRANSITION_INVOLVED_IN_METANEPHROS_MORPHOGENESIS
GOMF_C_X_C_CHEMOKINE_RECEPTOR_ACTIVITY, GOMF_C_X_C_CHEMOKINE_RECEPTOR_ACTIVITY
GOBP_PROSTANOID_BIOSYNTHETIC_PROCESS, GOBP_PROSTANOID_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_ENDOCRINE_PROCESS, GOBP_REGULATION_OF_ENDOCRINE_PROCESS
GOBP_ACETATE_ESTER_TRANSPORT, GOBP_ACETATE_ESTER_TRANSPORT
HP_HAND_CLENCHING, HP_HAND_CLENCHING
GOBP_NEGATIVE_REGULATION_OF_TRANSLATIONAL_ELONGATION, GOBP_NEGATIVE_REGULATION_OF_TRANSLATIONAL_ELONGATION
GOBP_NEGATIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION, GOBP_NEGATIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION
HP_KERATOSIS_PILARIS, HP_KERATOSIS_PILARIS
GOBP_LEUKOTRIENE_D4_METABOLIC_PROCESS, GOBP_LEUKOTRIENE_D4_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_ORGAN_GROWTH, GOBP_POSITIVE_REGULATION_OF_ORGAN_GROWTH
GOBP_SMOOTH_MUSCLE_CELL_MATRIX_ADHESION, GOBP_SMOOTH_MUSCLE_CELL_MATRIX_ADHESION
WP_SCFA_AND_SKELETAL_MUSCLE_SUBSTRATE_METABOLISM, WP_SCFA_AND_SKELETAL_MUSCLE_SUBSTRATE_METABOLISM
GOMF_GLUTATHIONE_HYDROLASE_ACTIVITY, GOMF_GLUTATHIONE_HYDROLASE_ACTIVITY
HP_PARIETAL_CORTICAL_ATROPHY, HP_PARIETAL_CORTICAL_ATROPHY
GOBP_TYPE_B_PANCREATIC_CELL_PROLIFERATION, GOBP_TYPE_B_PANCREATIC_CELL_PROLIFERATION
GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_STIMULUS, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_STIMULUS
REACTOME_FREE_FATTY_ACID_RECEPTORS, REACTOME_FREE_FATTY_ACID_RECEPTORS
HP_HYPOMINERALIZATION_OF_ENAMEL, HP_HYPOMINERALIZATION_OF_ENAMEL
GOBP_RETINAL_BLOOD_VESSEL_MORPHOGENESIS, GOBP_RETINAL_BLOOD_VESSEL_MORPHOGENESIS
chr12p11, chr12p11
GOBP_ACETYLCHOLINE_METABOLIC_PROCESS, GOBP_ACETYLCHOLINE_METABOLIC_PROCESS
HASEGAWA_TUMORIGENESIS_BY_RET_C634R, HASEGAWA_TUMORIGENESIS_BY_RET_C634R
GOBP_OTOLITH_DEVELOPMENT, GOBP_OTOLITH_DEVELOPMENT