NTI_TREM1_AND_LPS_MONOCYTE_UP, GSE9988_ANTI_TREM1_VS_ANTI_TREM1_AND_LPS_MONOCYTE_UP
GSE9988_LPS_VS_CTRL_TREATED_MONOCYTE_DN, GSE9988_LPS_VS_CTRL_TREATED_MONOCYTE_DN
GSE9988_LOW_LPS_VS_CTRL_TREATED_MONOCYTE_DN, GSE9988_LOW_LPS_VS_CTRL_TREATED_MONOCYTE_DN
GSE9988_ANTI_TREM1_AND_LPS_VS_CTRL_TREATED_MONOCYTES_DN, GSE9988_ANTI_TREM1_AND_LPS_VS_CTRL_TREATED_MONOCYTES_DN
GSE9988_LOW_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN, GSE9988_LOW_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN
GSE9988_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN, GSE9988_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN
GSE9988_ANTI_TREM1_VS_CTRL_TREATED_MONOCYTES_DN, GSE9988_ANTI_TREM1_VS_CTRL_TREATED_MONOCYTES_DN
GSE9988_ANTI_TREM1_AND_LPS_VS_VEHICLE_TREATED_MONOCYTES_DN, GSE9988_ANTI_TREM1_AND_LPS_VS_VEHICLE_TREATED_MONOCYTES_DN
GSE9988_ANTI_TREM1_VS_VEHICLE_TREATED_MONOCYTES_DN, GSE9988_ANTI_TREM1_VS_VEHICLE_TREATED_MONOCYTES_DN
GSE9988_ANTI_TREM1_VS_LPS_MONOCYTE_UP, GSE9988_ANTI_TREM1_VS_LPS_MONOCYTE_UP
GSE9988_ANTI_TREM1_VS_LOW_LPS_MONOCYTE_UP, GSE9988_ANTI_TREM1_VS_LOW_LPS_MONOCYTE_UP
GSE22140_GERMFREE_VS_SPF_MOUSE_CD4_TCELL_DN, GSE22140_GERMFREE_VS_SPF_MOUSE_CD4_TCELL_DN
GSE15330_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_VS_PRO_BCELL_UP, GSE15330_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_VS_PRO_BCELL_UP
GSE9988_ANTI_TREM1_VS_CTRL_TREATED_MONOCYTES_UP, GSE9988_ANTI_TREM1_VS_CTRL_TREATED_MONOCYTES_UP
GSE22196_HEALTHY_VS_OBESE_MOUSE_SKIN_GAMMADELTA_TCELL_DN, GSE22196_HEALTHY_VS_OBESE_MOUSE_SKIN_GAMMADELTA_TCELL_DN
GSE16266_CTRL_VS_LPS_STIM_MEF_UP, GSE16266_CTRL_VS_LPS_STIM_MEF_UP
GSE9988_LOW_LPS_VS_ANTI_TREM1_AND_LPS_MONOCYTE_DN, GSE9988_LOW_LPS_VS_ANTI_TREM1_AND_LPS_MONOCYTE_DN
GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP
GSE9988_ANTI_TREM1_VS_VEHICLE_TREATED_MONOCYTES_UP, GSE9988_ANTI_TREM1_VS_VEHICLE_TREATED_MONOCYTES_UP
GSE9988_LPS_VS_LPS_AND_ANTI_TREM1_MONOCYTE_DN, GSE9988_LPS_VS_LPS_AND_ANTI_TREM1_MONOCYTE_DN
GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_UP
GSE3982_EOSINOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP, GSE3982_EOSINOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP
MORF_PPP6C, MORF_PPP6C
GSE3982_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_UP, GSE3982_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_UP
GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_DN, GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_DN
GSE3039_CD4_TCELL_VS_B1_BCELL_DN, GSE3039_CD4_TCELL_VS_B1_BCELL_DN
GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN, GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN
GSE24634_TEFF_VS_TCONV_DAY7_IN_CULTURE_DN, GSE24634_TEFF_VS_TCONV_DAY7_IN_CULTURE_DN
GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION
GSE3982_EOSINOPHIL_VS_NKCELL_UP, GSE3982_EOSINOPHIL_VS_NKCELL_UP
GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
MODULE_177, MODULE_177
GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_UP, GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_UP
MCBRYAN_PUBERTAL_TGFB1_TARGETS_UP, MCBRYAN_PUBERTAL_TGFB1_TARGETS_UP
GSE17721_LPS_VS_POLYIC_12H_BMDC_DN, GSE17721_LPS_VS_POLYIC_12H_BMDC_DN
GSE34006_UNTREATED_VS_A2AR_AGONIST_TREATED_TREG_DN, GSE34006_UNTREATED_VS_A2AR_AGONIST_TREATED_TREG_DN
GSE3982_EFF_MEMORY_CD4_TCELL_VS_NKCELL_DN, GSE3982_EFF_MEMORY_CD4_TCELL_VS_NKCELL_DN
TRAVAGLINI_LUNG_B_CELL, TRAVAGLINI_LUNG_B_CELL
GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN, GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN
GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN
GSE360_L_DONOVANI_VS_T_GONDII_MAC_UP, GSE360_L_DONOVANI_VS_T_GONDII_MAC_UP
RBM15_TARGET_GENES, RBM15_TARGET_GENES
ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_1DY_DN, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_1DY_DN
MIR5698, MIR5698
GOBP_PROTEIN_MONOUBIQUITINATION, GOBP_PROTEIN_MONOUBIQUITINATION
MCBRYAN_PUBERTAL_BREAST_5_6WK_DN, MCBRYAN_PUBERTAL_BREAST_5_6WK_DN
THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_DN, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_DN
GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP
REACTOME_SIGNALING_BY_NTRKS, REACTOME_SIGNALING_BY_NTRKS
REACTOME_TRANS_GOLGI_NETWORK_VESICLE_BUDDING, REACTOME_TRANS_GOLGI_NETWORK_VESICLE_BUDDING
GREGORY_SYNTHETIC_LETHAL_WITH_IMATINIB, GREGORY_SYNTHETIC_LETHAL_WITH_IMATINIB
GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
GNF2_ICAM3, GNF2_ICAM3
BROWN_MYELOID_CELL_DEVELOPMENT_UP, BROWN_MYELOID_CELL_DEVELOPMENT_UP
GSE17721_4_VS_24H_CPG_BMDC_UP, GSE17721_4_VS_24H_CPG_BMDC_UP
GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_UP
GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MONOCYTE_UP, GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MONOCYTE_UP
MIR423_5P, MIR423_5P
GOBP_HISTONE_UBIQUITINATION, GOBP_HISTONE_UBIQUITINATION
MIR7109_5P, MIR7109_5P
GNF2_SELL, GNF2_SELL
DESCARTES_MAIN_FETAL_MYELOID_CELLS, DESCARTES_MAIN_FETAL_MYELOID_CELLS
REACTOME_GOLGI_ASSOCIATED_VESICLE_BIOGENESIS, REACTOME_GOLGI_ASSOCIATED_VESICLE_BIOGENESIS
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_7, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_7
GSE8515_CTRL_VS_IL1_4H_STIM_MAC_UP, GSE8515_CTRL_VS_IL1_4H_STIM_MAC_UP
MIR3184_5P, MIR3184_5P
GOCC_NEURON_PROJECTION_CYTOPLASM, GOCC_NEURON_PROJECTION_CYTOPLASM
MIR889_5P, MIR889_5P
GOBP_POSITIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBL
GNF2_HLA_C, GNF2_HLA_C
MIR3127_3P, MIR3127_3P
GOCC_AXON_CYTOPLASM, GOCC_AXON_CYTOPLASM
MIR6756_3P, MIR6756_3P
GSE12198_LOW_IL2_STIM_NK_CELL_VS_HIGH_IL2_STIM_NK_CELL_DN, GSE12198_LOW_IL2_STIM_NK_CELL_VS_HIGH_IL2_STIM_NK_CELL_DN
GOBP_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION
GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_UP, GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_UP
GOCC_PCG_PROTEIN_COMPLEX, GOCC_PCG_PROTEIN_COMPLEX
ZAMORA_NOS2_TARGETS_DN, ZAMORA_NOS2_TARGETS_DN
GOBP_AXO_DENDRITIC_TRANSPORT, GOBP_AXO_DENDRITIC_TRANSPORT
GOMF_PROTEIN_SERINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_SERINE_PHOSPHATASE_ACTIVITY
REACTOME_CARGO_RECOGNITION_FOR_CLATHRIN_MEDIATED_ENDOCYTOSIS, REACTOME_CARGO_RECOGNITION_FOR_CLATHRIN_MEDIATED_ENDOCYTOSIS
GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_UP, GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_UP
GSE3982_NEUTROPHIL_VS_BCELL_UP, GSE3982_NEUTROPHIL_VS_BCELL_UP
GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_60DY_UP, GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_60DY_UP
GOBP_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
MIR4687_3P, MIR4687_3P
GNF2_MSN, GNF2_MSN
HP_ANOREXIA, HP_ANOREXIA
GCM_BECN1, GCM_BECN1
GNF2_BNIP2, GNF2_BNIP2
WP_WNT_SIGNALING_PATHWAY_NETPATH, WP_WNT_SIGNALING_PATHWAY_NETPATH
HP_BONE_PAIN, HP_BONE_PAIN
GNF2_CD97, GNF2_CD97
MIR4524B_3P, MIR4524B_3P
GOCC_PHOSPHATASE_COMPLEX, GOCC_PHOSPHATASE_COMPLEX
HP_ABNORMAL_PULMONARY_VALVE_MORPHOLOGY, HP_ABNORMAL_PULMONARY_VALVE_MORPHOLOGY
GOBP_LYSOSOME_LOCALIZATION, GOBP_LYSOSOME_LOCALIZATION
GNATENKO_PLATELET_SIGNATURE, GNATENKO_PLATELET_SIGNATURE
GOBP_PLATELET_ACTIVATION, GOBP_PLATELET_ACTIVATION
GCM_SMARCC1, GCM_SMARCC1
MIR644A, MIR644A
GOBP_REGULATION_OF_VACUOLE_ORGANIZATION, GOBP_REGULATION_OF_VACUOLE_ORGANIZATION
MIR6081, MIR6081
GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP, GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP
GOBP_HISTONE_MONOUBIQUITINATION, GOBP_HISTONE_MONOUBIQUITINATION
GCM_DEAF1, GCM_DEAF1
MIR6884_3P, MIR6884_3P
GOBP_HISTONE_H2A_UBIQUITINATION, GOBP_HISTONE_H2A_UBIQUITINATION
GOMF_PROTEIN_TYROSINE_KINASE_BINDING, GOMF_PROTEIN_TYROSINE_KINASE_BINDING
MIR6755_3P, MIR6755_3P
GOBP_HISTONE_DEUBIQUITINATION, GOBP_HISTONE_DEUBIQUITINATION
GOBP_HOMOTYPIC_CELL_CELL_ADHESION, GOBP_HOMOTYPIC_CELL_CELL_ADHESION
JAZAG_TGFB1_SIGNALING_VIA_SMAD4_UP, JAZAG_TGFB1_SIGNALING_VIA_SMAD4_UP
GOBP_AUTOPHAGOSOME_MATURATION, GOBP_AUTOPHAGOSOME_MATURATION
WP_REGULATION_OF_TOLLLIKE_RECEPTOR_SIGNALING_PATHWAY, WP_REGULATION_OF_TOLLLIKE_RECEPTOR_SIGNALING_PATHWAY
GSE3565_DUSP1_VS_WT_SPLENOCYTES_POST_LPS_INJECTION_DN, GSE3565_DUSP1_VS_WT_SPLENOCYTES_POST_LPS_INJECTION_DN
MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER, MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER
GOBP_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GOBP_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
PID_NOTCH_PATHWAY, PID_NOTCH_PATHWAY
MIR6892_3P, MIR6892_3P
GOBP_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GOBP_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
HP_ABNORMALITY_OF_INCISOR_MORPHOLOGY, HP_ABNORMALITY_OF_INCISOR_MORPHOLOGY
GOBP_ANTEROGRADE_AXONAL_TRANSPORT, GOBP_ANTEROGRADE_AXONAL_TRANSPORT
MIR548Q, MIR548Q
GOBP_CELLULAR_PIGMENTATION, GOBP_CELLULAR_PIGMENTATION
VARELA_ZMPSTE24_TARGETS_UP, VARELA_ZMPSTE24_TARGETS_UP
GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP, GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP
YGACNNYACAR_UNKNOWN, YGACNNYACAR_UNKNOWN
MIR4633_3P_MIR6500_5P, MIR4633_3P_MIR6500_5P
MIR4764_3P, MIR4764_3P
FRASOR_TAMOXIFEN_RESPONSE_UP, FRASOR_TAMOXIFEN_RESPONSE_UP
GOBP_INTERLEUKIN_6_PRODUCTION, GOBP_INTERLEUKIN_6_PRODUCTION
MODULE_226, MODULE_226
MIR5189_3P, MIR5189_3P
GOBP_NATURAL_KILLER_CELL_MEDIATED_IMMUNITY, GOBP_NATURAL_KILLER_CELL_MEDIATED_IMMUNITY
GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_28DY_UP, GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_28DY_UP
RASHI_RESPONSE_TO_IONIZING_RADIATION_6, RASHI_RESPONSE_TO_IONIZING_RADIATION_6
GOBP_SYNAPTIC_VESICLE_LOCALIZATION, GOBP_SYNAPTIC_VESICLE_LOCALIZATION
MIR621, MIR621
GOBP_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GOBP_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
MIR4676_5P, MIR4676_5P
GOBP_PIGMENTATION, GOBP_PIGMENTATION
GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS, GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS
GOBP_VESICLE_TRANSPORT_ALONG_MICROTUBULE, GOBP_VESICLE_TRANSPORT_ALONG_MICROTUBULE
MIR575, MIR575
NAKAMURA_METASTASIS, NAKAMURA_METASTASIS
HP_ABNORMAL_PERICARDIUM_MORPHOLOGY, HP_ABNORMAL_PERICARDIUM_MORPHOLOGY
MIR490_3P, MIR490_3P
GOBP_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS, GOBP_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS
HP_VASCULITIS, HP_VASCULITIS
GOMF_LIPOPROTEIN_PARTICLE_RECEPTOR_ACTIVITY, GOMF_LIPOPROTEIN_PARTICLE_RECEPTOR_ACTIVITY
MIR6821_3P, MIR6821_3P
GOMF_RECEPTOR_TYROSINE_KINASE_BINDING, GOMF_RECEPTOR_TYROSINE_KINASE_BINDING
BIOCARTA_IL2RB_PATHWAY, BIOCARTA_IL2RB_PATHWAY
BIOCARTA_CTCF_PATHWAY, BIOCARTA_CTCF_PATHWAY
MIR515_3P, MIR515_3P
ABRAHAM_ALPC_VS_MULTIPLE_MYELOMA_UP, ABRAHAM_ALPC_VS_MULTIPLE_MYELOMA_UP
GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_NUCLEUS, GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_NUCLEUS
MIR4713_3P, MIR4713_3P
MIR33B_3P, MIR33B_3P
MIR6851_3P, MIR6851_3P
GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_UP, GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_UP
GSE14413_UNSTIM_VS_IFNB_STIM_L929_CELLS_UP, GSE14413_UNSTIM_VS_IFNB_STIM_L929_CELLS_UP
HP_MYOSITIS, HP_MYOSITIS
MIR519E_3P, MIR519E_3P
GOCC_RETROMER_COMPLEX, GOCC_RETROMER_COMPLEX
WP_3Q29_COPY_NUMBER_VARIATION_SYNDROME, WP_3Q29_COPY_NUMBER_VARIATION_SYNDROME
GSE13762_CTRL_VS_125_VITAMIND_DAY5_DC_DN, GSE13762_CTRL_VS_125_VITAMIND_DAY5_DC_DN
GOBP_PLATELET_AGGREGATION, GOBP_PLATELET_AGGREGATION
REACTOME_PLASMA_LIPOPROTEIN_CLEARANCE, REACTOME_PLASMA_LIPOPROTEIN_CLEARANCE
GOBP_REGULATION_OF_ERAD_PATHWAY, GOBP_REGULATION_OF_ERAD_PATHWAY
GOCC_TERTIARY_GRANULE_LUMEN, GOCC_TERTIARY_GRANULE_LUMEN
GNF2_TNFSF10, GNF2_TNFSF10
GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION, GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION
GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION
MIR4763_5P, MIR4763_5P
REACTOME_THROMBIN_SIGNALLING_THROUGH_PROTEINASE_ACTIVATED_RECEPTORS_PARS, REACTOME_THROMBIN_SIGNALLING_THROUGH_PROTEINASE_ACTIVATED_RECEPTORS_PARS
GOBP_SYNAPTIC_VESICLE_TRANSPORT, GOBP_SYNAPTIC_VESICLE_TRANSPORT
WP_COMPUTATIONAL_MODEL_OF_AEROBIC_GLYCOLYSIS, WP_COMPUTATIONAL_MODEL_OF_AEROBIC_GLYCOLYSIS
PID_IL8_CXCR2_PATHWAY, PID_IL8_CXCR2_PATHWAY
MIR3135A, MIR3135A
GOMF_CULLIN_FAMILY_PROTEIN_BINDING, GOMF_CULLIN_FAMILY_PROTEIN_BINDING
DAVIES_MULTIPLE_MYELOMA_VS_MGUS_DN, DAVIES_MULTIPLE_MYELOMA_VS_MGUS_DN
GOLUB_ALL_VS_AML_DN, GOLUB_ALL_VS_AML_DN
GOBP_NEGATIVE_REGULATION_OF_NATURAL_KILLER_CELL_MEDIATED_IMMUNITY, GOBP_NEGATIVE_REGULATION_OF_NATURAL_KILLER_CELL_MEDIATED_IMMUNITY
GOBP_HISTONE_H2A_MONOUBIQUITINATION, GOBP_HISTONE_H2A_MONOUBIQUITINATION
GOMF_TRANSLATION_REPRESSOR_ACTIVITY, GOMF_TRANSLATION_REPRESSOR_ACTIVITY
GOCC_BLOC_COMPLEX, GOCC_BLOC_COMPLEX
HP_PLEURITIS, HP_PLEURITIS
GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_MEDIATED_IMMUNITY, GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_MEDIATED_IMMUNITY
GOCC_BLOC_1_COMPLEX, GOCC_BLOC_1_COMPLEX
GOBP_ARF_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_ARF_PROTEIN_SIGNAL_TRANSDUCTION
HP_DECREASED_CIRCULATING_TOTAL_IGM, HP_DECREASED_CIRCULATING_TOTAL_IGM
HP_GINGIVAL_BLEEDING, HP_GINGIVAL_BLEEDING
PARK_TRETINOIN_RESPONSE_AND_RARA_PLZF_FUSION, PARK_TRETINOIN_RESPONSE_AND_RARA_PLZF_FUSION
REACTOME_SIGNALING_BY_RETINOIC_ACID, REACTOME_SIGNALING_BY_RETINOIC_ACID
HP_ENCEPHALITIS, HP_ENCEPHALITIS
BIOCARTA_GH_PATHWAY, BIOCARTA_GH_PATHWAY
GOBP_POSITIVE_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING, GOBP_POSITIVE_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING
REACTOME_G_PROTEIN_BETA_GAMMA_SIGNALLING, REACTOME_G_PROTEIN_BETA_GAMMA_SIGNALLING
GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_CYTOTOXICITY, GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_CYTOTOXICITY
GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX, GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX
HOXB7_TARGET_GENES, HOXB7_TARGET_GENES
GOCC_PRC1_COMPLEX, GOCC_PRC1_COMPLEX
GOBP_PIGMENT_ACCUMULATION, GOBP_PIGMENT_ACCUMULATION
MIR125A_3P, MIR125A_3P
HP_INCISOR_MACRODONTIA, HP_INCISOR_MACRODONTIA
HP_CHRONIC_INFECTION, HP_CHRONIC_INFECTION
GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY
GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN
GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_FISSION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_FISSION
MIR6872_3P, MIR6872_3P
GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY, GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY
LIAN_LIPA_TARGETS_6M, LIAN_LIPA_TARGETS_6M
GOMF_CLATHRIN_HEAVY_CHAIN_BINDING, GOMF_CLATHRIN_HEAVY_CHAIN_BINDING
GOBP_MITOCHONDRIAL_CALCIUM_ION_HOMEOSTASIS, GOBP_MITOCHONDRIAL_CALCIUM_ION_HOMEOSTASIS
GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
MIR6798_3P, MIR6798_3P
WP_OLIGODENDROCYTE_SPECIFICATION_AND_DIFFERENTIATION_LEADING_TO_MYELIN_COMPONENTS_FOR_CNS, WP_OLIGODENDROCYTE_SPECIFICATION_AND_DIFFERENTIATION_LEADING_TO
GOMF_CIS_TRANS_ISOMERASE_ACTIVITY, GOMF_CIS_TRANS_ISOMERASE_ACTIVITY
GOBP_PIGMENT_GRANULE_MATURATION, GOBP_PIGMENT_GRANULE_MATURATION
HP_POOR_COORDINATION, HP_POOR_COORDINATION
MIR181C_3P, MIR181C_3P
GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS
GSE15659_RESTING_VS_ACTIVATED_TREG_UP, GSE15659_RESTING_VS_ACTIVATED_TREG_UP
REACTOME_NUCLEAR_RECEPTOR_TRANSCRIPTION_PATHWAY, REACTOME_NUCLEAR_RECEPTOR_TRANSCRIPTION_PATHWAY
MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_23, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_23
RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_MODERATELY_DN, RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_MODERATELY_DN
WP_KISSPEPTINKISSPEPTIN_RECEPTOR_SYSTEM_IN_THE_OVARY, WP_KISSPEPTINKISSPEPTIN_RECEPTOR_SYSTEM_IN_THE_OVARY
GOBP_RAP_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_RAP_PROTEIN_SIGNAL_TRANSDUCTION
MIR365B_5P, MIR365B_5P
GNF2_SPI1, GNF2_SPI1
GOBP_RESPONSE_TO_TUMOR_CELL, GOBP_RESPONSE_TO_TUMOR_CELL
REACTOME_GAP_JUNCTION_DEGRADATION, REACTOME_GAP_JUNCTION_DEGRADATION
GOBP_ADAPTATION_OF_SIGNALING_PATHWAY, GOBP_ADAPTATION_OF_SIGNALING_PATHWAY
GOBP_TRIGLYCERIDE_METABOLIC_PROCESS, GOBP_TRIGLYCERIDE_METABOLIC_PROCESS
GOBP_INTERLEUKIN_4_PRODUCTION, GOBP_INTERLEUKIN_4_PRODUCTION
MIR31_3P, MIR31_3P
PID_BETA_CATENIN_DEG_PATHWAY, PID_BETA_CATENIN_DEG_PATHWAY
PID_IL8_CXCR1_PATHWAY, PID_IL8_CXCR1_PATHWAY
PID_BMP_PATHWAY, PID_BMP_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_NEURON_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_NEURON_DIFFERENTIATION
MIR365A_5P, MIR365A_5P
REACTOME_DISASSEMBLY_OF_THE_DESTRUCTION_COMPLEX_AND_RECRUITMENT_OF_AXIN_TO_THE_MEMBRANE, REACTOME_DISASSEMBLY_OF_THE_DESTRUCTION_COMPLEX_AND_RECRUITMEN
GOBP_PIGMENT_CELL_DIFFERENTIATION, GOBP_PIGMENT_CELL_DIFFERENTIATION
LI_PBMC_MENOMUNE_A_C_Y_W_135_AGE_18_45YO_3DY_UP, LI_PBMC_MENOMUNE_A_C_Y_W_135_AGE_18_45YO_3DY_UP
SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN, SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN
GOBP_SECRETORY_GRANULE_ORGANIZATION, GOBP_SECRETORY_GRANULE_ORGANIZATION
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_DN
HP_ABNORMALITY_OF_CIRCULATING_FIBRINOGEN, HP_ABNORMALITY_OF_CIRCULATING_FIBRINOGEN
CAVARD_LIVER_CANCER_MALIGNANT_VS_BENIGN, CAVARD_LIVER_CANCER_MALIGNANT_VS_BENIGN
GOBP_POSITIVE_REGULATION_OF_FILOPODIUM_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_FILOPODIUM_ASSEMBLY
REACTOME_NEF_MEDIATED_DOWNREGULATION_OF_MHC_CLASS_I_COMPLEX_CELL_SURFACE_EXPRESSION, REACTOME_NEF_MEDIATED_DOWNREGULATION_OF_MHC_CLASS_I_COMPLEX_CELL_S
REACTOME_P38MAPK_EVENTS, REACTOME_P38MAPK_EVENTS
GOBP_NEGATIVE_REGULATION_OF_CELL_KILLING, GOBP_NEGATIVE_REGULATION_OF_CELL_KILLING
GOBP_DENDRITIC_CELL_MIGRATION, GOBP_DENDRITIC_CELL_MIGRATION
GCM_RING1, GCM_RING1
PID_ALK1_PATHWAY, PID_ALK1_PATHWAY
MIR6804_5P, MIR6804_5P
GOCC_GLYCOGEN_GRANULE, GOCC_GLYCOGEN_GRANULE
GOBP_BETA_CATENIN_DESTRUCTION_COMPLEX_DISASSEMBLY, GOBP_BETA_CATENIN_DESTRUCTION_COMPLEX_DISASSEMBLY
REACTOME_SIGNALLING_TO_RAS, REACTOME_SIGNALLING_TO_RAS
HP_ABNORMAL_HAIR_WHORL, HP_ABNORMAL_HAIR_WHORL
GOBP_CELLULAR_RESPONSE_TO_ESTROGEN_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_ESTROGEN_STIMULUS
BIOCARTA_SODD_PATHWAY, BIOCARTA_SODD_PATHWAY
HOLLEMAN_DAUNORUBICIN_B_ALL_DN, HOLLEMAN_DAUNORUBICIN_B_ALL_DN
WP_IRON_METABOLISM_IN_PLACENTA, WP_IRON_METABOLISM_IN_PLACENTA
BIOCARTA_PML_PATHWAY, BIOCARTA_PML_PATHWAY
REACTOME_BETA_CATENIN_PHOSPHORYLATION_CASCADE, REACTOME_BETA_CATENIN_PHOSPHORYLATION_CASCADE
GOBP_FATTY_ACID_ELONGATION_SATURATED_FATTY_ACID, GOBP_FATTY_ACID_ELONGATION_SATURATED_FATTY_ACID
GOBP_TRANS_SYNAPTIC_SIGNALING_BY_LIPID, GOBP_TRANS_SYNAPTIC_SIGNALING_BY_LIPID
GOMF_3_OXO_ARACHIDOYL_COA_SYNTHASE_ACTIVITY, GOMF_3_OXO_ARACHIDOYL_COA_SYNTHASE_ACTIVITY
GOBP_NEURON_FATE_COMMITMENT, GOBP_NEURON_FATE_COMMITMENT
chr10q24, chr10q24
GOMF_CALMODULIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOMF_CALMODULIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
MIR517A_3P_MIR517B_3P_MIR517C_3P, MIR517A_3P_MIR517B_3P_MIR517C_3P
HP_NEUROGENIC_BLADDER, HP_NEUROGENIC_BLADDER
GAUSSMANN_MLL_AF4_FUSION_TARGETS_B_DN, GAUSSMANN_MLL_AF4_FUSION_TARGETS_B_DN
HP_MORNING_MYOCLONIC_JERKS, HP_MORNING_MYOCLONIC_JERKS
GOBP_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT, GOBP_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT
REACTOME_G_BETA_GAMMA_SIGNALLING_THROUGH_PI3KGAMMA, REACTOME_G_BETA_GAMMA_SIGNALLING_THROUGH_PI3KGAMMA
HP_ABNORMAL_CIRCULATING_FERRITIN_CONCENTRATION, HP_ABNORMAL_CIRCULATING_FERRITIN_CONCENTRATION
MODULE_454, MODULE_454
GOBP_G_PROTEIN_COUPLED_RECEPTOR_INTERNALIZATION, GOBP_G_PROTEIN_COUPLED_RECEPTOR_INTERNALIZATION
BAHD1_TARGET_GENES, BAHD1_TARGET_GENES
GOMF_EPHRIN_RECEPTOR_BINDING, GOMF_EPHRIN_RECEPTOR_BINDING
HP_ORCHITIS, HP_ORCHITIS
WP_COMMON_PATHWAYS_UNDERLYING_DRUG_ADDICTION, WP_COMMON_PATHWAYS_UNDERLYING_DRUG_ADDICTION
GOCC_INTEGRAL_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_INTEGRAL_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_ENDOPLASMIC_RETICULUM_ME
PID_IL4_2PATHWAY, PID_IL4_2PATHWAY
HP_IMPAIRED_T_CELL_FUNCTION, HP_IMPAIRED_T_CELL_FUNCTION
HP_OSTEOCHONDROSIS, HP_OSTEOCHONDROSIS
GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_APOPTOTIC_PROCESS
REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX_IN_CANCER, REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX_IN_CANCER
GOBP_DEVELOPMENTAL_PIGMENTATION, GOBP_DEVELOPMENTAL_PIGMENTATION
LIU_LIVER_CANCER, LIU_LIVER_CANCER
GOCC_M_BAND, GOCC_M_BAND
YANG_BREAST_CANCER_ESR1_DN, YANG_BREAST_CANCER_ESR1_DN
CHEN_HOXA5_TARGETS_6HR_UP, CHEN_HOXA5_TARGETS_6HR_UP
GOMF_TUMOR_NECROSIS_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, GOMF_TUMOR_NECROSIS_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY
MIR6090, MIR6090
GOBP_POSITIVE_REGULATION_OF_ATP_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_ATP_BIOSYNTHETIC_PROCESS
GOBP_CARNITINE_SHUTTLE, GOBP_CARNITINE_SHUTTLE
GOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY
GOBP_PIGMENT_GRANULE_ORGANIZATION, GOBP_PIGMENT_GRANULE_ORGANIZATION
REACTOME_INTERLEUKIN_2_FAMILY_SIGNALING, REACTOME_INTERLEUKIN_2_FAMILY_SIGNALING
LANDIS_ERBB2_BREAST_PRENEOPLASTIC_UP, LANDIS_ERBB2_BREAST_PRENEOPLASTIC_UP
PID_EPHA_FWDPATHWAY, PID_EPHA_FWDPATHWAY
GOBP_RETROGRADE_TRANS_SYNAPTIC_SIGNALING, GOBP_RETROGRADE_TRANS_SYNAPTIC_SIGNALING
MIR3615, MIR3615
GOMF_ARRESTIN_FAMILY_PROTEIN_BINDING, GOMF_ARRESTIN_FAMILY_PROTEIN_BINDING
GOBP_DENDRITIC_CELL_CHEMOTAXIS, GOBP_DENDRITIC_CELL_CHEMOTAXIS
REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOINOSITIDE_3_KINASE_PI3K, REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOINOSITIDE_3_KINASE_PI3K
GOBP_NEGATIVE_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION, GOBP_NEGATIVE_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION
GOBP_REGULATION_OF_LYSOSOME_ORGANIZATION, GOBP_REGULATION_OF_LYSOSOME_ORGANIZATION
GOMF_1_ALKYL_2_ACETYLGLYCEROPHOSPHOCHOLINE_ESTERASE_ACTIVITY, GOMF_1_ALKYL_2_ACETYLGLYCEROPHOSPHOCHOLINE_ESTERASE_ACTIVITY
SHIPP_DLBCL_VS_FOLLICULAR_LYMPHOMA_DN, SHIPP_DLBCL_VS_FOLLICULAR_LYMPHOMA_DN
ADCYAP1_TARGET_GENES, ADCYAP1_TARGET_GENES
CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_LD_MTX_UP, CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_LD_MTX_UP
REACTOME_REGULATION_OF_KIT_SIGNALING, REACTOME_REGULATION_OF_KIT_SIGNALING
IIZUKA_LIVER_CANCER_PROGRESSION_L0_L1_UP, IIZUKA_LIVER_CANCER_PROGRESSION_L0_L1_UP
GOMF_DEATH_RECEPTOR_ACTIVITY, GOMF_DEATH_RECEPTOR_ACTIVITY
GOMF_CALCIUM_CATION_ANTIPORTER_ACTIVITY, GOMF_CALCIUM_CATION_ANTIPORTER_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION, GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION
GOBP_PEPTIDYL_ARGININE_N_METHYLATION, GOBP_PEPTIDYL_ARGININE_N_METHYLATION
GOMF_PROTEIN_ARGININE_N_METHYLTRANSFERASE_ACTIVITY, GOMF_PROTEIN_ARGININE_N_METHYLTRANSFERASE_ACTIVITY
GOBP_THREONINE_METABOLIC_PROCESS, GOBP_THREONINE_METABOLIC_PROCESS
GOBP_IMMUNOGLOBULIN_V_D_J_RECOMBINATION, GOBP_IMMUNOGLOBULIN_V_D_J_RECOMBINATION
GOBP_HISTONE_H2A_K63_LINKED_UBIQUITINATION, GOBP_HISTONE_H2A_K63_LINKED_UBIQUITINATION
GOBP_NEGATIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION
WP_REGUCALCIN_IN_PROXIMAL_TUBULE_EPITHELIAL_KIDNEY_CELLS, WP_REGUCALCIN_IN_PROXIMAL_TUBULE_EPITHELIAL_KIDNEY_CELLS
HP_PULMONARY_ARTERY_DILATATION, HP_PULMONARY_ARTERY_DILATATION
GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY, GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY
HP_ABNORMAL_MAST_CELL_MORPHOLOGY, HP_ABNORMAL_MAST_CELL_MORPHOLOGY
GOBP_NEGATIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_CALCIUM_ION_CONCENTRATION, GOBP_NEGATIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_CALCIUM_ION_CONCENTRATIO
GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE_MEMBRANE, GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE_MEMBRANE
HP_SCHIZOPHRENIA, HP_SCHIZOPHRENIA
GOMF_PHOSPHATIDYLINOSITOL_KINASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_KINASE_ACTIVITY
REACTOME_SODIUM_CALCIUM_EXCHANGERS, REACTOME_SODIUM_CALCIUM_EXCHANGERS
REACTOME_INTERLEUKIN_6_SIGNALING, REACTOME_INTERLEUKIN_6_SIGNALING
GOBP_LYSOSOMAL_LUMEN_ACIDIFICATION, GOBP_LYSOSOMAL_LUMEN_ACIDIFICATION
GOBP_COENZYME_A_BIOSYNTHETIC_PROCESS, GOBP_COENZYME_A_BIOSYNTHETIC_PROCESS
GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATORY_SUBUNIT_BINDING, GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATORY_SUBUNIT_BINDING
MATZUK_PREOVULATORY_FOLLICLE, MATZUK_PREOVULATORY_FOLLICLE
GOCC_DIHYDROLIPOYL_DEHYDROGENASE_COMPLEX, GOCC_DIHYDROLIPOYL_DEHYDROGENASE_COMPLEX
GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE, GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE
WP_THIAMINE_METABOLIC_PATHWAYS, WP_THIAMINE_METABOLIC_PATHWAYS
MIR184, MIR184
GOBP_REGULATION_OF_ER_ASSOCIATED_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_ER_ASSOCIATED_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCE
HP_GASTROINTESTINAL_EOSINOPHILIA, HP_GASTROINTESTINAL_EOSINOPHILIA
GOBP_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION, GOBP_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION
GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT, GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT
GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_HYPOXIA, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_HYPOXIA
GOMF_HISTONE_ARGININE_N_METHYLTRANSFERASE_ACTIVITY, GOMF_HISTONE_ARGININE_N_METHYLTRANSFERASE_ACTIVITY
GOBP_REGULATION_OF_MITOCHONDRIAL_MRNA_STABILITY, GOBP_REGULATION_OF_MITOCHONDRIAL_MRNA_STABILITY
GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX_CLASS_III, GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX_CLASS_III
GOCC_CLATHRIN_COMPLEX, GOCC_CLATHRIN_COMPLEX
GOCC_PTW_PP1_PHOSPHATASE_COMPLEX, GOCC_PTW_PP1_PHOSPHATASE_COMPLEX
REACTOME_LOSS_OF_FUNCTION_OF_SMAD2_3_IN_CANCER, REACTOME_LOSS_OF_FUNCTION_OF_SMAD2_3_IN_CANCER
GOBP_SECRETORY_GRANULE_LOCALIZATION, GOBP_SECRETORY_GRANULE_LOCALIZATION
GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY
HP_JUVENILE_MYELOMONOCYTIC_LEUKEMIA, HP_JUVENILE_MYELOMONOCYTIC_LEUKEMIA
GOBP_ISOLEUCINE_METABOLIC_PROCESS, GOBP_ISOLEUCINE_METABOLIC_PROCESS
DONATO_CELL_CYCLE_TRETINOIN, DONATO_CELL_CYCLE_TRETINOIN
KUROKAWA_LIVER_CANCER_EARLY_RECURRENCE_DN, KUROKAWA_LIVER_CANCER_EARLY_RECURRENCE_DN
HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_DN, HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_DN
GOBP_REGULATION_OF_UNSATURATED_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_UNSATURATED_FATTY_ACID_BIOSYNTHETIC_PROCESS
MIR6862_3P, MIR6862_3P
HP_HYPERSPLENISM, HP_HYPERSPLENISM
GOBP_POSTSYNAPTIC_ACTIN_CYTOSKELETON_ORGANIZATION, GOBP_POSTSYNAPTIC_ACTIN_CYTOSKELETON_ORGANIZATION
HP_PHARYNGITIS, HP_PHARYNGITIS
GOBP_HISTONE_ARGININE_METHYLATION, GOBP_HISTONE_ARGININE_METHYLATION
GOBP_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY
MIR663B, MIR663B
GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT_UP, GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT_UP
GOMF_OPIOID_RECEPTOR_ACTIVITY, GOMF_OPIOID_RECEPTOR_ACTIVITY
REACTOME_DISPLACEMENT_OF_DNA_GLYCOSYLASE_BY_APEX1, REACTOME_DISPLACEMENT_OF_DNA_GLYCOSYLASE_BY_APEX1
GOBP_REGULATION_OF_GLYCOGEN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_GLYCOGEN_CATABOLIC_PROCESS
GOBP_REGULATION_OF_MICROVILLUS_LENGTH, GOBP_REGULATION_OF_MICROVILLUS_LENGTH
GOBP_SEQUESTERING_OF_IRON_ION, GOBP_SEQUESTERING_OF_IRON_ION