

GSE34006, UNTREATED, VS, A2AR, AGONIST, TREATED, TREG, UP, GSE34006, UNTREATED, VS, A2AR, AGONIST, TREATED, TREG, UP
MORF_TFR, MORF_TFR
GSE41176, WT, VS, TAK1_KO, ANTI_IJGM_STIM, BCELL_24H, UP, GSE41176, WT, VS, TAK1_KO, ANTI_IJGM_STIM, BCELL_24H, UP
MORF_PHB, MORF_PHB
GSE26018, LAIV, VS, TIV_F, FLU, VACCINE, DAY7, BCELL_DN, GSE26018, LAIV, VS, TIV_F, FLU, VACCINE, DAY7, BCELL_DN
MORF_TFR2IP, MORF_TFR2IP
WELCSH, BRCA1_TARGETS, UP, WELCSH, BRCA1_TARGETS, UP
GSE37416, 0H, VS, 24H_F, TULARENSIS, LVS, NEUTROPHIL_DN, GSE37416, 0H, VS, 24H_F, TULARENSIS, LVS, NEUTROPHIL_DN
MORF_BMI1, MORF_BMI1
GSE22886, JGM_MEMORY, BCELL_VS, BM_PLASMA, CELL_DN, GSE22886, JGM_MEMORY, BCELL_VS, BM_PLASMA, CELL_DN
MORF_BECN1, MORF_BECN1
GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX
GSE7548, NAIVE, VS, DAY7_PCC_IMMUNIZATION, CD4_TCELL_UP, GSE7548, NAIVE, VS, DAY7_PCC_IMMUNIZATION, CD4_TCELL_UP
GSE360, CTRL, VS, M, TUBERCULOSIS, MAC, UP, GSE360, CTRL, VS, M, TUBERCULOSIS, MAC, UP
GSE6269, FLU, VS, F, COLL, LINF, PBMC, UP, GSE6269, FLU, VS, F, COLL, LINF, PBMC, UP
GSE17721, CPG, VS, GARDIQUIMOD, 8H, BMDC, DN, GSE17721, CPG, VS, GARDIQUIMOD, 8H, BMDC, DN
MORF_JKBKG, MORF_JKBKG
MIR548M, MIR548M
CUI_DEVELOPING_HEART TRABECULAR_ ATRIAL_CARDIOMYOCYTE, CUI_DEVELOPING_HEART TRABECULAR_ ATRIAL_CARDIOMYOCYTE
GSE17721, POLYIC, VS, CPG, 6H, BMDC, DN, GSE17721, POLYIC, VS, CPG, 6H, BMDC, DN
MORF_PP2R5E, MORF_PP2R5E
REACTOME_SARS_COV_INFECTIONS, REACTOME_SARS_COV_INFECTIONS
GSE39152, CD103, NEG, VS, POS, MEMORY, CD8, TCELL_DN, GSE39152, CD103, NEG, VS, POS, MEMORY, CD8, TCELL_DN
GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING, GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING
GOBP_CYTOSKELETON_DEPENDENT_INTRACELLULAR_TRANSPORT, GOBP_CYTOSKELETON_DEPENDENT_INTRACELLULAR_TRANSPORT
REACTOME_TRANSPORT_TO_THE_GOLGI_AND_SUBSEQUENT_MODIFICATION, REACTOME_TRANSPORT_TO_THE_GOLGI_AND_SUBSEQUENT_MODIFICATION
MODULE_239, MODULE_239
GSE27786, NKCELL, VS, NKTCCELL_UP, GSE27786, NKCELL_VS, NKTCCELL_UP
MORF_FANCG, MORF_FANCG
PID_PDGRFB_PATHWAY, PID_PDGRFB_PATHWAY
GSE360_L_DONOVANI, VS, B, MALAY, LOW_DOSE, MAC, UP, GSE360_L_DONOVANI, VS, B, MALAY, LOW_DOSE, MAC, UP
GSE12845, JGD, NEG, VS, DARKZONE, GC, TONSIL, BCELL_UP, GSE12845, JGD, NEG, VS, DARKZONE, GC, TONSIL, BCELL_UP
GSE22611, UNSTIM, VS, 2H, MDP, STIM, MUTANT, NOD2, TRANSDUCED, HEK293T_CELL_UP, GSE22611, UNSTIM, VS, 2H, MDP, STIM, MUTANT, NOD2, TRANSDUCED, HEK293T_CELL_UP
REACTOME_SUMOYLATION_OF_DNA_DAMAGE_RESPONSE_AND_REPAIR_PROTEINS, REACTOME_SUMOYLATION_OF_DNA_DAMAGE_RESPONSE_AND_REPAIR_PROTEINS
REACTOME_ER_TO_GOLGI_ANTEROGRADE_TRANSPORT, REACTOME_ER_TO_GOLGI_ANTEROGRADE_TRANSPORT
GOCC_MEMBRANE_COAT, GOCC_MEMBRANE_COAT
GOBP_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GOBP_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
GSE22886, ICG, JGA, MEMORY, BCELL_VS, BM_PLASMA, CELL_DN, GSE22886, ICG, JGA, MEMORY, BCELL_VS, BM_PLASMA, CELL_DN
GOBP_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE, GOBP_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE
GSE39152, SPLEEN, CD103, NEG, VS, BRAIN, CD103, POS, MEMORY, CD8, TCELL_UP, GSE39152, SPLEEN, CD103, NEG, VS, BRAIN, CD103, POS, MEMORY, CD8, TCELL_UP
PID_ERBB1_DOWNSTREAM_PATHWAY, PID_ERBB1_DOWNSTREAM_PATHWAY
GOBP_PROTEIN_SUMOYLATION, GOBP_PROTEIN_SUMOYLATION
OCONNOR_PBMCM_MENVEJO, ACWYVAX_AGE_30_70YO_7DY_AFTER_SECOND_DOSE, VS, 7DY_AFTER_FIRST_DOSE, UP, OCONNOR_PBMCM_MENVEJO, ACWYVAX_AGE_30_70YO_7DY_AFTER_SECOND_DOSE, VS, 7DY_AFTER_FIRST_DOSE, UP
GSE43565, CTRL, VS, HCMV_INFECTION, NK_CELL_DN, GSE43565, CTRL, VS, HCMV_INFECTION, NK_CELL_DN
MORF_CSNK1D, MORF_CSNK1D
REACTOME_CLATHRIN_MEDIATED_ENDOCYTOSIS, REACTOME_CLATHRIN_MEDIATED_ENDOCYTOSIS
MORF_SS18, MORF_SS18
GSE12366, PLASMA, CELL, VS, MEMORY, BCELL_UP, GSE12366, PLASMA, CELL_VS, MEMORY, BCELL_UP
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE
HP_VASCULAR_DILATATION, HP_VASCULAR_DILATATION
DAZARD_RESPONSE_TO_UV_SCC_DN, DAZARD_RESPONSE_TO_UV_SCC_DN
GSE18884, BRAIN, VS, COLON_TUMORAL_MACROPHAGE_DN, GSE18884, BRAIN, VS, COLON_TUMORAL_MACROPHAGE_DN
GOMF_PHOSPHATASE_REGULATOR_ACTIVITY, GOMF_PHOSPHATASE_REGULATOR_ACTIVITY
MIR4514, MIR4514
AGCATT_A, MIR155, AGCATT_A, MIR155
REACTOME_TRANS_GOLGI_NETWORK_VESICLE_BUDDING, REACTOME_TRANS_GOLGI_NETWORK_VESICLE_BUDDING
HP_ABNORMALITY_OF_LATERAL_VENTRICLE, HP_ABNORMALITY_OF_LATERAL_VENTRICLE
SCHLOSSER_SERUM_RESPONSE, UP, SCHLOSSER_SERUM_RESPONSE, UP
MIR6511A_5P, MIR6511A_5P
LU_AGING, BRAIN, DN, LU_AGING, BRAIN, DN
GOBP_REGULATION_OF_ACTIN_FILAMENT_LENGTH, GOBP_REGULATION_OF_ACTIN_FILAMENT_LENGTH
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY
GOBP_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE, GOBP_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE
GSE19772, CTRL, VS, HCMV_INF, MONOCYTES_DN, GSE19772, CTRL, VS, HCMV_INF, MONOCYTES_DN
REACTOME_SARS_COV_2_INFECTION, REACTOME_SARS_COV_2_INFECTION
MIR940, MIR940
MIR6893_5P, MIR6893_5P
REACTOME_COPI_MEDIATED_ANTEROGRADE_TRANSPORT, REACTOME_COPI_MEDIATED_ANTEROGRADE_TRANSPORT
WP_ANGIOPETIN_LIKE_PROTEIN_8_REGULATORY_PATHWAY, WP_ANGIOPETIN_LIKE_PROTEIN_8_REGULATORY_PATHWAY
HP_MANDIBULAR_PROGNATHIA, HP_MANDIBULAR_PROGNATHIA
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE
REACTOME_EPH_EPHRIN_SIGNALING, REACTOME_EPH_EPHRIN_SIGNALING
REACTOME_SUMOYLATION_OF_CHROMATIN_ORGANIZATION_PROTEINS, REACTOME_SUMOYLATION_OF_CHROMATIN_ORGANIZATION_PROTEINS
SMARCA1_TARGET_GENES, SMARCA1_TARGET_GENES
MIR6884_5P, MIR6884_5P
WP_LEPTIN_SIGNALING_PATHWAY, WP_LEPTIN_SIGNALING_PATHWAY
HP_ABNORMAL_PROTEIN_GLYCOSYLATION, HP_ABNORMAL_PROTEIN_GLYCOSYLATION
REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS, REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS
GOBP_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
WP_FRAGILE_X_SYNDROME, WP_FRAGILE_X_SYNDROME
KEGG_INSULIN_SIGNALING_PATHWAY, KEGG_INSULIN_SIGNALING_PATHWAY
MIR6808_5P, MIR6808_5P
HP_OPEN_MOUTH, HP_OPEN_MOUTH
HP_MELANOMA, HP_MELANOMA
GOMF_ION_CHANNEL_BINDING, GOMF_ION_CHANNEL_BINDING
MORF_UBE2A, MORF_UBE2A
REACTOME_RAB_GERANYLGERANYLATION, REACTOME_RAB_GERANYLGERANYLATION
MORF_PML, MORF_PML
KEGG_N_GLYCAN_BIOSYNTHESIS, KEGG_N_GLYCAN_BIOSYNTHESIS
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_ASSEMBLY
GOBP_LYTIC_VASCULOC_ORGANIZATION, GOBP_LYTIC_VASCULOC_ORGANIZATION
GSE360, HIGH, VS, LOW, DOSE, B, MALAY, DC, DN, GSE360, HIGH, VS, LOW, DOSE, B, MALAY, DC, DN
GOBP_IJEL_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GOBP_IJEL_MEDIATED_UNFOLDED_PROTEIN_RESPONSE
HP_DILATION_OF_LATERAL_VENTRICLES, HP_DILATION_OF_LATERAL_VENTRICLES
REACTOME_TRANSLOCATION_OF_SLC2A4_GLUT4_TO_THE_PLASMA_MEMBRANE, REACTOME_TRANSLOCATION_OF_SLC2A4_GLUT4_TO_THE_PLASMA_MEMBRANE
GOCC_CLATHRIN_COATED_VESICLE, GOCC_CLATHRIN_COATED_VESICLE
MIR5586_5P, MIR5586_5P
GSE1791, CTRL, VS, NEUROMEDINU_IN_T_CELL_LINE_3H, UP, GSE1791, CTRL, VS, NEUROMEDINU_IN_T_CELL_LINE_3H, UP
MIR665, MIR665
MODULE_86, MODULE_86
GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT, GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT
GOBP_VESICLE_TRANSPORT_ALONG_MICROTUBULE, GOBP_VESICLE_TRANSPORT_ALONG_MICROTUBULE
REACTOME_GLYCOLYSIS, REACTOME_GLYCOLYSIS
GOCC_VESICLE_COAT, GOCC_VESICLE_COAT
EPPERT_LSC_R, EPPERTR_LSC_R
GSE3982, MAC, VS, NEUTROPHIL_LPS_STIM_DN, GSE3982, MAC, VS, NEUTROPHIL_LPS_STIM_DN
HP_LOW_POSTERIOR_HAIRLINE, HP_LOW_POSTERIOR_HAIRLINE
GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
GSE25088, CTRL, VS, ROSIGLITAZONE_STIM, STAT6, KO, MACROPHAGE_UP, GSE25088, CTRL, VS, ROSIGLITAZONE_STIM, STAT6, KO, MACROPHAGE_UP
MIR1228_3P, MIR1228_3P
MODULE_87, MODULE_87
HP_WIDE_INTERMAMILLARY_DISTANCE, HP_WIDE_INTERMAMILLARY_DISTANCE
MIR4733_5P, MIR4733_5P
HP_MACULE, HP_MACULE
GOBP_SYNAPTIC_VESICLE_TRANSPORT, GOBP_SYNAPTIC_VESICLE_TRANSPORT
GOBP_SYNAPTIC_VESICLE_LOCALIZATION, GOBP_SYNAPTIC_VESICLE_LOCALIZATION
MIR122_5P, MIR122_5P
REACTOME_RHO_GTPASE_CYCLE, REACTOME_RHO_GTPASE_CYCLE
WP_MELANOMA, WP_MELANOMA
GOCC_ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE
GOBP_VESICLE_CYTOSKELETAL_TRAFFICKING, GOBP_VESICLE_CYTOSKELETAL_TRAFFICKING
GOBP_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY, GOBP_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY
REACTOME_SUMOYLATION_OF_TRANSCRIPTION_COFACTORS, REACTOME_SUMOYLATION_OF_TRANSCRIPTION_COFACTORS
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MATURE_NEURONS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MATURE_NEURONS
HP_ATONIC_SEIZURE, HP_ATONIC_SEIZURE
GOCC_AP_TYPE_MEMBRANE_COAT_ADAPTOR_COMPLEX, GOCC_AP_TYPE_MEMBRANE_COAT_ADAPTOR_COMPLEX
WP_GLYCOGEN_SYNTHESIS_AND_DEGRADATION, WP_GLYCOGEN_SYNTHESIS_AND_DEGRADATION
BIOCARTA_PVK2_PATHWAY, BIOCARTA_PVK2_PATHWAY
GOBP_PROTEIN_K63_LINKED_UBIQUITINATION, GOBP_PROTEIN_K63_LINKED_UBIQUITINATION
GOCC_COPI_COATED_VESICLE, GOCC_COPI_COATED_VESICLE
GOBP_RESPONSE_TO_MONOSACCHARIDE, GOBP_RESPONSE_TO_MONOSACCHARIDE
GOCC_CLATHRIN_COAT, GOCC_CLATHRIN_COAT
GSE43955, TH17, ACT, CD4_TCELL_30H, UP, GSE43955, TH17, ACT, CD4_TCELL_30H, UP
HP_ECHOLALIA, HP_ECHOLALIA
REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE, REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE
HP_RECURRENT_FEVER, HP_RECURRENT_FEVER
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION
ACAACT_MIR82, ACAACT_MIR82
REACTOME_SUMOYLATION_OF_INTRACELLULAR_RECEPTORS, REACTOME_SUMOYLATION_OF_INTRACELLULAR_RECEPTORS
GSE26488, WT, VS, HDACT, DELTA_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION, UP, GSE26488, WT, VS, HDACT, DELTA_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION, UP
BIOCARTA_FMLP_PATHWAY, BIOCARTA_FMLP_PATHWAY
REACTOME_RHO_GTPASES_ACTIVATE_WASPS_AND_WAVES, REACTOME_RHO_GTPASES_ACTIVATE_WASPS_AND_WAVES
BIOCARTA_TCR_PATHWAY, BIOCARTA_TCR_PATHWAY
KEGG_WNT_SIGNALING_PATHWAY, KEGG_WNT_SIGNALING_PATHWAY
MIR409_3P, MIR409_3P
HP_PLACIOCEPHALY, HP_PLACIOCEPHALY
SASSON_RESPONSE_TO_GONADOTROPHINS, UP, SASSON_RESPONSE_TO_GONADOTROPHINS, UP
GOCC_DYNEIN_COMPLEX, GOCC_DYNEIN_COMPLEX
GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
GSE36476, YOUNG, VS, OLD, DONOR, MEMORY_CD4_TCELL_40H_TSTT_ACT_UP, GSE36476, YOUNG, VS, OLD, DONOR, MEMORY_CD4_TCELL_40H_TSTT_ACT_UP
GSE29618, PRE, VS, DAY7_POST_LAIV_F, FLU, VACCINE, MONOCYTE_UP, GSE29618, PRE, VS, DAY7_POST_LAIV_F, FLU, VACCINE, MONOCYTE_UP
HP_TYPE_I_TRANSFERRIN_ISOFORM_PROFILE, HP_TYPE_I_TRANSFERRIN_ISOFORM_PROFILE
REACTOME_INSERTION_OF_TAIL_ANCHORED_PROTEINS_INTO_THE_ENDOPLASMIC_RETICULUM_MEMBRANE, REACTOME_INSERTION_OF_TAIL_ANCHORED_PROTEINS_INTO_THE_ENDOPLASMIC_RETICULUM_MEMBRANE
MIR3118, MIR3118
HP_ABNORMALITY_OF_THE_INTRINSIC_PATHWAY, HP_ABNORMALITY_OF_THE_INTRINSIC_PATHWAY
HP_REDUCED_ANTITHROMBIN_III_ACTIVITY, HP_REDUCED_ANTITHROMBIN_III_ACTIVITY
KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY, KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY
HP_ABNORMALITY_OF_THE_EXTRINSIC_PATHWAY, HP_ABNORMALITY_OF_THE_EXTRINSIC_PATHWAY
WP_CARDIAC_HYPERTROPHIC_RESPONSE, WP_CARDIAC_HYPERTROPHIC_RESPONSE
SASSON_RESPONSE_TO_FORSKOLIN, UP, SASSON_RESPONSE_TO_FORSKOLIN, UP
MIR4752, MIR4752
MIR299_3P, MIR299_3P
KLEIN_PRIMARY_EFFUSION_LYMPHOMA, UP, KLEIN_PRIMARY_EFFUSION_LYMPHOMA, UP
GOBP_CELLULAR_PIGMENTATION, GOBP_CELLULAR_PIGMENTATION
REACTOME_EPH_EPHRIN_MEDIATED_REPULSION_OF_CELLS, REACTOME_EPH_EPHRIN_MEDIATED_REPULSION_OF_CELLS
BIOCARTA_NDKDYNAMIN_PATHWAY, BIOCARTA_NDKDYNAMIN_PATHWAY
MIR5189_3P, MIR5189_3P
GSE18281, PERIMEDULLARY_CORTICAL_REGION_VS, WHOLE_CORTX_THYMUS_DN, GSE18281, PERIMEDULLARY_CORTICAL_REGION_VS, WHOLE_CORTX_THYMUS_DN
REACTOME_SIGNALING_BY_PTFRS, REACTOME_SIGNALING_BY_PTFRS
GOBP_ENDOPLASMIC_RETICULUM_CALCIIUM_ION_HOMEOSTASIS, GOBP_ENDOPLASMIC_RETICULUM_CALCIIUM_ION_HOMEOSTASIS
GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT, GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT
GOCC_ROUGH_ENDOPLASMIC_RETICULUM, GOCC_ROUGH_ENDOPLASMIC_RETICULUM
HP_BROAD_TOE, HP_BROAD_TOE
KEGG_VASOPRESSIN_REGULATED_WATER_REABSORPTION, KEGG_VASOPRESSIN_REGULATED_WATER_REABSORPTION
MODULE_366, MODULE_366
GOCC_CLATHRIN_COATED_PIT, GOCC_CLATHRIN_COATED_PIT
GOBP_CELLULAR_GLUCOSE_HOMEOSTASIS, GOBP_CELLULAR_GLUCOSE_HOMEOSTASIS
GOBP_REGULATION_OF_PROTEIN_MATURATION, GOBP_REGULATION_OF_PROTEIN_MATURATION
REACTOME_CALNEXIN_CALRETICULIN_CYCLE, REACTOME_CALNEXIN_CALRETICULIN_CYCLE
GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
HP_ANKLE_FLEXION_CONTRACTURE, HP_ANKLE_FLEXION_CONTRACTURE
REACTOME_NEGATIVE_REGULATION_OF_MAPK_PATHWAY, REACTOME_NEGATIVE_REGULATION_OF_MAPK_PATHWAY
REACTOME_TBC_RABGAPS, REACTOME_TBC_RABGAPS
GOCC_COPI_COATED_VESICLE_MEMBRANE, GOCC_COPI_COATED_VESICLE_MEMBRANE
PID_ADPS_2PATHWAY, PID_ADPS_2PATHWAY
GOBP_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION, GOBP_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION
BURTON_ADIPOGENESIS_11, BURTON_ADIPOGENESIS_11
GOBP_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GOBP_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
PID_ARF_3PATHWAY, PID_ARF_3PATHWAY
GOBP_PIGMENT_GNANULE_LOCALIZATION, GOBP_PIGMENT_GNANULE_LOCALIZATION
MIR301A_5P, MIR301A_5P
REACTOME_COPI_INDEPENDENT_GOLGI_TO_ER_RETROGRADE_TRAFFIC, REACTOME_COPI_INDEPENDENT_GOLGI_TO_ER_RETROGRADE_TRAFFIC
LXR_DR4_Q3, LXR_DR4_Q3
BIOCARTA_PTINS_PATHWAY, BIOCARTA_PTINS_PATHWAY
MIR301B_5P, MIR301B_5P
MIR6769B_3P, MIR6769B_3P
GOCC_CLATHRIN_VESICLE_COAT, GOCC_CLATHRIN_VESICLE_COAT
MIR4723_3P, MIR4723_3P
MIR1383, MIR1383
WP_G_PROTEIN_SIGNALING_PATHWAYS, WP_G_PROTEIN_SIGNALING_PATHWAYS
AHR_01_AHR_01
HP_AORTIC_ROOT_ANEURYSM, HP_AORTIC_ROOT_ANEURYSM
GOBP_PIGMENTATION, GOBP_PIGMENTATION
GOMF_CLATHRIN_BINDING, GOMF_CLATHRIN_BINDING
GOCC_COPI_VESICLE_COAT, GOCC_COPI_VESICLE_COAT
CHEOK_RESPONSE_TO_HD_MTX_DN, CHEOK_RESPONSE_TO_HD_MTX_DN
GOBP_SNARE_COMPLEX_ASSEMBLY, GOBP_SNARE_COMPLEX_ASSEMBLY
HP_WEAK_VOICE, HP_WEAK_VOICE
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MATURATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MATURATION
GSE24972, WT, VS, IRRF_KO, MARGINAL_ZONE_SPLEEN, BCELL_DN, GSE24972, WT, VS, IRRF_KO, MARGINAL_ZONE_SPLEEN, BCELL_DN
MIR3665, MIR3665
GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY
NAGY_STAGA_COMPONENTS_HUMAN, NAGY_STAGA_COMPONENTS_HUMAN
WP_HEDGEHOG_SIGNALING_PATHWAY, WP_HEDGEHOG_SIGNALING_PATHWAY
GOBP_MAINTENANCE_OF_CELL_POLARITY, GOBP_MAINTENANCE_OF_CELL_POLARITY
REACTOME_RAF_ACTIVATION, REACTOME_RAF_ACTIVATION
GOMF_UBIQUITIN_SPECIFIC_PROTEASE_BINDING, GOMF_UBIQUITIN_SPECIFIC_PROTEASE_BINDING
GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORT
GOBP_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ENDOPLASMIC_RETICULUM, GOBP_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ENDOPLASMIC_RETICULUM
GOBP_ESTABLISHMENT_OF_PIGMENT_GNANULE_LOCALIZATION, GOBP_ESTABLISHMENT_OF_PIGMENT_GNANULE_LOCALIZATION
GOCC_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX, GOCC_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX
VARELA_ZMPST24_TARGETS_UP, VARELA_ZMPST24_TARGETS_UP
GOBP_POSITIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION
HP_GROWTH_HORMONE_EXCESS, HP_GROWTH_HORMONE_EXCESS
GOCC_CYTOPLASMIC_DYNEIN_COMPLEX, GOCC_CYTOPLASMIC_DYNEIN_COMPLEX
HP_ENTROPION, HP_ENTROPION
GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSPORT
HP_THROMBOEMBOLISM, HP_THROMBOEMBOLISM
GOBP_REGULATION_OF_LYSOSOMAL_LUMEN_PH, GOBP_REGULATION_OF_LYSOSOMAL_LUMEN_PH
GOCC_LUMENAL_SIDE_OF_MEMBRANE, GOCC_LUMENAL_SIDE_OF_MEMBRANE
GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
HP_OLIVOPONTOCEREBELLAR_HYPOPLASIA, HP_OLIVOPONTOCEREBELLAR_HYPOPLASIA
GOCC_STAGA_COMPLEX, GOCC_STAGA_COMPLEX
REACTOME_SIGNALING_BY_CTNNB1_PHOSPHO_SITE_MUTANTS, REACTOME_SIGNALING_BY_CTNNB1_PHOSPHO_SITE_MUTANTS
GOBP_PROTEIN_FOLDING_IN_ENDOPLASMIC_RETICULUM, GOBP_PROTEIN_FOLDING_IN_ENDOPLASMIC_RETICULUM
GOMF_CHANNEL_INHIBITOR_ACTIVITY, GOMF_CHANNEL_INHIBITOR_ACTIVITY
GOBP_INTEGRATED_STRESS_RESPONSE_SIGNALING, GOBP_INTEGRATED_STRESS_RESPONSE_SIGNALING
REACTOME_PROCESSING_AND_ACTIVATION_OF_SUMO, REACTOME_PROCESSING_AND_ACTIVATION_OF_SUMO
GOCC_CLATHRIN_COATED_VESICLE_MEMBRANE, GOCC_CLATHRIN_COATED_VESICLE_MEMBRANE
HP_DECREASED_TESTICULAR_SIZE, HP_DECREASED_TESTICULAR_SIZE
HP_ABNORMAL_THYROID_STIMULATING_HORMONE_LEVEL, HP_ABNORMAL_THYROID_STIMULATING_HORMONE_LEVEL
REACTOME_BETA_CATENIN_PHOSPHORYLATION_CASCADE, REACTOME_BETA_CATENIN_PHOSPHORYLATION_CASCADE
GOMF_PHOSPHATASE_ACTIVATOR_ACTIVITY, GOMF_PHOSPHATASE_ACTIVATOR_ACTIVITY
ARID3B_TARGET_GENES, ARID3B_TARGET_GENES
GOBP_HYALURONAN_METABOLIC_PROCESS, GOBP_HYALURONAN_METABOLIC_PROCESS
GOBP_RECEPTOR_CATABOLIC_PROCESS, GOBP_RECEPTOR_CATABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_RYANODINE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_RYANODINE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY
REACTOME_GPIB_DX_V_ACTIVATION_SIGNALLING, REACTOME_GPIB_DX_V_ACTIVATION_SIGNALLING
MIR744_3P, MIR744_3P
REACTOME_DISASSEMBLY_OF_THE_DESTRUCTION_COMPLEX_AND_RECRUITMENT_OF_AXIN_TO_THE_MEMBRANE, REACTOME_DISASSEMBLY_OF_THE_DESTRUCTION_COMPLEX_AND_RECRUITMENT_OF_AXIN_TO_THE_MEMBRANE
REACTOME_WNT5A_DEPENDENT_INTERNALIZATION_OF_FZD2_FZD5_AND_ROR2, REACTOME_WNT5A_DEPENDENT_INTERNALIZATION_OF_FZD2_FZD5_AND_ROR2
GOBP_REGULATION_OF_RNA_EXPORT_FROM_NUCLEUS, GOBP_REGULATION_OF_RNA_EXPORT_FROM_NUCLEUS
BIOCARTA_NOS1_PATHWAY, BIOCARTA_NOS1_PATHWAY
GOCC_ENDOLYSOSOME_MEMBRANE, GOCC_ENDOLYSOSOME_MEMBRANE
HP_SMALL_HAND, HP_SMALL_HAND
REACTOME_DARPP_32_EVENTS, REACTOME_DARPP_32_EVENTS
GOBP_EMBRYONIC_AXIS_SPECIFICATION, GOBP_EMBRYONIC_AXIS_SPECIFICATION
GOBP_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GOBP_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY
CAMPS_COLON_CANCER_COPY_NUMBER_UP, CAMPS_COLON_CANCER_COPY_NUMBER_UP
GOBP_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS, GOBP_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS
WP_FACTORS_AND_PATHWAYS_AFFECTING_INSULINLIKE_GROWTH_FACTOR_IGF1AKT_SIGNALING, WP_FACTORS_AND_PATHWAYS_AFFECTING_INSULINLIKE_GROWTH_FACTOR_IGF1AKT_SIGNALING
GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE_BY_P53_CLASS_MEDIATOR, GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE_BY_P53_CLASS_MEDIATOR
REACTOME_PLATELET_ACTIVATION_BY_ID1, REACTOME_PLATELET_ACTIVATION_BY_ID1
HP_LOW_HANGING_COLUMELLA, HP_LOW_HANGING_COLUMELLA
MIR4280, MIR4280
BIOCARTA_VIP_PATHWAY, BIOCARTA_VIP_PATHWAY
GOMF_ADENYLATE_CYCLASE_BINDING, GOMF_ADENYLATE_CYCLASE_BINDING
WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY, WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY
MIR6750_5P, MIR6750_5P
REACTOME_PLASMA_LIPOPROTEIN_CLEARANCE, REACTOME_PLASMA_LIPOPROTEIN_CLEARANCE
GOBP_POSITIVE_REGULATION_OF_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING, GOBP_POSITIVE_REGULATION_OF_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING
MIR822_5P, MIR822_5P
chr3q23, chr3q23
GOBP_MINUS_END_DIRECTED_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE, GOBP_MINUS_END_DIRECTED_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE
GOCC_LUMENAL_SIDE_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_LUMENAL_SIDE_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
GOBP_UV_DAMAGE_EXCISION_REPAIR, GOBP_UV_DAMAGE_EXCISION_REPAIR
GOBP_NEGATIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
TERAMOTO_OPN_TARGETS_CLUSTER_6, TERAMOTO_OPN_TARGETS_CLUSTER_6
HP_ADRENAL_HYPERPLASIA, HP_ADRENAL_HYPERPLASIA
PID_ENDOTHELIN_PATHWAY, PID_ENDOTHELIN_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_TRANSCRIPTION_BY_COMPETITIVE_PROMOTER_BINDING, GOBP_NEGATIVE_REGULATION_OF_TRANSCRIPTION_BY_COMPETITIVE_PROMOTER_BINDING
GOBP_LYSOSOMAL_LUMEN_ACIDIFICATION, GOBP_LYSOSOMAL_LUMEN_ACIDIFICATION
GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_LIPOPROTEIN_PARTICLE_RECEPTOR_BINDING, GOMF_LIPOPROTEIN_PARTICLE_RECEPTOR_BINDING
GOBP_POSITIVE_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA1_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA1_PRODUCTION
HP_HYPOFLEXIA_OF_UPPER_LIMBS, HP_HYPOFLEXIA_OF_UPPER_LIMBS
GOCC_TRANSLOCON_COMPLEX, GOCC_TRANSLOCON_COMPLEX
HP_BROAD_FEMORAL_NECK, HP_BROAD_FEMORAL_NECK
GOBP_REGULATION_OF_COATED_ADHESION_DEPENDENT_CELL_SPREADING, GOBP_REGULATION_OF_COATED_ADHESION_DEPENDENT_CELL_SPREADING
GOCC_CLATHRIN_COAT_OF_COATED_PIT, GOCC_CLATHRIN_COAT_OF_COATED_PIT
GOMF_EPIDERMAL_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_EPIDERMAL_GROWTH_FACTOR_RECEPTOR_BINDING
GOBP_REGULATION_OF_CALCINEURIN_MEDIATED_SIGNALING, GOBP_REGULATION_OF_CALCINEURIN_MEDIATED_SIGNALING
GSE1971, CTRL, VS, NEUROMEDINU_IN_T_CELL_LINE_12H, DN, GSE1971, CTRL, VS, NEUROMEDINU_IN_T_CELL_LINE_12H, DN
REACTOME_VLDLR_INTERNALISATION_AND_DEGRADATION, REACTOME_VLDLR_INTERNALISATION_AND_DEGRADATION
GOBP_AMYLOID_PRECURSOR_PROTEIN_CATABOLIC_PROCESS, GOBP_AMYLOID_PRECURSOR_PROTEIN_CATABOLIC_PROCESS
MODULE_395, MODULE_395
GOBP_REGULATION_OF_JRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GOBP_REGULATION_OF_JRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE
GOMF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_BINDING, GOMF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_BINDING
GOMF_PROTEIN_PHOSPHATASE_ACTIVATOR_ACTIVITY, GOMF_PROTEIN_PHOSPHATASE_ACTIVATOR_ACTIVITY
GOBP_NUCLEAR_MEMBRANE_ORGANIZATION, GOBP_NUCLEAR_MEMBRANE_ORGANIZATION
GOCC_DYNEIN_LIGHT_CHAIN_BINDING, GOMF_DYNEIN_LIGHT_CHAIN_BINDING
REACTOME_SEMA4D_MEDIATED_INHIBITION_OF_CELL_ATTACHMENT_AND_MIGRATION, REACTOME_SEMA4D_MEDIATED_INHIBITION_OF_CELL_ATTACHMENT_AND_MIGRATION
GOBP_DETERMINATION_OF_LEFT_RIGHT_ASYMMETRY_IN_LATERAL_MESODERM, GOBP_DETERMINATION_OF_LEFT_RIGHT_ASYMMETRY_IN_LATERAL_MESODERM
WP_ENDOTHELIN_PATHWAYS, WP_ENDOTHELIN_PATHWAYS
GOMF_PROTEIN_PHOSPHATASE_2A_BINDING, GOMF_PROTEIN_PHOSPHATASE_2A_BINDING
GOCC_ANCHORED_COMPONENT_OF_SYNAPTIC_VESICLE_MEMBRANE, GOCC_ANCHORED_COMPONENT_OF_SYNAPTIC_VESICLE_MEMBRANE
BIOCARTA_RAB_PATHWAY, BIOCARTA_RAB_PATHWAY
LJUKA_LIVER_CANCER_PROGRESSION_10_11_11_UP, LJUKA_LIVER_CANCER_PROGRESSION_10_11_11_UP
GOBP_NEGATIVE_REGULATION_OF_CALCIIUM_ION_TRANSPORT_INTO_CYTOSOL, GOBP_NEGATIVE_REGULATION_OF_CALCIIUM_ION_TRANSPORT_INTO_CYTOSOL
HINATA_NFKB_MATRIX, HINATA_NFKB_MATRIX
HP_DEFORMED_TARSAL_BONIS, HP_DEFORMED_TARSAL_BONIS
HP_EEG_WITH_SPIKE_WAVE_COMPLEXES_3_5_HZ, HP_EEG_WITH_SPIKE_WAVE_COMPLEXES_3_5_HZ
GOBP_ATTACHMENT_OF_GPI_ANCHOR_TO_PROTEIN, GOBP_ATTACHMENT_OF_GPI_ANCHOR_TO_PROTEIN
GOCC_SAGA_COMPLEX, GOCC_SAGA_COMPLEX
GOBP_AMYLOID_PRECURSOR_PROTEIN_METABOLIC_PROCESS, GOBP_AMYLOID_PRECURSOR_PROTEIN_METABOLIC_PROCESS
REACTOME_SIGNALING_BY_WNT_IN_CANCER, REACTOME_SIGNALING_BY_WNT_IN_CANCER