

	MORF_RPA2, MORF_RPA2 GSE29618_BCELL_VS_MDC_DN, GSE29618_BCELL_VS_MDC_DN MORF_TTPR, MORF_TTPR GSE22886_NAIVE_CD4_TCELL_VS_NKCELL_DN, GSE22886_NAIVE_CD4_TCELL_VS_NKCELL_DN GSE41176_WT_VS_TAK1_KO_ANT1JGM_STIM_BCELL_24H_UP, GSE41176_WT_VS_TAK1_KO_ANT1JGM_STIM_BCELL_24H_UP GSE360_DC_VS_MAC_B_MALAYI_LOW_DOSE_UP, GSE360_DC_VS_MAC_B_MALAYI_LOW_DOSE_UP GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN GSE45365_NK_CELL_VS_BCELL_MCMV_INFECTION_UP, GSE45365_NK_CELL_VS_BCELL_MCMV_INFECTION_UP MORF_SP3, MORF_SP3 GSE20727_CTRL_VS_ROS_INHIBITOR_TREATED_DC_DN, GSE20727_CTRL_VS_ROS_INHIBITOR_TREATED_DC_DN MORF_BECN1, MORF_BECN1 GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS MODULE_239, MODULE_239 GSE34205_HEALTHY_VS_FLU_INF_INFANT_PBMCDN, GSE34205_HEALTHY_VS_FLU_INF_INFANT_PBMCDN HALLMARK_PROTEIN_SECRETION, HALLMARK_PROTEIN_SECRETION GSE22886_TH1_VS_TH2_48H_ACT_DN, GSE22886_TH1_VS_TH2_48H_ACT_DN WELCSH_BRCA1_TARGETS_UP, WELCSH_BRCA1_TARGETS_UP GO_PROTEIN_SUMOYLATION, GO_PROTEIN_SUMOYLATION MORF_PHB, MORF_PHB MORF_RPA1, MORF_RPA1 GO_UBIQUITIN_LIKE_PROTEIN_BINDING, GO_UBIQUITIN_LIKE_PROTEIN_BINDING GO_CELL_CYCLE_G2_M_PHASE_TRANSITION, GO_CELL_CYCLE_G2_M_PHASE_TRANSITION GSE45365_CTRL_VS_MCMV_INFECTION_NK_CELL_DN, GSE45365_CTRL_VS_MCMV_INFECTION_NK_CELL_DN GNF2_DENR, GNF2_DENR GO_POSITIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS LAIHO_COLORECTAL_CANCER_SERRATED_UP, LAIHO_COLORECTAL_CANCER_SERRATED_UP YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_10, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_10 MORF_BAG5, MORF_BAG5 GSE43863_NAIVE_VS_TFH_CD4_EFF_TCELL_D6_1CMV_UP, GSE43863_NAIVE_VS_TFH_CD4_EFF_TCELL_D6_1CMV_UP MORF_CSNK1D, MORF_CSNK1D GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_UP GSE1460_DP_THYMOCYTE_VS_THYMIC_STROMAL_CELL_DN, GSE1460_DP_THYMOCYTE_VS_THYMIC_STROMAL_CELL_DN GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP, GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_DN, GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_DN GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_MAC_DN, GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_MAC_DN TAKAO_RESPONSE_TO_UVB_RADIATION_DN, TAKAO_RESPONSE_TO_UVB_RADIATION_DN MORF_SS18, MORF_SS18 GSE3920_UNTREATED_VS_IFNA_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_UNTREATED_VS_IFNA_TREATED_ENDOTHELIAL_CELL_UP GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_MAC_UP, GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_MAC_UP GO_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT, GO_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT TARTE_PLASMA_CELL_VS_B_LYMPHOCYTE_UP, TARTE_PLASMA_CELL_VS_B_LYMPHOCYTE_UP GO_VACUOLE_ORGANIZATION, GO_VACUOLE_ORGANIZATION GSE17721_ALL_VS_24H_PAM3CSK4_BMDC_DN, GSE17721_ALL_VS_24H_PAM3CSK4_BMDC_DN GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN MORF_BMI1, MORF_BMI1 MORF_JKBKG, MORF_JKBKG PID_PDGFRRB_PATHWAY, PID_PDGFRRB_PATHWAY GSE22886_JGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_DN, GSE22886_JGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_DN GO_NUCLEUS_ORGANIZATION, GO_NUCLEUS_ORGANIZATION MORF_FANCCG, MORF_FANCCG GO_MEMBRANE_BUDDING, GO_MEMBRANE_BUDDING REACTOME_TRANS_GOLGI_NETWORK_VESICLE_BUDDING, REACTOME_TRANS_GOLGI_NETWORK_VESICLE_BUDDING GSE18893_CTRL_VS_TNF_TREATED_TCONV_2H_UP, GSE18893_CTRL_VS_TNF_TREATED_TCONV_2H_UP GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_DN, GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_DN KEGG_LYSOSOME, KEGG_LYSOSOME GO_NUCLEAR_ENVELOPE_ORGANIZATION, GO_NUCLEAR_ENVELOPE_ORGANIZATION HALLMARK_PIK_AKT_MTOR_SIGNALING, HALLMARK_PIK_AKT_MTOR_SIGNALING GSE12845_IKD_NEG_BLOOD_VS_DARKZONE_GC_TONSIL_BCELL_UP, GSE12845_IKD_NEG_BLOOD_VS_DARKZONE_GC_TONSIL_BCELL_UP GO_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GO_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE GSE360_DC_VS_MAC_UP, GSE360_DC_VS_MAC_UP GSE5099_UNSTIM_VS_MCSF_TREATED_MONOCYTE_DAY3_DN, GSE5099_UNSTIM_VS_MCSF_TREATED_MONOCYTE_DAY3_DN KEGG_OOCYTE_MEIOSIS, KEGG_OOCYTE_MEIOSIS GSE12366_PLASMA_CELL_VS_MEMORY_BCELL_UP, GSE12366_PLASMA_CELL_VS_MEMORY_BCELL_UP GSE12366_PLASMA_CELL_VS_NAIVE_BCELL_UP, GSE12366_PLASMA_CELL_VS_NAIVE_BCELL_UP GSE17721_CTRL_VS_POLYIC_1H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_1H_BMDC_UP REACTOME_SIGNALING_BY_EGFR_IN_CANCER, REACTOME_SIGNALING_BY_EGFR_IN_CANCER PID_ERBB1_DOWNSTREAM_PATHWAY, PID_ERBB1_DOWNSTREAM_PATHWAY REACTOME_GOLGI_ASSOCIATED_VESICLE_BIOGENESIS, REACTOME_GOLGI_ASSOCIATED_VESICLE_BIOGENESIS GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_MEMORY_CD8_TCELL_UP, GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_MEMORY_CD8_TCELL_UP GO_GOLGI_ORGANIZATION, GO_GOLGI_ORGANIZATION GSE360_LOW_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_MAC_UP, GSE360_LOW_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_MAC_UP REACTOME_NGF_SIGNALING_VIA_TRKA_FROM_THE_PLASMA_MEMBRANE, REACTOME_NGF_SIGNALING_VIA_TRKA_FROM_THE_PLASMA_MEMBRANE FAELT_B_CLL_WITH_VH3_21_UP, FAELT_B_CLL_WITH_VH3_21_UP GNF2_MSN, GNF2_MSN GCM_TPR, GCM_TPR AGCATT_A_MIR155, AGCATT_A_MIR155 MORF_USP5, MORF_USP5 CATGTAA_MIR496, CATGTAA_MIR496 GO_COATED_MEMBRANE, GO_COATED_MEMBRANE GO_TRANS_GOLGI_NETWORK, GO_TRANS_GOLGI_NETWORK GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ER, GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ER GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PBMCDN, GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PBMCDN REACTOME_MEMBRANE_TRAFFICKING, REACTOME_MEMBRANE_TRAFFICKING MODULE_206, MODULE_206 DACOSTA_UV_RESPONSE_VIA_ERCC3_TTD_UP, DACOSTA_UV_RESPONSE_VIA_ERCC3_TTD_UP IKEDA_MIR1_TARGETS_UP, IKEDA_MIR1_TARGETS_UP GO_MAINTENANCE_OF_LOCATION_IN_CELL, GO_MAINTENANCE_OF_LOCATION_IN_CELL GSE360_DC_VS_MAC_B_MALAYI_LOW_DOSE_DN, GSE360_DC_VS_MAC_B_MALAYI_LOW_DOSE_DN GSE3982_MAC_VS_TH2_UP, GSE3982_MAC_VS_TH2_UP ACAACTT_MIR382, ACAACTT_MIR382 GO_SINGLE_ORGANISM_MEMBRANE_BUDDING, GO_SINGLE_ORGANISM_MEMBRANE_BUDDING REACTOME_ASPARAGINE_N_LINKED_GLYCOSYLATION, REACTOME_ASPARAGINE_N_LINKED_GLYCOSYLATION GO_REGULATION_OF_CHROMATIN_ORGANIZATION, GO_REGULATION_OF_CHROMATIN_ORGANIZATION GSE3982_EOSINOPHIL_VS_TH2_UP, GSE3982_EOSINOPHIL_VS_TH2_UP SESTO_RESPONSE_TO_UV_C2, SESTO_RESPONSE_TO_UV_C2 GO_VESICLE_TARGETING, GO_VESICLE_TARGETING GO_LOCALIZATION_WITHIN_MEMBRANE, GO_LOCALIZATION_WITHIN_MEMBRANE GSE13306_TREG_VS_TCONV_LAMINA_PROPRIA_UP, GSE13306_TREG_VS_TCONV_LAMINA_PROPRIA_UP GO_ORGANELLE_DISASSEMBLY, GO_ORGANELLE_DISASSEMBLY GO_HISTONE_UBIQUITINATION, GO_HISTONE_UBIQUITINATION GO_MAINTENANCE_OF_LOCATION, GO_MAINTENANCE_OF_LOCATION BAE_BRCA1_TARGETS_UP, BAE_BRCA1_TARGETS_UP GNF2_PAK2, GNF2_PAK2 GO_GOLGI_STACK, GO_GOLGI_STACK GO_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT, GO_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT REACTOME_GLOBAL_GENOMIC_NER_GG_NER, REACTOME_GLOBAL_GENOMIC_NER_GG_NER FAELT_B_CLL_WITH_VH3_21_DN, FAELT_B_CLL_WITH_VH3_21_DN LU_AGING_BRAIN_DN, LU_AGING_BRAIN_DN SCHLOSSER_SERUM_RESPONSE_UP, SCHLOSSER_SERUM_RESPONSE_UP GO_REGULATION_OF_PROTEIN_POLYMERIZATION, GO_REGULATION_OF_PROTEIN_POLYMERIZATION GO_COATED_VESICLE_MEMBRANE, GO_COATED_VESICLE_MEMBRANE GO_REGULATION_OF_ACTIN_FILAMENT_LENGTH, GO_REGULATION_OF_ACTIN_FILAMENT_LENGTH GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN, GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN GSE20715_WT_VS_TLR4_KO_6H_OZONE_LUNG_DN, GSE20715_WT_VS_TLR4_KO_6H_OZONE_LUNG_DN AHR_I01, AHR_I01 CGGAARNGGCNG_UNKNOWN, CGGAARNGGCNG_UNKNOWN WANG_ESOPHAGUS_CANCER_VS_NORMAL_DN, WANG_ESOPHAGUS_CANCER_VS_NORMAL_DN GO_AUTOPHAGOSOME, GO_AUTOPHAGOSOME GO_EXOCYTIC_VESICLE, GO_EXOCYTIC_VESICLE PID_ARF_3PATHWAY, PID_ARF_3PATHWAY GO_CYTOPLASMIC_SEQUESTERING_OF_PROTEIN, GO_CYTOPLASMIC_SEQUESTERING_OF_PROTEIN GO_REGULATION_OF_PLASMA_MEMBRANE_ORGANIZATION, GO_REGULATION_OF_PLASMA_MEMBRANE_ORGANIZATION GO_MEMBRANE_DISASSEMBLY, GO_MEMBRANE_DISASSEMBLY MORF_PML, MORF_PML MISSIAGLIA_REGULATED_BY_METHYLATION_UP, MISSIAGLIA_REGULATED_BY_METHYLATION_UP FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN, FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TREG_DN, GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TREG_DN GO_NEGATIVE_REGULATION_OF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY GSE22025_PROGESTERONE_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN, GSE22025_PROGESTERONE_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN SCHAEFFER_PROSTATE_DEVELOPMENT_6HR_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_6HR_UP GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_DN, GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_DN EPPERT_LSC_R, EPPERT_LSC_R WATANABE_RECTAL_CANCER_RADIOOTHERAPY_RESPONSE_DN, WATANABE_RECTAL_CANCER_RADIOOTHERAPY_RESPONSE_DN REACTOME_SIGNALING_BY_ERBB2, REACTOME_SIGNALING_BY_ERBB2 GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_UP, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_UP GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS POMEROY_MEDULLOBLASTOMA_DESMOPLASIC_VS_CLASSIC_UP, POMEROY_MEDULLOBLASTOMA_DESMOPLASIC_VS_CLASSIC_UP KEGG_SNARE_INTERACTIONS_IN_VESICULAR_TRANSPORT, KEGG_SNARE_INTERACTIONS_IN_VESICULAR_TRANSPORT BIOCARTA_PTDINS_PATHWAY, BIOCARTA_PTDINS_PATHWAY CHOW_RASSF1_TARGETS_DN, CHOW_RASSF1_TARGETS_DN GO_VESICLE_COAT, GO_VESICLE_COAT GSE19401_PAM2CSK4_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP, GSE19401_PAM2CSK4_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_UP GO_VESICLE_COATING, GO_VESICLE_COATING KLEIN_PRIMARY_EFFUSION_LYMPHOMA_UP, KLEIN_PRIMARY_EFFUSION_LYMPHOMA_UP GO_SNAP_RECEPTOR_ACTIVITY, GO_SNAP_RECEPTOR_ACTIVITY PID_HEDGEHOG_GLI_PATHWAY, PID_HEDGEHOG_GLI_PATHWAY REACTOME_SIGNALING_BY_FGFR_IN_DISEASE, REACTOME_SIGNALING_BY_FGFR_IN_DISEASE GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY3_UP, GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY3_UP GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_40H_TSST_ACT_UP, GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_40H_TSST_ACT_UP GO_GOLGI_CISTERNA, GO_GOLGI_CISTERNA GO_VESICLE_CYTOSKELETAL_TRAFFICKING, GO_VESICLE_CYTOSKELETAL_TRAFFICKING PID_AJDISS_2PATHWAY, PID_AJDISS_2PATHWAY GO_BINDING_BRIDGING, GO_BINDING_BRIDGING GO_SEX_CHROMOSOME, GO_SEX_CHROMOSOME GO_GPI_ANCHOR_METABOLIC_PROCESS, GO_GPI_ANCHOR_METABOLIC_PROCESS GO_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY, GO_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY GO_ION_CHANNEL_BINDING, GO_ION_CHANNEL_BINDING SIG_INSULIN_RECEPTOR_PATHWAY_IN_CARDIAC_MYOCYTES, SIG_INSULIN_RECEPTOR_PATHWAY_IN_CARDIAC_MYOCYTES REACTOME_TRANSPORT_OF_MATURE_MRNA_DERIVED_FROM_AN_INTRONLESS_TRANSCRIPT, REACTOME_TRANSPORT_OF_MATURE_MRNA_DERIVED_FROM_AN_INTRONLESS_TRANSCRIPT EINAV_INTERFERON_SIGNATURE_IN_CANCER, EINAV_INTERFERON_SIGNATURE_IN_CANCER REACTOME_POST_TRANSLATIONAL_PROTEIN_MODIFICATION, REACTOME_POST_TRANSLATIONAL_PROTEIN_MODIFICATION CAGTCAC_MIR134, CAGTCAC_MIR134 REACTOME_SIGNAL_TRANSDUCTION_BY_L1, REACTOME_SIGNAL_TRANSDUCTION_BY_L1 GO_CELLULAR_PIGMENTATION, GO_CELLULAR_PIGMENTATION GO_SIGNAL_RELEASE, GO_SIGNAL_RELEASE REACTOME_LYSOSOME_VESICLE_BIOGENESIS, REACTOME_LYSOSOME_VESICLE_BIOGENESIS BIOCARTA_CREB_PATHWAY, BIOCARTA_CREB_PATHWAY REACTOME_PKB_MEDIATED_EVENTS, REACTOME_PKB_MEDIATED_EVENTS CAFFAREL_RESPONSE_TO_THC_24HR_5_UP, CAFFAREL_RESPONSE_TO_THC_24HR_5_UP GO_REGULATION_OF_DEPHOSPHORYLATION, GO_REGULATION_OF_DEPHOSPHORYLATION KEGG_PANCREATIC_CANCER, KEGG_PANCREATIC_CANCER GO_AP_TYPE_MEMBRANE_COAT_ADAPTOR_COMPLEX, GO_AP_TYPE_MEMBRANE_COAT_ADAPTOR_COMPLEX GO_MACROMITOPHAGY, GO_MACROMITOPHAGY REACTOME_DOWNSTREAM_SIGNAL_TRANSDUCTION, REACTOME_DOWNSTREAM_SIGNAL_TRANSDUCTION REACTOME_SIGNALING_BY_PDGF, REACTOME_SIGNALING_BY_PDGF LXR_DR4_Q3, LXR_DR4_Q3 GO_REGULATION_OF_PROTEIN_PHOSPHATASE_TYPE_2A_ACTIVITY, GO_REGULATION_OF_PROTEIN_PHOSPHATASE_TYPE_2A_ACTIVITY GO_SNARE_COMPLEX, GO_SNARE_COMPLEX GO_REGULATION_OF_ENDOCYTOSIS, GO_REGULATION_OF_ENDOCYTOSIS GO_TRNA_TRANSPORT, GO_TRNA_TRANSPORT SPIRA_SMOKERS_LUNG_CANCER_UP, SPIRA_SMOKERS_LUNG_CANCER_UP ATCMTNCCGY_UNKNOWN, ATCMTNCCGY_UNKNOWN KEGG_N_GLYCAN_BIOSYNTHESIS, KEGG_N_GLYCAN_BIOSYNTHESIS NIKOLSKY_BREAST_CANCER_8Q23_Q24_AMPLICON, NIKOLSKY_BREAST_CANCER_8Q23_Q24_AMPLICON GO_AUTOPHAGOSOME_ORGANIZATION, GO_AUTOPHAGOSOME_ORGANIZATION GO_ENDOCYTIC_VESICLE_MEMBRANE, GO_ENDOCYTIC_VESICLE_MEMBRANE chr8q24, chr8q24 REACTOME_SIGNALING_BY_FGFR, REACTOME_SIGNALING_BY_FGFR GO_LIPOPROTEIN_BIOSYNTHETIC_PROCESS, GO_LIPOPROTEIN_BIOSYNTHETIC_PROCESS BIOCARTA_FMLP_PATHWAY, BIOCARTA_FMLP_PATHWAY GCCAGT_MIR193A_MIR193B, GCCAGT_MIR193A_MIR193B HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_UP, HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_UP GO_RECEPTOR_METABOLIC_PROCESS, GO_RECEPTOR_METABOLIC_PROCESS GO_INTRA_GOLGI_VESICLE_MEDIATED_TRANSPORT, GO_INTRA_GOLGI_VESICLE_MEDIATED_TRANSPORT GO_CLATHRIN_COATED_ENDOCYTIC_VESICLE_MEMBRANE, GO_CLATHRIN_COATED_ENDOCYTIC_VESICLE_MEMBRANE YAP1_UP, YAP1_UP REACTOME_COPI_MEDIATED_TRANSPORT, REACTOME_COPI_MEDIATED_TRANSPORT CCTGAGT_MIR510, CCTGAGT_MIR510 GO_MEMBRANE_INVAGINATION, GO_MEMBRANE_INVAGINATION GUO_HEX_TARGETS_UP, GUO_HEX_TARGETS_UP GO_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING, GO_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING LIN_MELANOMA_COPY_NUMBER_UP, LIN_MELANOMA_COPY_NUMBER_UP GO_TRANSCRIPTION_FACTOR_TTFC_COMPLEX, GO_TRANSCRIPTION_FACTOR_TTFC_COMPLEX BIOCARTA_RAB_PATHWAY, BIOCARTA_RAB_PATHWAY GO_NEGATIVE_REGULATION_OF_PROTEIN_COMPLEX_ASSEMBLY, GO_NEGATIVE_REGULATION_OF_PROTEIN_COMPLEX_ASSEMBLY SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX4_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX4_DN GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_COMPLEX, GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_COMPLEX BARIS_THYROID_CANCER_DN, BARIS_THYROID_CANCER_DN GO_COPI_COATED_VESICLE_MEMBRANE, GO_COPI_COATED_VESICLE_MEMBRANE LUCAS_HNF4A_TARGETS_UP, LUCAS_HNF4A_TARGETS_UP REACTOME_GAP_JUNCTION_TRAFFICKING, REACTOME_GAP_JUNCTION_TRAFFICKING GSE3982_EOSINOPHIL_VS_TH1_UP, GSE3982_EOSINOPHIL_VS_TH1_UP GO_GOLGI_ASSOCIATED_VESICLE_MEMBRANE, GO_GOLGI_ASSOCIATED_VESICLE_MEMBRANE GO_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR, GO_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR GO_POSITIVE_REGULATION_OF_TRANSLATIONAL_INITIATION, GO_POSITIVE_REGULATION_OF_TRANSLATIONAL_INITIATION REACTOME_RECYCLING_PATHWAY_OF_L1, REACTOME_RECYCLING_PATHWAY_OF_L1 AAGTCCA_MIR422B_MIR422A, AAGTCCA_MIR422B_MIR422A NAGY_TTFC_COMPONENTS_HUMAN, NAGY_TTFC_COMPONENTS_HUMAN REACTOME_FORMATION_OF_INCISION_COMPLEX_IN_GG_NER, REACTOME_FORMATION_OF_INCISION_COMPLEX_IN_GG_NER GO_GOLGI_CISTERNA_MEMBRANE, GO_GOLGI_CISTERNA_MEMBRANE GO_NEGATIVE_REGULATION_OF_ACTIN_FILAMENT_POLYMERIZATION, GO_NEGATIVE_REGULATION_OF_ACTIN_FILAMENT_POLYMERIZATION PID_CDC42_PATHWAY, PID_CDC42_PATHWAY GO_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION, GO_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION PID_AR_NONGENOMIC_PATHWAY, PID_AR_NONGENOMIC_PATHWAY GO_NEGATIVE_REGULATION_OF_HISTONE_MODIFICATION, GO_NEGATIVE_REGULATION_OF_HISTONE_MODIFICATION GO_NEGATIVE_REGULATION_OF_ENDOCYTOSIS, GO_NEGATIVE_REGULATION_OF_ENDOCYTOSIS AGGCAG_MIR18A, AGGCAG_MIR18A REACTOME_PLATELET_SENSITIZATION_BY_LDL, REACTOME_PLATELET_SENSITIZATION_BY_LDL GO_CLATHRIN_COATED_VESICLE, GO_CLATHRIN_COATED_VESICLE GCM_AIP, GCM_AIP RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP, RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP GO_PIGMENTATION, GO_PIGMENTATION KEGG_THYROID_CANCER, KEGG_THYROID_CANCER GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE, GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE GO_APOPTOTIC_CELL_CLEARANCE, GO_APOPTOTIC_CELL_CLEARANCE GO_CLATHRIN_COAT, GO_CLATHRIN_COAT GO_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY, GO_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY GO_POSITIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION GENTILE_UV_RESPONSE_CLUSTER_D9, GENTILE_UV_RESPONSE_CLUSTER_D9 GO_PROTEIN_PHOSPHATASE_2A_BINDING, GO_PROTEIN_PHOSPHATASE_2A_BINDING REACTOME_GAP_JUNCTION_DEGRADATION, REACTOME_GAP_JUNCTION_DEGRADATION REACTOME_INSULIN_RECEPTOR_SIGNALING_CASCADE, REACTOME_INSULIN_RECEPTOR_SIGNALING_CASCADE REACTOME_POST_TRANSLATIONAL_MODIFICATION_SYNTHESIS_OF_GPI_ANCHOR, REACTOME_POST_TRANSLATIONAL_MODIFICATION_SYNTHESIS_OF_GPI_ANCHOR VALK_AML_CLUSTER_I2, VALK_AML_CLUSTER_I2 GO_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY, GO_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY IIZUKA_LIVER_CANCER_PROGRESSION_L1_G1_DN, IIZUKA_LIVER_CANCER_PROGRESSION_L1_G1_DN GO_AUTOPHAGOSOME_MEMBRANE, GO_AUTOPHAGOSOME_MEMBRANE GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_16h_TSST_ACT_UP, GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_16h_TSST_ACT_UP GO_COPI_VESICLE_COAT, GO_COPI_VESICLE_COAT PID_A6B1_A6B4_INTEGRIN_PATHWAY, PID_A6B1_A6B4_INTEGRIN_PATHWAY GO_NUCLEOPHAGY, GO_NUCLEOPHAGY GO_CLATHRIN_BINDING, GO_CLATHRIN_BINDING GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE, GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE GO_CLATHRIN_COATED_ENDOCYTIC_VESICLE, GO_CLATHRIN_COATED_ENDOCYTIC_VESICLE GO_ARF_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY, GO_ARF_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY TCATATGA_MIR376A_MIR376B, TCATATGA_MIR376A_MIR376B REACTOME_DOWNSTREAM_SIGNALING_OF_ACTIVATED_FGFR, REACTOME_DOWNSTREAM_SIGNALING_OF_ACTIVATED_FGFR LEE_METASTASIS_AND_RNA_PROCESSING_UP, LEE_METASTASIS_AND_RNA_PROCESSING_UP GO_INORGANIC_ION_IMPORT_INTO_CELL, GO_INORGANIC_ION_IMPORT_INTO_CELL GO_CLATHRIN_COAT_OF_ENDOCYTIC_VESICLE, GO_CLATHRIN_COAT_OF_ENDOCYTIC_VESICLE GO_NUCLEOTIDE_EXCISION_REPAIR_COMPLEX, GO_NUCLEOTIDE_EXCISION_REPAIR_COMPLEX GO_NEGATIVE_REGULATION_OF_CHROMATIN_MODIFICATION, GO_NEGATIVE_REGULATION_OF_CHROMATIN_MODIFICATION GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN, GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN GO_KINASE_INHIBITOR_ACTIVITY, GO_KINASE_INHIBITOR_ACTIVITY GO_REGULATION_OF_HISTONE_DEACETYLATION, GO_REGULATION_OF_HISTONE_DEACETYLATION GO_RETROGRADE_TRANSPORT_VESICLE_RECYCLING_WITHIN_GOLGI, GO_RETROGRADE_TRANSPORT_VESICLE_RECYCLING_WITHIN_GOLGI BIOCARTA_ARAP_PATHWAY, BIOCARTA_ARAP_PATHWAY GO_GTP_DEPENDENT_PROTEIN_BINDING, GO_GTP_DEPENDENT_PROTEIN_BINDING GO_PHOSPHATIDYLINOSITOL_3_KINASE_BINDING, GO_PHOSPHATIDYLINOSITOL_3_KINASE_BINDING GO_BARBED_END_ACTIN_FILAMENT_CAPPING, GO_BARBED_END_ACTIN_FILAMENT_CAPPING REACTOME_ENERGY_DEPENDENT_REGULATION_OF_MTOR_BY_LKB1_AMPK, REACTOME_ENERGY_DEPENDENT_REGULATION_OF_MTOR_BY_LKB1_AMPK GO_REGULATION_OF_CELL_MATURATION, GO_REGULATION_OF_CELL_MATURATION GO_ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE, GO_ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_I2, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_I2 GO_HISTONE_H2A_UBIQUITINATION, GO_HISTONE_H2A_UBIQUITINATION KEGG_GLYCOSYLPHOSPHATIDYLINOSITOL_GPIANCHOR_BIOSYNTHESIS, KEGG_GLYCOSYLPHOSPHATIDYLINOSITOL_GPIANCHOR_BIOSYNTHESIS GO_PHOSPHOPROTEIN_BINDING, GO_PHOSPHOPROTEIN_BINDING MODULE_395, MODULE_395 GO_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS, GO_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS REACTOME_CALNEXIN_CALRETICULIN_CYCLE, REACTOME_CALNEXIN_CALRETICULIN_CYCLE GO_PROTEIN_LOCALIZATION_TO_GOLGI_APPARATUS, GO_PROTEIN_LOCALIZATION_TO_GOLGI_APPARATUS DANG_MYC_TARGETS_DN, DANG_MYC_TARGETS_DN GO_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT, GO_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE, REACTOME_N_GLYCAN_TRIMMING_IN_THE_ER_AND_CALNEXIN_CALRETICULIN_CYCLE GO_REGULATION_OF_NUCLEOBASE_CONTAINING_COMPOUND_TRANSPORT, GO_REGULATION_OF_NUCLEOBASE_CONTAINING_COMPOUND_TRANSPORT BIOCARTA_TCR_PATHWAY, BIOCARTA_TCR_PATHWAY BIOCARTA_GPCR_PATHWAY, BIOCARTA_GPCR_PATHWAY REACTOME_SOS_MEDIATED_SIGNALING, REACTOME_SOS_MEDIATED_SIGNALING REACTOME_SYNTHESIS_OF_SUBSTRATES_IN_N_GLYCAN_BIOSYTHESIS, REACTOME_SYNTHESIS_OF_SUBSTRATES_IN_N_GLYCAN_BIOSYTHESIS chr7q22, chr7q22 GO_ORGAN_GROWTH, GO_ORGAN_GROWTH GO_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION, GO_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION REACTOME_CTNNB1_PHOSPHORYLATION_CASCADE, REACTOME_CTNNB1_PHOSPHORYLATION_CASCADE GO_PROTEIN_LOCALIZATION_TO_CENTROSOME, GO_PROTEIN_LOCALIZATION_TO_CENTROSOME GO_CYTOPLASMIC_DYNEIN_COMPLEX, GO_CYTOPLASMIC_DYNEIN_COMPLEX GO_POLYSACCHARIDE_CATABOLIC_PROCESS, GO_POLYSACCHARIDE_CATABOLIC_PROCESS GO_POST_EMBRYONIC_DEVELOPMENT, GO_POST_EMBRYONIC_DEVELOPMENT GO_R_SMAD_BINDING, GO_R_SMAD_BINDING RAMPON_ENRICHED_LEARNING_ENVIRONMENT_LATE_UP, RAMPON_ENRICHED_LEARNING_ENVIRONMENT_LATE_UP REACTOME_GRB2_EVENTS_IN_ERBB2_SIGNALING, REACTOME_GRB2_EVENTS_IN_ERBB2_SIGNALING GO_MUSCLE_CELL_CELLULAR_HOMEOSTASIS, GO_MUSCLE_CELL_CELLULAR_HOMEOSTASIS
F_RAB1A, MORF_RAB1A	