ELL_AND_IL12_TREATED_MELANOMA_DAY3_UP, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP
GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP
GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN
GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_DN, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_DN
GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN, GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN
GSE32423_IL7_VS_IL7_IL4_MEMORY_CD8_TCELL_UP, GSE32423_IL7_VS_IL7_IL4_MEMORY_CD8_TCELL_UP
GSE369_SOCS3_KO_VS_WT_LIVER_DN, GSE369_SOCS3_KO_VS_WT_LIVER_DN
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_12H_CD4_TCELL_UP, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_12H_CD4_TCELL_UP
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_DN
GSE3982_DC_VS_BASOPHIL_UP, GSE3982_DC_VS_BASOPHIL_UP
GSE13493_DP_VS_CD8POS_THYMOCYTE_UP, GSE13493_DP_VS_CD8POS_THYMOCYTE_UP
MIR6893_5P, MIR6893_5P
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP
GSE3982_DC_VS_NKCELL_UP, GSE3982_DC_VS_NKCELL_UP
MIR6808_5P, MIR6808_5P
GOBP_VACUOLAR_TRANSPORT, GOBP_VACUOLAR_TRANSPORT
GOBP_REGULATION_OF_CELL_PROJECTION_ASSEMBLY, GOBP_REGULATION_OF_CELL_PROJECTION_ASSEMBLY
MIR520F_5P, MIR520F_5P
GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP, GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP
GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THYMUS_DN, GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THYMUS_DN
MANNE_COVID19_NONICU_VS_HEALTHY_DONOR_PLATELETS_UP, MANNE_COVID19_NONICU_VS_HEALTHY_DONOR_PLATELETS_UP
GSE11864_CSF1_PAM3CYS_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN, GSE11864_CSF1_PAM3CYS_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN
GOBP_ORGANELLE_FUSION, GOBP_ORGANELLE_FUSION
GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_UP, GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_UP
GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP, GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP
GSE14769_UNSTIM_VS_20MIN_LPS_BMDM_DN, GSE14769_UNSTIM_VS_20MIN_LPS_BMDM_DN
GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP
MIR513B_3P, MIR513B_3P
HP_TAPERED_FINGER, HP_TAPERED_FINGER
MIR3920, MIR3920
GOBP_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_ALPHA_BETA_T_CELL_ACTIVATION
MIR4318, MIR4318
ZNF669_TARGET_GENES, ZNF669_TARGET_GENES
GSE13522_CTRL_VS_T_CRUZI_BRAZIL_STRAIN_INF_SKIN_DN, GSE13522_CTRL_VS_T_CRUZI_BRAZIL_STRAIN_INF_SKIN_DN
SUNG_METASTASIS_STROMA_UP, SUNG_METASTASIS_STROMA_UP
WP_PANCREATIC_ADENOCARCINOMA_PATHWAY, WP_PANCREATIC_ADENOCARCINOMA_PATHWAY
HP_ABNORMAL_METABOLISM, HP_ABNORMAL_METABOLISM
GOBP_LYSOSOMAL_TRANSPORT, GOBP_LYSOSOMAL_TRANSPORT
MIR197_3P, MIR197_3P
MIR671_5P, MIR671_5P
HP_HALLUCINATIONS, HP_HALLUCINATIONS
MIR202_5P, MIR202_5P
GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
MIR935, MIR935
BROWNE_HCMV_INFECTION_20HR_DN, BROWNE_HCMV_INFECTION_20HR_DN
MIR512_3P, MIR512_3P
GOBP_ORGANELLE_MEMBRANE_FUSION, GOBP_ORGANELLE_MEMBRANE_FUSION
MIR4793_3P, MIR4793_3P
REACTOME_ONCOGENIC_MAPK_SIGNALING, REACTOME_ONCOGENIC_MAPK_SIGNALING
HP_ABNORMAL_VENOUS_MORPHOLOGY, HP_ABNORMAL_VENOUS_MORPHOLOGY
HP_ABNORMALITY_OF_HAIR_TEXTURE, HP_ABNORMALITY_OF_HAIR_TEXTURE
GOCC_VESICLE_TETHERING_COMPLEX, GOCC_VESICLE_TETHERING_COMPLEX
MIR6806_3P, MIR6806_3P
HP_2_3_TOE_SYNDACTYLY, HP_2_3_TOE_SYNDACTYLY
MIR3928_5P, MIR3928_5P
PID_FCER1_PATHWAY, PID_FCER1_PATHWAY
GOBP_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_ALPHA_BETA_T_CELL_DIFFERENTIATION
MIR4727_5P, MIR4727_5P
GOBP_MEMBRANE_FUSION, GOBP_MEMBRANE_FUSION
MIR5585_3P, MIR5585_3P
MIR4494, MIR4494
HP_EXCESSIVE_SALIVATION, HP_EXCESSIVE_SALIVATION
MIR4499, MIR4499
GGGATGC_MIR3245P, GGGATGC_MIR3245P
MIR584_3P, MIR584_3P
GOBP_POSITIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY
GUO_HEX_TARGETS_UP, GUO_HEX_TARGETS_UP
GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_0, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_0
GSE29949_CD8_POS_DC_SPLEEN_VS_DC_BRAIN_DN, GSE29949_CD8_POS_DC_SPLEEN_VS_DC_BRAIN_DN
GOBP_LAMELLIPODIUM_ORGANIZATION, GOBP_LAMELLIPODIUM_ORGANIZATION
WP_ONCOSTATIN_M_SIGNALING_PATHWAY, WP_ONCOSTATIN_M_SIGNALING_PATHWAY
MIR3179, MIR3179
GOMF_OMEGA_PEPTIDASE_ACTIVITY, GOMF_OMEGA_PEPTIDASE_ACTIVITY
HP_X_LINKED_RECESSIVE_INHERITANCE, HP_X_LINKED_RECESSIVE_INHERITANCE
MIR5088_3P, MIR5088_3P
REACTOME_FOXO_MEDIATED_TRANSCRIPTION, REACTOME_FOXO_MEDIATED_TRANSCRIPTION
WENDT_COHESIN_TARGETS_UP, WENDT_COHESIN_TARGETS_UP
MIR7515, MIR7515
MIR141_5P, MIR141_5P
ZNF34_TARGET_GENES, ZNF34_TARGET_GENES
GOMF_PALMITOYLTRANSFERASE_ACTIVITY, GOMF_PALMITOYLTRANSFERASE_ACTIVITY
REACTOME_REGULATION_OF_TP53_EXPRESSION_AND_DEGRADATION, REACTOME_REGULATION_OF_TP53_EXPRESSION_AND_DEGRADATION
GSE29618_PRE_VS_DAY7_FLU_VACCINE_MONOCYTE_UP, GSE29618_PRE_VS_DAY7_FLU_VACCINE_MONOCYTE_UP
GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX
HP_LONG_PALPEBRAL_FISSURE, HP_LONG_PALPEBRAL_FISSURE
GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP
MIR3912_5P, MIR3912_5P
GOBP_PROTEIN_TARGETING_TO_VACUOLE, GOBP_PROTEIN_TARGETING_TO_VACUOLE
ZSCAN18_TARGET_GENES, ZSCAN18_TARGET_GENES
GSE24574_BCL6_HIGH_TFH_VS_TCONV_CD4_TCELL_DN, GSE24574_BCL6_HIGH_TFH_VS_TCONV_CD4_TCELL_DN
REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS, REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS
GOBP_POSITIVE_REGULATION_OF_DEPHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_DEPHOSPHORYLATION
HP_CONGENITAL_HIP_DISLOCATION, HP_CONGENITAL_HIP_DISLOCATION
GOBP_LAMELLIPODIUM_ASSEMBLY, GOBP_LAMELLIPODIUM_ASSEMBLY
GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN
BIOCARTA_FCER1_PATHWAY, BIOCARTA_FCER1_PATHWAY
JOHNSTONE_PARVB_TARGETS_1_DN, JOHNSTONE_PARVB_TARGETS_1_DN
ZHAN_MULTIPLE_MYELOMA_CD1_DN, ZHAN_MULTIPLE_MYELOMA_CD1_DN
MIR6500_3P, MIR6500_3P
HP_PREMATURE_SKIN_WRINKLING, HP_PREMATURE_SKIN_WRINKLING
MIR556_3P, MIR556_3P
MIR4769_3P, MIR4769_3P
MIR6817_5P, MIR6817_5P
GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPHOSPHORYLATION
MIR4252, MIR4252
ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_3DY_DN, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_3DY_DN
XU_GH1_EXOGENOUS_TARGETS_DN, XU_GH1_EXOGENOUS_TARGETS_DN
GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE
MIR374C_3P, MIR374C_3P
GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS
MIR500B_3P, MIR500B_3P
HNF3_Q6, HNF3_Q6
MIR215_5P, MIR215_5P
GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_III
BIOCARTA_KERATINOCYTE_PATHWAY, BIOCARTA_KERATINOCYTE_PATHWAY
PID_MYC_PATHWAY, PID_MYC_PATHWAY
MIR4509, MIR4509
HP_TOE_CLINODACTYLY, HP_TOE_CLINODACTYLY
HP_ABNORMAL_FEMORAL_HEAD_MORPHOLOGY, HP_ABNORMAL_FEMORAL_HEAD_MORPHOLOGY
HP_ABNORMAL_T_CELL_SUBSET_DISTRIBUTION, HP_ABNORMAL_T_CELL_SUBSET_DISTRIBUTION
GOBP_AUTOPHAGOSOME_MATURATION, GOBP_AUTOPHAGOSOME_MATURATION
MORF_DMPK, MORF_DMPK
MIR192_5P, MIR192_5P
GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT, GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT
PID_INSULIN_PATHWAY, PID_INSULIN_PATHWAY
HP_CHRONIC_CONSTIPATION, HP_CHRONIC_CONSTIPATION
GOBP_ACTIVATION_OF_JUN_KINASE_ACTIVITY, GOBP_ACTIVATION_OF_JUN_KINASE_ACTIVITY
GOMF_TRANSITION_METAL_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_TRANSITION_METAL_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_S_ACYLTRANSFERASE_ACTIVITY, GOMF_S_ACYLTRANSFERASE_ACTIVITY
MIR7703, MIR7703
REACTOME_NRAGE_SIGNALS_DEATH_THROUGH_JNK, REACTOME_NRAGE_SIGNALS_DEATH_THROUGH_JNK
CHEN_LIVER_METABOLISM_QTL_CIS, CHEN_LIVER_METABOLISM_QTL_CIS
HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE, HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE
GSE43863_NAIVE_VS_MEMORY_TH1_CD4_TCELL_D150_LCMV_DN, GSE43863_NAIVE_VS_MEMORY_TH1_CD4_TCELL_D150_LCMV_DN
GOBP_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION, GOBP_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION
HP_DISPROPORTIONATE_TALL_STATURE, HP_DISPROPORTIONATE_TALL_STATURE
HP_PERIORBITAL_FULLNESS, HP_PERIORBITAL_FULLNESS
MIR28_3P, MIR28_3P
ZHANG_ANTIVIRAL_RESPONSE_TO_RIBAVIRIN_DN, ZHANG_ANTIVIRAL_RESPONSE_TO_RIBAVIRIN_DN
GSE10422_WT_VS_BAFF_TRANSGENIC_LN_BCELL_DN, GSE10422_WT_VS_BAFF_TRANSGENIC_LN_BCELL_DN
GOCC_MICROVILLUS, GOCC_MICROVILLUS
HP_MALE_REPRODUCTIVE_SYSTEM_NEOPLASM, HP_MALE_REPRODUCTIVE_SYSTEM_NEOPLASM
MIR3614_5P, MIR3614_5P
HP_DOWN_SLOPING_SHOULDERS, HP_DOWN_SLOPING_SHOULDERS
MORF_BCL2L11, MORF_BCL2L11
PAX8_B, PAX8_B
GOBP_NEGATIVE_REGULATION_OF_COLD_INDUCED_THERMOGENESIS, GOBP_NEGATIVE_REGULATION_OF_COLD_INDUCED_THERMOGENESIS
GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_DN, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_DN
MORF_BMPR2, MORF_BMPR2
GOBP_PEPTIDYL_CYSTEINE_MODIFICATION, GOBP_PEPTIDYL_CYSTEINE_MODIFICATION
MIR4740_5P, MIR4740_5P
HP_ABNORMALITY_OF_LOWER_LIMB_EPIPHYSIS_MORPHOLOGY, HP_ABNORMALITY_OF_LOWER_LIMB_EPIPHYSIS_MORPHOLOGY
GOBP_REGULATION_OF_LIPID_CATABOLIC_PROCESS, GOBP_REGULATION_OF_LIPID_CATABOLIC_PROCESS
GOBP_RESPONSE_TO_INTERFERON_ALPHA, GOBP_RESPONSE_TO_INTERFERON_ALPHA
MCCABE_HOXC6_TARGETS_CANCER_DN, MCCABE_HOXC6_TARGETS_CANCER_DN
GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION
GOBP_PHOSPHOLIPID_CATABOLIC_PROCESS, GOBP_PHOSPHOLIPID_CATABOLIC_PROCESS
BIOCARTA_SPPA_PATHWAY, BIOCARTA_SPPA_PATHWAY
GOCC_ORGANELLE_MEMBRANE_CONTACT_SITE, GOCC_ORGANELLE_MEMBRANE_CONTACT_SITE
REACTOME_INACTIVATION_OF_CSF3_G_CSF_SIGNALING, REACTOME_INACTIVATION_OF_CSF3_G_CSF_SIGNALING
GOBP_T_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_T_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE
GOBP_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GOBP_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY
MIR4715_3P, MIR4715_3P
GOBP_PEPTIDYL_L_CYSTEINE_S_PALMITOYLATION, GOBP_PEPTIDYL_L_CYSTEINE_S_PALMITOYLATION
GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY
HP_ABNORMAL_PULMONARY_VALVE_MORPHOLOGY, HP_ABNORMAL_PULMONARY_VALVE_MORPHOLOGY
HP_MASK_LIKE_FACIES, HP_MASK_LIKE_FACIES
HP_COARSE_HAIR, HP_COARSE_HAIR
GOBP_REGULATION_OF_FILOPODIUM_ASSEMBLY, GOBP_REGULATION_OF_FILOPODIUM_ASSEMBLY
GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT, GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT
GOBP_FIBROBLAST_APOPTOTIC_PROCESS, GOBP_FIBROBLAST_APOPTOTIC_PROCESS
MIR4638_3P, MIR4638_3P
GOBP_BROWN_FAT_CELL_DIFFERENTIATION, GOBP_BROWN_FAT_CELL_DIFFERENTIATION
LANDIS_BREAST_CANCER_PROGRESSION_UP, LANDIS_BREAST_CANCER_PROGRESSION_UP
GOMF_LIGASE_ACTIVITY_FORMING_CARBON_SULFUR_BONDS, GOMF_LIGASE_ACTIVITY_FORMING_CARBON_SULFUR_BONDS
GOBP_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT, GOBP_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT
GOBP_T_CELL_SELECTION, GOBP_T_CELL_SELECTION
HP_URTICARIA, HP_URTICARIA
GOBP_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GOBP_T_HELPER_1_TYPE_IMMUNE_RESPONSE
MIR5002_3P, MIR5002_3P
GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS
MIR146B_3P, MIR146B_3P
MIR1231, MIR1231
GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY
BERNARD_PPAPDC1B_TARGETS_DN, BERNARD_PPAPDC1B_TARGETS_DN
GOBP_REGULATION_OF_SUPEROXIDE_ANION_GENERATION, GOBP_REGULATION_OF_SUPEROXIDE_ANION_GENERATION
GOBP_COPPER_ION_TRANSPORT, GOBP_COPPER_ION_TRANSPORT
GOMF_CARBOXYLIC_ESTER_HYDROLASE_ACTIVITY, GOMF_CARBOXYLIC_ESTER_HYDROLASE_ACTIVITY
REACTOME_CD28_CO_STIMULATION, REACTOME_CD28_CO_STIMULATION
HP_ABNORMALITY_OF_THE_2ND_TOE, HP_ABNORMALITY_OF_THE_2ND_TOE
GOBP_MORPHOGENESIS_OF_AN_EPITHELIAL_SHEET, GOBP_MORPHOGENESIS_OF_AN_EPITHELIAL_SHEET
MIR7109_3P, MIR7109_3P
GOBP_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY, GOBP_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY
TERAO_AOX4_TARGETS_HG_UP, TERAO_AOX4_TARGETS_HG_UP
HP_PROMINENT_SUPERFICIAL_BLOOD_VESSELS, HP_PROMINENT_SUPERFICIAL_BLOOD_VESSELS
BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP
BALDWIN_PRKCI_TARGETS_UP, BALDWIN_PRKCI_TARGETS_UP
GOBP_PROTEIN_PALMITOYLATION, GOBP_PROTEIN_PALMITOYLATION
GOBP_REGULATION_OF_HOMOTYPIC_CELL_CELL_ADHESION, GOBP_REGULATION_OF_HOMOTYPIC_CELL_CELL_ADHESION
ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN, ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN
MIR4515, MIR4515
HP_DELAYED_CLOSURE_OF_THE_ANTERIOR_FONTANELLE, HP_DELAYED_CLOSURE_OF_THE_ANTERIOR_FONTANELLE
GOBP_ER_NUCLEUS_SIGNALING_PATHWAY, GOBP_ER_NUCLEUS_SIGNALING_PATHWAY
HP_UVEITIS, HP_UVEITIS
WP_FBXL10_ENHANCEMENT_OF_MAPERK_SIGNALING_IN_DIFFUSE_LARGE_BCELL_LYMPHOMA, WP_FBXL10_ENHANCEMENT_OF_MAPERK_SIGNALING_IN_DIFFUSE_LARGE_BCELL_LYMPHOMA
GOBP_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS
MIR339_5P, MIR339_5P
GOMF_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOMF_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY
WP_4HYDROXYTAMOXIFEN_DEXAMETHASONE_AND_RETINOIC_ACIDS_REGULATION_OF_P27_EXPRESSION, WP_4HYDROXYTAMOXIFEN_DEXAMETHASONE_AND_RETINOIC_ACIDS_REGULATION_OF_P27_EXPRESS
HP_RETINAL_HEMORRHAGE, HP_RETINAL_HEMORRHAGE
GOMF_PROTEIN_DEACETYLASE_ACTIVITY, GOMF_PROTEIN_DEACETYLASE_ACTIVITY
GOBP_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING, GOBP_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING
MIR4711_3P, MIR4711_3P
MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER, MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER
HP_APLASIA_HYPOPLASIA_OF_THE_ABDOMINAL_WALL_MUSCULATURE, HP_APLASIA_HYPOPLASIA_OF_THE_ABDOMINAL_WALL_MUSCULATURE
HP_MILD_HEARING_IMPAIRMENT, HP_MILD_HEARING_IMPAIRMENT
ZHU_SKIL_TARGETS_UP, ZHU_SKIL_TARGETS_UP
HP_SCHIZOPHRENIA, HP_SCHIZOPHRENIA
GOBP_ENDOSOMAL_VESICLE_FUSION, GOBP_ENDOSOMAL_VESICLE_FUSION
WP_P38_MAPK_SIGNALING_PATHWAY, WP_P38_MAPK_SIGNALING_PATHWAY
HP_EXCESSIVE_WRINKLED_SKIN, HP_EXCESSIVE_WRINKLED_SKIN
BIOCARTA_AGR_PATHWAY, BIOCARTA_AGR_PATHWAY
HP_EEG_WITH_MULTIFOCAL_SLOW_ACTIVITY, HP_EEG_WITH_MULTIFOCAL_SLOW_ACTIVITY
HP_CHRONIC_PULMONARY_OBSTRUCTION, HP_CHRONIC_PULMONARY_OBSTRUCTION
GOMF_NON_MEMBRANE_SPANNING_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOMF_NON_MEMBRANE_SPANNING_PROTEIN_TYROSINE_KINASE_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS
GSE33292_DN3_THYMOCYTE_VS_TCF1_KO_TCELL_LYMPHOMA_DN, GSE33292_DN3_THYMOCYTE_VS_TCF1_KO_TCELL_LYMPHOMA_DN
HP_APLASIA_INVOLVING_BONES_OF_THE_EXTREMITIES, HP_APLASIA_INVOLVING_BONES_OF_THE_EXTREMITIES
WP_NEURAL_CREST_CELL_MIGRATION_DURING_DEVELOPMENT, WP_NEURAL_CREST_CELL_MIGRATION_DURING_DEVELOPMENT
BIOCARTA_P38MAPK_PATHWAY, BIOCARTA_P38MAPK_PATHWAY
XU_RESPONSE_TO_TRETINOIN_UP, XU_RESPONSE_TO_TRETINOIN_UP
GOBP_MITOCHONDRION_DISTRIBUTION, GOBP_MITOCHONDRION_DISTRIBUTION
HP_CUBITUS_VALGUS, HP_CUBITUS_VALGUS
HP_ABNORMAL_PROXIMAL_PHALANX_MORPHOLOGY_OF_THE_HAND, HP_ABNORMAL_PROXIMAL_PHALANX_MORPHOLOGY_OF_THE_HAND
MIR1225_5P, MIR1225_5P
GOBP_VIRAL_BUDDING_VIA_HOST_ESCRT_COMPLEX, GOBP_VIRAL_BUDDING_VIA_HOST_ESCRT_COMPLEX
REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION, REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION
HP_SHORTENING_OF_ALL_DISTAL_PHALANGES_OF_THE_FINGERS, HP_SHORTENING_OF_ALL_DISTAL_PHALANGES_OF_THE_FINGERS
REACTOME_VEGFR2_MEDIATED_VASCULAR_PERMEABILITY, REACTOME_VEGFR2_MEDIATED_VASCULAR_PERMEABILITY
GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS, GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_D
WP_KISSPEPTINKISSPEPTIN_RECEPTOR_SYSTEM_IN_THE_OVARY, WP_KISSPEPTINKISSPEPTIN_RECEPTOR_SYSTEM_IN_THE_OVARY
HP_CURLY_HAIR, HP_CURLY_HAIR
GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION
HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7DY_TOP_100_DEG_EX_VIVO_UP, HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7D
GOBP_POSITIVE_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY
HP_THICK_HAIR, HP_THICK_HAIR
GOBP_REGULATION_OF_ACTIN_NUCLEATION, GOBP_REGULATION_OF_ACTIN_NUCLEATION
GOBP_VITAMIN_TRANSMEMBRANE_TRANSPORT, GOBP_VITAMIN_TRANSMEMBRANE_TRANSPORT
WP_ALPHA_6_BETA_4_SIGNALING_PATHWAY, WP_ALPHA_6_BETA_4_SIGNALING_PATHWAY
MIR6816_3P, MIR6816_3P
HP_PES_VALGUS, HP_PES_VALGUS
GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION, GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION
HP_ABNORMALITY_OF_FIBULA_MORPHOLOGY, HP_ABNORMALITY_OF_FIBULA_MORPHOLOGY
GOBP_SUPEROXIDE_METABOLIC_PROCESS, GOBP_SUPEROXIDE_METABOLIC_PROCESS
MIR6753_3P, MIR6753_3P
GOCC_TORC1_COMPLEX, GOCC_TORC1_COMPLEX
HP_BLOODY_DIARRHEA, HP_BLOODY_DIARRHEA
WINTER_HYPOXIA_DN, WINTER_HYPOXIA_DN
MIR7107_3P, MIR7107_3P
HP_SEBORRHEIC_DERMATITIS, HP_SEBORRHEIC_DERMATITIS
GOBP_TRIGLYCERIDE_BIOSYNTHETIC_PROCESS, GOBP_TRIGLYCERIDE_BIOSYNTHETIC_PROCESS
HP_RETICULOCYTOSIS, HP_RETICULOCYTOSIS
HP_ABNORMAL_MORPHOLOGY_OF_THE_ABDOMINAL_MUSCULATURE, HP_ABNORMAL_MORPHOLOGY_OF_THE_ABDOMINAL_MUSCULATURE
WP_UNFOLDED_PROTEIN_RESPONSE, WP_UNFOLDED_PROTEIN_RESPONSE
KASLER_HDAC7_TARGETS_1_DN, KASLER_HDAC7_TARGETS_1_DN
HP_MODERATE_SENSORINEURAL_HEARING_IMPAIRMENT, HP_MODERATE_SENSORINEURAL_HEARING_IMPAIRMENT
GOMF_MAP_KINASE_KINASE_KINASE_ACTIVITY, GOMF_MAP_KINASE_KINASE_KINASE_ACTIVITY
MIR4771, MIR4771
HP_INCISOR_MACRODONTIA, HP_INCISOR_MACRODONTIA
GOBP_T_HELPER_1_CELL_DIFFERENTIATION, GOBP_T_HELPER_1_CELL_DIFFERENTIATION
GOMF_ADENYLYLTRANSFERASE_ACTIVITY, GOMF_ADENYLYLTRANSFERASE_ACTIVITY
GOBP_REGULATION_OF_PROTEIN_DEACETYLATION, GOBP_REGULATION_OF_PROTEIN_DEACETYLATION
MORF_MAP2K7, MORF_MAP2K7
MIR656_5P, MIR656_5P
HP_SLENDER_TOE, HP_SLENDER_TOE
WP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, WP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION
REACTOME_CD209_DC_SIGN_SIGNALING, REACTOME_CD209_DC_SIGN_SIGNALING
GOBP_CELLULAR_RESPONSE_TO_CADMIUM_ION, GOBP_CELLULAR_RESPONSE_TO_CADMIUM_ION
GOMF_CALMODULIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOMF_CALMODULIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
GOBP_SUPEROXIDE_ANION_GENERATION, GOBP_SUPEROXIDE_ANION_GENERATION
HP_JUVENILE_ASEPTIC_NECROSIS, HP_JUVENILE_ASEPTIC_NECROSIS
HP_PROMINENT_NASOLABIAL_FOLD, HP_PROMINENT_NASOLABIAL_FOLD
HP_ABNORMAL_PELVIS_BONE_MORPHOLOGY, HP_ABNORMAL_PELVIS_BONE_MORPHOLOGY
HP_BULL_S_EYE_MACULOPATHY, HP_BULL_S_EYE_MACULOPATHY
GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
PIEPOLI_LGI1_TARGETS_UP, PIEPOLI_LGI1_TARGETS_UP
GOBP_POSITIVE_REGULATION_OF_PROTEIN_ACETYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_ACETYLATION
GOBP_IRON_ION_TRANSMEMBRANE_TRANSPORT, GOBP_IRON_ION_TRANSMEMBRANE_TRANSPORT
HP_GENERALIZED_JOINT_LAXITY, HP_GENERALIZED_JOINT_LAXITY
KIM_HYPOXIA, KIM_HYPOXIA
BACOLOD_RESISTANCE_TO_ALKYLATING_AGENTS_UP, BACOLOD_RESISTANCE_TO_ALKYLATING_AGENTS_UP
GOBP_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN
MIR4662A_3P, MIR4662A_3P
GOCC_ESCRT_I_COMPLEX, GOCC_ESCRT_I_COMPLEX
HP_VARICOSE_VEINS, HP_VARICOSE_VEINS
REACTOME_FOXO_MEDIATED_TRANSCRIPTION_OF_CELL_DEATH_GENES, REACTOME_FOXO_MEDIATED_TRANSCRIPTION_OF_CELL_DEATH_GENES
GOBP_GLYCEROPHOSPHOLIPID_CATABOLIC_PROCESS, GOBP_GLYCEROPHOSPHOLIPID_CATABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_GLUCOSE_IMPORT, GOBP_POSITIVE_REGULATION_OF_GLUCOSE_IMPORT
HP_ABNORMAL_CIRCULATING_IGE_LEVEL, HP_ABNORMAL_CIRCULATING_IGE_LEVEL
SHIN_B_CELL_LYMPHOMA_CLUSTER_2, SHIN_B_CELL_LYMPHOMA_CLUSTER_2
HP_ABNORMAL_NATURAL_KILLER_CELL_PHYSIOLOGY, HP_ABNORMAL_NATURAL_KILLER_CELL_PHYSIOLOGY
HP_HEMIMEGALENCEPHALY, HP_HEMIMEGALENCEPHALY
GOBP_PROSTANOID_METABOLIC_PROCESS, GOBP_PROSTANOID_METABOLIC_PROCESS
GOBP_REGULATION_OF_RECEPTOR_INTERNALIZATION, GOBP_REGULATION_OF_RECEPTOR_INTERNALIZATION
GOBP_POSITIVE_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT
GOMF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN_PHOSPHOSERINE_BINDING, GOMF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN_PHOSPHOSERINE_BINDING
MIR3690, MIR3690
CGTCTTA_MIR208, CGTCTTA_MIR208
CAO_BLOOD_FLUMIST_AGE_05_14YO_30DY_DN, CAO_BLOOD_FLUMIST_AGE_05_14YO_30DY_DN
GOBP_NEURON_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_NEURON_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS
GOMF_RDNA_BINDING, GOMF_RDNA_BINDING
GOBP_REGULATION_OF_SUPEROXIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_SUPEROXIDE_METABOLIC_PROCESS
MIR5196_3P, MIR5196_3P
HP_ABNORMALITY_OF_THE_NASOLABIAL_REGION, HP_ABNORMALITY_OF_THE_NASOLABIAL_REGION
GOBP_NEUROTROPHIN_SIGNALING_PATHWAY, GOBP_NEUROTROPHIN_SIGNALING_PATHWAY
GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL
HP_HEMATEMESIS, HP_HEMATEMESIS
HP_VENOUS_INSUFFICIENCY, HP_VENOUS_INSUFFICIENCY
GOBP_PROTEIN_INSERTION_INTO_ER_MEMBRANE_BY_STOP_TRANSFER_MEMBRANE_ANCHOR_SEQUENCE, GOBP_PROTEIN_INSERTION_INTO_ER_MEMBRANE_BY_STOP_TRANSFER_MEMBRANE_ANCHOR_SEQUEN
GOCC_EMC_COMPLEX, GOCC_EMC_COMPLEX
GOMF_IRON_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_IRON_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
REACTOME_REGULATION_OF_FOXO_TRANSCRIPTIONAL_ACTIVITY_BY_ACETYLATION, REACTOME_REGULATION_OF_FOXO_TRANSCRIPTIONAL_ACTIVITY_BY_ACETYLATION
GOBP_POSITIVE_T_CELL_SELECTION, GOBP_POSITIVE_T_CELL_SELECTION
HP_COOMBS_POSITIVE_HEMOLYTIC_ANEMIA, HP_COOMBS_POSITIVE_HEMOLYTIC_ANEMIA
HP_FOLLICULITIS, HP_FOLLICULITIS
HP_PSYCHOMOTOR_DETERIORATION, HP_PSYCHOMOTOR_DETERIORATION
MIR5011_3P, MIR5011_3P
GOMF_K48_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_K48_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING
GOBP_REGULATION_OF_BROWN_FAT_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_BROWN_FAT_CELL_DIFFERENTIATION
HP_NON_MIDLINE_CLEFT_LIP, HP_NON_MIDLINE_CLEFT_LIP
GOBP_REGULATION_OF_BEHAVIOR, GOBP_REGULATION_OF_BEHAVIOR
WP_TYPE_II_DIABETES_MELLITUS, WP_TYPE_II_DIABETES_MELLITUS
HP_ABNORMAL_PROPORTION_OF_CD8_POSITIVE_T_CELLS, HP_ABNORMAL_PROPORTION_OF_CD8_POSITIVE_T_CELLS
GOBP_ELASTIC_FIBER_ASSEMBLY, GOBP_ELASTIC_FIBER_ASSEMBLY
REACTOME_SYNTHESIS_OF_VERY_LONG_CHAIN_FATTY_ACYL_COAS, REACTOME_SYNTHESIS_OF_VERY_LONG_CHAIN_FATTY_ACYL_COAS
GOMF_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY, GOMF_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY
GOMF_PROTEIN_KINASE_C_ACTIVITY, GOMF_PROTEIN_KINASE_C_ACTIVITY
GOBP_EPIBOLY, GOBP_EPIBOLY
HP_PERIPHERAL_VISUAL_FIELD_LOSS, HP_PERIPHERAL_VISUAL_FIELD_LOSS
GOBP_FOREBRAIN_NEURON_DEVELOPMENT, GOBP_FOREBRAIN_NEURON_DEVELOPMENT
HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR, HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR
HP_INCREASED_VERTEBRAL_HEIGHT, HP_INCREASED_VERTEBRAL_HEIGHT
GOBP_CELLULAR_RESPONSE_TO_IRON_ION, GOBP_CELLULAR_RESPONSE_TO_IRON_ION
HP_APLASIA_OF_METACARPAL_BONES, HP_APLASIA_OF_METACARPAL_BONES
FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_TO_336HR_SUSTAINED_UP, FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_TO_336HR_SUSTAINED_UP
GOBP_TORC2_SIGNALING, GOBP_TORC2_SIGNALING
LEE_AGING_CEREBELLUM_DN, LEE_AGING_CEREBELLUM_DN
GOBP_SPHINGOMYELIN_CATABOLIC_PROCESS, GOBP_SPHINGOMYELIN_CATABOLIC_PROCESS
GOBP_GLYCOSYLCERAMIDE_CATABOLIC_PROCESS, GOBP_GLYCOSYLCERAMIDE_CATABOLIC_PROCESS
GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT, GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT
HP_RIGHT_BUNDLE_BRANCH_BLOCK, HP_RIGHT_BUNDLE_BRANCH_BLOCK
HP_FIRST_DORSAL_INTEROSSEI_MUSCLE_WEAKNESS, HP_FIRST_DORSAL_INTEROSSEI_MUSCLE_WEAKNESS
KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_4NQO_IN_OLD, KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_4NQO_IN_OLD
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_OLFACTORY_MICROVILLAR_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_OLFACTORY_MICROVILLAR_CELLS
GOCC_CORVET_COMPLEX, GOCC_CORVET_COMPLEX
GOBP_NEGATIVE_REGULATION_OF_ACTIN_NUCLEATION, GOBP_NEGATIVE_REGULATION_OF_ACTIN_NUCLEATION
GOMF_LYS48_SPECIFIC_DEUBIQUITINASE_ACTIVITY, GOMF_LYS48_SPECIFIC_DEUBIQUITINASE_ACTIVITY
GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS
GOBP_NEURON_MATURATION, GOBP_NEURON_MATURATION
HP_HEMORRHAGE_OF_THE_EYE, HP_HEMORRHAGE_OF_THE_EYE
GOBP_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_PLATELET_AGGREGATION, GOBP_NEGATIVE_REGULATION_OF_PLATELET_AGGREGATION
GOBP_PROTEIN_K11_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K11_LINKED_DEUBIQUITINATION
MIR769_5P, MIR769_5P
BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_UP, BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_UP
GOBP_REGULATION_OF_PLATELET_AGGREGATION, GOBP_REGULATION_OF_PLATELET_AGGREGATION
GOBP_REGULATION_OF_NEUTROPHIL_DEGRANULATION, GOBP_REGULATION_OF_NEUTROPHIL_DEGRANULATION
GOCC_HOPS_COMPLEX, GOCC_HOPS_COMPLEX
RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_UP, RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_UP
MIR615_3P, MIR615_3P
GOCC_ESCRT_III_COMPLEX, GOCC_ESCRT_III_COMPLEX
WP_SOMATROPH_AXIS_GH_AND_ITS_RELATIONSHIP_TO_DIETARY_RESTRICTION_AND_AGING, WP_SOMATROPH_AXIS_GH_AND_ITS_RELATIONSHIP_TO_DIETARY_RESTRICTION_AND_AGING
HP_BUNDLE_BRANCH_BLOCK, HP_BUNDLE_BRANCH_BLOCK
HP_SECONDARY_AMENORRHEA, HP_SECONDARY_AMENORRHEA
GOBP_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_REGULATION_OF_FEEDING_BEHAVIOR
BIOCARTA_IL2_PATHWAY, BIOCARTA_IL2_PATHWAY
GOBP_LONG_CHAIN_FATTY_ACID_TRANSPORT, GOBP_LONG_CHAIN_FATTY_ACID_TRANSPORT
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_SUMOYLATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_SUMOYLATION
GOMF_ARACHIDONATE_COA_LIGASE_ACTIVITY, GOMF_ARACHIDONATE_COA_LIGASE_ACTIVITY
HP_CHYLOTHORAX, HP_CHYLOTHORAX
GOMF_PHOSPHATIDYL_PHOSPHOLIPASE_B_ACTIVITY, GOMF_PHOSPHATIDYL_PHOSPHOLIPASE_B_ACTIVITY
GOMF_LONG_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY, GOMF_LONG_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY
GOBP_CELLULAR_RESPONSE_TO_LEUCINE, GOBP_CELLULAR_RESPONSE_TO_LEUCINE
GOCC_SCAR_COMPLEX, GOCC_SCAR_COMPLEX
RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_SMAD4_UP, RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_SMAD4_UP
GOBP_CEREBELLAR_PURKINJE_CELL_DIFFERENTIATION, GOBP_CEREBELLAR_PURKINJE_CELL_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_ACTIVATION, GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_ACTIVATION
GOBP_REGULATION_OF_HISTONE_H3_K27_METHYLATION, GOBP_REGULATION_OF_HISTONE_H3_K27_METHYLATION
REACTOME_SMALL_INTERFERING_RNA_SIRNA_BIOGENESIS, REACTOME_SMALL_INTERFERING_RNA_SIRNA_BIOGENESIS
GOBP_REGULATION_OF_INORGANIC_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_REGULATION_OF_INORGANIC_ANION_TRANSMEMBRANE_TRANSPORT
HP_BILATERAL_CLEFT_LIP_AND_PALATE, HP_BILATERAL_CLEFT_LIP_AND_PALATE
HP_INCREASED_B_CELL_COUNT, HP_INCREASED_B_CELL_COUNT
GOBP_POSITIVE_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT
KOBAYASHI_EGFR_SIGNALING_6HR_UP, KOBAYASHI_EGFR_SIGNALING_6HR_UP
MIR6814_3P_MIR6872_5P, MIR6814_3P_MIR6872_5P
HP_CERVICAL_SPINAL_CANAL_STENOSIS, HP_CERVICAL_SPINAL_CANAL_STENOSIS
GOBP_REGULATION_OF_BONE_DEVELOPMENT, GOBP_REGULATION_OF_BONE_DEVELOPMENT
GTF3C1_TARGET_GENES, GTF3C1_TARGET_GENES
GOBP_GLUCOSYLCERAMIDE_METABOLIC_PROCESS, GOBP_GLUCOSYLCERAMIDE_METABOLIC_PROCESS
GOBP_CLEAVAGE_FURROW_FORMATION, GOBP_CLEAVAGE_FURROW_FORMATION
GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION
GOMF_RNA_POLYMERASE_III_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOMF_RNA_POLYMERASE_III_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BI
YAN_ESCAPE_FROM_ANOIKIS, YAN_ESCAPE_FROM_ANOIKIS
MORF_CTSB, MORF_CTSB
HP_RECURRENT_HYPOGLYCEMIA, HP_RECURRENT_HYPOGLYCEMIA
GOCC_TORC2_COMPLEX, GOCC_TORC2_COMPLEX
HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_INNER_EAR, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_INNER_EAR
GOMF_POLYNUCLEOTIDE_ADENYLYLTRANSFERASE_ACTIVITY, GOMF_POLYNUCLEOTIDE_ADENYLYLTRANSFERASE_ACTIVITY
GOBP_CD4_POSITIVE_OR_CD8_POSITIVE_ALPHA_BETA_T_CELL_LINEAGE_COMMITMENT, GOBP_CD4_POSITIVE_OR_CD8_POSITIVE_ALPHA_BETA_T_CELL_LINEAGE_COMMITMENT
HP_CONE_CONE_ROD_DYSTROPHY, HP_CONE_CONE_ROD_DYSTROPHY
GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS
BIOCARTA_ERK_PATHWAY, BIOCARTA_ERK_PATHWAY
GOBP_POSITIVE_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_POSITIVE_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM
HP_APLASIA_HYPOPLASIA_OF_THE_PROXIMAL_PHALANGES_OF_THE_HAND, HP_APLASIA_HYPOPLASIA_OF_THE_PROXIMAL_PHALANGES_OF_THE_HAND
GOBP_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM
IKEDA_MIR30_TARGETS_DN, IKEDA_MIR30_TARGETS_DN
WP_OMEGA9_FA_SYNTHESIS, WP_OMEGA9_FA_SYNTHESIS
MIR6514_5P, MIR6514_5P
GOBP_POSITIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS
HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_DN, HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_DN
HP_MULTIPLE_SUTURE_CRANIOSYNOSTOSIS, HP_MULTIPLE_SUTURE_CRANIOSYNOSTOSIS
GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING
GOBP_LOCOMOTOR_RHYTHM, GOBP_LOCOMOTOR_RHYTHM
REACTOME_MAPK1_ERK2_ACTIVATION, REACTOME_MAPK1_ERK2_ACTIVATION
GOBP_PEPTIDYL_SERINE_AUTOPHOSPHORYLATION, GOBP_PEPTIDYL_SERINE_AUTOPHOSPHORYLATION
GOBP_NATURAL_KILLER_CELL_DEGRANULATION, GOBP_NATURAL_KILLER_CELL_DEGRANULATION
SMCHD1_TARGET_GENES, SMCHD1_TARGET_GENES
GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_DOPAMINERGIC, GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_DOPAMINERGIC
GOBP_THIAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_THIAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
KREPPEL_CD99_TARGETS_DN, KREPPEL_CD99_TARGETS_DN
DESCARTES_MAIN_FETAL_OLIGODENDROCYTES, DESCARTES_MAIN_FETAL_OLIGODENDROCYTES
GOBP_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS, GOBP_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS
GOBP_ICOSANOID_TRANSPORT, GOBP_ICOSANOID_TRANSPORT
REACTOME_ACTIVATION_OF_RAC1, REACTOME_ACTIVATION_OF_RAC1
HP_FOLLICULAR_HYPERPLASIA, HP_FOLLICULAR_HYPERPLASIA
HP_ELEVATED_PROPORTION_OF_CD4_NEGATIVE_CD8_NEGATIVE_ALPHA_BETA_REGULATORY_T_CELLS, HP_ELEVATED_PROPORTION_OF_CD4_NEGATIVE_CD8_NEGATIVE_ALPHA_BETA_REGULATORY_T_CELLS
FOURATI_BLOOD_TWINRIX_AGE_65_83YO_POOR_RESPONDERS_VS_RESPONDERS_0DY_NETWORK_INFERENCE_UP, FOURATI_BLOOD_TWINRIX_AGE_65_83YO_POOR_RESPONDERS_VS_RESPONDERS_0DY_NETWORK_
GOBP_RESPONSE_TO_MOLECULE_OF_FUNGAL_ORIGIN, GOBP_RESPONSE_TO_MOLECULE_OF_FUNGAL_ORIGIN
LIU_IL13_MEMORY_MODEL_DN, LIU_IL13_MEMORY_MODEL_DN
HP_PAROXYSMAL_SUPRAVENTRICULAR_TACHYCARDIA, HP_PAROXYSMAL_SUPRAVENTRICULAR_TACHYCARDIA
HP_DECREASED_SERUM_COMPLEMENT_C4, HP_DECREASED_SERUM_COMPLEMENT_C4
GOBP_POSITIVE_REGULATION_OF_HISTONE_H3_K27_METHYLATION, GOBP_POSITIVE_REGULATION_OF_HISTONE_H3_K27_METHYLATION
GOBP_POSITIVE_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION, GOBP_POSITIVE_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION
GOMF_ADP_RIBOSE_DIPHOSPHATASE_ACTIVITY, GOMF_ADP_RIBOSE_DIPHOSPHATASE_ACTIVITY
MIR3619_3P, MIR3619_3P
HP_ABNORMAL_SARCOMERE_MORPHOLOGY, HP_ABNORMAL_SARCOMERE_MORPHOLOGY
GOBP_FOLIC_ACID_TRANSPORT, GOBP_FOLIC_ACID_TRANSPORT
HP_SCHIZENCEPHALY, HP_SCHIZENCEPHALY
HP_PULMONARY_VENOUS_HYPERTENSION, HP_PULMONARY_VENOUS_HYPERTENSION
GOBP_GLYCERALDEHYDE_3_PHOSPHATE_METABOLIC_PROCESS, GOBP_GLYCERALDEHYDE_3_PHOSPHATE_METABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_GLIAL_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_GLIAL_CELL_APOPTOTIC_PROCESS
HP_HEMERALOPIA, HP_HEMERALOPIA
REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_AND_CELL_SPREADING_BY_IPP_COMPLEX_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_AND_CELL_SPREADING_BY_
WP_NEUROINFLAMMATION, WP_NEUROINFLAMMATION
GOBP_ICOSANOID_BIOSYNTHETIC_PROCESS, GOBP_ICOSANOID_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_HISTONE_DEACETYLATION, GOBP_REGULATION_OF_HISTONE_DEACETYLATION
GOBP_ICOSANOID_SECRETION, GOBP_ICOSANOID_SECRETION
GOBP_T_CELL_LINEAGE_COMMITMENT, GOBP_T_CELL_LINEAGE_COMMITMENT
WP_TYPE_III_INTERFERON_SIGNALING, WP_TYPE_III_INTERFERON_SIGNALING
HP_BROAD_CLAVICLES, HP_BROAD_CLAVICLES
GOBP_REGULATION_OF_POSTSYNAPTIC_NEUROTRANSMITTER_RECEPTOR_INTERNALIZATION, GOBP_REGULATION_OF_POSTSYNAPTIC_NEUROTRANSMITTER_RECEPTOR_INTERNALIZATION
HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS, HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS
GOBP_REGULATION_OF_GLIAL_CELL_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_GLIAL_CELL_APOPTOTIC_PROCESS
HP_HYPERCONVEX_FINGERNAILS, HP_HYPERCONVEX_FINGERNAILS
GOBP_NEGATIVE_REGULATION_OF_CELL_SIZE, GOBP_NEGATIVE_REGULATION_OF_CELL_SIZE
TURJANSKI_MAPK14_TARGETS, TURJANSKI_MAPK14_TARGETS
GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_MORPHOGENESIS, GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_MORPHOGENESIS
MIR6082, MIR6082
MIR377_5P, MIR377_5P
GOBP_POSITIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION
HP_ABNORMAL_PERIFOLLICULAR_MORPHOLOGY, HP_ABNORMAL_PERIFOLLICULAR_MORPHOLOGY
LIU_CMYB_TARGETS_DN, LIU_CMYB_TARGETS_DN
MIR933, MIR933
HP_CHORIORETINAL_DYSTROPHY, HP_CHORIORETINAL_DYSTROPHY
GOBP_MITOCHONDRIAL_RIBOSOME_ASSEMBLY, GOBP_MITOCHONDRIAL_RIBOSOME_ASSEMBLY
GOBP_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION, GOBP_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION
SOBOLEV_PBMC_PANDEMRIX_AGE_18_64YO_RESPONDERS_VS_NONRESPONDERS_7DY_UP, SOBOLEV_PBMC_PANDEMRIX_AGE_18_64YO_RESPONDERS_VS_NONRESPONDERS_7DY_UP
GOBP_ACTIVATION_OF_PROTEIN_KINASE_B_ACTIVITY, GOBP_ACTIVATION_OF_PROTEIN_KINASE_B_ACTIVITY
GOBP_REGULATION_OF_KERATINOCYTE_MIGRATION, GOBP_REGULATION_OF_KERATINOCYTE_MIGRATION
GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEACETYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEACETYLATION
GOBP_IODIDE_TRANSPORT, GOBP_IODIDE_TRANSPORT
LAU_APOPTOSIS_CDKN2A_DN, LAU_APOPTOSIS_CDKN2A_DN
HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION, HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION
HP_ABNORMAL_PROPORTION_OF_CLASS_SWITCHED_MEMORY_B_CELLS, HP_ABNORMAL_PROPORTION_OF_CLASS_SWITCHED_MEMORY_B_CELLS
HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II, HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II
GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY
GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_HYDROGEN_PEROXIDE, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_HYDROGEN_PEROXIDE
GOBP_RNA_3_URIDYLATION, GOBP_RNA_3_URIDYLATION
GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_NEURON_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_NEURON_INTRINS
GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATIONAL_INITIATION, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATIONAL_INITIATION
GOCC_NATA_COMPLEX, GOCC_NATA_COMPLEX
GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H4_K20_SPECIFIC, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H4_K20_SPECIFIC
GOMF_CYSTEINE_TYPE_EXOPEPTIDASE_ACTIVITY, GOMF_CYSTEINE_TYPE_EXOPEPTIDASE_ACTIVITY
DPPA3_TARGET_GENES, DPPA3_TARGET_GENES
GOMF_ANGIOTENSIN_RECEPTOR_BINDING, GOMF_ANGIOTENSIN_RECEPTOR_BINDING
WP_CALORIC_RESTRICTION_AND_AGING, WP_CALORIC_RESTRICTION_AND_AGING
GOMF_P_TYPE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_P_TYPE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
BIOCARTA_S1P_PATHWAY, BIOCARTA_S1P_PATHWAY
GOCC_CELL_TRAILING_EDGE, GOCC_CELL_TRAILING_EDGE
GOBP_MUSCLE_ATROPHY, GOBP_MUSCLE_ATROPHY
WP_SRF_AND_MIRS_IN_SMOOTH_MUSCLE_DIFFERENTIATION_AND_PROLIFERATION, WP_SRF_AND_MIRS_IN_SMOOTH_MUSCLE_DIFFERENTIATION_AND_PROLIFERATION
GOBP_POSITIVE_REGULATION_OF_MEMBRANE_POTENTIAL, GOBP_POSITIVE_REGULATION_OF_MEMBRANE_POTENTIAL
GOBP_MAGNESIUM_ION_TRANSPORT, GOBP_MAGNESIUM_ION_TRANSPORT
GOBP_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_KILLING_OF_CELLS_OF_OTHER_ORGANISM
HP_RECURRENT_CORNEAL_EROSIONS, HP_RECURRENT_CORNEAL_EROSIONS
HP_INTELLECTUAL_DISABILITY_BORDERLINE, HP_INTELLECTUAL_DISABILITY_BORDERLINE