

GCM_MYST2, GCM_MYST2

GSE13484_UNSTM_VS_3H_YF17D_VACCINE_STIM_PBMIC_UP, GSE13484_UNSTM_VS_3H_YF17D_VACCINE_STIM_PBMIC_UP

GSE36026_NORMA1_VS_STAPH_AUREUS_INF_IL1R_KO_SKIN_DN, GSE36026_NORMA1_VS_STAPH_AUREUS_INF_IL1R_KO_SKIN_DN

GSE2706_UNSTM_VS_8H_LPS_AND_R848_DC_UP, GSE2706_UNSTM_VS_8H_LPS_AND_R848_DC_UP

GSE14000_UNSTM_VS_16H_LPS_DC_UP, GSE14000_UNSTM_VS_16H_LPS_DC_UP

GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_DN, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_DN

GSE2409_PDI_LIGATION_VS_CTRL_IN_ACT_TCELL_LINE_DN, GSE2409_PDI_LIGATION_VS_CTRL_IN_ACT_TCELL_LINE_DN

GSE9960_HEALTHY_VS_GRAM_NEG_SEIPS_PBMIC_UP, GSE9960_HEALTHY_VS_GRAM_NEG_SEIPS_PBMIC_UP

GSE13484_12H_UNSTM_VS_YF17D_VACCINE_STIM_PBMIC_UP, GSE13484_12H_UNSTM_VS_YF17D_VACCINE_STIM_PBMIC_UP

GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP, GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP

GCM_LBENZ, GCM_LBENZ

GSE13484_UNSTM_VS_12H_YF17D_VACCINE_STIM_PBMIC_UP, GSE13484_UNSTM_VS_12H_YF17D_VACCINE_STIM_PBMIC_UP

GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_2H_DN, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_2H_DN

GSE16755_CTRL_VS_IFNA_TREATED_MAC_UP, GSE16755_CTRL_VS_IFNA_TREATED_MAC_UP

GSE7598_NAIVE_VS_DAY7_PCC_IMMUNIZATION_CD4_TCELL_UP, GSE7598_NAIVE_VS_DAY7_PCC_IMMUNIZATION_CD4_TCELL_UP

GSE1035_ICELL_VS_MYELOID_DN, GSE1035_ICELL_VS_MYELOID_DN

GSE13484_3H_UNSTM_VS_YF17D_VACCINE_STIM_PBMIC_UP, GSE13484_3H_UNSTM_VS_YF17D_VACCINE_STIM_PBMIC_UP

GSE36476_CTRL_VS_T8ST_ACT_40H_MEMORY_CD4_TCELL_YOUNG_UP, GSE36476_CTRL_VS_T8ST_ACT_40H_MEMORY_CD4_TCELL_YOUNG_UP

GSE360_10W_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_DC_UP, GSE360_10W_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_DC_UP

GSE2706_UNSTM_VS_8H_LPS_DC_UP, GSE2706_UNSTM_VS_8H_LPS_DC_UP

GSE14000_4H_VS_16H_LPS_DC_UP, GSE14000_4H_VS_16H_LPS_DC_UP

GSE19300_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_UP, GSE19300_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_UP

GSE37533_PPARG2_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_UP, GSE37533_PPARG2_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_UP

GSE42021_TREG_VS_TCONV_PLN_DN, GSE42021_TREG_VS_TCONV_PLN_DN

GSE1432_1H_VS_6H_JFNG_MICROGLIA_UP, GSE1432_1H_VS_6H_JFNG_MICROGLIA_UP

GSE42021_CD24H_VS_CD24INT_TREG_THYMUS_UP, GSE42021_CD24H_VS_CD24INT_TREG_THYMUS_UP

GSE10325_CD4_TCELL_VS_MYELOID_UP, GSE10325_CD4_TCELL_VS_MYELOID_UP

GCM_ZNF198, GCM_ZNF198

GSE19300_STIM_VS_STIM_AND_IFNAB_24H_CD8_T_CELL_UP, GSE19300_STIM_VS_STIM_AND_IFNAB_24H_CD8_T_CELL_UP

GSE18281_SUBCAPSULAR_VS_PERIMEDULLARY_CORTICAL_REGION_OF_THYMUS_UP, GSE18281_SUBCAPSULAR_VS_PERIMEDULLARY_CORTICAL_REGION_OF_THYMUS_UP

GSE36476_CTRL_VS_T8ST_ACT_16H_MEMORY_CD4_TCELL_OLD_UP, GSE36476_CTRL_VS_T8ST_ACT_16H_MEMORY_CD4_TCELL_OLD_UP

GSE2903_PFE_VS_DAY7_ILU_VACCINE_MDC_DN, GSE2903_PFE_VS_DAY7_ILU_VACCINE_MDC_DN

GSE19401_PLN_VS_PFEYERS_PATCH_FOLLICULAR_DC_UP, GSE19401_PLN_VS_PFEYERS_PATCH_FOLLICULAR_DC_UP

GSE27899_CD11C_INT_F480_HI_MACROPHAGE_VS_CD11C_ING_F480_INT_DC_DN, GSE27899_CD11C_INT_F480_HI_MACROPHAGE_VS_CD11C_ING_F480_INT_DC_DN

GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_DC_DN, GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_DC_DN

GSE3962_MAST_CELL_VS_TH2_UP, GSE3962_MAST_CELL_VS_TH2_UP

GSE33701_MULTIPOTENT_PROGENTOR_VS_RAG2_KO_NK_CELL_UP, GSE33701_MULTIPOTENT_PROGENTOR_VS_RAG2_KO_NK_CELL_UP

GSE3982_CTRL_VS_LPS_IH_NEUTROPHIL_UP, GSE3982_CTRL_VS_LPS_IH_NEUTROPHIL_UP

GSE360_1_DONOVANI_VS_M_TUBERCULOSIS_DC_UP, GSE360_1_DONOVANI_VS_M_TUBERCULOSIS_DC_UP

GCM_SUFU, GCM_SUFU

MIR5138_3P, MIR5138_3P

GSE42021_TREG_PLN_VS_CD24LO_TREG_THYMUS_UP, GSE42021_TREG_PLN_VS_CD24LO_TREG_THYMUS_UP

GSE360_CTRL_VS_T_GONDIL_CD_UP, GSE360_CTRL_VS_T_GONDIL_CD_UP

GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_1CMV_ARMSTRONG_DN, GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_1CMV_ARMSTRONG_DN

MORF_RV171, MORF_RV171

GNF2_CD53, GNF2_CD53

ZSCAN5C_TARGET_GENES, ZSCAN5C_TARGET_GENES

GSE27786_ICELL_VS_CD8_TCELL_DN, GSE27786_ICELL_VS_CD8_TCELL_DN

MIR5867, MIR5867

PATEL_SKIN_OF_BODY_ZOSTAVAX_AGE_70_93YO_VZV_CHALLENGE_3DY_UP, PATEL_SKIN_OF_BODY_ZOSTAVAX_AGE_70_93YO_VZV_CHALLENGE_3DY_UP

GSE1448_CTRL_VS_ANT1_VALPHA2_DP_THYMOCYTE_DN, GSE1448_CTRL_VS_ANT1_VALPHA2_DP_THYMOCYTE_DN

MIR3692_5P, MIR3692_5P

MIR645, MIR645

IL_RESPONSE_TO_FSH_DN, IL_RESPONSE_TO_FSH_DN

GOBP_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY, GOBP_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY

GSE19300_STIM_VS_STIM_AND_IFNAB_72H_CD8_T_CELL_UP, GSE19300_STIM_VS_STIM_AND_IFNAB_72H_CD8_T_CELL_UP

MODULE_277, MODULE_277

MIR6757_3P, MIR6757_3P

MIR5583_5P, MIR5583_5P

CHOL_ATL_STAGE_PREDICTOR, CHOL_ATL_STAGE_PREDICTOR

GSE42021_CD24INT_VS_CD24LOW_TREG_THYMUS_UP, GSE42021_CD24INT_VS_CD24LOW_TREG_THYMUS_UP

MODULE_243, MODULE_243

MIR3129_3P, MIR3129_3P

GNF2_TYK2, GNF2_TYK2

MIR935, MIR935

HP_ABNORMAL_LARYNX_MORPHOLOGY, HP_ABNORMAL_LARYNX_MORPHOLOGY

BUSSLINGER_GASTRIC_X_CELLS, BUSSLINGER_GASTRIC_X_CELLS

MIR1224_3P, MIR1224_3P

ROSS_AML_WITH_CBF, MYH11_FUSION, ROSS_AML_WITH_CBF, MYH11_FUSION

CTCAAG, MIR43, CTCAAG, MIR43

HP_CONFUSION, HP_CONFUSION

GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_CELL, GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_CELL

GSE19300_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_UP, GSE19300_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_UP

HP_BRBK_REFLEXES, HP_BRBK_REFLEXES

MIR7843_5P, MIR7843_5P

PARK_HSC_MARKERS, PARK_HSC_MARKERS

GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_6H_IFNA_STIM_UP, GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_6H_IFNA_STIM_UP

GCM_CSNK1D, GCM_CSNK1D

HP_BONE_MARROW_HYPOCELLULARITY, HP_BONE_MARROW_HYPOCELLULARITY

GOBP_HISTONE_H3_ACETYLATION, GOBP_HISTONE_H3_ACETYLATION

WP_WNT_SIGNALING_PATHWAY_AND_PLURIPOTENCY, WP_WNT_SIGNALING_PATHWAY_AND_PLURIPOTENCY

GOCC_DNA_REPAIR_COMPLEX, GOCC_DNA_REPAIR_COMPLEX

GOMF_MIRNA_BINDING, GOMF_MIRNA_BINDING

GCM_RAP2A, GCM_RAP2A

BOYAULT_LIVER_CANCER_SUBCLASS_G12_UP, BOYAULT_LIVER_CANCER_SUBCLASS_G12_UP

GOBP_PEPTIDE_N_ACETYLTRANSFERASE_ACTIVITY, GOMF_PEPTIDE_N_ACETYLTRANSFERASE_ACTIVITY

HP_KIRAT1B, HP_KIRAT1B

GOMF_PEPTIDE_LYSINE_N_ACETYLTRANSFERASE_ACTIVITY, GOMF_PEPTIDE_LYSINE_N_ACETYLTRANSFERASE_ACTIVITY

GOBP_PEPTIDYL_TYROSINE_DEPHOSPHORYLATION, GOBP_PEPTIDYL_TYROSINE_DEPHOSPHORYLATION

HP_MUCOSAL_TELANGIECTASIAE, HP_MUCOSAL_TELANGIECTASIAE

MIR3116, MIR3116

MIR431_5P, MIR431_5P

BCAT_BILD_ET_AL_DN, BCAT_BILD_ET_AL_DN

GOBP_PEPTIDYL_CYSINE_MODIFICATION, GOBP_PEPTIDYL_CYSINE_MODIFICATION

HP_PREMATURE_OVARIAN_INSUFFICIENCY, HP_PREMATURE_OVARIAN_INSUFFICIENCY

HP_CRANIAL_HYPEROSTOSIS, HP_CRANIAL_HYPEROSTOSIS

MIR654_5P, MIR654_5P

HP_BLEPHARITIS, HP_BLEPHARITIS

GNF2_RAD23A, GNF2_RAD23A

MIR541_3P, MIR541_3P

MIR6881_5P, MIR6881_5P

AIZARANI_LIVER_C12_NK_NKT_CELLS_4, AIZARANI_LIVER_C12_NK_NKT_CELLS_4

ASTON_MAJOR_DEPRESSIVE_DISORDER_UP, ASTON_MAJOR_DEPRESSIVE_DISORDER_UP

HP_FRECKLING, HP_FRECKLING

MIR937_5P, MIR937_5P

MIR8055, MIR8055

MODULE_281, MODULE_281

GCM_FANCL, GCM_FANCL

GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_DN, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_DN

GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_2_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_2_PRODUCTION

GNF2_SPTA1, GNF2_SPTA1

MIR477_3P, MIR477_3P

GOBP_ACTIVATION_OF_MAPKK_ACTIVITY, GOBP_ACTIVATION_OF_MAPKK_ACTIVITY

SIG_REGULATION_OF_THE_ACTIN_CYTOSKELETON_BY_RHO_GTPASES, SIG_REGULATION_OF_THE_ACTIN_CYTOSKELETON_BY_RHO_GTPASES

MIR6866_5P, MIR6866_5P

WP_RAC1PAK1P38MAP2_PATHWAY, WP_RAC1PAK1P38MAP2_PATHWAY

MIR676_3P, MIR676_3P

GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ORGANIZATION, GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ORGANIZATION

HP_MIXED_HEARING_IMPAIRMENT, HP_MIXED_HEARING_IMPAIRMENT

GOCC_DNA_REPLICATION_FACTOR_A_COMPLEX, GOCC_DNA_REPLICATION_FACTOR_A_COMPLEX

GOBP_HISTONE_H4_K16_ACETYLATION, GOBP_HISTONE_H4_K16_ACETYLATION

PID_INSULIN_PATHWAY, PID_INSULIN_PATHWAY

PID_P38_MKK3_6PATHWAY, PID_P38_MKK3_6PATHWAY

HP_PORILODERMA, HP_PORILODERMA

MODULE_284, MODULE_284

HP_ABNORMAL_PLATELET_VOLUME, HP_ABNORMAL_PLATELET_VOLUME

HP_ENTROPION, HP_ENTROPION

MIR634_3P, MIR634_3P

GOMF_PROTEIN_DISULFIDE_OXIDOREDUCTASE_ACTIVITY, GOMF_PROTEIN_DISULFIDE_OXIDOREDUCTASE_ACTIVITY

MIR450B_3P, MIR450B_3P

GGTAACC, MIR4099P, GGTAACC, MIR4099P

GENTILE_UV_RESPONSE_CLUSTER_D8, GENTILE_UV_RESPONSE_CLUSTER_D8

BROCAARTA_WNT_PATHWAY, BROCAARTA_WNT_PATHWAY

GOMF_PHOSPHATIDIC_ACID_BINDING, GOMF_PHOSPHATIDIC_ACID_BINDING

REACTOME_PEPTIDE_HORMONE_METABOLISM, REACTOME_PEPTIDE_HORMONE_METABOLISM

HP_PTERYGLUM, HP_PTERYGLUM

GNF2_ATM, GNF2_ATM

MODULE_300, MODULE_300

GOBP_HISTONE_H3_K9_METHYLATION, GOBP_HISTONE_H3_K9_METHYLATION

GOMF_METALLOAMINOPEPTIDASE_ACTIVITY, GOMF_METALLOAMINOPEPTIDASE_ACTIVITY

MIR799_3P, MIR799_3P

HP_ABNORMAL_MEGAKARYOCYTE_MORPHOLOGY, HP_ABNORMAL_MEGAKARYOCYTE_MORPHOLOGY

GTACAGG, MIR486, GTACAGG, MIR486

GOBP_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION, GOBP_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION

WP_TRANSCRIPTION_FACTOR_REGULATION_IN_ADIPONECISIS, WP_TRANSCRIPTION_FACTOR_REGULATION_IN_ADIPONECISIS

MCCLUNG_DELTA_FOSB_TARGETS_8WK, MCCLUNG_DELTA_FOSB_TARGETS_8WK

GOBP_RETINOIC_ACID_RECEPTOR_SIGNALING_PATHWAY, GOBP_RETINOIC_ACID_RECEPTOR_SIGNALING_PATHWAY

QI_PBMIC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_T_CELL_RESPONSES_EXPANSION_AND_CONTRACTION_ID_NOT_INFORMATIVE_OF_LONG_TERM_RESPONSES_POSITIVE, QI_PBMIC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_T_CELL_RESPONSES_EXPANSION_AND_CONTRACTION_ID_NOT_INFORMATIVE_OF_LONG_TERM_RESPONSES_POSITIVE

HP_LONG_NOSE, HP_LONG_NOSE

MODULE_166, MODULE_166

MIR769_3P, MIR769_3P

GOCC_NUCLEOTIDE_EXCISION_REPAIR_COMPLEX, GOCC_NUCLEOTIDE_EXCISION_REPAIR_COMPLEX

BROCAARTA_ETC_PATHWAY, BROCAARTA_ETC_PATHWAY

GOBP_POSITIVE_REGULATION_OF_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY

PID_P38_ALPHA_BETA_PATHWAY, PID_P38_ALPHA_BETA_PATHWAY

FLECHNER_BIOPSY_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN, FLECHNER_BIOPSY_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN

GOCC_SEPTIN_COMPLEX, GOCC_SEPTIN_COMPLEX

GOBP_SNO_5_RNA_METABOLIC_PROCESS, GOBP_SNO_5_RNA_METABOLIC_PROCESS

MIR5586_3P, MIR5586_3P

MIR3680_5P, MIR3680_5P

GOBP_WNT_SIGNALING_PATHWAY_CALCULUM_MODULATING_PATHWAY, GOBP_WNT_SIGNALING_PATHWAY_CALCULUM_MODULATING_PATHWAY

GOBP_PRL_MIRNA_PROCESSING, GOBP_PRL_MIRNA_PROCESSING

GOMF_RIBOSOMAL_LARGE_SUBUNIT_BINDING, GOMF_RIBOSOMAL_LARGE_SUBUNIT_BINDING

GOBP_HISTONE_H3_K9_DEMETHYLATION, GOBP_HISTONE_H3_K9_DEMETHYLATION

GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K9_SPECIFIC, GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K9_SPECIFIC

REACTOME_STING_MEDIATED_INDUCION_OF_HOST_IMMUNE_RESPONSES, REACTOME_STING_MEDIATED_INDUCION_OF_HOST_IMMUNE_RESPONSES

MODULE_362, MODULE_362

REACTOME_TRANSCRIPTIONAL_REGULATION_BY_MECTP2, REACTOME_TRANSCRIPTIONAL_REGULATION_BY_MECTP2

GOBP_INCLUSION_BODY_ASSEMBLY, GOBP_INCLUSION_BODY_ASSEMBLY

HP_DIFFUSE_WHITE_MATTER_ABNORMALITIES, HP_DIFFUSE_WHITE_MATTER_ABNORMALITIES

MIR702, MIR702

HP_ULNAR_DEVIATION_OF_THE_HAND, HP_ULNAR_DEVIATION_OF_THE_HAND

GOBP_PEPTIDYL_LYSINE_DIMETHYLATION, GOBP_PEPTIDYL_LYSINE_DIMETHYLATION

GOBP_RESPONSE_TO_LITHIUM_ION, GOBP_RESPONSE_TO_LITHIUM_ION

GOCC_H3_HISTONE_ACETYLTRANSFERASE_COMPLEX, GOCC_H3_HISTONE_ACETYLTRANSFERASE_COMPLEX

GOBP_PROTEIN_PALMITOYLATION, GOBP_PROTEIN_PALMITOYLATION

MIR375_3P, MIR375_3P

MODULE_395, MODULE_395

CACGTTT, MIR302A, CACGTTT, MIR302A

MANN_RESPONSE_TO_AMFPOSTINE_UP, MANN_RESPONSE_TO_AMFPOSTINE_UP

WP_PDKAKTMTOR_VITD3_SIGNALLING, WP_PDKAKTMTOR_VITD3_SIGNALLING

GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY

GOBP_POSITIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY, GOBP_POSITIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY

HP_GIANT_PLATELETS, HP_GIANT_PLATELETS

GOMF_PYRIMIDINE_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PYRIMIDINE_NUCLEOTIDE_SUGAR_TRANSMEMBRANE_TRANSPORTER_ACTIVITY

MODULE_399, MODULE_399

STAMBOLSKY_RESPONSE_TO_VITAMIN_D3_UP, STAMBOLSKY_RESPONSE_TO_VITAMIN_D3_UP

REACTOME_BMAL1_CLOCK_NPAS2_ACTIVATES_CIRCADIAN_GENE_EXPRESSION, REACTOME_BMAL1_CLOCK_NPAS2_ACTIVATES_CIRCADIAN_GENE_EXPRESSION

REACTOME_IRF3_MEDIATED_INDUCION_OF_TYPE_1_IFN, REACTOME_IRF3_MEDIATED_INDUCION_OF_TYPE_1_IFN

MIR9903, MIR9903

MORF_MYC, MORF_MYC

HP_SYSTEMIC_LUPUS_ERYTHEMATOSUS, HP_SYSTEMIC_LUPUS_ERYTHEMATOSUS

GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION

WILLIAMS_FSR2_TARGETS_UP, WILLIAMS_FSR2_TARGETS_UP

REACTOME_TRANSPORT_OF_NUCLEOTIDE_SUGARS, REACTOME_TRANSPORT_OF_NUCLEOTIDE_SUGARS

GOBP_PEPTIDYL_L_CYSINE_S_PALMITOYLATION, GOBP_PEPTIDYL_L_CYSINE_S_PALMITOYLATION

GOMF_QUATERNARY_AMMONIUM_GROUP_BINDING, GOMF_QUATERNARY_AMMONIUM_GROUP_BINDING

HP_APLASIA_HYPOPLASIA_INVOLVING_THE_SKELETAL_MUSCULATURE, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_SKELETAL_MUSCULATURE

GOBP_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GOBP_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY

GOBP_REGULATION_OF_RETINOIC_ACID_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_RETINOIC_ACID_RECEPTOR_SIGNALING_PATHWAY

HP_VARICOSE_VEINS, HP_VARICOSE_VEINS

HP_ABNORMALITY_OF_LYSOSOMAL_METABOLISM, HP_ABNORMALITY_OF_LYSOSOMAL_METABOLISM

HP_FLAT_NASAL_ALAE, HP_FLAT_NASAL_ALAE

GOBP_NEGATIVE_REGULATION_OF_HISTONE_METHYLATION, GOBP_NEGATIVE_REGULATION_OF_HISTONE_METHYLATION

HP_ANKYLOBLEPHARON, HP_ANKYLOBLEPHARON

HP_ABNORMAL_PLATELET_MEMBRANE_PROTEIN_EXPRESSION, HP_ABNORMAL_PLATELET_MEMBRANE_PROTEIN_EXPRESSION

GOBP_POSITIVE_REGULATION_OF_NMDA_GLUTAMATE_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_NMDA_GLUTAMATE_RECEPTOR_ACTIVITY

GOBP_REGULATION_OF_HISTONE_H4_K16_ACETYLATION, GOBP_REGULATION_OF_HISTONE_H4_K16_ACETYLATION

GOMF_U2_SNRNA_BINDING, GOMF_U2_SNRNA_BINDING

GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMATIN, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMATIN

GOMF_RIBBLE_DNA_BINDING, GOMF_RIBBLE_DNA_BINDING

HP_SCLEROSIS_OF_SKULL_BASE, HP_SCLEROSIS_OF_SKULL_BASE

HP_PAIN_IN_HEAD_AND_NECK_REGION, HP_PAIN_IN_HEAD_AND_NECK_REGION

YORDY_RECIPROCAL_REGULATION_BY_ETSI1_AND_SPI100_DN, YORDY_RECIPROCAL_REGULATION_BY_ETSI1_AND_SPI100_DN

HP_SUBCORTICAL_HETEROTOPIA, HP_SUBCORTICAL_HETEROTOPIA

GOMF_PHOSPHATIDYLETHANOLAMINE_BINDING, GOMF_PHOSPHATIDYLETHANOLAMINE_BINDING

GOBP_REGULATION_OF_INCLUSION_BODY_ASSEMBLY, GOBP_REGULATION_OF_INCLUSION_BODY_ASSEMBLY

HP_VENOUS_INSUFFICIENCY, HP_VENOUS_INSUFFICIENCY

PRDM5_TARGET_GENES, PRDM5_TARGET_GENES

GOBP_REGULATION_OF_HISTONE_H3_K9_ACETYLATION, GOBP_REGULATION_OF_HISTONE_H3_K9_ACETYLATION

GOBP_SPERMATID_NUCLEUS_DIFFERENTIATION, GOBP_SPERMATID_NUCLEUS_DIFFERENTIATION

GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION

HP_THYROID_AGENESIS, HP_THYROID_AGENESIS

REACTOME_NRH2_NRH3_REGULATE_GENE_EXPRESSION_TO_LIMIT_CHOLESTEROL_UPTAKE, REACTOME_NRH2_NRH3_REGULATE_GENE_EXPRESSION_TO_LIMIT_CHOLESTEROL_UPTAKE

GOBP_WHITE_FAT_CELL_DIFFERENTIATION, GOBP_WHITE_FAT_CELL_DIFFERENTIATION

MIR548A0_3P, MIR548A0_3P

GOBP_REGULATION_OF_HISTONE_H3_K9_METHYLATION, GOBP_REGULATION_OF_HISTONE_H3_K9_METHYLATION

WP_EBV_LMP1_SIGNALING, WP_EBV_LMP1_SIGNALING

GOBP_RNA_DECAPPING, GOBP_RNA_DECAPPING

HP_THYROID_DYSGENESIS, HP_THYROID_DYSGENESIS

GOBP_NEGATIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_STRESS, GOBP_NEGATIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_STRESS

HP_ABNORMAL_CRY, HP_ABNORMAL_CRY

GNF2_PCAF, GNF2_PCAF

WIKMAN_ASBESTOS_LUNG_CANCER_UP, WIKMAN_ASBESTOS_LUNG_CANCER_UP

GOCC_INTRINSIC_COMPONENT_OF_NUCLEAR_INNER_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_NUCLEAR_INNER_MEMBRANE

DAVIES_MULTIPLE_MYELOMA_VS_MGUS_UP, DAVIES_MULTIPLE_MYELOMA_VS_MGUS_UP

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_7DY_TOP_100_DEG_EX_VIVO_UP

GOCC_SEMAPHORIN_RECEPTOR_COMPLEX, GOCC_SEMAPHORIN_RECEPTOR_COMPLEX

MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_VACCINATED_VS_CANDIN_PLACEBO_BCG_PRIMED_1DY_DN, MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_VACCINATED_VS_CANDIN_PLACEBO_BCG_PRIMED_1DY_DN

GOBP_NEGATIVE_REGULATION_OF_INCLUSION_BODY_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_INCLUSION_BODY_ASSEMBLY

GOBP_MICROTUBULE_ANCHORING_AT_CENTROSOME, GOBP_MICROTUBULE_ANCHORING_AT_CENTROSOME

REACTOME_RORA_ACTIVATES_GENE_EXPRESSION, REACTOME_RORA_ACTIVATES_GENE_EXPRESSION

GOBP_ENDOPLASMIC_RETICULUM_LOCALIZATION, GOBP_ENDOPLASMIC_RETICULUM_LOCALIZATION

HP_AORTIC_ATHEROSCLEROTIC_LESION, HP_AORTIC_ATHEROSCLEROTIC_LESION

MIR371A_3P, MIR371A_3P

HP_FOAM_CELLS, HP_FOAM_CELLS

HP_APLASIA_HYPOPLASIA_OF_THE_5TH_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_5TH_METACARPAL

HP_VISUAL_FIXATION_INSTABILITY, HP_VISUAL_FIXATION_INSTABILITY

GOBP_URIDYLYLTRANSFERASE_ACTIVITY, GOMF_URIDYLYLTRANSFERASE_ACTIVITY

MEISSNER_NPC_ICP_WITH_H3K4ME3, MEISSNER_NPC_ICP_WITH_H3K4ME3

GOBP_POSITIVE_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY

GOBP_CAVEOLIN_MEDIATED_ENDOCYTOSIS, GOBP_CAVEOLIN_MEDIATED_ENDOCYTOSIS

GSE12707_AT1H1_HYPOMORPH_VS_WT_THYMUS_DN, GSE12707_AT1H1_HYPOMORPH_VS_WT_THYMUS_DN

HP_MACULAR_THICKENING, HP_MACULAR_THICKENING

GOCC_EXOCYST, GOCC_EXOCYST

MIR227_3P, MIR227_3P

MEISSNER_ES_ICP_WITH_H3K4ME3, MEISSNER_ES_ICP_WITH_H3K4ME3

REACTOME_AQUAPORIN_MEDIATED_TRANSPORT, REACTOME_AQUAPORIN_MEDIATED_TRANSPORT

GOBP_REGULATION_OF_LYASE_ACTIVITY, GOBP_REGULATION_OF_LYASE_ACTIVITY

GOBP_SEMAPHORIN_PLEXIN_SIGNALING_PATHWAY_INVOLVED_IN_AXON_GUIDANCE, GOBP_SEMAPHORIN_PLEXIN_SIGNALING_PATHWAY_INVOLVED_IN_AXON_GUIDANCE

HP_ULCERATIVE_COLITIS, HP_ULCERATIVE_COLITIS

HP_ABNORMAL_OSSIFICATION_INVOLVING_THE_FEMORAL_HEAD_AND_NECK, HP_ABNORMAL_OSSIFICATION_INVOLVING_THE_FEMORAL_HEAD_AND_NECK

GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY

GOBP_POSITIVE_REGULATION_OF_CHROMATIN_BINDING, GOBP_POSITIVE_REGULATION_OF_CHROMATIN_BINDING

GOBP_SPHINGOSINE_BIOSYNTHETIC_PROCESS, GOBP_SPHINGOSINE_BIOSYNTHETIC_PROCESS

GOCC_AP_3_ADAPTOR_COMPLEX, GOCC_AP_3_ADAPTOR_COMPLEX

HP_PROGRESSIVE_EXTRAPYRAMIDAL_MOVEMENT_DISORDER, HP_PROGRESSIVE_EXTRAPYRAMIDAL_MOVEMENT_DISORDER

GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPOLYMERIZATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPOLYMERIZATION