

Y30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN,GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN

GSE41867_DAY6_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN,GSE41867_DAY6_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN
GSE12845_IJD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_DN,GSE12845_IJD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_DN
GSE11961_FOLLICULAR_BCELL_VS_MEMORY_BCELL_DAY7_DN,GSE11961_FOLLICULAR_BCELL_VS_MEMORY_BCELL_DAY7_DN
GSE11721_CTRL_VS_POLYIC_H1BMDCL_UP,GSE11721_CTRL_VS_POLYIC_H1BMDCL_UP
GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP,GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GSE42021_TREG_PLN_VS_CD24HL_TREG_THYMUS_UP,GSE42021_TREG_PLN_VS_CD24HL_TREG_THYMUS_UP
GOBP_NCRNA_TRANSCRIPTION,GOBP_NCRNA_TRANSCRIPTION
GSE16385_ROSIGLITAZONE_IPNG_TNF_VS_IL4_STIM_MACROPHAGE_DN,GSE16385_ROSIGLITAZONE_IPNG_TNF_VS_IL4_STIM_MACROPHAGE_DN
GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_UP,GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_UP
GSE339_CDRP5_VS_CD438DN_DC_IN_CULTURE_DN,GSE339_CDRP5_VS_CD438DN_DC_IN_CULTURE_DN
GSE1448_ANTI_VALPHA2_VS_VBETA5_DP_THYMOCYTE_DN,GSE1448_ANTI_VALPHA2_VS_VBETA5_DP_THYMOCYTE_DN
GOBP_SNRNA_TRANSCRIPTION,GOBP_SNRNA_TRANSCRIPTION
MIR2054,MIR2054
MIR3679_3P,MIR3679_3P
HP_TRIANGULAR_FACE,HP_TRIANGULAR_FACE
MIR1264,MIR1264
GSE27786_ERYTHROBLAST_VS_NEUTROPHIL_UP,GSE27786_ERYTHROBLAST_VS_NEUTROPHIL_UP
GSE41176_WT_VS_TAK1_KO_ANTI_IKM_STIM_BCELL_1H_UP,GSE41176_WT_VS_TAK1_KO_ANTI_IKM_STIM_BCELL_1H_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_30H_UP,GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_30H_UP
MIR5801_3P,MIR5801_3P
GOBP_ENDOPLASMIC_RETICULUM_ORGANIZATION,GOBP_ENDOPLASMIC_RETICULUM_ORGANIZATION
CGGAARNGGCNG_UNKNOWN,CGGAARNGGCNG_UNKNOWN
MIR4772_3P,MIR4772_3P
GOMF_RNA_POLYMERASE_BINDING,GOMF_RNA_POLYMERASE_BINDING
GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN,GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN
MIR297,MIR297
RITTTNNNTTTCM_UNKNOWN,RITTTNNNTTTCM_UNKNOWN
MIR3171,MIR3171
MIR6806_5P,MIR6806_5P
GOMF_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY,GOMF_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
HP_RETINAL_DETACHMENT,HP_RETINAL_DETACHMENT
MIR217_3P,MIR217_3P
MIR4714_5P,MIR4714_5P
HP_SHORT_PALM,HP_SHORT_PALM
MIR5007_3P,MIR5007_3P
CRX_DN.V1_DN,CRX_DN.V1_DN
MIR4540,MIR4540
GOBP_MANNOSYLATION,GOBP_MANNOSYLATION
HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_PYRAMIDAL_TRACT,HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_PYRAMIDAL_TRACT
MIR4748,MIR4748
CCAGGTT,MIR490,CCAGGTT,MIR490
GOMF_RNA_POLYMERASE_IL_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY,GOMF_RNA_POLYMERASE_IL_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
MIR6824_5P,MIR6824_5P
GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN,GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN
FIGUEROA_AML_METHYLATION_CLUSTER_1_DN,FIGUEROA_AML_METHYLATION_CLUSTER_1_DN
HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_CORTICOSPINAL_TRACT,HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_CORTICOSPINAL_TRACT
GOCC_TRANSCRIPTION_FACTOR_THFD_COMPLEX,GOCC_TRANSCRIPTION_FACTOR_THFD_COMPLEX
MIR3161,MIR3161
GENTILE_UV_RESPONSE_CLUSTER_D5,GENTILE_UV_RESPONSE_CLUSTER_D5
MIR8060,MIR8060
GAZDA_DIAMOND_BLACKFAN_ANEMIA_ERYTHROID_UP,GAZDA_DIAMOND_BLACKFAN_ANEMIA_ERYTHROID_UP
GOBP_PROTEIN_LOCALIZATION_TO_ENDOSOME,GOBP_PROTEIN_LOCALIZATION_TO_ENDOSOME
REACTOME_RND2_GTPASE_CYCLE,REACTOME_RND2_GTPASE_CYCLE
HP_ABNORMALLY_LARGE_GLOBE,HP_ABNORMALLY_LARGE_GLOBE
WP_COPPER_HOMEOSTASIS,WP_COPPER_HOMEOSTASIS
MIR6502_3P,MIR6502_3P
GOBP_INTRA_S_DNA_DAMAGE_CHECKPOINT,GOBP_INTRA_S_DNA_DAMAGE_CHECKPOINT
GOBP_PROTEIN_INSERTION INTO_ER_MEMBRANE,GOBP_PROTEIN_INSERTION INTO_ER_MEMBRANE
REACTOME_RND3_GTPASE_CYCLE,REACTOME_RND3_GTPASE_CYCLE
GOBP_DNA_REPLICATION_CHECKPOINT,GOBP_DNA_REPLICATION_CHECKPOINT
MIR6780B_3P,MIR6780B_3P
GOBP_PROTEIN_MANNOSYLATION,GOBP_PROTEIN_MANNOSYLATION
ANDERSON_BLOOD_CN54GP40_ADJUVANTED_WITH_GLA_AE_AGE_18_45YO_6HR_UP,ANDERSON_BLOOD_CN54GP40_ADJUVANTED_WITH_GLA_AE_AGE_18_45YO_6HR_UP
KEGC_BASAL_TRANSCRIPTION_FACTORS_TRANSCRIPTION_FACTORS
HP_BEAKING_OF_VERTEBRAL_BODIES,HP_BEAKING_OF_VERTEBRAL_BODIES
WP_ALZHEIMERS_DISEASE,WP_ALZHEIMERS_DISEASE
GOMF_CARBOXY_LYASE_ACTIVITY,GOMF_CARBOXY_LYASE_ACTIVITY
GTACGGA,MIR378,GTACGGA,MIR378
MIR710B_3P,MIR710B_3P
GOBP_MITOTIC_DNA_REPLICATION_CHECKPOINT,GOBP_MITOTIC_DNA_REPLICATION_CHECKPOINT
GOBP_ISOPRENOID_METABOLIC_PROCESS,GOBP_ISOPRENOID_METABOLIC_PROCESS
MIR1911_3P,MIR1911_3P
WP_PI3KAKTMTOR_SIGNALING_PATHWAY_AND_THERAPEUTIC_OPPORTUNITIES,WP_PI3KAKTMTOR_SIGNALING_PATHWAY_AND_THERAPEUTIC_OPPORTUNITIES
SIG_PIP3_SIGNALING_IN_B_LYMPHOCTES,SIG_PIP3_SIGNALING_IN_B_LYMPHOCTES
HP_LOWER_LIMB_ASYMMETRY,HP_LOWER_LIMB_ASYMMETRY
HP_NEURONAL_LOSS_IN_CENTRAL_NERVOUS_SYSTEM,HP_NEURONAL_LOSS_IN_CENTRAL_NERVOUS_SYSTEM
GOBP_ERYTHROCYTE_DEVELOPMENT,GOBP_ERYTHROCYTE_DEVELOPMENT
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAGENTA_DN,GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAGENTA_DN
MODULE_528,MODULE_528
SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_UP,SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_UP
GROSS_HIF1A_TARGETS_DN,GROSS_HIF1A_TARGETS_DN
GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_SINGLE_STRAND_ANNHEALING,GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_SINGLE_STRAND_ANNHEALING
GOBP_SMALL_NUCLEOLAR_RIBONUCLEOPROTEIN_COMPLEX_ASSEMBLY,GOBP_SMALL_NUCLEOLAR_RIBONUCLEOPROTEIN_COMPLEX_ASSEMBLY
NKX25_O1,NKX25_O1
MIR6501_5P,MIR6501_5P
MODULE_471,MODULE_471
HP_ABNORMAL_MUSCLE_FIBER_PROTEIN_EXPRESSION,HP_ABNORMAL_MUSCLE_FIBER_PROTEIN_EXPRESSION
GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_BY_EIF2_ALPHA_PHOSPHORYLATION,GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_BY_EIF2_ALPHA_PHOSPHORYLATION
HP_APLASIA_HYPOPLASIA_INVOLVING_THE_CARPAL_BONES,HP_APLASIA_HYPOPLASIA_INVOLVING_THE_CARPAL_BONES
HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY,HP_ABNORMALITY_OF_UPPER_LIMB_EPIPHYSIS_MORPHOLOGY
WP_PKCGAMMA_CALCIIUM_SIGNALING_PATHWAY_IN_ATAXIA,WP_PKCGAMMA_CALCIIUM_SIGNALING_PATHWAY_IN_ATAXIA
LMDX1B_TARGET_GENES,LMDX1B_TARGET_GENES
HP_SCLEROSIS_OF_HAND_BONE,HP_SCLEROSIS_OF_HAND_BONE
MEF2_O4,MEF2_O4
GOBP_ISOPRENOID_BIOSYNTHETIC_PROCESS,GOBP_ISOPRENOID_BIOSYNTHETIC_PROCESS
MIR382_5P,MIR382_5P
MODULE_432,MODULE_432
MIR4529_3P,MIR4529_3P
GOBP_MANGANESE_ION_TRANSMEMBRANE_TRANSPORT,GOBP_MANGANESE_ION_TRANSMEMBRANE_TRANSPORT
GOMF_MANGANESE_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY,GOMF_MANGANESE_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
PALOMERO_GS1_SENSITIVITY_UP,PALOMERO_GS1_SENSITIVITY_UP
GOBP_DOLCHIL_DIPHOSPHATE_BIOSYNTHETIC_PROCESS,GOBP_DOLCHIL_DIPHOSPHATE_BIOSYNTHETIC_PROCESS
HARALAMBIEVA_PBMC_FLUARX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LATE_GENE_EXPR_INDIVID_GENE_MODELS,PRED_PEAK_B_CELL_ELISPOT_RESP_NEGATIVE,HARALAMBIEVA_PBMC_FLUARX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LATE_GENE_EXPR_INDIVID_GENE_MODELS,PRED
GOBP_NEGATIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS
GOMF_RNA_POLYMERASE_III_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY,GOMF_RNA_POLYMERASE_III_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
GOCC_PALMITOYLTRANSFERASE_COMPLEX,GOCC_PALMITOYLTRANSFERASE_COMPLEX
MIR128B_3P,MIR128B_3P
GOBP_IRON_IMPORT_INTO_CELL,GOBP_IRON_IMPORT_INTO_CELL
GOMF_G_PROTEIN_COUPLED_SEROTONIN_RECEPTOR_BINDING,GOMF_G_PROTEIN_COUPLED_SEROTONIN_RECEPTOR_BINDING
GOBP_CAMP_CATABOLIC_PROCESS,GOBP_CAMP_CATABOLIC_PROCESS
HP_PROGRESSIVE_SPASTIC_PARAPARESIS,HP_PROGRESSIVE_SPASTIC_PARAPARESIS
GOBP_NEURAL_FOLD_FORMATION,GOBP_NEURAL_FOLD_FORMATION
TERAMOTO_OPN_TARGETS_CLUSTER_1,TERAMOTO_OPN_TARGETS_CLUSTER_1
chr18q12,chr18q12
HP_ULNAR_DEVIATION_OF_FINGER,HP_ULNAR_DEVIATION_OF_FINGER
HP_BUPHTHALMOS,HP_BUPHTHALMOS
HP_DILATATED_INTERNAL_AUDITORY_CANAL,HP_DILATATED_INTERNAL_AUDITORY_CANAL
HP_ABNORMAL_BRAIN_CHOLINE_LEVEL_BY_MRS,HP_ABNORMAL_BRAIN_CHOLINE_LEVEL_BY_MRS
chr18q21,chr18q21
HP_JUVENILE_CATARACT,HP_JUVENILE_CATARACT
GOMF_INTERLEUKIN_17_RECEPTOR_ACTIVITY,GOMF_INTERLEUKIN_17_RECEPTOR_ACTIVITY
HP_HEMIPLEGIA,HP_HEMIPLEGIA
HP_ABNORMAL_COCHLEA_MORPHOLOGY,HP_ABNORMAL_COCHLEA_MORPHOLOGY
WP_GAMMAGLUTAMYL_CYCLE_FOR_THE_BIOSYNTHESIS_AND_DEGRADATION_OF_GLUTATHIONE_INCLUDING_DISEASES,WP_GAMMAGLUTAMYL_CYCLE_FOR_THE_BIOSYNTHESIS_AND_DEGRADATION_OF_GLUTATHIONE_INCLUDING_DISEASES
HP_POST_PARTUM_HEMORRHAGE,HP_POST_PARTUM_HEMORRHAGE
HP_SHOULDER_DISLOCATION,HP_SHOULDER_DISLOCATION
GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGNALING_PATHWAY,GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGNALING_PATHWAY
REACTOME_SYNTHESIS_OF_DOLICHYL_PHOSPHATE,REACTOME_SYNTHESIS_OF_DOLICHYL_PHOSPHATE
GOME_3_5_CYCLIC_AMP_PHOSPHODIESTERASE_ACTIVITY,GOME_3_5_CYCLIC_AMP_PHOSPHODIESTERASE_ACTIVITY
GOBP_PROTEIN_C_LINKED_GLYCOSYLATION,GOBP_PROTEIN_C_LINKED_GLYCOSYLATION
GOBP_NEGATIVE_REGULATION_OF_CERAMIDE_BIOSYNTHETIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_CERAMIDE_BIOSYNTHETIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION,GOBP_POSITIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION
REACTOME_RELEASE_OF_HH_NP_FROM_THE_SECRETING_CELL,REACTOME_RELEASE_OF_HH_NP_FROM_THE_SECRETING_CELL
GOBP_NEGATIVE_REGULATION_OF_MYELINATION,GOBP_NEGATIVE_REGULATION_OF_MYELINATION
REACTOME_GLUTATHIONE_SYNTHESIS_AND_RECYCLING,REACTOME_GLUTATHIONE_SYNTHESIS_AND_RECYCLING
HP_ABNORMAL_EPIPHYSIS_MORPHOLOGY_OF_THE_PHALANGES_OF_THE_HAND,HP_ABNORMAL_EPIPHYSIS_MORPHOLOGY_OF_THE_PHALANGES_OF_THE_HAND
WP_NRF2ARE_REGULATION,WP_NRF2ARE_REGULATION
GOME_ARYL_HYDROCARBON_RECEPTOR_BINDING,GOME_ARYL_HYDROCARBON_RECEPTOR_BINDING
HP_COCHLEAR_MALFORMATION,HP_COCHLEAR_MALFORMATION
GOBP_MULTICELLULAR_ORGANISMAL_IRON_ION_HOMEOSTASIS,GOBP_MULTICELLULAR_ORGANISMAL_IRON_ION_HOMEOSTASIS
GOMF_INTERLEUKIN_6_RECEPTOR_BINDING,GOMF_INTERLEUKIN_6_RECEPTOR_BINDING
REACTOME_PECAM1_INTERACTIONS,REACTOME_PECAM1_INTERACTIONS
GOBP_KERATINOCYTE_APOPTOTIC_PROCESS,GOBP_KERATINOCYTE_APOPTOTIC_PROCESS
HP_SCLEROSIS_OF_FINGER_PHALANX,HP_SCLEROSIS_OF_FINGER_PHALANX
GOBP_THREONINE_METABOLIC_PROCESS,GOBP_THREONINE_METABOLIC_PROCESS
GOBP_HISTONE_DEPHOSPHORYLATION,GOBP_HISTONE_DEPHOSPHORYLATION
GOBP_DENTATE_GYRUS_DEVELOPMENT,GOBP_DENTATE_GYRUS_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_NON_CANONICAL_WNT_SIGNALING_PATHWAY,GOBP_POSITIVE_REGULATION_OF_NON_CANONICAL_WNT_SIGNALING_PATHWAY
GOBP_INNER_DYNEIN_ARM_ASSEMBLY,GOBP_INNER_DYNEIN_ARM_ASSEMBLY
GOBP_DETECTION_OF_LIGHT_STIMULUS,GOBP_DETECTION_OF_LIGHT_STIMULUS
HP_NEUROMUSCULAR_DYSPHAGIA,HP_NEUROMUSCULAR_DYSPHAGIA
REACTOME_SIGNALING_BY_NOTCH1_T_7_9_NOTCH1_M1580_K2555_TRANSLOCATION_MUTANT,REACTOME_SIGNALING_BY_NOTCH1_T_7_9_NOTCH1_M1580_K2555_TRANSLOCATION_MUTANT
GOBP_PHOTOTRANSDUCTION_VISIBLE_LIGHT,GOBP_PHOTOTRANSDUCTION_VISIBLE_LIGHT
HP_SHORT_THORAX,HP_SHORT_THORAX
WIEMANN_TELOMERE_SHORTENING_AND_CHRONIC_LIVER_DAMAGE_DN,WIEMANN_TELOMERE_SHORTENING_AND_CHRONIC_LIVER_DAMAGE_DN
BIOCARTA_TURBY_PATHWAY,BIOCARTA_TURBY_PATHWAY
HP_LACRIMAL_DUCT_APLASIA,HP_LACRIMAL_DUCT_APLASIA
GOBP_THREONINE_CATABOLIC_PROCESS,GOBP_THREONINE_CATABOLIC_PROCESS
GOBP_FAT_PAD_DEVELOPMENT,GOBP_FAT_PAD_DEVELOPMENT
MIR3677_3P,MIR3677_3P
HP_WIDE_PUBIC_SYMPHYSIS,HP_WIDE_PUBIC_SYMPHYSIS
HP_HEAD_BANGING,HP_HEAD_BANGING
GOBP_REGULATION_OF_NADP_METABOLIC_PROCESS,GOBP_REGULATION_OF_NADP_METABOLIC_PROCESS
GOBP_MESENODERM_DEVELOPMENT,GOBP_MESENODERM_DEVELOPMENT
HP_PROLONGED_BLEEDING_FOLLOWING_PROCEDURE,HP_PROLONGED_BLEEDING_FOLLOWING_PROCEDURE
GOBP_MANGANESE_ION_TRANSPORT,GOBP_MANGANESE_ION_TRANSPORT
ZHONG_PFC_HES1_PFC_C1_NPC_ZHONG_PFC_HES1_PFC_C1_NPC
HP_IMPAIRED_RISTOCETIN_INDUCED_PLATELET_AGGREGATION,HP_IMPAIRED_RISTOCETIN_INDUCED_PLATELET_AGGREGATION
HP_BARREL_SHAPED_CHEST,HP_BARREL_SHAPED_CHEST
HP_CONE_SHAPED_EPIPHYSES_OF_THE_PHALANGES_OF_THE_HAND,HP_CONE_SHAPED_EPIPHYSES_OF_THE_PHALANGES_OF_THE_HAND
HP_ABNORMAL_ANTERIOR_CHAMBER_MORPHOLOGY,HP_ABNORMAL_ANTERIOR_CHAMBER_MORPHOLOGY
HP_REDUCED_PANCREATIC_BETA_CELLS,HP_REDUCED_PANCREATIC_BETA_CELLS
HP_ABNORMALITY_OF_THE_GLENOID_FOSSA,HP_ABNORMALITY_OF_THE_GLENOID_FOSSA
GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_PHOSPHORYLATION,GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_PHOSPHORYLATION
GOMF_GLUTAMATE_BINDING,GOMF_GLUTAMATE_BINDING
HP_ONCOSTATIN_M_MEDIATED_SIGNALING_PATHWAY,GOBP_ONCOSTATIN_M_MEDIATED_SIGNALING_PATHWAY
HP_CORNEL_DYSTROPHY,HP_CORNEL_DYSTROPHY
HOEK_PBMC_INACTIVATED_INFLUENZA_ADULT_3DY_UP,HOEK_PBMC_INACTIVATED_INFLUENZA_ADULT_3DY_UP
HP_SHALLOW_ANTERIOR_CHAMBER,HP_SHALLOW_ANTERIOR_CHAMBER
HP_ABNORMALITY_OF_THE_EPIPHYSES_OF_THE_FEET,HP_ABNORMALITY_OF_THE_EPIPHYSES_OF_THE_FEET
GOMF_CADMIUM_ION_BINDING,GOMF_CADMIUM_ION_BINDING
HP_ANAL_FISSURE,HP_ANAL_FISSURE
GOBP_CILIARY_NEUROTROPHIC_FACTOR_MEDIATED_SIGNALING_PATHWAY,GOBP_CILIARY_NEUROTROPHIC_FACTOR_MEDIATED_SIGNALING_PATHWAY
WP_FTO_OBESITY_VARIANT_MECHANISM,WP_FTO_OBESITY_VARIANT_MECHANISM
HAY_BONE_MARROW_CD34_POS_MEP,HAY_BONE_MARROW_CD34_POS_MEP
GOMF_CILIARY_NEUROTROPHIC_FACTOR_RECEPTOR_ACTIVITY,GOMF_CILIARY_NEUROTROPHIC_FACTOR_RECEPTOR_ACTIVITY
REACTOME_TANDEM_PORE_DOMAIN_POTASSIUM_CHANNELS,REACTOME_TANDEM_PORE_DOMAIN_POTASSIUM_CHANNELS
GOMF_ONCOSTATIN_M_RECEPTOR_ACTIVITY,GOMF_ONCOSTATIN_M_RECEPTOR_ACTIVITY
CHESLER_BRAIN_DMW150_QTL_TRANS_CHESLER_BRAIN_DMW150_QTL_TRANS
GOMF_CILIARY_NEUROTROPHIC_FACTOR_RECEPTOR_BINDING,GOBP_CILIARY_NEUROTROPHIC_FACTOR_RECEPTOR_BINDING
GOBP_POINTED_END_ACTIN_FILAMENT_CAPPING,GOBP_POINTED_END_ACTIN_FILAMENT_CAPPING
HP_HYPOPLASTIC_ILIAC_WING,HP_HYPOPLASTIC_ILIAC_WING
GOBP_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_PRODUCTION,GOBP_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_PRODUCTION