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GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP
 GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP
 GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP
 GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP
 GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP
 GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_6H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_6H_UP
 GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP
 GSE9988_LOW_LPS_VS_CTRL_TREATED_MONOCYTE_DN, GSE9988_LOW_LPS_VS_CTRL_TREATED_MONOCYTE_DN
 GSE9988_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN, GSE9988_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN
 GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP
 GSE14000_4H_VS_16H_LPS_DC_DN, GSE14000_4H_VS_16H_LPS_DC_DN
 GSE14000_UNSTIM_VS_16H_LPS_DC_UP, GSE14000_UNSTIM_VS_16H_LPS_DC_UP
 GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_UP, GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_UP
 GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_DN, GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_DN
 GSE7768_OVA_WITH_LPS_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_WITH_LPS_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP
 GSE35543_IN_VITRO_ITREG_VS_CONVERTED_EX_ITREG_UP, GSE35543_IN_VITRO_ITREG_VS_CONVERTED_EX_ITREG_UP
 GSE37416_CTRL_VS_6H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_6H_F_TULARENSIS_LVS_NEUTROPHIL_UP
 GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP
 GSE2706_UNSTIM_VS_8H_LPS_DC_UP, GSE2706_UNSTIM_VS_8H_LPS_DC_UP
 GOBP_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GOBP_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
  BLANCO_MELO_COVID19_SARS_COV_2_POS_PATIENT_LUNG_TISSUE_DN, BLANCO_MELO_COVID19_SARS_COV_2_POS_PATIENT_LUNG_TISSUE_DN
 GSE2706_UNSTIM_VS_2H_LPS_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_DC_UP
 HP_JOINT_HYPERFLEXIBILITY, HP_JOINT_HYPERFLEXIBILITY
  GSE2706_UNSTIM_VS_2H_R848_DC_UP, GSE2706_UNSTIM_VS_2H_R848_DC_UP
  HP_ABNORMAL_FEMORAL_NECK_HEAD_MORPHOLOGY, HP_ABNORMAL_FEMORAL_NECK_HEAD_MORPHOLOGY
  MIR6852 3P, MIR6852 3P
  HP_PREMATURELY_AGED_APPEARANCE, HP_PREMATURELY_AGED_APPEARANCE
 GSE21546_WT_VS_SAP1A_KO_DP_THYMOCYTES_DN, GSE21546_WT_VS_SAP1A_KO_DP_THYMOCYTES_DN
 GOMF_RIBONUCLEASE_ACTIVITY, GOMF_RIBONUCLEASE_ACTIVITY
 WEI_MIR34A_TARGETS, WEI_MIR34A_TARGETS
 HP_SLENDER_FINGER, HP_SLENDER_FINGER
 GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_TURQUOISE_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_TURQUOISE_DN
 HP_SMALL_NAIL, HP_SMALL_NAIL
  HP_APLASIA_HYPOPLASIA_INVOLVING_THE_PELVIS, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_PELVIS
 MIR3176, MIR3176
 GOMF_DEMETHYLASE_ACTIVITY, GOMF_DEMETHYLASE_ACTIVITY
 MIR3922_3P, MIR3922_3P
 HP_ABNORMAL_PATELLA_MORPHOLOGY, HP_ABNORMAL_PATELLA_MORPHOLOGY
 HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_HAND, HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_HAND
 HP_THROMBOCYTOSIS, HP_THROMBOCYTOSIS
 GOCC PRERIBOSOME SMALL SUBUNIT PRECURSOR, GOCC PRERIBOSOME SMALL SUBUNIT PRECURSOR
 HP_APLASIA_HYPOPLASIA_OF_THE_FEMUR, HP_APLASIA_HYPOPLASIA_OF_THE_FEMUR
 HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_UPPER_LIMBS, HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_UPPER_LIMBS
 HP_RADIOULNAR_SYNOSTOSIS, HP_RADIOULNAR_SYNOSTOSIS
 WP_RETT_SYNDROME_CAUSING_GENES, WP_RETT_SYNDROME_CAUSING_GENES
 GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP
 GAO_STOMACH_24W_C1_PROCRPOS_MULTIPOTENT_PROGENITOR, GAO_STOMACH_24W_C1_PROCRPOS_MULTIPOTENT_PROGENITOR
 GOMF ORGANIC ACID TRANSMEMBRANE TRANSPORTER ACTIVITY, GOMF ORGANIC ACID TRANSMEMBRANE TRANSPORTER ACTIVITY
 FIGUEROA_AML_METHYLATION_CLUSTER_4_UP, FIGUEROA_AML_METHYLATION_CLUSTER_4_UP
 NUTT_GBM_VS_AO_GLIOMA_DN, NUTT_GBM_VS_AO_GLIOMA_DN
 GOMF_ACETYLTRANSFERASE_ACTIVITY, GOMF_ACETYLTRANSFERASE_ACTIVITY
 MIR3909, MIR3909
 GSE13547_CTRL_VS_ANTI_IGM_STIM_ZFX_KO_BCELL_12H_UP, GSE13547_CTRL_VS_ANTI_IGM_STIM_ZFX_KO_BCELL_12H_UP
 HP_INCREASED_BONE_MINERAL_DENSITY, HP_INCREASED_BONE_MINERAL_DENSITY
  MIR7114 3P, MIR7114 3P
 HP_METATARSUS_ADDUCTUS, HP_METATARSUS_ADDUCTUS
 GOCC_MRNA_CLEAVAGE_FACTOR_COMPLEX, GOCC_MRNA_CLEAVAGE_FACTOR_COMPLEX
 GOBP_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION, GOBP_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION
 MIR3976, MIR3976
 GOCC_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX
 GOBP_DEMETHYLATION, GOBP_DEMETHYLATION
 HP_APLASIA_HYPOPLASIA_INVOLVING_THE_FEMORAL_HEAD_AND_NECK, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_FEMORAL_HEAD_AND_NECK
 SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5, SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5
  FERREIRA_EWINGS_SARCOMA_UNSTABLE_VS_STABLE_DN, FERREIRA_EWINGS_SARCOMA_UNSTABLE_VS_STABLE_DN
 HP_ABNORMAL_CIRCULATING_ASPARTATE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_ASPARTATE_FAMILY_AMINO_ACID_CONCENTRATION
  IZADPANAH_STEM_CELL_ADIPOSE_VS_BONE_DN, IZADPANAH_STEM_CELL_ADIPOSE_VS_BONE_DN
 MIR6727_3P, MIR6727_3P
 MIR1180_5P, MIR1180_5P
 MIR500A_3P, MIR500A_3P
 GOBP_REGULATION_OF_RRNA_PROCESSING, GOBP_REGULATION_OF_RRNA_PROCESSING
 HP_ABNORMALITY_OF_EARLOBE, HP_ABNORMALITY_OF_EARLOBE
 HP_NAIL_DYSPLASIA, HP_NAIL_DYSPLASIA
 MIR4763_5P, MIR4763_5P
 SNAPC4_TARGET_GENES, SNAPC4_TARGET_GENES
 GOCC_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX, GOCC_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX
 MIR3189_3P, MIR3189_3P
 HP_IMPAIRED_SMOOTH_PURSUIT, HP_IMPAIRED_SMOOTH_PURSUIT
 GOBP_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS
 GCM_PPM1D, GCM_PPM1D
 GOBP_NEGATIVE_REGULATION_OF_SMOOTHENED_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_SMOOTHENED_SIGNALING_PATHWAY
 HP_ABSENT_FINGERNAIL, HP_ABSENT_FINGERNAIL
 GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK, GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK
 GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_DN, GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_DN
 HP APLASIA HYPOPLASIA OF THE PHALANGES OF THE TOES, HP APLASIA HYPOPLASIA OF THE PHALANGES OF THE TOES
 GOBP_DICARBOXYLIC_ACID_TRANSPORT, GOBP_DICARBOXYLIC_ACID_TRANSPORT
 VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN, VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN
 GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_DN
 HP_MILD_GLOBAL_DEVELOPMENTAL_DELAY, HP_MILD_GLOBAL_DEVELOPMENTAL_DELAY
 GOMF_LYSINE_ACETYLATED_HISTONE_BINDING, GOMF_LYSINE_ACETYLATED_HISTONE_BINDING
 GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_FROM_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICISTRONIC_TRICITRONIC_TRICITRONIC_TRICITRONIC_TRICITRONIC_TRICISTRONIC_TRICITRONIC_TRICITRONIC_TRICITRONIC_TRICITRONIC_TRICITRO
  BIOCARTA_CPSF_PATHWAY, BIOCARTA_CPSF_PATHWAY
 GOBP_HISTONE_H2A_ACETYLATION, GOBP_HISTONE_H2A_ACETYLATION
 GOBP_PIRNA_METABOLIC_PROCESS, GOBP_PIRNA_METABOLIC_PROCESS
 HP_ABNORMAL_DELIVERY, HP_ABNORMAL_DELIVERY
 HP_PROGRESSIVE_GAIT_ATAXIA, HP_PROGRESSIVE_GAIT_ATAXIA
 HP_THORACOLUMBAR_SCOLIOSIS, HP_THORACOLUMBAR_SCOLIOSIS
 GOBP_PHOTOPERIODISM, GOBP_PHOTOPERIODISM
 HP_ABNORMAL_CIRCULATING_FATTY_ACID_ANION_CONCENTRATION, HP_ABNORMAL_CIRCULATING_FATTY_ACID_ANION_CONCENTRATION
 GOMF_CYTOSKELETAL_ANCHOR_ACTIVITY, GOMF_CYTOSKELETAL_ANCHOR_ACTIVITY
 GOMF_5S_RRNA_BINDING, GOMF_5S_RRNA_BINDING
 MIR3158_3P, MIR3158_3P
 GOBP_CELLULAR_SENESCENCE, GOBP_CELLULAR_SENESCENCE
 GOMF_RNA_POLYMERASE_I_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOMF_RNA_POLYMERASE_I_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING
 REACTOME_POLB_DEPENDENT_LONG_PATCH_BASE_EXCISION_REPAIR, REACTOME_POLB_DEPENDENT_LONG_PATCH_BASE_EXCISION_REPAIR
 GOBP_RNA_INTERFERENCE, GOBP_RNA_INTERFERENCE
 MIR758 5P, MIR758 5P
 SA_B_CELL_RECEPTOR_COMPLEXES, SA_B_CELL_RECEPTOR_COMPLEXES
  MIR424_3P, MIR424_3P
 GOMF_L_ASCORBIC_ACID_BINDING, GOMF_L_ASCORBIC_ACID_BINDING
 HP_ABNORMAL_EYELID_PHYSIOLOGY, HP_ABNORMAL_EYELID_PHYSIOLOGY
 GOBP_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE
 HP_APLASIA_HYPOPLASIA_OF_THE_PUBIC_BONE, HP_APLASIA_HYPOPLASIA_OF_THE_PUBIC_BONE
 chr17q12, chr17q12
 GOMF_SOLUTE_ANION_ANTIPORTER_ACTIVITY, GOMF_SOLUTE_ANION_ANTIPORTER_ACTIVITY
 MODULE_372, MODULE_372
 HP SEVERE POSTNATAL GROWTH RETARDATION, HP SEVERE POSTNATAL GROWTH RETARDATION
 GOBP_MITOTIC_CHROMOSOME_CONDENSATION, GOBP_MITOTIC_CHROMOSOME_CONDENSATION
 MIR1915_5P, MIR1915_5P
 GOCC_CHROMATIN_SILENCING_COMPLEX, GOCC_CHROMATIN_SILENCING_COMPLEX
 GOCC_CONDENSIN_COMPLEX, GOCC_CONDENSIN_COMPLEX
 HP_ABNORMAL_ORAL_FRENULUM_MORPHOLOGY, HP_ABNORMAL_ORAL_FRENULUM_MORPHOLOGY
 GOBP_CARDIOLIPIN_METABOLIC_PROCESS, GOBP_CARDIOLIPIN_METABOLIC_PROCESS
 HP_CALCIFICATION_OF_CARTILAGE, HP_CALCIFICATION_OF_CARTILAGE
 GOBP_ASPARTATE_TRANSMEMBRANE_TRANSPORT, GOBP_ASPARTATE_TRANSMEMBRANE_TRANSPORT
 GOBP_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION, GOBP_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION
 GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
GOBP_POSITIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GOBP_POSITIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
WP_ATR_SIGNALING, WP_ATR_SIGNALING
HP_NONPROGRESSIVE_CEREBELLAR_ATAXIA, HP_NONPROGRESSIVE_CEREBELLAR_ATAXIA
HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANX_OF_THE_5TH_TOE, HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANX_OF_THE_5TH_TOE
GOMF_DICARBOXYLIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_DICARBOXYLIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
 GOBP_PHOSPHATIDYLSERINE_METABOLIC_PROCESS, GOBP_PHOSPHATIDYLSERINE_METABOLIC_PROCESS
GOBP_SULFUR_COMPOUND_TRANSPORT, GOBP_SULFUR_COMPOUND_TRANSPORT
GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY
MIR6798_3P, MIR6798_3P
GOBP_CHROMOSOME_CONDENSATION, GOBP_CHROMOSOME_CONDENSATION
GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING, GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING
  HP_ABNORMALITY_OF_THE_ANTIHELIX, HP_ABNORMALITY_OF_THE_ANTIHELIX
  MIR663B, MIR663B
  HP_BREECH_PRESENTATION, HP_BREECH_PRESENTATION
  GOMF_NADPLUS_KINASE_ACTIVITY, GOMF_NADPLUS_KINASE_ACTIVITY
  HP_PROTRUSIO_ACETABULI, HP_PROTRUSIO_ACETABULI
  GOMF_FLAP_ENDONUCLEASE_ACTIVITY, GOMF_FLAP_ENDONUCLEASE_ACTIVITY
  HP_BILATERAL_TALIPES_EQUINOVARUS, HP_BILATERAL_TALIPES_EQUINOVARUS
  HP_ABSENT_EARLOBE, HP_ABSENT_EARLOBE
 HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR, HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR
 GOBP_NEGATIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY
 GOMF_GLYCINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_GLYCINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
 REACTOME_SMALL_INTERFERING_RNA_SIRNA_BIOGENESIS, REACTOME_SMALL_INTERFERING_RNA_SIRNA_BIOGENESIS
 GOBP_FOREBRAIN_NEURON_FATE_COMMITMENT, GOBP_FOREBRAIN_NEURON_FATE_COMMITMENT
 GOMF_RNA_2_O_METHYLTRANSFERASE_ACTIVITY, GOMF_RNA_2_O_METHYLTRANSFERASE_ACTIVITY
  GOCC_EARLY_PHAGOSOME, GOCC_EARLY_PHAGOSOME
  HP_CERVICAL_KYPHOSIS, HP_CERVICAL_KYPHOSIS
  GOCC_RDNA_HETEROCHROMATIN, GOCC_RDNA_HETEROCHROMATIN
  GOBP VITAMIN K METABOLIC PROCESS, GOBP VITAMIN K METABOLIC PROCESS
  GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K4_SPECIFIC, GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K4_SPECIFIC
 GOCC_H3_HISTONE_ACETYLTRANSFERASE_COMPLEX, GOCC_H3_HISTONE_ACETYLTRANSFERASE_COMPLEX
  HP_ABNORMALITY_OF_THE_PUBIC_BONE, HP_ABNORMALITY_OF_THE_PUBIC_BONE
 GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_CELL_DIFFERENTIATION
 GOBP_COLLAGEN_ACTIVATED_TYROSINE_KINASE_RECEPTOR_SIGNALING_PATHWAY, GOBP_COLLAGEN_ACTIVATED_TYROSINE_KINASE_RECEPTOR_SIGNALING_PATHWAY
GOBP_PURINE_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION, GOBP_POSITIVE_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION
GOBP_3_PHOSPHOADENOSINE_5_PHOSPHOSULFATE_BIOSYNTHETIC_PROCESS, GOBP_3_PHOSPHOADENOSINE_5_PHOSPHOSULFATE_BIOSYNTHETIC_PROCESS
 REACTOME_TRANSPORT_AND_SYNTHESIS_OF_PAPS, REACTOME_TRANSPORT_AND_SYNTHESIS_OF_PAPS
 HP_FLAT_CORNEA, HP_FLAT_CORNEA
 GOMF_SULFUR_COMPOUND_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_SULFUR_COMPOUND_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
 HP_LARGE_EARLOBE, HP_LARGE_EARLOBE
 GSE21546_UNSTIM_VS_ANTI_CD3_STIM_SAP1A_KO_AND_ELK1_KO_DP_THYMOCYTES_DN, GSE21546_UNSTIM_VS_ANTI_CD3_STIM_SAP1A_KO_AND_ELK1_KO_DP_THYMOCYTES_DN
 HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT, HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT
 GOBP_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION
 GOBP_REGULATION_OF_INTESTINAL_CHOLESTEROL_ABSORPTION, GOBP_REGULATION_OF_INTESTINAL_CHOLESTEROL_ABSORPTION
 HP_NYSTAGMUS_INDUCED_HEAD_NODDING, HP_NYSTAGMUS_INDUCED_HEAD_NODDING
 HP_SHORT_LINGUAL_FRENULUM, HP_SHORT_LINGUAL_FRENULUM
 OKAWA_NEUROBLASTOMA_1P36_31_DELETION, OKAWA_NEUROBLASTOMA_1P36_31_DELETION
 HP_PERIVENTRICULAR_NODULAR_HETEROTOPIA, HP_PERIVENTRICULAR_NODULAR_HETEROTOPIA
 HP_TOENAIL_DYSPLASIA, HP_TOENAIL_DYSPLASIA
 GOBP_HEMATOPOIETIC_STEM_CELL_HOMEOSTASIS, GOBP_HEMATOPOIETIC_STEM_CELL_HOMEOSTASIS
 GOBP_REGULATION_OF_INTESTINAL_ABSORPTION, GOBP_REGULATION_OF_INTESTINAL_ABSORPTION
 GOBP_NUCLEAR_BODY_ORGANIZATION, GOBP_NUCLEAR_BODY_ORGANIZATION
 CARDOSO_RESPONSE_TO_GAMMA_RADIATION_AND_3AB, CARDOSO_RESPONSE_TO_GAMMA_RADIATION_AND_3AB
  HP_BROAD_ALVEOLAR_RIDGES, HP_BROAD_ALVEOLAR_RIDGES
 GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION
 GOBP_TELOMERE_ASSEMBLY, GOBP_TELOMERE_ASSEMBLY
 GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS, GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS
 GOBP_HISTONE_H3_K4_DEMETHYLATION_TRIMETHYL_H3_K4_SPECIFIC, GOBP_HISTONE_H3_K4_DEMETHYLATION_TRIMETHYL_H3_K4_SPECIFIC
 GOBP NUCLEAR RETENTION OF PRE MRNA AT THE SITE OF TRANSCRIPTION, GOBP NUCLEAR RETENTION OF PRE MRNA AT THE SITE OF TRANSCRIPTION
 GOMF_RRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY, GOMF_RRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY
 GOBP_PRIMITIVE_HEMOPOIESIS, GOBP_PRIMITIVE_HEMOPOIESIS
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NSTIM VS 4H LPS DC UP, GSE14000 UNSTIM VS 4H LPS DC UP

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MIR10399_3P, MIR10399_3P
HP_INCOMPREHENSIBLE_SPEECH, HP_INCOMPREHENSIBLE_SPEECH
HP_ABNORMAL_CERVICAL_CURVATURE, HP_ABNORMAL_CERVICAL_CURVATURE
HP_SIRENOMELIA, HP_SIRENOMELIA
HP_ABNORMAL_RIB_OSSIFICATION, HP_ABNORMAL_RIB_OSSIFICATION
GOMF_1_PHOSPHATIDYLINOSITOL_4_PHOSPHATE_5_KINASE_ACTIVITY, GOMF_1_PHOSPHATIDYLINOSITOL_4_PHOSPHATE_5_KINASE_ACTIVITY
HP_LABIAL_HYPERTROPHY, HP_LABIAL_HYPERTROPHY
HP_LOW_ALKALINE_PHOSPHATASE, HP_LOW_ALKALINE_PHOSPHATASE
GOMF_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATE_PHOSPHATASE_ACTIVITY
WHITE_NEUROBLASTOMA_WITH_1P36.3_DELETION, WHITE_NEUROBLASTOMA_WITH_1P36.3_DELETION
GOBP_CELLULAR_RESPONSE_TO_BRAIN_DERIVED_NEUROTROPHIC_FACTOR_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_BRAIN_DERIVED_NEUROTROPHIC_FACTOR_STIMULUS
MPHOSPH8_TARGET_GENES, MPHOSPH8_TARGET_GENES
GOBP_XENOPHAGY, GOBP_XENOPHAGY
KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_4NQO_IN_OLD, KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_4NQO_IN_OLD
GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MEMBRANE, GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MEMBRANE
HP_CONSPICUOUSLY_HAPPY_DISPOSITION, HP_CONSPICUOUSLY_HAPPY_DISPOSITION
GOBP_TAURINE_TRANSPORT, GOBP_TAURINE_TRANSPORT
GOBP EMBRYONIC CLEAVAGE, GOBP EMBRYONIC CLEAVAGE
GOBP_REGULATION_OF_ARACHIDONIC_ACID_SECRETION, GOBP_REGULATION_OF_ARACHIDONIC_ACID_SECRETION
MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_30, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_30
GOBP_PRE_B_CELL_DIFFERENTIATION, GOBP_PRE_B_CELL_DIFFERENTIATION
GOBP_REGULATION_OF_SEQUESTERING_OF_ZINC_ION, GOBP_REGULATION_OF_SEQUESTERING_OF_ZINC_ION
GOBP_ADENOSINE_TO_INOSINE_EDITING, GOBP_ADENOSINE_TO_INOSINE_EDITING
GOMF_O_ACETYLTRANSFERASE_ACTIVITY, GOMF_O_ACETYLTRANSFERASE_ACTIVITY
chr17q23, chr17q23
GOBP_CDP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_CDP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS
GOCC_ZYMOGEN_GRANULE, GOCC_ZYMOGEN_GRANULE
HP_INTELLECTUAL_DISABILITY_BORDERLINE, HP_INTELLECTUAL_DISABILITY_BORDERLINE
HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_TOES, HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_TOES
REACTOME_BETA_OXIDATION_OF_PRISTANOYL_COA, REACTOME_BETA_OXIDATION_OF_PRISTANOYL_COA
GOMF VERY LONG CHAIN FATTY ACID COA LIGASE ACTIVITY, GOMF VERY LONG CHAIN FATTY ACID COA LIGASE ACTIVITY
GOMF_BIOACTIVE_LIPID_RECEPTOR_ACTIVITY, GOMF_BIOACTIVE_LIPID_RECEPTOR_ACTIVITY
GOBP_UV_PROTECTION, GOBP_UV_PROTECTION
MIR95 3P. MIR95 3P
HP_ABNORMAL_HUMERUS_MORPHOLOGY, HP_ABNORMAL_HUMERUS_MORPHOLOGY
GOBP_VOLUNTARY_MUSCULOSKELETAL_MOVEMENT, GOBP_VOLUNTARY_MUSCULOSKELETAL_MOVEMENT
GOBP_RESPONSE_TO_FUNGICIDE, GOBP_RESPONSE_TO_FUNGICIDE
REACTOME_FLT3_SIGNALING_THROUGH_SRC_FAMILY_KINASES, REACTOME_FLT3_SIGNALING_THROUGH_SRC_FAMILY_KINASES
GOBP_BONE_RESORPTION, GOBP_BONE_RESORPTION
GOCC_POSTSYNAPTIC_DENSITY_INTRACELLULAR_COMPONENT, GOCC_POSTSYNAPTIC_DENSITY_INTRACELLULAR_COMPONENT
FIGUEROA_AML_METHYLATION_CLUSTER_2_DN, FIGUEROA_AML_METHYLATION_CLUSTER_2_DN
HP_DACRYOCYSTITIS, HP_DACRYOCYSTITIS
GOBP_CELL_PROLIFERATION_IN_BONE_MARROW, GOBP_CELL_PROLIFERATION_IN_BONE_MARROW
HP_INCREASED_MUSCLE_GLYCOGEN_CONTENT, HP_INCREASED_MUSCLE_GLYCOGEN_CONTENT
HP_THYROID_AGENESIS, HP_THYROID_AGENESIS
MIR6814_3P_MIR6872_5P, MIR6814_3P_MIR6872_5P
HP_CYSTIC_RENAL_DYSPLASIA, HP_CYSTIC_RENAL_DYSPLASIA
GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS
GOMF_7S_RNA_BINDING, GOMF_7S_RNA_BINDING
GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATASE_ACTIVITY
STREICHER_LSM1_TARGETS_UP, STREICHER_LSM1_TARGETS_UP
NIKOLSKY_BREAST_CANCER_10Q22_AMPLICON, NIKOLSKY_BREAST_CANCER_10Q22_AMPLICON
GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_2, GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_2
GOBP_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY, GOBP_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY
GOCC_APICAL_CORTEX, GOCC_APICAL_CORTEX
MUNSHI_MULTIPLE_MYELOMA_DN, MUNSHI_MULTIPLE_MYELOMA_DN
HP_DECREASED_CALVARIAL_OSSIFICATION, HP_DECREASED_CALVARIAL_OSSIFICATION
GOBP_PHOSPHATIDYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLGLYCEROL_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_STEM_CELL_DIVISION, GOBP_REGULATION_OF_STEM_CELL_DIVISION
NIKOLSKY_BREAST_CANCER_21Q22_AMPLICON, NIKOLSKY_BREAST_CANCER_21Q22_AMPLICON
CHANG_POU5F1_TARGETS_UP, CHANG_POU5F1_TARGETS_UP
HP_DUMBBELL_SHAPED_LONG_BONE, HP_DUMBBELL_SHAPED_LONG_BONE
MIR3677_3P, MIR3677_3P
HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION, HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION
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GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MICROTUBULE, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MICROTUBULE GOBP_NEGATIVE_REGULATION_OF_HISTONE_DEACETYLATION, GOBP_NEGATIVE_REGULATION_OF_HISTONE_DEACETYLATION GOBP_POSITIVE_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS, GOBP_POSITIVE_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS GOBP_POSITIVE_REGULATION_OF_CELL_PROLIFERATION_IN_BONE_MARROW, GOBP_POSITIVE_REGULATION_OF_CELL_PROLIFERATION_IN_BONE_MARROW REACTOME_RUNX1_REGULATES_ESTROGEN_RECEPTOR_MEDIATED_TRANSCRIPTION, REACTOME_RUNX1_REGULATES_ESTROGEN_RECEPTOR_MEDIATED_TRANSCRIPTION

HP_ABNORMALLY_OSSIFIED_VERTEBRAE, HP_ABNORMALLY_OSSIFIED_VERTEBRAE GOBP_ANTEROGRADE_AXONAL_PROTEIN_TRANSPORT, GOBP_ANTEROGRADE_AXONAL_PROTEIN_TRANSPORT HP_LIMITED_ELBOW_FLEXION, HP_LIMITED_ELBOW_FLEXION HP_POSTPRANDIAL_HYPERGLYCEMIA, HP_POSTPRANDIAL_HYPERGLYCEMIA HP_CHRONIC_PAIN, HP_CHRONIC_PAIN HP_CALCIFIC_STIPPLING, HP_CALCIFIC_STIPPLING GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_1_ALPHA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_1_ALPHA_PRODUCTION GOMF_SPHINGOSINE_1_PHOSPHATE_RECEPTOR_ACTIVITY, GOMF_SPHINGOSINE_1_PHOSPHATE_RECEPTOR_ACTIVITY HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_UP, HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_UP GOMF_CHLORIDE_ION_BINDING, GOMF_CHLORIDE_ION_BINDING GOBP_SEROTONIN_SECRETION, GOBP_SEROTONIN_SECRETION GOBP_ENDOPLASMIC_RETICULUM_LOCALIZATION, GOBP_ENDOPLASMIC_RETICULUM_LOCALIZATION GOBP_UTERUS_DEVELOPMENT, GOBP_UTERUS_DEVELOPMENT HP_NEONATAL_SHORT_LIMB_SHORT_STATURE, HP_NEONATAL_SHORT_LIMB_SHORT_STATURE GOBP_LENS_FIBER_CELL_DIFFERENTIATION, GOBP_LENS_FIBER_CELL_DIFFERENTIATION GOBP_INTERLEUKIN_1_ALPHA_PRODUCTION, GOBP_INTERLEUKIN_1_ALPHA_PRODUCTION HP_DELAYED_ERUPTION_OF_PERMANENT_TEETH, HP_DELAYED_ERUPTION_OF_PERMANENT_TEETH

HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD, HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD

REACTOME_TRANSPORT_OF_FATTY_ACIDS, REACTOME_TRANSPORT_OF_FATTY_ACIDS

GOBP_C_TERMINAL_PROTEIN_LIPIDATION, GOBP_C_TERMINAL_PROTEIN_LIPIDATION

GOBP_CARDIOLIPIN_BIOSYNTHETIC_PROCESS, GOBP_CARDIOLIPIN_BIOSYNTHETIC_PROCESS

GOBP_REGULATION_OF_DEACETYLASE_ACTIVITY, GOBP_REGULATION_OF_DEACETYLASE_ACTIVITY

GOMF_CARNITINE_O_ACYLTRANSFERASE_ACTIVITY, GOMF_CARNITINE_O_ACYLTRANSFERASE_ACTIVITY

GOBP_REGULATION_OF_ATP_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_ATP_BIOSYNTHETIC_PROCESS

HP_ABSENT_OR_MINIMALLY_OSSIFIED_VERTEBRAL_BODIES, HP_ABSENT_OR_MINIMALLY_OSSIFIED_VERTEBRAL_BODIES

HP_LETHAL_SKELETAL_DYSPLASIA, HP_LETHAL_SKELETAL_DYSPLASIA

HP_EEG_WITH_FOCAL_SPIKE_WAVES, HP_EEG_WITH_FOCAL_SPIKE_WAVES

GOBP_OSTEOBLAST_DEVELOPMENT, GOBP_OSTEOBLAST_DEVELOPMENT

HAY_BONE_MARROW_CD34_POS_MEP, HAY_BONE_MARROW_CD34_POS_MEP

HP_THICK_CEREBRAL_CORTEX, HP_THICK_CEREBRAL_CORTEX

REACTOME_SYNTHESIS_OF_PG, REACTOME_SYNTHESIS_OF_PG

HP_TRICHORRHEXIS_NODOSA, HP_TRICHORRHEXIS_NODOSA

HP_THYROID_DYSGENESIS, HP_THYROID_DYSGENESIS

HP_OPTIC_NERVE_DYSPLASIA, HP_OPTIC_NERVE_DYSPLASIA

GOBP_UTERUS_MORPHOGENESIS, GOBP_UTERUS_MORPHOGENESIS

MIR1226_5P, MIR1226_5P

ROAZ_01, ROAZ_01

chr10q26, chr10q26