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|  | GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP   |
|  | GSE21033_CTRL_VS_POLYIC_STIM_DC_3H_DN, GSE21033_CTRL_VS_POLYIC_STIM_DC_3H_DN   |
|  | GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP   |
|  | GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP   |
|  | GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP   |
|  | GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP   |
|  | GSE14000_UNSTIM_VS_4H_LPS_DC_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_UP   |
|  | GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP   |
|  | GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP   |
|  | GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_DN, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_DN   |
|  | GSE41176_UNSTIM_VS_ANTL_IGM_STIM_BCELL_3H_DN, GSE41176_UNSTIM_VS_ANTL_IGM_STIM_BCELL_3H_DN   |
|  | GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP   |
|  | GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_DN, GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_DN   |
|  | WP_DNA_REPAIR_PATHWAYS_FULL_NETWORK, WP_DNA_REPAIR_PATHWAYS_FULL_NETWORK   |
|  | GSE1432_CTRL_VS_IFNG_6H_MICROGLIA_UP, GSE1432_CTRL_VS_IFNG_6H_MICROGLIA_UP   |
|  | GSE2706_UNSTIM_VS_2H_R848_DC_UP, GSE2706_UNSTIM_VS_2H_R848_DC_UP   |
|  | GSE42021_CD24HL_VS_CD24LOW_TREG_THYMUS_UP, GSE42021_CD24HL_VS_CD24LOW_TREG_THYMUS_UP   |
|  | MORF_PP2SC, MORF_PP2SC   |
|  | GSE37534_UNTREATED_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3 TRASDUCED_UP, GSE37534_UNTREATED_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3 TRASDUCED_UP                             |
|  | CSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_DN, GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_DN   |
|  | MIR6501_3P, MIR6501_3P   |
|  | WAKASUGI_HAVE_ZNF143_BINDING_SITES, WAKASUGI_HAVE_ZNF143_BINDING_SITES   |
|  | GSE45365_WT_VS_IFNAR_KO_BCELL_MCMV_INFECTION_UP, GSE45365_WT_VS_IFNAR_KO_BCELL_MCMV_INFECTION_UP   |
|  | HP_NEOPLASM_OF_THE_PANCREAS, HP_NEOPLASM_OF_THE_PANCREAS   |
|  | MIR6716_5P, MIR6716_5P   |
|  | MIR1304_5P, MIR1304_5P   |
|  | HP_NEOPLASM_OF_THE_BREAST, HP_NEOPLASM_OF_THE_BREAST   |
|  | GOBP_MATURATION_OF_5_8S_RRNA, GOBP_MATURATION_OF_5_8S_RRNA   |
|  | GSE21546_WT_VS_SAPIA_KO_DP_THYMOCYTES_DN, GSE21546_WT_VS_SAPIA_KO_DP_THYMOCYTES_DN   |
|  | HP_NEOPLASM_OF_THE_PITUITARY_GLAND, HP_NEOPLASM_OF_THE_PITUITARY_GLAND   |
|  | GSE19401_PAM2CSK4_VS_RETINOIC_ACID_STIM_FOLLICULAR_DC_DN, GSE19401_PAM2CSK4_VS_RETINOIC_ACID_STIM_FOLLICULAR_DC_DN   |
|  | MIR550A_3P, MIR550A_3P   |
|  | REACTOME_RESOLUTION_OF_ABASIC_SITES_AP_SITES, REACTOME_RESOLUTION_OF_ABASIC_SITES_AP_SITES   |
|  | MIR3922_3P, MIR3922_3P   |
|  | HP_BREAST_CARCINOMA, HP_BREAST_CARCINOMA   |
|  | HP_MALIGNANT_NEOPLASM_OF_THE_CENTRAL_NERVOUS_SYSTEM, HP_MALIGNANT_NEOPLASM_OF_THE_CENTRAL_NERVOUS_SYSTEM   |
|  | HP_NEOPLASM_OF_THE_RECTUM, HP_NEOPLASM_OF_THE_RECTUM   |
|  | REACTOME_RESOLUTION_OF_AP_SITES_VIA_THE_MULTIPLE_NUCLEOTIDE_PATCH_REPLACEMENT_PATHWAY, REACTOME_RESOLUTION_OF_AP_SITES_VIA_THE_MULTIPLE_NUCLEOTIDE_PATCH_REPLACEMENT_PATHWAY                         |
|  | HP_DYSGRAPHIA, HP_DYSGRAPHIA   |
|  | GOBP_TELOMERE_MAINTENANCE_VIA_SEMI_CONSERVATIVE_REPLICATION, GOBP_TELOMERE_MAINTENANCE_VIA_SEMI_CONSERVATIVE_REPLICATION   |
|  | HP_BASAL_CELL_CARCINOMA, HP_BASAL_CELL_CARCINOMA   |
|  | MIR3176, MIR3176   |
|  | GOBP_BASE_EXCISION_REPAIR, GOBP_BASE_EXCISION_REPAIR   |
|  | SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5, SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5   |
|  | HP_PITUITARY_ADENOMA, HP_PITUITARY_ADENOMA   |
|  | HP_PARESTHESIA, HP_PARESTHESIA   |
|  | HP_GONADAL_NEOPLASM, HP_GONADAL_NEOPLASM   |
|  | WP_BASE_EXCISION_REPAIR, WP_BASE_EXCISION_REPAIR   |
|  | MIR3188, MIR3188   |
|  | GOBP_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_15U_RRNA, GOBP_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_15U_RRNA               |
|  | KEGG_BASE_EXCISION_REPAIR, KEGG_BASE_EXCISION_REPAIR   |
|  | GOBP_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA, GOBP_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA   |
|  | MIR6774_5P, MIR6774_5P   |
|  | GOMF_ENDODEOXYRIBONUCLEASE_ACTIVITY, GOMF_ENDODEOXYRIBONUCLEASE_ACTIVITY   |
|  | HP_POLYCYSTIC_KIDNEY_DYSPLASIA, HP_POLYCYSTIC_KIDNEY_DYSPLASIA   |
|  | HP_GLIOMA, HP_GLIOMA   |
|  | GOCC_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR, GOCC_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR   |
|  | HP_NEOPLASM_OF_THE_RESPIRATORY_SYSTEM, HP_NEOPLASM_OF_THE_RESPIRATORY_SYSTEM   |
|  | MIR635, MIR635   |
|  | HP_TRUNCAL_ATAXIA, HP_TRUNCAL_ATAXIA   |
|  | GOBP_CENTRIOLE_ASSEMBLY, GOBP_CENTRIOLE_ASSEMBLY   |
|  | HP_AGNOSIA, HP_AGNOSIA   |
|  | HP_MIGRAINE, HP_MIGRAINE   |
|  | GOMF_REGULATORY_RNA_BINDING, GOMF_REGULATORY_RNA_BINDING   |
|  | MIR5695, MIR5695   |
|  | PAX8_B, PAX8_B   |
|  | HP_OVARIAN_NEOPLASM, HP_OVARIAN_NEOPLASM   |
|  | HP_PRIMITIVE_NEUROECTODERMAL_TUMOR, HP_PRIMITIVE_NEUROECTODERMAL_TUMOR   |
|  | GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN   |
|  | ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_DN, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_DN   |
|  | GOBP_MITOTIC_RECOMBINATION, GOBP_MITOTIC_RECOMBINATION   |
|  | GOBP_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION, GOBP_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION   |
|  | KEGG_LONG_TERM_POTENTIATION, KEGG_LONG_TERM_POTENTIATION   |
|  | GOBP_REGULATION_OF_TELOMERE_CAPPING, GOBP_REGULATION_OF_TELOMERE_CAPPING   |
|  | GOCC_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR, GOCC_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR   |
|  | HP_AMAUROSIS_FUGAX, HP_AMAUROSIS_FUGAX   |
|  | MIR5581_5P, MIR5581_5P   |
|  | MODULE_285, MODULE_285   |
|  | GOMF_MISMATCHED_DNA_BINDING, GOMF_MISMATCHED_DNA_BINDING   |
|  | HP_DECREASED_LACRIMATION, HP_DECREASED_LACRIMATION   |
|  | GOBP_REGULATION_OF_CENTRIOLE_REPLICATION, GOBP_REGULATION_OF_CENTRIOLE_REPLICATION   |
|  | HP_ASTROCYTOMA, HP_ASTROCYTOMA   |
|  | HP_ESOPHAGITIS, HP_ESOPHAGITIS   |
|  | MORF_JAG1, MORF_JAG1   |
|  | BHAT_ESR1_TARGETS_NOT_VIA_AKT1_DN, BHAT_ESR1_TARGETS_NOT_VIA_AKT1_DN   |
|  | RYAAAKNNNNNNNTTGW_UNKNOWN, RYAAAKNNNNNNNTTGW_UNKNOWN   |
|  | GOCC_NUCLEAR_EXOSOME_RNASE_COMPLEX, GOCC_NUCLEAR_EXOSOME_RNASE_COMPLEX   |
|  | HP_PANCREATIC_ADENOCARCINOMA, HP_PANCREATIC_ADENOCARCINOMA   |
|  | GOMF_EXODEOXYRIBONUCLEASE_ACTIVITY, GOMF_EXODEOXYRIBONUCLEASE_ACTIVITY   |
|  | HP_PROSTATE_NEOPLASM, HP_PROSTATE_NEOPLASM   |
|  | MODULE_366, MODULE_366   |
|  | HP_ABNORMALITY_OF_THE_ENDOMETRIUM, HP_ABNORMALITY_OF_THE_ENDOMETRIUM   |
|  | HP_CEREBRAL_WHITE_MATTER_ATROPHY, HP_CEREBRAL_WHITE_MATTER_ATROPHY   |
|  | MYOGNF1_01, MYOGNF1_01   |
|  | HP_DEATH_IN_EARLY_ADULTHOOD, HP_DEATH_IN_EARLY_ADULTHOOD   |
|  | GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS_VIA_SOMATIC_MUTATION, GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS_VIA_SOMATIC_MUTATION   |
|  | GOMF_HYDROLASE_ACTIVITY_HYDROLYZING_N_GLYCOSYL_COMPOUNDS, GOMF_HYDROLASE_ACTIVITY_HYDROLYZING_N_GLYCOSYL_COMPOUNDS   |
|  | GOMF_FOUR_WAY_JUNCTION_DNA_BINDING, GOMF_FOUR_WAY_JUNCTION_DNA_BINDING   |
|  | GOMF_OXIDIZED_DNA_BINDING, GOMF_OXIDIZED_DNA_BINDING   |
|  | MIR1178_5P, MIR1178_5P   |
|  | HP_ANGIOFIBROMAS, HP_ANGIOFIBROMAS   |
|  | HP_ACIDEMIA, HP_ACIDEMIA   |
|  | WP_4HYDROXYTAMOXIFEN_DEXAMETHASONE_AND_RETINOIC_ACIDS_REGULATION_OF_P27_EXPRESSION, WP_4HYDROXYTAMOXIFEN_DEXAMETHASONE_AND_RETINOIC_ACIDS_REGULATION_OF_P27_EXPRESSION                               |
|  | GOBP_MITOCHONDRIAL_RNA_PROCESSING, GOBP_MITOCHONDRIAL_RNA_PROCESSING   |
|  | GOBP_TRNA_WOBBLE_BASE_MODIFICATION, GOBP_TRNA_WOBBLE_BASE_MODIFICATION   |
|  | HP_INCREASED_CIRCULATING_PROLACTIN_CONCENTRATION, HP_INCREASED_CIRCULATING_PROLACTIN_CONCENTRATION   |
|  | HP_CARDIAC_DIVERTICULUM, HP_CARDIAC_DIVERTICULUM   |
|  | GOBP_PEROXISOMAL_MEMBRANE_TRANSPORT, GOBP_PEROXISOMAL_MEMBRANE_TRANSPORT   |
|  | HP_SPASTIC_DYSARTHRIA, HP_SPASTIC_DYSARTHRIA   |
|  | REACTOME_SYNTHESIS_OF_PIP3_AT_THE_EARLY_ENDOSOME_MEMBRANE, REACTOME_SYNTHESIS_OF_PIP3_AT_THE_EARLY_ENDOSOME_MEMBRANE   |
|  | GOBP_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY   |
|  | GOMF_GLUCOSYLTRANSFERASE_ACTIVITY, GOMF_GLUCOSYLTRANSFERASE_ACTIVITY   |
|  | PETRETTO_HEART_MASS_QTL_CIS_DN, PETRETTO_HEART_MASS_QTL_CIS_DN   |
|  | WANG_RECURRENT_LIVER_CANCER_UP, WANG_RECURRENT_LIVER_CANCER_UP   |
|  | WP_CAMKK2_PATHWAY, WP_CAMKK2_PATHWAY   |
|  | HP_ADENOMA_SEBACEUM, HP_ADENOMA_SEBACEUM   |
|  | HP_ABNORMALITY_OF_THE_PERITONEUM, HP_ABNORMALITY_OF_THE_PERITONEUM   |
|  | GOBP_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS, GOBP_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS   |
|  | WILLERT_WNT_SIGNALING, WILLERT_WNT_SIGNALING   |
|  | MIR548AD_3P, MIR548AD_3P   |
|  | GOMF_MISMATCH_REPAIR_COMPLEX_BINDING, GOMF_MISMATCH_REPAIR_COMPLEX_BINDING   |
|  | HP_GLIOMASTOMA_MULTIFORME, HP_GLIOMASTOMA_MULTIFORME   |
|  | GOBP_TELOMERE_MAINTENANCE_VIA_RECOMBINATION, GOBP_TELOMERE_MAINTENANCE_VIA_RECOMBINATION   |
|  | HP_ALACRIMA, HP_ALACRIMA   |
|  | GOCC_MISMATCH_REPAIR_COMPLEX, GOCC_MISMATCH_REPAIR_COMPLEX   |
|  | HP_NON_HODGKIN_LYMPHOMA, HP_NON_HODGKIN_LYMPHOMA   |
|  | MIR5684, MIR5684   |
|  | NIKOLSKY_BREAST_CANCER_12Q24_AMPLICON, NIKOLSKY_BREAST_CANCER_12Q24_AMPLICON   |
|  | GOBP_DETERMINATION_OF_ADULT_LIFESPAN, GOBP_DETERMINATION_OF_ADULT_LIFESPAN   |
|  | HP_STRIAE_DISTENSAE, HP_STRIAE_DISTENSAE   |
|  | GOMF_LYSINE_ACETYLATED_HISTONE_BINDING, GOMF_LYSINE_ACETYLATED_HISTONE_BINDING   |
|  | GOBP_POSITIVE_REGULATION_OF_P38MAPK_CASCADE, GOBP_POSITIVE_REGULATION_OF_P38MAPK_CASCADE   |
|  | GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION   |
|  | MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_8_MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_8  |
|  | GOBP_PYRIMIDINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS, GOBP_PYRIMIDINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS   |
|  | GOBP_PPTIDYL_GLUTAMIC_ACID_MODIFICATION, GOBP_PPTIDYL_GLUTAMIC_ACID_MODIFICATION   |
|  | GOMF_UBIQUITIN_UBIQUITIN_LIGASE_ACTIVITY, GOMF_UBIQUITIN_UBIQUITIN_LIGASE_ACTIVITY   |
|  | GOBP_PYRIMIDINE_NUCLEOTIDE_CATABOLIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOTIDE_CATABOLIC_PROCESS   |
|  | GOBP_HISTONE_H3_K36_DEMETHYLATION, GOBP_HISTONE_H3_K36_DEMETHYLATION   |
|  | GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K36_SPECIFIC, GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K36_SPECIFIC   |
|  | HP_HYPOCHROMIC_ANEMIA, HP_HYPOCHROMIC_ANEMIA   |
|  | LIU_TARGETS_OF_VMYB_VS_CMYB_DN, LIU_TARGETS_OF_VMYB_VS_CMYB_DN   |
|  | GOBP_PROGESTERONE_RECEPTOR_SIGNALING_PATHWAY, GOBP_PROGESTERONE_RECEPTOR_SIGNALING_PATHWAY   |
|  | GOBP_MITOCHONDRIAL_RNA_MODIFICATION, GOBP_MITOCHONDRIAL_RNA_MODIFICATION   |
|  | HP_NEPHROGENIC_DIABETES_INSIPIDUS, HP_NEPHROGENIC_DIABETES_INSIPIDUS   |
|  | GOMF_DEOXYRIBONUCLEOTIDE_BINDING, GOMF_DEOXYRIBONUCLEOTIDE_BINDING   |
|  | GOBP_MITOCHONDRIAL_TRNA_PROCESSING, GOBP_MITOCHONDRIAL_TRNA_PROCESSING   |
|  | VALK_AML_CLUSTER_13, VALK_AML_CLUSTER_13   |
|  | MIR596, MIR596   |
|  | GOBP_ACTIVATION_INDUCED_CELL_DEATH_OF_T_CELLS, GOBP_ACTIVATION_INDUCED_CELL_DEATH_OF_T_CELLS   |
|  | GOBP_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS, GOBP_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS   |
|  | GOMF_PROTEIN_TYROSINE_THREONINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_TYROSINE_THREONINE_PHOSPHATASE_ACTIVITY   |
|  | HP_PROGRESSIVE_ENCEPHALOPATHY, HP_PROGRESSIVE_ENCEPHALOPATHY   |
|  | GOBP_TELOMERIC_D_LOOP_DISASSEMBLY, GOBP_TELOMERIC_D_LOOP_DISASSEMBLY   |
|  | REACTOME_DISEASES_OF_DNA_REPAIR, REACTOME_DISEASES_OF_DNA_REPAIR   |
|  | GOBP_CONSTITUTIVE_SECRETORY_PATHWAY, GOBP_CONSTITUTIVE_SECRETORY_PATHWAY   |
|  | GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAGENTA_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAGENTA_DN   |
|  | GOBP_MULTICELLULAR_ORGANISM_AGING, GOBP_MULTICELLULAR_ORGANISM_AGING   |
|  | HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II, HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II   |
|  | GOBP_MEIOTIC_MISMATCH_REPAIR, GOBP_MEIOTIC_MISMATCH_REPAIR   |
|  | HP_CHYLOTHORAX, HP_CHYLOTHORAX   |
|  | GOMF_MAP_KINASE_KINASE_ACTIVITY, GOMF_MAP_KINASE_KINASE_ACTIVITY   |
|  | KOBAYASHI_EGFR_SIGNALING_6HR_UP, KOBAYASHI_EGFR_SIGNALING_6HR_UP   |
|  | HP_MULTIPLE_ROWS_OF_EYELASHES, HP_MULTIPLE_ROWS_OF_EYELASHES   |
|  | MIR4781_3P, MIR4781_3P   |
|  | GOBP_CHONDROCYTE_DEVELOPMENT_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS, GOBP_CHONDROCYTE_DEVELOPMENT_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS   |
|  | REACTOME_COMPETING_ENDOGENOUS_RNAS_CERNAS_REGULATE_PTEEN_TRANSLATION, REACTOME_COMPETING_ENDOGENOUS_RNAS_CERNAS_REGULATE_PTEEN_TRANSLATION   |
|  | HP_LIPOMA, HP_LIPOMA   |
|  | HP_ABNORMALITY_OF_THE_SUBUNGUAL_REGION, HP_ABNORMALITY_OF_THE_SUBUNGUAL_REGION   |
|  | BOYAUULT_LIVER_CANCER_SUBCLASS_G56_UP, BOYAUULT_LIVER_CANCER_SUBCLASS_G56_UP   |
|  | YAMANAKA_GLIOMASTOMA_SURVIVAL_UP, YAMANAKA_GLIOMASTOMA_SURVIVAL_UP   |
|  | GOBP_NEGATIVE_REGULATION_OF_RNA_POLYMERASE_II_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOBP_NEGATIVE_REGULATION_OF_RNA_POLYMERASE_II_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING         |
|  | HP_ABNORMAL_LYMPHATIC_VESSEL_MORPHOLOGY, HP_ABNORMAL_LYMPHATIC_VESSEL_MORPHOLOGY   |
|  | GSE12839_CTRL_VS_IL12_TREATED_PBMCDN, GSE12839_CTRL_VS_IL12_TREATED_PBMCDN   |
|  | GOBP_PROTEIN_HEXAMERIZATION, GOBP_PROTEIN_HEXAMERIZATION   |
|  | GOBP_PROTEIN_DEGLUTAMYLATION, GOBP_PROTEIN_DEGLUTAMYLATION   |
|  | GOBP_NEGATIVE_REGULATION_OF_PROTEIN_K63_LINKED_UBIQUITINATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_K63_LINKED_UBIQUITINATION   |
|  | GOMF_DNA_N_GLYCOSYLASE_ACTIVITY, GOMF_DNA_N_GLYCOSYLASE_ACTIVITY   |
|  | BIOCARTA_CPSF_PATHWAY, BIOCARTA_CPSF_PATHWAY   |
|  | GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_HEME_PROTEIN_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_HEME_PROTEIN_AS_ACCEPTOR   |
|  | GOBP_ARYL_HYDROCARBON_RECEPTOR_BINDING, GOMF_ARYL_HYDROCARBON_RECEPTOR_BINDING   |
|  | REACTOME_REGULATION_OF_PTEEN_MRNA_TRANSLATION, REACTOME_REGULATION_OF_PTEEN_MRNA_TRANSLATION   |
|  | HP_MICROCYTIC_ANEMIA, HP_MICROCYTIC_ANEMIA   |
|  | GOBP_REGULATION_OF_NITRIC_OXIDE_SYNTHASE_ACTIVITY, GOBP_REGULATION_OF_NITRIC_OXIDE_SYNTHASE_ACTIVITY   |
|  | HP_DOUBLE_AORTIC_ARCH, HP_DOUBLE_AORTIC_ARCH   |
|  | GOBP_DE_NOVO_IMP_BIOSYNTHETIC_PROCESS, GOBP_DE_NOVO_IMP_BIOSYNTHETIC_PROCESS   |
|  | GOMF_RNA_POLYMERASE_I_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_I_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY   |
|  | HP_HEMIMEGALENCEPHALY, HP_HEMIMEGALENCEPHALY   |
|  | HP_ABNORMAL_SALIVARY_GLAND_MORPHOLOGY, HP_ABNORMAL_SALIVARY_GLAND_MORPHOLOGY   |
|  | GOBP_POSITIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION   |
|  | REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION, REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION   |
|  | GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MATRIX, GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MATRIX   |
|  | HP_NEOPLASM_OF_THE_SMALL_INTESTINE, HP_NEOPLASM_OF_THE_SMALL_INTESTINE   |
|  | GOMF_UDP_XYLOSYLTRANSFERASE_ACTIVITY, GOMF_UDP_XYLOSYLTRANSFERASE_ACTIVITY   |
|  | GOBP_FATTY_ACID_BETA_OXIDATION_USING_ACYL_COA_DEHYDROGENASE, GOBP_FATTY_ACID_BETA_OXIDATION_USING_ACYL_COA_DEHYDROGENASE   |
|  | GOMF_NADH_PYROPHOSPHATASE_ACTIVITY, GOMF_NADH_PYROPHOSPHATASE_ACTIVITY   |
|  | HP_TALIPES_CALCANEVALGUS, HP_TALIPES_CALCANEVALGUS   |
|  | HP_THICKENED_EARS, HP_THICKENED_EARS   |
|  | GOBP_REGULATION_OF_NEUROTRANSMITTER_RECEPTOR_LOCALIZATION_TO_POSTSYNAPTIC_SPECIALIZATION_MEMBRANE, GOBP_REGULATION_OF_NEUROTRANSMITTER_RECEPTOR_LOCALIZATION_TO_POSTSYNAPTIC_SPECIALIZATION_MEMBRANE |
|  | GOMF_INTERLEUKIN_1_RECEPTOR_ACTIVITY, GOMF_INTERLEUKIN_1_RECEPTOR_ACTIVITY   |
|  | REACTOME_SIGNALING_BY_MAPK_MUTANTS, REACTOME_SIGNALING_BY_MAPK_MUTANTS   |
|  | HP_CAESARIAN_SECTION, HP_CAESARIAN_SECTION   |
|  | GOBP_ACTIN_POLYMERIZATION_DEPENDENT_CELL_MOTILITY, GOBP_ACTIN_POLYMERIZATION_DEPENDENT_CELL_MOTILITY   |
|  | GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING, GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING   |
|  | GOCC_SHELTERIN_COMPLEX, GOCC_SHELTERIN_COMPLEX   |
|  | GOMF_CALCIIUM_DEPENDENT_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY, GOMF_CALCIIUM_DEPENDENT_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY   |
|  | GOBP_POSITIVE_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE   |
|  | GOBP_MITOCHONDRIAL_DNA_REPAIR, GOBP_MITOCHONDRIAL_DNA_REPAIR   |
|  | HP_MYELOPATHY, HP_MYELOPATHY   |
|  | WP_NOCGMPKKG_MEDIATED_NEUROPROTECTION, WP_NOCGMPKKG_MEDIATED_NEUROPROTECTION   |
|  | GOBP_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE   |
|  | HP_INCREASED_BLOOD_UREA_NITROGEN, HP_INCREASED_BLOOD_UREA_NITROGEN   |
|  | KIM_MYCL1_AMPLIFICATION_TARGETS_UP, KIM_MYCL1_AMPLIFICATION_TARGETS_UP   |
|  | HOECKERKORP_CD44_TARGETS_DIRECT_DN, HOECKERKORP_CD44_TARGETS_DIRECT_DN   |
|  | GOBP_MEMBRANOUS_SEPTUM_MORPHOGENESIS, GOBP_MEMBRANOUS_SEPTUM_MORPHOGENESIS   |
|  | HP_ABNORMALITY_OF_CHORIORETINAL_PIGMENTATION, HP_ABNORMALITY_OF_CHORIORETINAL_PIGMENTATION   |
|  | GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATASE_ACTIVITY   |
|  | GOMF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_BINDING, GOMF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_BINDING   |