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| | GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP, GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP, GSE32986_CURDLAN_LOWDOSE_VS_CURDLAN_HIGHDOSE_STIM_DC_UP, GSE32986_CURDLAN_LOWDOSE_VS_CURDLAN_HIGHDOSE_STIM_DC_UP, GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_UP, GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_UP, GSE369_IFNG_KO_VS_WT_LIVER_UP, GSE369_IFNG_KO_VS_WT_LIVER_UP, GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP, GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP, GSE45365_NK_CELL_VS_CD11B_DC_MCMV_INFECTION_DN, GSE45365_NK_CELL_VS_CD11B_DC_MCMV_INFECTION_DN, GSE411_UNSTIM_VS_100MIN_IL6_STIM_S0CS3_KO_MACROPHAGE_UP, GSE411_UNSTIM_VS_100MIN_IL6_STIM_S0CS3_KO_MACROPHAGE_UP, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_UP, GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_UP, GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_DN, GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_DN, GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UP, GSE27786_NKCELL_VS_NEUTROPHIL_DN, GSE27786_NKCELL_VS_NEUTROPHIL_DN, GSE12845_ICD_NEG_BLOOD_VS_NAIVE_TONSI_BCELL_DN, GSE12845_ICD_NEG_BLOOD_VS_NAIVE_TONSI_BCELL_DN, MIR525_5P, MIR525_5P, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_UP, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_UP, MIR520A_5P, MIR520A_5P, GSE27786_NKCELL_VS_MONO_MAC_DN, GSE27786_NKCELL_VS_MONO_MAC_DN, MIR3663_3P, MIR3663_3P, ZHENG_CORD_BLOOD_C10_MULTILYMPHOID_PROGENITOR, ZHENG_CORD_BLOOD_C10_MULTILYMPHOID_PROGENITOR, MIR623, MIR623, WP_CHEMOKINE_SIGNALING_PATHWAY, WP_CHEMOKINE_SIGNALING_PATHWAY, MIR922, MIR922, WP_FRAGILE_X_SYNDROME, WP_FRAGILE_X_SYNDROME, HP_LOWER_LIMB_MUSCLE_WEAKNESS, HP_LOWER_LIMB_MUSCLE_WEAKNESS, MIR1179, MIR1179, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_BCELL_UP, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_BCELL_UP, PID_AR_PATHWAY, PID_AR_PATHWAY, KEGG_CHEMOKINE_SIGNALING_PATHWAY, KEGG_CHEMOKINE_SIGNALING_PATHWAY, BASSO_CD40_SIGNALING_DN, BASSO_CD40_SIGNALING_DN, GSE22886_TH1_VS_TH2_12H_ACT_DN, GSE22886_TH1_VS_TH2_12H_ACT_DN, GSE13411_NAIVE_VS_ICM_MEMORY_BCELL_UP, GSE13411_NAIVE_VS_ICM_MEMORY_BCELL_UP, HP_MITRAL_VALVE_PROLAPSE, HP_MITRAL_VALVE_PROLAPSE, WAMUNYOKOLI_OVARIAN_CANCER_GRADES_1_2_UP, WAMUNYOKOLI_OVARIAN_CANCER_GRADES_1_2_UP, HP_ABNORMAL_MITRAL_VALVE_MORPHOLOGY, HP_ABNORMAL_MITRAL_VALVE_MORPHOLOGY, MIR17_3P, MIR17_3P, GOCC_CELL_DIVISION_SITE, GOCC_CELL_DIVISION_SITE, MIR4697_3P, MIR4697_3P, HP_BROAD_BASED_GAIT, HP_BROAD_BASED_GAIT, MIR4299, MIR4299, AR_Q2, AR_Q2, GSE21063_CTRL_VS_ANTL_ICM_STIM_BCELL_8H_UP, GSE21063_CTRL_VS_ANTL_ICM_STIM_BCELL_8H_UP, PID_BCR_PATHWAY, PID_BCR_PATHWAY, MIR4526, MIR4526, MIR3660, MIR3660, MIR4463, MIR4463, AGTCAGC_MIR345, AGTCAGC_MIR345, PID_FAK_PATHWAY, PID_FAK_PATHWAY, MIR7156_3P, MIR7156_3P, GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP, GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP, HP_LIMITED_ELBOW_MOVEMENT, HP_LIMITED_ELBOW_MOVEMENT, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_UP, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_UP, REACTOME_DDX58_IFIH1_MEDIATED_INDUCTION_OF_INTERFERON_ALPHA_BETA, REACTOME_DDX58_IFIH1_MEDIATED_INDUCTION_OF_INTERFERON_ALPHA_BETA, GSE1112_OT1_CD8A8_VS_HY_CD8AA_THYMOCYTE_RT0C_CULTURE_DN, GSE1112_OT1_CD8A8_VS_HY_CD8AA_THYMOCYTE_RT0C_CULTURE_DN, GAGCGTCG_MIR844, GAGCGTCG_MIR844, WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY, WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY, REACTOME_NEGATIVE_REGULATORS_OF_DDX58_IFIH1_SIGNALING, REACTOME_NEGATIVE_REGULATORS_OF_DDX58_IFIH1_SIGNALING, HP_ELBOW_FLEXION_CONTRACTURE, HP_ELBOW_FLEXION_CONTRACTURE, BCAT_BILD_ET_AL_DN, BCAT_BILD_ET_AL_DN, MIR4708_5P, MIR4708_5P, MIR3201, MIR3201, AR_Q2, AR_Q2, MIR683_3P, MIR683_3P, CHUNG_BLISTER_CYTOTOXICITY_DN, CHUNG_BLISTER_CYTOTOXICITY_DN, DCA_UP_V1_DN, DCA_UP_V1_DN, MIR550B_3P, MIR550B_3P, MIR3622A_3P, MIR3622B_3P, MIR3622A_3P, MIR3622B_3P, MIR1225_3P, MIR1225_3P, GSE20366_TREG_VS_TCONV_UP, GSE20366_TREG_VS_TCONV_UP, MIR1238_3P, MIR1238_3P, HP_PRIMITIVE_NEUROECTODERMAL_TUMOR, HP_PRIMITIVE_NEUROECTODERMAL_TUMOR, MIR3140_5P, MIR3140_5P, MIR4639_5P, MIR4639_5P, WEIGEL_OXIDATIVE_STRESS_RESPONSE, WEIGEL_OXIDATIVE_STRESS_RESPONSE, WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_INTERFERON_INDUCTION, WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_INTERFERON_INDUCTION, NIKOLSKY_BREAST_CANCER_6P24_P22_AMPICON, NIKOLSKY_BREAST_CANCER_6P24_P22_AMPICON, GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZLI_Y_STRAIN_INF_DN, GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZLI_Y_STRAIN_INF_DN, MIR4717_3P, MIR4717_3P, WP_SIGNALING_OF_HEPATOCYTE_GROWTH_FACTOR_RECEPTOR, WP_SIGNALING_OF_HEPATOCYTE_GROWTH_FACTOR_RECEPTOR, MIR640, MIR640, GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, AR_Q2, AR_Q2, PID_IL2_PATHWAY, PID_IL2_PATHWAY, BIOCARTA_MET_PATHWAY, BIOCARTA_MET_PATHWAY, HP_LANGUAGE_IMPAIRMENT, HP_LANGUAGE_IMPAIRMENT, MIR4791, MIR4791, MIR6818_5P, MIR6818_5P, HP_TRIGONOCEPHALY, HP_TRIGONOCEPHALY, MIR431_5P, MIR431_5P, GRADE_COLON_AND_RECTAL_CANCER_DN, GRADE_COLON_AND_RECTAL_CANCER_DN, MIR219A_2_3P, MIR219A_2_3P, SEITZ_NEOPLASTIC_TRANSFORMATION_BY_SP_DELETION_UP, SEITZ_NEOPLASTIC_TRANSFORMATION_BY_SP_DELETION_UP, HP_ABNORMAL_SPERMATOGENESIS, HP_ABNORMAL_SPERMATOGENESIS, MIR4633_3P, MIR6500_5P, MIR4633_3P, MIR6500_5P, HP_VASCULAR_TORTUOSITY, HP_VASCULAR_TORTUOSITY, WP_PDGF_PATHWAY, WP_PDGF_PATHWAY, WP_TYPE_1_INTERFERON_INDUCTION_AND_SIGNALING_DURING_SARSCOV2_INFECTION, WP_TYPE_1_INTERFERON_INDUCTION_AND_SIGNALING_DURING_SARSCOV2_INFECTION, MIR6745, MIR6745, HP_NEUROFIBRILLARY_TANGLES, HP_NEUROFIBRILLARY_TANGLES, BIOCARTA_CXCR4_PATHWAY, BIOCARTA_CXCR4_PATHWAY, REACTOME_TICAM1_DEPENDENT_ACTIVATION_OF_IRF3_IRF7, REACTOME_TICAM1_DEPENDENT_ACTIVATION_OF_IRF3_IRF7, MIR6508_3P, MIR6508_3P, MORF_PDPK1, MORF_PDPK1, GOBP_ANTEROGRADE_AXONAL_TRANSPORT, GOBP_ANTEROGRADE_AXONAL_TRANSPORT, MIR3099_3P, MIR3099_3P, GOBP_NEGATIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS, BAE_BRCA1_TARGETS_DN, BAE_BRCA1_TARGETS_DN, DASU_IL6_SIGNALING_SCAR_UP, DASU_IL6_SIGNALING_SCAR_UP, MIR6798_3P, MIR6798_3P, HP_PROTRUDING_TONGUE, HP_PROTRUDING_TONGUE, ZHONG_PFC_C7_ORG_UNDERGOING_NEURONAL_DIFFERENTIATION, ZHONG_PFC_C7_ORG_UNDERGOING_NEURONAL_DIFFERENTIATION, REACTOME_ACTIVATION_OF_IRF3_IRF7_MEDIATED_BY_TBK1_IKK_EPSILON, REACTOME_ACTIVATION_OF_IRF3_IRF7_MEDIATED_BY_TBK1_IKK_EPSILON, REACTOME_DAPI2_INTERACTIONS, REACTOME_DAPI2_INTERACTIONS, GOBP_BONE_REMODELING, GOBP_BONE_REMODELING, BIOCARTA_EGF_PATHWAY, BIOCARTA_EGF_PATHWAY, WP_EPO_RECEPTOR_SIGNALING, WP_EPO_RECEPTOR_SIGNALING, MIR6793_5P, MIR6793_5P, HP_ABNORMAL_PANCREAS_MORPHOLOGY, HP_ABNORMAL_PANCREAS_MORPHOLOGY, GOBP_CELLULAR_RESPONSE_TO_CALCIUM_ION, GOBP_CELLULAR_RESPONSE_TO_CALCIUM_ION, PID_RAS_PATHWAY, PID_RAS_PATHWAY, MORLEMU_MYC_LYMPHOMA_BY_ONSET_TIME_DN, MORLEMU_MYC_LYMPHOMA_BY_ONSET_TIME_DN, HP_LIMITED_HIP_MOVEMENT, HP_LIMITED_HIP_MOVEMENT, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP, SAGIV_CD24_TARGETS_UP, SAGIV_CD24_TARGETS_UP, GOBP_RESPONSE_TO_NERVE_GROWTH_FACTOR, GOBP_RESPONSE_TO_NERVE_GROWTH_FACTOR, GOMF_ACTININ_BINDING, GOMF_ACTININ_BINDING, GOCC_DENDRITIC_SHAFT, GOCC_DENDRITIC_SHAFT, GOMF_PHOSPHOTYROSINE_RESIDUE_BINDING, GOMF_PHOSPHOTYROSINE_RESIDUE_BINDING, NUNODA_RESPONSE_TO_DASATINIB_IMATINIB_DN, NUNODA_RESPONSE_TO_DASATINIB_IMATINIB_DN, HP_SUBARACHNOID_HEMORRHAGE, HP_SUBARACHNOID_HEMORRHAGE, PID_TCR_RAS_PATHWAY, PID_TCR_RAS_PATHWAY, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSPORT, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSPORT, HP_LIMITATION_OF_NECK_MOTION, HP_LIMITATION_OF_NECK_MOTION, GOBP_RESPONSE_TO_DIETARY_EXCESS, GOBP_RESPONSE_TO_DIETARY_EXCESS, GOBERT_CORE_OLIGODENDROCYTE_DIFFERENTIATION, GOBERT_CORE_OLIGODENDROCYTE_DIFFERENTIATION, HP_SYNOVITIS, HP_SYNOVITIS, GOCC_FILAMENTOUS_ACTIN, GOCC_FILAMENTOUS_ACTIN, HP_HIGH_HYPERMETROPIA, HP_HIGH_HYPERMETROPIA, GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, HP_CYSTIC_HYDROMA, HP_CYSTIC_HYDROMA, BIOCARTA_CDMAC_PATHWAY, BIOCARTA_CDMAC_PATHWAY, GOBP_REGULATION_OF_NEUTROPHIL_ACTIVATION, GOBP_REGULATION_OF_NEUTROPHIL_ACTIVATION, LIU_IL13_PRIMING_MODEL, LIU_IL13_PRIMING_MODEL, GOBP_TELOMERIC_LOOP_DISASSEMBLY, GOBP_TELOMERIC_LOOP_DISASSEMBLY, CHEOK_RESPONSE_TO_MERCAPTOPURINE_DN, CHEOK_RESPONSE_TO_MERCAPTOPURINE_DN, GSE43863_TH1_VS_TFH_EFFECTOR_CD4_TCELL_UP, GSE43863_TH1_VS_TFH_EFFECTOR_CD4_TCELL_UP, GOBP_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION, BIOCARTA_CD40_PATHWAY, BIOCARTA_CD40_PATHWAY, GOBP_GALACTOSE_CATABOLIC_PROCESS, GOBP_GALACTOSE_CATABOLIC_PROCESS, GOBP_CELLULAR_RESPONSE_TO_D5RNA, GOBP_CELLULAR_RESPONSE_TO_D5RNA, MIR8074, MIR8074, WP_MAPK_PATHWAY_IN_CONGENITAL_THYROID_CANCER, WP_MAPK_PATHWAY_IN_CONGENITAL_THYROID_CANCER, TIAN_BHLHA15_TARGETS, TIAN_BHLHA15_TARGETS, CUL1_TCF21_TARGETS_DN, CUL1_TCF21_TARGETS_DN, REACTOME_PTK6_REGULATES_RHO_GTPASES, REACTOME_PTK6_REGULATES_RHO_GTPASES, MIR911_3P, MIR911_3P, REACTOME_FRUCTOSE_METABOLISM, REACTOME_FRUCTOSE_METABOLISM, GOBP_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE, HP_UNDETECTABLE_ELECTRORETINOGRAM, HP_UNDETECTABLE_ELECTRORETINOGRAM, FLOTHO_PEDIATRIC_ALL_THERAPY_RESPONSE_DN, FLOTHO_PEDIATRIC_ALL_THERAPY_RESPONSE_DN, HP_LIVER_ABSCESS, HP_LIVER_ABSCESS, WP_TLR4_SIGNALING_AND_TOLERANCE, WP_TLR4_SIGNALING_AND_TOLERANCE, GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY, GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY, GOBP_STEM_CELL_DIVISION, GOBP_STEM_CELL_DIVISION, GOBP_CYTOPLASMIC_SEQUESTERING_OF_PROTEIN, GOBP_CYTOPLASMIC_SEQUESTERING_OF_PROTEIN, WP_SARS_CORONAVIRUS_AND_INNATE_IMMUNITY, WP_SARS_CORONAVIRUS_AND_INNATE_IMMUNITY, REACTOME_TRAF3_DEPENDENT_IRF_ACTIVATION_PATHWAY, REACTOME_TRAF3_DEPENDENT_IRF_ACTIVATION_PATHWAY, HP_ADVANCED_ERUPTION_OF_TEETH, HP_ADVANCED_ERUPTION_OF_TEETH, BIOCARTA_TNFR2_PATHWAY, BIOCARTA_TNFR2_PATHWAY, HP_PECTUS_EXCAVATUM_OF_INFERIOR_STERNUM, HP_PECTUS_EXCAVATUM_OF_INFERIOR_STERNUM, HP_REDUCED_FACTOR_XII_ACTIVITY, HP_REDUCED_FACTOR_XII_ACTIVITY, GOBP_HISTONE_H3_K4_DEMETHYLATION_TRIMETHYL_H3_K4_SPECIFIC, GOBP_HISTONE_H3_K4_DEMETHYLATION_TRIMETHYL_H3_K4_SPECIFIC, GOBP_AUTOCRINE_SIGNALING, GOBP_AUTOCRINE_SIGNALING, WP_CANONICAL_NFKB_PATHWAY, WP_CANONICAL_NFKB_PATHWAY, GOBP_GOLGI_TO_ENDOSOME_TRANSPORT, GOBP_GOLGI_TO_ENDOSOME_TRANSPORT, GOMF_PHOSPHATIDYLINOSITOL_3_4_BISPHOSPHATE_BINDING, GOMF_PHOSPHATIDYLINOSITOL_3_4_BISPHOSPHATE_BINDING, GOBP_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE, GOBP_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE, HP_HYPOGLYCORRHACHIA, HP_HYPOGLYCORRHACHIA, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_ENDOSOME, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_ENDOSOME, HP_IRIS_HYPOPIGMENTATION, HP_IRIS_HYPOPIGMENTATION, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS, HP_AUTOIMMUNE_THROMBOCYTOPENIA, HP_AUTOIMMUNE_THROMBOCYTOPENIA, WORSCHICH_TUMOR_EVASION_AND_TOLEROGENICITY_UP, WORSCHICH_TUMOR_EVASION_AND_TOLEROGENICITY_UP, CUL1_DEVELOPING_HEART_5TH_WEEK_VENTRICULAR_CARDIOMYOCYTE_CUL1_DEVELOPING_HEART_5TH_WEEK_VENTRICULAR_CARDIOMYOCYTE, TURASHVILL_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_DN, TURASHVILL_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_DN, HP_BROAD_METATARSAL, HP_BROAD_METATARSAL, GOBP_NEUROMUSCULAR_PROCESS, GOBP_NEUROMUSCULAR_PROCESS, GOBP_NEUROTROPHIN_TRK_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEUROTROPHIN_TRK_RECEPTOR_SIGNALING_PATHWAY, REACTOME_GALACTOSE_CATABOLISM, REACTOME_GALACTOSE_CATABOLISM, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME, ZN774_TARGET_GENECYCLING_TARGETS, ZN774_TARGET_GENECYCLING_TARGETS, GRANDVAUX_IRF3_TARGETS_DN, GRANDVAUX_IRF3_TARGETS_DN, MIR6865_3P, MIR6865_3P, CDP_Q1, CDP_Q1, ZHAN_LATE_DIFFERENTIATION_GENES_DN, ZHAN_LATE_DIFFERENTIATION_GENES_DN, GOBP_CYTOPLASMIC_SEQUESTERING_OF_TRANSCRIPTION_FACTOR, GOBP_CYTOPLASMIC_SEQUESTERING_OF_TRANSCRIPTION_FACTOR, WILLIAMS_ESR1_TARGETS_DN, WILLIAMS_ESR1_TARGETS_DN, HP_AMEGAKARYOCYTIC_THROMBOCYTOPENIA, HP_AMEGAKARYOCYTIC_THROMBOCYTOPENIA, GOCC_MULTIVESICULAR_BODY_MEMBRANE, GOCC_MULTIVESICULAR_BODY_MEMBRANE, REACTOME_INTEGRIN_SIGNALING, REACTOME_INTEGRIN_SIGNALING, REACTOME_NF_KB_IS_ACTIVATED_AND_SIGNALS_SURVIVAL, REACTOME_NF_KB_IS_ACTIVATED_AND_SIGNALS_SURVIVAL, HP_SCANNING_SPEECH, HP_SCANNING_SPEECH, GOBP_NEGATIVE_REGULATION_OF_NEURON_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_NEURON_DIFFERENTIATION, GOCC_GOLGI_LUMEN, GOCC_GOLGI_LUMEN, GOBP_CAMP_CATABOLIC_PROCESS, GOBP_CAMP_CATABOLIC_PROCESS, GOBP_SYNAPTIC_GROWTH_AT_NEUROMUSCULAR_JUNCTION, GOBP_SYNAPTIC_GROWTH_AT_NEUROMUSCULAR_JUNCTION, GOBP_HINDBRAIN_RADIAL_GLIA_GUIDED_CELL_MIGRATION, GOBP_HINDBRAIN_RADIAL_GLIA_GUIDED_CELL_MIGRATION, GOBP_RESPONSE_TO_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR, GOBP_RESPONSE_TO_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR, GOBP_REGULATION_OF_SPONTANEOUS_SYNAPTIC_TRANSMISSION, GOBP_REGULATION_OF_SPONTANEOUS_SYNAPTIC_TRANSMISSION, HP_APLASIA_HYPOPLASIA_OF_THE_EPIGLOTTIS, HP_APLASIA_HYPOPLASIA_OF_THE_EPIGLOTTIS, GOBP_ADIPONECTIN_ACTIVATED_SIGNALING_PATHWAY, GOBP_ADIPONECTIN_ACTIVATED_SIGNALING_PATHWAY, GOBP_RESPONSE_TO_GLYCOPROTEIN, GOBP_RESPONSE_TO_GLYCOPROTEIN, GOCC_PINKSOME, GOCC_PINKSOME, GOCC_PHOSPHOLIPID_TRANSLOCATING_ATPASE_COMPLEX, GOCC_PHOSPHOLIPID_TRANSLOCATING_ATPASE_COMPLEX, GOMF_MANNOSYL_OLIGOSACCHARIDE_1_2_ALPHA_MANNOSIDASE_ACTIVITY, GOMF_MANNOSYL_OLIGOSACCHARIDE_1_2_ALPHA_MANNOSIDASE_ACTIVITY, HP_DECREASED_LEVEL_OF_GABA_IN_SERUM, HP_DECREASED_LEVEL_OF_GABA_IN_SERUM, HP_DIFFUSE_PALMOPLANTAR_KERATODERMA, HP_DIFFUSE_PALMOPLANTAR_KERATODERMA, WP_ESTROGEN_SIGNALING_PATHWAY, WP_ESTROGEN_SIGNALING_PATHWAY, HP_SCALING_SKIN, HP_SCALING_SKIN, GOCC_POSTSYNAPTIC_ENDOCYTIC_ZONE, GOCC_POSTSYNAPTIC_ENDOCYTIC_ZONE, BIOCARTA_SPRY_PATHWAY, BIOCARTA_SPRY_PATHWAY, MORF_FLT1, MORF_FLT1, GOBP_PURINE_CONTAINING_COMPOUND_SALVAGE, GOBP_PURINE_CONTAINING_COMPOUND_SALVAGE, GOBP_SUCKLING_BEHAVIOR, GOBP_SUCKLING_BEHAVIOR, GOBP_LOOP_OF_HENLE_DEVELOPMENT, GOBP_LOOP_OF_HENLE_DEVELOPMENT, GOBP_RECEPTOR_RECYCLING, GOBP_RECEPTOR_RECYCLING, GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, HP_REDUCED_FACTOR_XIII_ACTIVITY, HP_REDUCED_FACTOR_XIII_ACTIVITY, GOMF_SUMO_POLYMER_BINDING, GOMF_SUMO_POLYMER_BINDING, GOBP_S_SHAPED_BODY_MORPHOGENESIS, GOBP_S_SHAPED_BODY_MORPHOGENESIS, GOMF_1_NISOPHOSPHATIDIC_ACID_BINDING, GOMF_1_NISOPHOSPHATIDIC_ACID_BINDING, HP_FATIGUABLE_WEAKNESS_OF_PROXIMAL_LIMB_MUSCLES, HP_FATIGUABLE_WEAKNESS_OF_PROXIMAL_LIMB_MUSCLES, GOBP_NEGATIVE_REGULATION_OF_STEROID_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_STEROID_METABOLIC_PROCESS, GOBP_DIET_INDUCED_THERMOGENESIS, GOBP_DIET_INDUCED_THERMOGENESIS, HP_RECURRENT_STAPHYLOCOCCAL_INFECTIONS, HP_RECURRENT_STAPHYLOCOCCAL_INFECTIONS, GOMF_CARBOHYDRATE_CATION_SYMPORTER_ACTIVITY, GOMF_CARBOHYDRATE_CATION_SYMPORTER_ACTIVITY, GOCC_MICROFIBRIL, GOCC_MICROFIBRIL, WP_TTS_REGULATE_MIRNAS_RELATED_TO_CARDIAC_HYPERTROPHY, WP_TTS_REGULATE_MIRNAS_RELATED_TO_CARDIAC_HYPERTROPHY, MOTAMED_RESPONSE_TO_ANDROGEN_DN, MOTAMED_RESPONSE_TO_ANDROGEN_DN, GOBP_TRIGEMINAL_NERVE_DEVELOPMENT, GOBP_TRIGEMINAL_NERVE_DEVELOPMENT, GOBP_VERY_LONG_CHAIN_FATTY_ACID_CATABOLIC_PROCESS, GOBP_VERY_LONG_CHAIN_FATTY_ACID_CATABOLIC_PROCESS, REACTOME_PLATELET_AGGREGATION_PLUG_FORMATION, REACTOME_PLATELET_AGGREGATION_PLUG_FORMATION, MIR8086, MIR8086, GOCC_PERINUCLEOLAR_COMPARTMENT, GOCC_PERINUCLEOLAR_COMPARTMENT, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MICROTUBULE, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MICROTUBULE, GOBP_UBIQUITIN_DEPENDENT_GLYCOPROTEIN_BREAKDOWN_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_GLYCOPROTEIN_BREAKDOWN_PATHWAY, REACTOME_NEGATIVE_REGULATION_OF_TCF_DEPENDENT_SIGNALING_BY_DVL_INTERACTING_PROTEINS, REACTOME_NEGATIVE_REGULATION_OF_TCF_DEPENDENT_SIGNALING_BY_DVL_INTERACTING_PROTEINS, GOBP_REGULATION_OF_ADIPOSE_TISSUE_DEVELOPMENT, GOBP_REGULATION_OF_ADIPOSE_TISSUE_DEVELOPMENT, HP_INTRAMUSCULAR_HEMATOMA, HP_INTRAMUSCULAR_HEMATOMA, HP_PERIPHERAL_ARTERIOVENOUS_FISTULA, HP_PERIPHERAL_ARTERIOVENOUS_FISTULA, REACTOME_COMPETING_ENDOGENOUS_RNAS_CERNAS_REGULATE_PTEIN_TRANSLATION, REACTOME_COMPETING_ENDOGENOUS_RNAS_CERNAS_REGULATE_PTEIN_TRANSLATION, GOBP_POSITIVE_REGULATION_OF_EPIDERMAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_EPIDERMAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, WP_OSTEOPONTIN_SIGNALING, WP_OSTEOPONTIN_SIGNALING |
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