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YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_13, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_13
REACTOME_MITOCHONDRIAL_BIOGENESIS, REACTOME_MITOCHONDRIAL_BIOGENESIS
WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_WITH_H4K20ME1_MARK, WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_WITH_H4K20ME1_MARK
ALCALA_APOPTOSIS, ALCALA_APOPTOSIS
GROSS_HYPOXIA_VIA_HIF1A_UP, GROSS_HYPOXIA_VIA_HIF1A_UP
KEGG_PEROXISOME, KEGG_PEROXISOME
REACTOME_FANCONI_ANEMIA_PATHWAY, REACTOME_FANCONI_ANEMIA_PATHWAY
WELCSH_BRCA1_TARGETS_UP, WELCSH_BRCA1_TARGETS_UP
REACTOME_HDR_THROUGH_HOMOLOGOUS_RECOMBINATION_HRR, REACTOME_HDR_THROUGH_HOMOLOGOUS_RECOMBINATION_HRR
REACTOME_AUTOPHAGY, REACTOME_AUTOPHAGY
BLALOCK_ALZHEIMERS_DISEASE_INCIPIENT_DN, BLALOCK_ALZHEIMERS_DISEASE_INCIPIENT_DN
RODRIGUES_DCC_TARGETS_DN, RODRIGUES_DCC_TARGETS_DN
REACTOME_SARS_COV_2_INFECTION, REACTOME_SARS_COV_2_INFECTION
REACTOME_CRISTAE_FORMATION, REACTOME_CRISTAE_FORMATION
REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT, REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT
REACTOME_CITRIC_ACID_CYCLE_TCA_CYCLE, REACTOME_CITRIC_ACID_CYCLE_TCA_CYCLE
REACTOME_AMYLOID_FIBER_FORMATION, REACTOME_AMYLOID_FIBER_FORMATION
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION, KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION
REACTOME_SIGNALING_BY_TGFB_FAMILY_MEMBERS, REACTOME_SIGNALING_BY_TGFB_FAMILY_MEMBERS
REACTOME_CYTOSOLIC_SENSORS_OF_PATHOGEN_ASSOCIATED_DNA, REACTOME_CYTOSOLIC_SENSORS_OF_PATHOGEN_ASSOCIATED_DNA
REACTOME_DNA_DOUBLE_STRAND_BREAK_RESPONSE, REACTOME_DNA_DOUBLE_STRAND_BREAK_RESPONSE
REACTOME_TOLL_LIKE_RECEPTOR_9_TLR9_CASCADE, REACTOME_TOLL_LIKE_RECEPTOR_9_TLR9_CASCADE
REACTOME_PEROXISOMAL_PROTEIN_IMPORT, REACTOME_PEROXISOMAL_PROTEIN_IMPORT
REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX, REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX
REACTOME DDX58_IFIH1_MEDIATED_INDUCTION_OF_INTERFERON_ALPHA_BETA, REACTOME_DDX58_IFIH1_MEDIATED_INDUCTION_OF_INTERFERON_ALPHA_BETA
REACTOME_REGULATED_NECROSIS, REACTOME_REGULATED_NECROSIS
REACTOME_SARS_COV_1_INFECTION, REACTOME_SARS_COV_1_INFECTION
REACTOME_DNA_DAMAGE_BYPASS, REACTOME_DNA_DAMAGE_BYPASS
REACTOME_CLASS_I_PEROXISOMAL_MEMBRANE_PROTEIN_IMPORT, REACTOME_CLASS_I_PEROXISOMAL_MEMBRANE_PROTEIN_IMPORT
REACTOME_PROTEIN_UBIQUITINATION, REACTOME_PROTEIN_UBIQUITINATION
WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_36HR, WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_36HR
WP_7Q1123_COPY_NUMBER_VARIATION_SYNDROME, WP_7Q1123_COPY_NUMBER_VARIATION_SYNDROME
REACTOME_DUAL_INCISION_IN_TC_NER, REACTOME_DUAL_INCISION_IN_TC_NER
WP_EICOSANOID_METABOLISM_VIA_CYCLO_OXYGENASES_COX, WP_EICOSANOID_METABOLISM_VIA_CYCLO_OXYGENASES_COX
REACTOME_FLT3_SIGNALING, REACTOME_FLT3_SIGNALING
KEGG_LYSINE_DEGRADATION, KEGG_LYSINE_DEGRADATION
REACTOME_TRANSCRIPTIONAL_ACTIVITY_OF_SMAD2_SMAD3_SMAD4_HETEROTRIMER, REACTOME_TRANSCRIPTIONAL_ACTIVITY_OF_SMAD2_SMAD3_SMAD4_HETEROTRIMER
REACTOME_TRANSLATION_OF_SARS_COV_2_STRUCTURAL_PROTEINS, REACTOME_TRANSLATION_OF_SARS_COV_2_STRUCTURAL_PROTEINS
REACTOME_DUAL_INCISION_IN_GG_NER, REACTOME_DUAL_INCISION_IN_GG_NER
REACTOME_FATTY_ACID_METABOLISM, REACTOME_FATTY_ACID_METABOLISM
DESERT_PERIVENOUS_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP, DESERT_PERIVENOUS_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP
REACTOME_TRANSLESION_SYNTHESIS_BY_Y_FAMILY_DNA_POLYMERASES_BYPASSES_LESIONS_ON_DNA_TEMPLATE, REACTOME_TRANSLESION_SYNTHESIS_BY_Y_FAMILY_DNA_POLYMERASES_BYPASSES_LESION
KEGG_TRYPTOPHAN_METABOLISM, KEGG_TRYPTOPHAN_METABOLISM
REACTOME_FORMATION_OF_INCISION_COMPLEX_IN_GG_NER, REACTOME_FORMATION_OF_INCISION_COMPLEX_IN_GG_NER
REACTOME_NOTCH1_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION, REACTOME_NOTCH1_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION
REACTOME_MYD88_INDEPENDENT_TLR4_CASCADE, REACTOME_MYD88_INDEPENDENT_TLR4_CASCADE
RUAN_RESPONSE_TO_TNF_DN, RUAN_RESPONSE_TO_TNF_DN
KEGG_FATTY_ACID_METABOLISM, KEGG_FATTY_ACID_METABOLISM
REACTOME CYCLIN D ASSOCIATED EVENTS IN G1, REACTOME CYCLIN D ASSOCIATED EVENTS IN G1
REACTOME E3 UBIQUITIN LIGASES UBIQUITINATE TARGET PROTEINS, REACTOME E3 UBIQUITIN LIGASES UBIQUITINATE TARGET PROTEINS
REACTOME_MITOCHONDRIAL_IRON_SULFUR_CLUSTER_BIOGENESIS, REACTOME_MITOCHONDRIAL_IRON_SULFUR_CLUSTER_BIOGENESIS
WP_EICOSANOID_METABOLISM_VIA_LIPO_OXYGENASES_LOX, WP_EICOSANOID_METABOLISM_VIA_LIPO_OXYGENASES_LOX
KEGG_BUTANOATE_METABOLISM, KEGG_BUTANOATE_METABOLISM
REACTOME_SELECTIVE_AUTOPHAGY, REACTOME_SELECTIVE_AUTOPHAGY
PID_ERBB1_INTERNALIZATION_PATHWAY, PID_ERBB1_INTERNALIZATION_PATHWAY
KEGG_PROPANOATE_METABOLISM, KEGG_PROPANOATE_METABOLISM
WP_FATTY_ACID_BIOSYNTHESIS, WP_FATTY_ACID_BIOSYNTHESIS
KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS, KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS
REACTOME_INTERLEUKIN_3_INTERLEUKIN_5_AND_GM_CSF_SIGNALING, REACTOME_INTERLEUKIN_3_INTERLEUKIN_5_AND_GM_CSF_SIGNALING
WP_MITOCHONDRIAL_CIII_ASSEMBLY, WP_MITOCHONDRIAL_CIII_ASSEMBLY
RASHI_RESPONSE_TO_IONIZING_RADIATION_5, RASHI_RESPONSE_TO_IONIZING_RADIATION_5
CHO_NR4A1_TARGETS, CHO_NR4A1_TARGETS
REACTOME_SIGNALING_BY_NOTCH3, REACTOME_SIGNALING_BY_NOTCH3
REACTOME_SIGNALING_BY_PTK6, REACTOME_SIGNALING_BY_PTK6
REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL, REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL
REACTOME_SIGNALING_BY_ERBB2, REACTOME_SIGNALING_BY_ERBB2
REACTOME_FORMATION_OF_ATP_BY_CHEMIOSMOTIC_COUPLING, REACTOME_FORMATION_OF_ATP_BY_CHEMIOSMOTIC_COUPLING
MAHAJAN_RESPONSE_TO_IL1A_DN, MAHAJAN_RESPONSE_TO_IL1A_DN
KIM_LIVER_CANCER_POOR_SURVIVAL_DN, KIM_LIVER_CANCER_POOR_SURVIVAL_DN
LANDIS_ERBB2_BREAST_PRENEOPLASTIC_DN, LANDIS_ERBB2_BREAST_PRENEOPLASTIC_DN
REACTOME_METABOLISM_OF_STEROIDS, REACTOME_METABOLISM_OF_STEROIDS
REACTOME_CARGO_RECOGNITION_FOR_CLATHRIN_MEDIATED_ENDOCYTOSIS, REACTOME_CARGO_RECOGNITION_FOR_CLATHRIN_MEDIATED_ENDOCYTOSIS
HASLINGER_B_CLL_WITH_17P13_DELETION, HASLINGER_B_CLL_WITH_17P13_DELETION
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ΓΕΙΝ_LOCALIZATION, REACTOME_PROTEIN_LOCALIZATION