

RXRA_BOUND_WITH_H4K20ME1_MARK, WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_WITH_H4K20ME1_MARK

- KEGG_PEROXISOME, KEGG_PEROXISOME
- WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_36HR, WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_36HR
- NAKAYAMA_SOFT_TISSUE_TUMORS_PCA2_DN, NAKAYAMA_SOFT_TISSUE_TUMORS_PCA2_DN
- KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION, KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION
- CHIANG_LIVER_CANCER_SUBCLASS_PROLIFERATION_DN, CHIANG_LIVER_CANCER_SUBCLASS_PROLIFERATION_DN
- RASHI_RESPONSE_TO_IONIZING_RADIATION_5, RASHI_RESPONSE_TO_IONIZING_RADIATION_5
- REACTOME_BRANCHED_CHAIN_AMINO_ACID_CATABOLISM, REACTOME_BRANCHED_CHAIN_AMINO_ACID_CATABOLISM
- MARTINEZ_RESPONSE_TO TRABECTEDIN, MARTINEZ_RESPONSE_TO TRABECTEDIN
- MOOTHA_FFA_OXYDATION, MOOTHA_FFA_OXYDATION
- REACTOME_NUCLEAR_RECEPTOR_TRANSCRIPTION_PATHWAY, REACTOME_NUCLEAR_RECEPTOR_TRANSCRIPTION_PATHWAY
- WP_MITOCHONDRIAL_LCFATTY_ACID_BETAOXIDATION, WP_MITOCHONDRIAL_LCFATTY_ACID_BETAOXIDATION
- OZEN_MIR125B1_TARGETS, OZEN_MIR125B1_TARGETS
- WP_GALANIN_RECEPTOR_PATHWAY, WP_GALANIN_RECEPTOR_PATHWAY
- VERNOCHET_ADIPOGENESIS, VERNOCHE_T_ADIPOGENESIS
- REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION, REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION
- REACTOME_METABOLISM_OF_WATER_SOLUBLE_VITAMINS_AND_COFACTORS, REACTOME_METABOLISM_OF_WATER_SOLUBLE_VITAMINS_AND_COFACTORS
- REACTOME_TRIGLYCERIDE_CATABOLISM, REACTOME_TRIGLYCERIDE_CATABOLISM