RXRA_BOUND_WITH_H4K20ME1_MARK, WAKABAYASHI_ADIPOGENESIS_PPARG_RXRA_BOUND_WITH_H4K20ME1_MARK 🤘 `REACTOME_ABC_FAMILY_PROTEINS_MEDIATED_TRANSPORT, REACTOME_ABC_FAMILY_PROTEINS_MEDIATED_TRANSPORT GO NEGATIVE REGULATION OF PROTEIN AUTOPHOSPHORYLATION, GO NEGATIVE REGULATION OF PROTEIN AUTOPH

/ GSE4748_CTRL_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_DN, GSE4748_CTRL_VS_LPS_AND_CYANOBACTERIUI / GO_COFACTOR_BIOSYNTHETIC_PROCESS, GO_COFACTOR_BIOSYNTHETIC_PROCESS GO_BROWN_FAT_CELL_DIFFERENTIATION, GO_BROWN_FAT_CELL_DIFFERENTIATION BRUECKNER_TARGETS_OF_MIRLET7A3_DN, BRUECKNER_TARGETS_OF_MIRLET7A3_DN

NAKAYAMA_SOFT_TISSUE_TUMORS_PCA2_DN, NAKAYAMA_SOFT_TISSUE_TUMORS_PCA2_DN