75 GO terms clustered by 'binary_cut' deacetylase phospholipase succinatecoa oacyltransferase ligase activity kinase histone dna nadph heavy clathrin chain binding sialic acid heparin derivative carbohydrate binding proteoglycan compound sulfate chondroitin nfkappabinducing **Similarity** activity kinase activator inhibitor atpase 0.4 0.3 calciumdependent 0.2 binding phospholipid phosphatidylinositol 0.1 binding gammatubulin dynactin atpase voltagegated channel transfer lipid phospholipid activity antiporter rotational mechanism coregulator transcription binding tfiiicclass complex rna polymerase complex binding mutlalpha repair mismatch mutsalpha growth fibroblast binding histone binding acetyltransferase binding anion amino acid glutamate binding receptor tyrosine kinase semaphorin neuropilin laminin binding folding chaperone binding rdna mirna translation elongation activity