

Record: AKID

name AKID

description

AKID is a Deep Neural Network-based method that identifies kinase-specific phosphorylation events at single-kinase level in any given Eukaryotic proteome without prior functional annotation. It automatically detects kinase domains, their residues responsible for target specificity and finally their target peptides.

softwareVersion 1.0

url <http://akid.bio.uniroma2.it>

citation(pmid) <https://www.ncbi.nlm.nih.gov/m/pubmed/30016513/> 

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operatingSystem Linux