

HaDeX: analysis of data from hydrogen deuterium exchange-mass spectrometry

Choose state :

☒ gg_Nucb2_CaCl2

☐ gg_Nucb2_EDTA

Choose range :

0

407

Name	Value
Length	407
Covered	87.22%
Cys	0

Protein name

db_Nucb2

Reconstructed sequence

xxxxxxxxxxxxxxxxxVPIDIDKTKVKGEHVEGEKIENPDTGLYYDEYLRQVIDVLETDKH
FREKLQTADIEEIKSGKLSRELDLVSHHVTRLDELKRQEVARLRMLIAKAKMDSVQDTGI
DHQALLKQFEHLNHQNPDTFEPKDLDMLIKAATSDLENYDKTRHEEFKKYEMxxxxxx
xxxxxxLDEEKRQREESKFGEMxxxxxxxxxxxxxxxxxxKEVWEEADGLDPNEFDPKTF
FKLHDVNNDRFLDEQELEAxFTKELEKVYDPKNEEDDMVEMEEERLxxxxHVMNEVD
INKDRLVTLEEFRLATEKKEFLEPDSWETLDQQQLFTEDELKEFESHISQQEDELRRKAE
ELQKQKEELQRQHDQLAQEQELQQVVKQMEQKKLQQANPPAGPAGELK

