

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac region		
Q61666	HIRA_MOUSE	Mus musculus	Protein HIRA	36.743721	T541	S111;S548;T554;S556;T575;S583;S608;S609;S610;S612;S659;S673	40997131;40885482	MKLLKPTWVNHNGKPIFSVDIHPDG TKFATGGQGODSGKVVWVWNSPVL QEDDEKDENIPKMLCQMDNHLACV NCVRWSNSGMYLASGGDDKLMVW KRATYIGPSTVFGSSGKLANVEQWR CVSILRSHSGDVMVAVSPHDAWL ASCSVDNTVVIWNAVKFPEILATLRG HSGLVKGLTWDVPVKYIASQADDRS LKVVRTLDWQLETSITKPFDECGGT THVLRLSWSPDGHYLVSAHAMNNS GPTAQIHEREGWKTNDMDFVGHKAV TVVKFNPKIFKKKQKNGSSTKPSCPY CCCAVGSKDRSLSVWLTCLKRPLVVI HELFDKSIDISWTLNGLGILVCSM DGSVAFLDFSQDELGDPLSEEEKSRI HQSTYGKSLAIMTEAQLSTAVIENPE MLKYQRRQQQLDQKNATTRETS SASSVTGVVNGESLEDIRKNLLKKQ VETRTADGRRRITPLCIAQLDTGDFS TAFNSIPLSSSLAGTMLSSPSGQQL LPLDSSTPSFGASKPCTEPAATSAR PTGESVSKDSMNATSTPAASSPSVLT TPSKIEPMKAFDSRFTERSKATPGAP SLTSVIPTAVERLKEQNLVKELRSRE LESSSDSDEKVHLAKPSSLKRKLEL EVETVEKKKGRPRKDSRLLPMSLS VQSPAALSTEKEAMCLSA PALALKLP IPGPQRAFTLQVSSDPSMYIEVENEV TTVGGIRLSRLKCNREGKEWETVLS SRVLTAAAGSDVVCACEKRMLSVF STCGRLLPPILLPSPISTLHCTGYPV MALTAATLSVWDVHRQVVVKEE SLHSILSGSDMTVSQILLTOHGIPVM NLSDGKAYCFNPSLSTWNLVSDKQ DSLQCADFRNLSLPSODAMLCSGFL AIIQGRTSNSGRQAARLFSVPHVVQ QETTLAYLENQVAAALTLQSSHEYR HWLLLYARYLVNEGFEYRLREICKD LLGPVHCSTGSQWESTVVGLRKRKEL LKELLPVIGQNLRFQRLFTECQEQL DILRDK	True	False	1.925	4.544	0.606	False	False	False	False	False	False

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