

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
088622	PARG_MOUSE	Mus musculus	Poly(ADP-ribose) glycohydrolase	13.597734	NaN	S135;T137;S195;T197;S256;S259;S281;S286;S293;S297;S311	28528544;25153642	MSAGPGWEPCTKRPRWGAAGTSAP TASDSRSFFPGRQRRVLDPKDAPVQF RVPPSSPACVSGRAGPHRGNATSFV FKQKTITTWMDTKGPKTAESESKEN NNTRIDSMSSVQKDNFYPHKVEK LENVPQLNLDKSPTEKSSQYLNQQQ TASVCKWQNEGKHAEQLLASEPPA GTPLPKQLSNANIGQSPHTDDHSDT DHEEDRDNOQFLTPIKLANTKPTVG DGQARSNCKCSGSRQSVKDCGTGCQ QEEVDVLPESPLSDVGAEDIGTGPK NDNKLGTQESSLGDSPPEKESEPE SPMDVDNSKNSCQDSEADEETSPV FDEQDDRSSQTANKLSSCQAREADG DLRKRYLTKGSEVRLHFQFEGENNA GTSDLNAKPSGNSSSLNVECRSSKQ HGKRDSKITDHFMRISKSEDRRKEQ CEVRHQRTERRKIPKYIPNLPPEKKW LGTPIEMRMKMPRCGIHLPSLRPSAS HTVTVRVDLLRAGEVPKPFPTHYKD LWDNKHVKMPCEQONLYPVEDENG ERTAGSRWELIQTALLNKFTRPQNL KDAILKYNVAYSKKWDFALVDFWD KVLEEAQAQHLYSILPDMVKIALCL PNICTQPIPLLKQKMNHSVTMSQEQ IASLLANAFFCTFPRRNAMKSEYSS YPDINFNRLFEGRSSRKPEKLTLC YFRRVTEKKPTGLVTFTRQSLEDFPE WERCEKPLTRLHVTYEGTIEGNRG MLQVDFANRFVGGGVTGAGLVQEEI RFLINPELIVSRFTEVLDHNECLIIT GTEQYSEYTYAETYRWARSHEDGS EKDDWQRRCTEIVDAIDALHFRRYLD QFVPEKVRRELNKAYCGFLRPGVPS ENLSAVATGNWGCGAFGGDARLKA LIQILAAAAAERDVVYFTFGDSELMR DIYSMHTFLTERKLDVGKVKLLLR YNEECRNCSTPGPDIKLYPFIYHAVE SSAETDMPGQKAGT