

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q86W56	PARG_HUMAN	Homo sapiens	Poly(ADP-ribose) glycohydrolase	22.892873	S26	S22;S68;S133;S137;T139;S197;T199;S261;S264;S286;S291;S298;S302;S316;S448	30379171;23301498;37858678	MNAGPGCEPCTKRPRWGAATTSPA ASDARSFPSRQRRVLDPKDAHVQFR VPPSSPACVPRAGQHRGSATSLVF KQKTITSWMDTKGIKTAESLDSK ENNNTRIESMMSSVQKDNFYQHNV EKLENVSQSLDKSPTKSTQYLNQ HQTAAAMCKWONEGKHTEQLESEP QTVTLVPEQFSNANIDRSPONDDHS DTDSEENRDNQQLTTVKLANAKQT TEDEQAREAKSHQKCSKCDPGE CASCQQDEIDVVPESPLSDVGS GTGPKNDNKLTRQESCLGNSPPFEK ESEPEPMDVDNSKNSCQDSEADE ETSPGFDEQEDGSSQTANKPSRFQ ARDADIEFRKRYSTKGGEVRLHFQF EGGESRTGMNDLNAKLPGNISSLNV ECRNSKQHGKKDSKITDHFMRLLPK AEDRRKEQWETKHQRTERKIPKYVP PHLSPDKKWLGTPIEEMRRMPRCGI RLPLLRPSANHTVTIRVDLLRAGEVP KPFPTHYKDLWDNKHVKMPCSEON LYPVEDENGERTAGSRWELIQTALL NKFTRPQNLKDAILKYNVAYSKKWD FTALIDFWDKVL EEAQHLYSILP DMVKIALCLPNICTQPIPLLKQKMN HSITMSQEQLASLLANAFCTFPRRN AKMKSEYSSYPDINFNRLFEGRSSR KPEKLTLCFYFRRVTEKKPTGLVTF TRQSLDFPEWERCEKPLTRLHVTY EGTIEENGQGM LQVDFANRFVGGG VTSAGLVQEIRFLINPELIHSRLFTE VLDHNECLIITGTEQYSEYTGAEY RWSRSHEDGSEDDWQRRCTEIVAI DALHFRRYLDQFVPEKMRRRELNKAY CGFLRPGVSSENLSAVATGNWCCG AFGGDARLKALIQILAAAAAERDVVY FTFGDSELMRDIYSMHIFLTERKLT GDVYKLLLRYYNEECRNCSTPGPDIK LYPFIYHAVESCAETADHSGQRTGT	True	False	4.57	4.638	4.067	1.353	1.105