

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q8TEW0	PARD3_HUMAN	Homo sapiens	Partitioning defective 3 homolog	14.605553	S437	S25;T91;S156;S174;S383;T489;S692;S695;S715;S728;S809;S827;S837;S852;S873;S962;S971;S973;S1046;S792	30379171;34725712;23301498;34019948;39534244;39302247;35254053	MKVTVCFGRTVVPCGDGHMKVFLIQQAVTRYRKAIAKDPNYWVHRLVLEHGGGILDDLDLDCVADDKDRLVAVFDEQDPHHGGDGTASSTGTQSP EIFGSELGTNNVSAFQPYQATSEIEVTPSVLRANMPLHVRRSSDPALIGLSTSVSDSNFSSEEPSRKNPTRWSTTAGFLKQNTAGSPKTCDRKKDENYRSLPRDTSNWSNQFQDNARSSLSASHPMVGVKWLKQEQDEDGTEEDNSRVEPVGHADTGLEHIPNFSLDDMVKLV EVPNDDGGPLGIHVVPFSAARGGRTLGLLVKRLKGGKAEHENLFREND CIVRINDGDLNRNRRFEQAQHMFRAQAMRTPHWFHVVPAANKEQYEQLSSEKNNYSSRFSQYIDNRSVNSAGLHTVQAPRLNHPPEQIDSHSRLPHSAHPSGKPPSAPASAPQNVFSTTVSSGYNTKKIGKRLNIQLKKGTEGLGFSITSRDVTIGGSAPIYVKNILPRGAAIQDGRKAGDRLIEVNGVDLVGKSQEEVVSLLRSTKMEGTVSLVFRQEDAFHPR ELSRIMGESGKYQLSPTVNMPQDDT VIIIEDDRLPVLPPHLSQSSSSSHDDVGFVTADAGTWAKAAISDSADC SLPDVPVLAFOREGFGROSMSEKRTKQFSDASQLDFVKTRKSKSMDLGIAD ETKLNTVDDQKAGSPSRDVGPSLGLKKSSSLESQTAAVEVTLNGDIPFHRPRPRIIRGRGCNESFRAAIDKSYDKP AVDDDDDEGMETLEEDTEESSRSGRE SVSTASDQPSHSLERQMNGNQEKGDKTDRKKDKTGKEKKDRDKEKDKMKAKKMLKGLGDMFRFGKHRKDKIEKTGKIKQESFTSEEIRRMKQEQERIQAKTREFRERQARERDYAEIQDFHRTFGCDDDELMYGGVSSYEGSMALNARPQSPREGHMMDALYAQVKKPRNSKPSVDSNRSTPSNHDRIQRLRQEFQQAQKQEDVEDRRRTYSFEQPWPNARPATQSGRHSVSVVEVQMQRQRQEEERSSQQAQRYSSLPRQSRKNASSVSQDSWEQNYSPGEGFQSAKENPRYSSYQGSRNGLGGHGFNARVMLETQELLRQEQRRKEQMKKQPPSEGSPSNYDSYKKVQDPSYAPPKGPF RQDVPPSPSQVARLNLQTPEKGRPFYS	True	True	4.671	2.058	1.359	1.293	1.502