

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8TEW0	PAR3_HUMAN	Homo sapiens	Partitioning defective 3 homolog	6.61901	S437	S25;T91;S156;S174;S383;T489;S692;S695;S715;S728;S809;S827;S837;S852;S873;S962;S971;S973;S1046;S792	34725712;34019948;35254053;23301498;30379171	MKVTVCFGRTRVVPCGDGHMKVF SLIQQAVTRYRKAIAKDPNYWQVHR LEHGDGGILDLDLDCDVADDDKDR VAVFDEQDPHHGGDGTSSASSTGT PEIFGSELGTNNVSAFQPYQATSEI VTPSVLRANMPLHVRSSDPALIGLS TSVSDSNFSSEEPSRKNPTRWSTTA GFLKQNTAGSPKTCDRKKDENYRSL PRDTSNWSNQFQRDNARSSLSASH PMVGKWLEKQEDEDGTEEDNSRV EPVGHADTGLEHIPNFSLDDMVKLV EVPNDGGPLGIHVVPFSARGGRTL LLVKRLEKGGKAEHENLFREND RINDGDLRNRREFEAQHMFRQAMR TPIWFHVVPAANKEQYEQLSQSEK NNYSSRFSPDSQYIDNRSVNSAGL HTVQRAPRLNHPPEQIDSHSRLPHS AHPSGKPPSAPASAPQNVFSTTVSSG YNTKKIGKRLNIQLKKGTEGLGFSIT SRDVTIGGSAPIYVKNILPRGAAIQDG RLKAGDRLIEVNGVDLVGKSQEEVV SLLRSTKMEGTVSLLVFRQEDAFHP RELNAEPSQMQIPKETAEDDIVLT PDGTREFLTFEVPLNDSGSAGLGVS VKGNRSKENHADLGIFVKSIIINGGAA SKDGRLRVNDQLIAVNGESLLGKTN QDAMETLRRSMSTEGNKRGMQLIV ARRISKCNELKSPGSPGPELPIETAL DDRERRISHSLYSGIEGLDESPSRNA ALSRIMGESGKYQLSPTVNMPQDDT VIEDDRLPVLPPHLSQSSSSSHDD VGFVTADAGTWAKAASDSADCSLS PDVDPVLAFAQREGFGGRQSMSEKRTK QFSDASQLDFVKTREKSKSMDLGIAD ETKLNVTDDQKAGSPSRDVGPSLGL KKSSSLESLSQTAVAQVTLNGDIPFHR PRPRIIRGRGCNESFRAAIDKSYDKP AVDDDDDEGMETLEEDTESSRSGRE SVSTASDQPSHSRLERQMNNGNQEKG DKTDRKKDKTGKEKKKDRDKEKDK MKAKKGMMLKGLGDMFRFGKHRKD DKIEKTGKIKIQESFTSEERIRMKQ EQERIQAKTREFRERQARERDYAEIQ DFHRTFGCDEL MYGGVSSYEGSM ALNARQSPREGHMMDALYAQVKK PRNSKPSPVDSNRSTPSNHDRIQRL RQEFQAKQDEDVEDRRRTYSFEQP WPNARPATQSGRHSVSVEVQMQRQ RQEERESSQQAQRQYSSLPRQSRKN ASSVSQDSWEQNYSPGEGFQSAKE NPRYSSYQGSRNGLGGHGFNARV MLETQELLRQEQRRKEQOMKKQPP SEGSPNYSYKKVQDPSYAPPKGP RQDVPPSPSQVARLNLQTPEKGRF FYS

