

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O75170	PP6R2_HUMAN	Homo sapiens	Serine/threonine-protein phosphatase 6 regulatory subunit 2	13.989682	T887;T899;T903;T904;S907;S919;S920	S289;S771;S828	35254053;28657654;29351928;33214551;35132862;35138101;34019948;30620550	MFWKFDLNTTSHVDKLLDKEHVTL QELMDEDDILQECKAQNKLLDFLC RQQCMEELVSLITQDPPDMEEKVR FKYPNTACELLTCDVPQISDRLGGE SLLSLLYDFLDHEPPLNPLLASFFSK TIGNLIARKTEQVITFLKKKDKFISLV LKHIGTSALMDLLLRLVSCVEPAGLR QDVLHWNEEKVIQRLVELIHPSQD EDRQSNASQTLCDIVRLGRDQGSQ QEALPDPLLTALESQDCVEQLLKN MFDGDRTESCLVSGTQVLLTLETR RVGTEGLVDSFSQGLERSYAVSSSVL HGIEPRLKDFHQLLLNPCKKAILTT IGVLEEPLGNARLHGARLMAALLHT NTPSINQELCRLNTMDLLDLFFKY TWNNFLHFQVELCIAAILSHAAREE RTEASGSESRVEPPHENGNSLETP QPAASLPDNTMVTHLFQKCLVQRI LEAWEANDHTQAAGGMRRGNMGH LTRIANAVVQNLERGPVQTHISEVIR GLPADCRGRWESFVEETLTETNRRN TVDLVSTHHLHSSSEDEDIEGAFP ELSLQQAFSDYQIQMTANFVDQFG FNDEEFADQDDNINAPFDRIAEINF NIDADEDSPSAAALFEACCSDRIQPF DDEDEDIWEDSDTRCAARVMARPR FGAPHASESCSKNGPERGGQDGKA SLEAHRDAPGAGAPPAPGKKEAPPV EGDSEGAMWTAVFDEPANSTPTAP GVVRDVGSSVWAAGTSAPEEKGWA KFTDFQPFCCSESGPRCSPVDTEC SHAEGSRSQGPEKASQASYFAVSPA SPCAWNVCVTRKAPLLASDSSSSGG SHSEGDQKAASAMDAVSRGPGRE APPLPTVARTEEAVGRVGCADSRLLS PACPAPKEVTAAPAVAVPEATVAITT ALSKAGPAIPTAVSSALAVAVPLGPI MAVTAAPAMVATLGTVTKDGKTDAP PEGAALNGPV

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