

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q5TCZ1	SPD2A_HUMAN	Homo sapiens	SH3 and PX domain-containing protein 2A	29.030064	T747	T256;S406;S421;S547;S567;S593;S644;T731;S767;S769;S819;T829;S1002;S1016;S1017;S1038	29237092	MLAYCVQDATVVDVEKRRNPSKHVY YIINVTWSDSTSQTYYRRYSKFFDLQ MQLLDKFFIEGGQKDPKQRIIFLPG KILFRRSHIRDVAVKRLKPIDEYCRAL VRLPPHISQCDEVFRFFEARPEDEVN PPKEDYGSKRKSVWLSWAESPCK DVTGADATAAEPMILEQYVVVSNYKK QENSELSLQAGEVVDVIEKNESGW WVSTSEEQGWVWPATYLEAQNQTR DDSDINTSKTGEVSKRRKAHLRLD RRWTLGGMVNRQHSREEKYVTVP YTSQSKDEIGFEKGVTVVIRKNLEG WWYIRYLGKEGWAPASYLKKAKDDL PTRKKNLAGPVEIIGNIMEISNLLNK KASGDKETPPAEGEGHEAPIAKKEIS LPILCNASNGSAVGPDRTVSRLAQ GSPAVARIAPQRAQISSPNLRTRPPP RRESSLGFQLPKPPEPPSVEVEYTI AEFQSCISDGISFRGGQKAEVIDKNS GGWVYVQIGEKKEGWAPASYIDKRKK PNLSRRTSLTRPKVPPAPPSPKPE AEEGPTGASESQDSPRKLKYEPEY DIPAFGFDSEPELSEEPVEDRASGER RPAQPHRPSPASSLQRRARFKVGESS EDVALEEEETIYENEGFRPYAEDTL SARSSGSDSPGSSSLSLTRKNSPKS GSPKSSLLKKAEKNAQAEMGKN HSSASFSSITINTCCSSSSSSSSSL SKTSGDLKPRSASDAGIRGTPKVR KADANAGLTSCPRAKPSVRPKPFL NRAESQSQEKMDISTLRRQLRPTGQ LRGGLKGSKSESELPPQTASEAPSE GSRSSSDLITLPTTTPCPTKKEWE GPATSYMTC SAYQKVQDSEISFPAGV EVQVLEKQESGWVYVRFGELEGWA PSHYLVLDENEQDPDPSGKELDTVPA KGRQNEGKSDSLEKIERRVQALNTV NQSKKATPIPSKPPGGFGKTSGTPA VKMRNGVRQVAVRPQSVFVSPPKD NNLSCALRRNESLTATDGLRGVRRN SSFSTARSAAAEAKGRLAERAASQG SDSPLLPAQRNSIPVSPVRPKPIEK QFIHNNLKDVVYSIADYEGDEETAG FQEGVSMEVLERNPNGWVYWCQILD GVKPFKGVVPSNYLEKKN	True	False	4.659	3.33	1.324	1.23	1.01	2.112	1.671

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