

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasm reticulum
P23246	SFPQ_HUMAN	Homo sapiens	Splicing factor, proline- and glutamine-rich	38.481322	S496;T652;S654;S660	S8;S33;S273;S283;T293;T368;S374;S379;S496;S626;T687;T691	27655845;34725712;36580660;30397120;37217939;40596516;21740066;38665916;39534244;39302247;34019948;32870666;22121020;35289036;30444036;35083852;20305658;33214551;41018975;35254053;40930100;32119511;33465208	MSRDRFRSRGGGGGFHRRGGGGGRGGLHDFRSPPPGMLNQNRRGPMGPGPGQSGPKPIPPPPPHQQQQPPQPPPPQPPPHQPPPHQPHQQQPPPPPPQDSSKPVVAQGPAPGPGVGSAPPASSAPPATPPTSGAPPGSGPGPTPTPPAVTSAPPGAPPPTPPSSGVP TTPPQAGGPPPPAAVPGPGPKQ GPGPGPKGGKMPGGPKPGGGPGL STPGGHPKPPHRGGGEPGRGRQHH PPYHQHHQGGPPGGPGRSEKIS DSEGFKANLSLLRRPGEKTYTQRCLFVGNLPADITEDEFKRLFAKYGEPGEVFINKGKGGFKLESRALAEIAKA ELDDTPMRGRQLRVRFATHAAALSV RNLSPPYVSNELLEAFSQFGPIERAV VIVDDRGRSTGKGIVEFASKPAARKA FERCSEGVFLTTTPRPVIVEPLEQL DDEDGLPEKLAQKNPMYQKERETPP RFAQHGTFEYEYSQRWKSLEMEK QQREQVEKNMKDAKLESEMEDA YHEHQANLLRQDLMRRQEELRRME ELHNQEMQKRKEMQLRQEEERRR REEEMMIRQREMEEQMRQRREESY SRMGYMDPRERDMRMGGGGAMN MGDPYGGGQKFPPLGGGGGIGYEA NPGVPPATMSGSMGSDMRTERFG QGGAGPVGGQGPGRMGPGTPAGYGRGEEYEGPNKKPRF	True	False	4.545	5.0	2.843	1.501