

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	exti regi
Q9ES52	SHIP1_MOUSE	Mus musculus	Phosphatidylinositol 3,4,5-trisphosphate 5-phosphatase 1	25.685855	NaN	S246;T918;S935;T945;T964;S967;S972;T1021	39627609	MPAMVPGWNHGNITRSKAEELLSR AGKDGSLVRASESIPRAYALCVLFR NCVYTYRILPNEDDKFTVQASEGVP MRFFTKLDQLIDFYKKENMGLVTHL QYPVPLEEEDDAIDEAEEDTVESVMSP PELPPRNIPMSAGPSEAKDLPLATEN PRAPEVTRLSESETLQRLQSMDS GLPEEHLKAIQDYLTQLLDSDFLK TGSSNLPHLKKLMSLLCKELHGEVI RTLPSLESLQRLFDQQLSPGLRPRPQ VPGEASPTMVAKLSQLTSLSSIED KVKSLLHEGSESTNRRSLIPVTFEV KSESLGIPQKMHKVDVESGKLIVKK SKDGEDKFYSHKKILQIKSQFLN KLVILVETEKEKILRKEYVFADSKKR EGFCQLLQOMKNKHSEQPEPDMITI FIGTWNMGNAPPPKITSWFLSKGQ GKTRDSDADYIPHDYVIGTQEDPLG EKEWLELLRHSLQEVTSMTFKTVAI HTLWNIRIVVLAKPEHENRISHICTD NVKTGIANTLGNKGAVGVSMFNG TSLGFVNSHLTSGSEKKLRRNQNY MNILRFLALGDKKLSPFNITHRFTHL FWLGDLYRVELPTWEAEAIQKIKQ QQYSDLLAHDQLLLERKDQKVFLHF EEEEITFAPTYRFERLTDKYAYTKQ KATGMKYNLPSWCDRVLWKSYPV HVVCQSYGSTSDIMSDHSPVFATF EAGVTSQFVSKNGPVTVDSDGQIEF LACYATLKTQSQTKFYLFHSSCLES FVKSQEGENEEGSEGLVVRFGETL PKLPHSDPEYLLDQHILISIKSSDSD ESYGEGCIALRLETTAQHPYITPLT HHGEMTGFRGEIKLQTSQGMRE KLYDFVKTERDESSGMKCLKNLTSH DPMRQWEPGRVPACGVSSLNEMI NPNYIGMGPFQPLHGKSTLSPDQ LTAWSYDQLPKDSSLGPRGEGPPT PPSQPPLSPKFFSSSTANRGPCPRVQ EARPGDLGKVEALLQEDLLLTKPEM FENPLYGSVSSFPKLVPRKEQESPK MLRKEPPPCPDPGISSPSIVLPKAE VESVKGTSKQAPVVLGPTPRIRSFT CSSSAEGRMTSGDKSQGKPKASASS QAPVPVKRPVKSRSSEMSQQTTPIPA PRPPLPVKSPAVLQHQHSGGRDYRD NTELPHHGKHRQEGLLGRYAMQ	True	False	4.706	2.726	1.707	1.414	1.283	3.846	1.96