

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q15020	SART3_HUMAN	Homo sapiens	Spliceosome associated factor 3, U4/U6 recycling protein	20.600146	T946	S10;S16;S215;S650;T657;S769;S795;S852	33214551;31492838;38665916;35254053;32119511;40914422;28411811;34019948;28510447	MATAAETSASEPEAESKAGPKADGE EDEVKAARTRRKVLRAVAATYKT MGPAWDQEEGVSESDGDEYAMAS SAESSPGEYEWYDEEEKNQLEIE RLEEQLSINVYDYNCHVDLIRLLRLE GELTKVRMARQKMSIIFPLTEELWL EWLHDEISMAQDGLDREHVDLFE KAVKDYICPNIWLEYGQYSVGGIGQK GGLEKVRVFERALSSVGLHMTKGL ALWEAYREFESAIVEAARLEKVVHSLF RRQLAIPLYDMEATFAEYEEWSEDPI PESVIQNYNKALQQLEKYKPYEEALL QAEAPRLAEYQAYIDFEMKIGDPARI QLIFERALVENCLVPDLWIRYSQYLD ROLKVKDLVLSVHNRAIRNCPWTVA LWSRYLLAMERHGVDHQVISVTFEK ALNAGFIQATDYVEIWQAYLDYLRRL VDFKQDSSKELEELRAAFTRALEYK QEVEERFNEGDPSCVIMQNWARIE ARLCNNMQKARELWDSIMTRGNAK YANMWLEYNLERAHGDTQHCRKA LHRAVQCTSDYPEHVCEVLLTMERT EGSLEDWDIAVQKTETRLARVNEQR MKAAEKEAALVQEEKAEQRKRA RAEKKALKKKKIRGPEKRGADDD EKEKGDDEEEQPSKRRRVENSIPAA GETONVEVAAGPAGKCAAVDVEPPS KQKEKAASLKRDMPKVLHDSKDSI TVFVSNLPYSMQEPDTKLRPLFEAC GEVVQIRPIFSNRGDFRGYCYVEFKE EKSAQALEMDRKSVEGRPMFVSPC VDKSKNPDFKVFYRSTSLKHKLFIS GLPFSCTKEELEIICKAHGTVKDLRL VTNRAGKPKGLAYVEYENESQASQA VMKMDGMTIKENIIVAISSPPORK VPEKPETRKAPGGPMLLPQTYGARG KGRTQLSLLPRALQRPSAAAPQAEN GPAAPAVAAPAATEAPKMSNADFA KLFLRK	True	False	1.282	5.0	0.947	0.697	False