

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
043290	SNUT1_HUMAN	Homo sapiens	U4/U6.U5 tri-snRNP-associated protein 1	9.141658	T258	T189;S321;S348;T392;T430;S448;S474;S486;S521;S591;S596;S598;S621;T695;S761;T764;S789	34229054;37217939;32522876;34019948;38253038;28510447;38665916	MGSSKKHRGEKEAAGTTAAAGTGG ATEQPPRHREHKKHKHSRGGSGGS GGERRKRSRERGGGSGRRGAEA EARSSTHGRERSQAEPSEKRVKREK RDDGYEAAASSKTSSGDASSLSIET NKLRAKLGKPLEVNAIKKEAGTKE EPVTADVINPMALRQREELREKLA AKEKRLLNQKLGKIKTLGEDDPWLD DTAAWIERSRQLQEKDLAEKRAKL LEEMDQEFVSTLVEEFGQRRQDL YSARDLQGLTVEHAIDSFREGETMIL TLKDKGVLQEEEDLVNVNLDKER AEKNVELRKKKPDYLPYAEDESVD LAQQKPRSILSKYDEELEGEPHSFR LEQGGTADGLRERELEEIRAKRLQ AQSLSTVGPRLASEYLTPEEMVTFKK TKRRVKKIRKKEKVVRADDLPLG DQTQDGFSGSRLRGRGRRRVSEVE EEKEPVPQPLPSDDTRVENMDSIDE EEGGAPPPGSPQVLEEDAELELQK QLEKGRRLRQLQQLRDSGEKVV EIVKKLESRORGWEEDEDPERKGA VFNATSEFCRTLGEIPTYGLAGNREE QEELMDFERDEERSANGGSESDGE ENIGWSTVNLDEEKQQDFASSTT ILDEEPIVNRGLAAALLCQNKGLLE TTVQKVARVKAPNKSLPSAVYCIEDK MAIDDKYSRREERYRGFTQDFKEKDG YKPDVKIEYVDETGKRLTPKEAFRQL SHRFHGKSGKMKTERRMKKLDEE ALLKMSSTDTPLGTVALLEKQKA QKTPYIVLSGSGKSMNANTITK	True	True	5.0	5.0	1.428	1.128	4.48