

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
Q78PY7	SND1_MOUSE	Mus musculus	Staphylococcal nuclease domain-containing protein 1	25.700107	NaN	T103;T235;T240;S426;S645;T779;S785;S909	37453647;39627609	MASSAQSSGSSGGPAVPTVQRGIVK MVLSGCAIIVRGQPRGGPPPERQINL SNIRAGNLARRAAATQPDGKDPDE PWAFFPAREFLRKKLIGKEVCFTIENK TPQGREYGMIIYLGKDTNGENIAESL VAEGLATRREGMRANNPEQNRLSE CEEQAKASKKGMWSENGSHTIRD LKYTIENPRHFFVDSHHQKPVNAIEH VRDGSVVRALLPGHHLVTVMLSGI KCPTFRRETDGSETPEPFAAEAKFFT ESRLLQRDVQHILESCHNQNLGTL HPNGNITELLKKEGFARCVDSIAV YTRGAEKLRAAERFAKERRLRWRD YVPPTANLDQKDKQFVAKVMQVLN ADAIIVKLNsgDYKTIHLSSIRPPRE GDNIQDKNKKLRPLYDIPYMFARE FLRKKLIGKKVNVTVDIRPASPATE TVPAFSERTCATVTIGGINIAEALVSK GLATVIRYQDDQRRSHYDELLAA EARAIKNGKGLHKKKEVPIHRVADIS GDTQKAKQFLPFLQRAGRSEAVVEY VFGSRLKLYLPKETCLITFLLAGIEC PRGARNLPLVQEGEPFSEATLFT KELVLQREVEVEVESMDKAGNFIGW LHMDGANLSVLLVEQALSKVHFTAE RSAYYKPLLSAEEAAKQRKEKVVWAH YEERPVEEVMPVLEEKERSASYKPV FVTEITDDLHFYVQDVETGTQLEKL MENMRNDISSHPVVEGSYAPRRGE FCIAKFVDGEWYRARVEKVESPAKV HVFYIDYGNREILPSTRLGTLPAPFST RVLPAQATEYAFAFIQVPQDEDARTD AVDSVVRDIQNTQCLLNVEHLSASC PHVTLOFADSKGDVGLGLVKEGLVM VEVRKEKQFQKVITEYLNQAQESAKS ARLNLWRYGDFRADDDEFQYSR	True	False	3.879	5.0	4.411	1.604	1.024	1.473