

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	extracellular region
Q9Y6K5	OAS3_HUMAN	Homo sapiens	2'-5'-oligoadenylate synthase 3	16.682301	S461	T365	30379171;37217939	MDLYSTPAAALDRFVARRLQPRKEF VEKARRALGALAAALRERGGRLGAA APRVLKTVKGSSGRGTALKGGCDS ELVIFLDCFKSYVDQRARRAEILSEM RASLESWWQNPVPLRLTFPEQSV GALQFRLTSVDLEDWMDVSLVPAF NVLGQAGSGVKPKQVYSTLLNSGC QGGEHAACFTELRRNFVNIRPAKLN NLILLVKHWYHQVCLQGLWKETLPP VYALELLTIFAWEQGCKKDAFSLAE GLRTVLGLIQHQHLVCFWTVNYGF EDPAVGQFLQRLKRPVILDPAFP TWDLGNAAWHWDLLAQEAASCY DHPFCFLRGMGDPVQSWKGPGLPRA GCSGLGHPQLDPNQKTPENSKSLN AVYPRAGSKPPSCPAPGPTGAASIVP SVPGMALDLSQIPTKELDRFIQDHLK PSPQFQEQVKKAIIDILRCLHENCVH KASRVSKGGSFGRGTDLRDGCDEL IIFLNCFTDYKDQGPRAEILDEMRA QLESWWQDQVPSLSLQFPEQNVPE ALQFQLVSTALKSWTDVSLLPAFDA VGQLSSGKPNPQVYSRLTSGCQE GEHKACFAELRRNFMNIRPVKLN ILLVKHWYRQVAAQNKKGKGPAPASL PPAYALELLTIFAWEQGCRQDCFN AQGFRTVLGLVQHQQLCVYWTVN YSTEDPAMRMHLLGQLRKRPLVLD PADPTWNVGHGSWELLAQEAALG MQACFLSRDGTSPVQPDVMPALLY QTPAGDLDFKIFELQPNRQFLAQQ NKAVDTICSLKENCFRNSPIKVIK VKGSSAKGTALRGRSDADLVVFLS CFSQFTEQGNKRAEIISEIRAQLEAC QQRQFEVKFEVSKWENPRVLSFSL TSQTMLDQSVDFVLPALDQALQV SGSRPSSQVYVDLIHYSNAGEYSTC FTELQRDFIISRPTKLSLIRLVKHWY QQCTKISKGRGSLPPQHGLELLTVYA WEQGGKDSQFNMAEGFRTVLELVT QYRQLCIYWTINYNADKTVGDFLK QQLQKPRPIILDPAADPTGNLGHNAR WDLAKEAAACTSALCCMGRNGIPI QPWPVKA	True	True	4.781	5.0	1.96	1.837	1.269	4.602	4.422