

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	es
Q92625	ANS1A_HUMAN	Homo sapiens	Ankyrin repeat and SAM domain-containing protein 1A	18.419428	NaN	T318;S507;S620;S622;S624;S626;S628;S647;S661;S663;S666;S677;S887	30379171;23301498	MGKEQELLEAARTGHLPVAVKLLSG KRLSSGFGGGGGGGGGGGGGGGGG GGGGLGSSSHPLSSLLSMWRGPNV NCVDSTGYTPLHHAALNGHKDVVE VLLRNDALTNVADSKGCYPLHLAAW KGDAQIVRLLIHQGPSHTRVNEQNN DNETALHCAAQYGHTEVVVLEEL TDPTMRNNKFETPLDLAALYGRLEV VKMLLNAHPNLLSCNTKKHTPLHL AARNGHKAVVQVLLDAGMDSNYQT EMGSALHEAALFGKTDVVQILLAAG TDVNIKDNHGLTALDTRVRELPSQKS QQIAALIEDHMTGKRSTKEVDKTPPP QPPLISSMDSISQKSQGDVEKAVTEL IIDFDANAE EEGPYEALYNAISCHSL DSMASGRSSDQDSTNKEAEAAGVK PAGVPRERPPPAKPPPEEEEEEDI DKKYFPLTASEVLSMRPRIHGSAARE EDEHPYELLLTAETKKVVLVDGKTK DHRRSSSSRSQDSAEGQDGOVPEQ FSGLLHGSSPVEVQDPFQLLCTA GQSHPDGSPQQGACHKASMQLLEET GVHAPGASQPSALDQSKRVGYLTGL PTTNSRSHPETLHTASPHPGGAE GDRSGARSRAPPTSKPKAELKLSRSL SKSDSLLTCSPTEDATMGRSESEL SNCSIGKKRLEKSPFASEWDEIEKI MSSIGEGIDFSQERQKISGLRTLEQS VGEWLESIGLQYESKLLNGFDDV HFLGSNVMEEQDLRDIGISDPQHRR KLLQAARSLPKVKALGYDGNSPPSV PSWLDLGLQDYVHSFLSSGYSSIDT VKNLWELELVNVLKVQLLGHKRRII ASLADRPYEEPPQKPPRFSQLRCQD LLSQTSPLSQNDSCGRSADLLLP GDTGRRRHDSLHDPAAAPRAERFRI QEEHREAKLTLRPPSLAAPYAPVQS WQHQPPEKLIFESCGYEANYLGSMLI KDLRGTESTQDACAKMRKSTEHMK KIPTIILSITYKGVKFIDASNKNVIAEH EIRNISCAAQDPEDLCTFAYITKDLQ TSHHYCHVFSTVDVNLTYEIIITLGO AFEVAYQALQAQKSRATGASAAEM IETKSSKPVKPRVGVKSALEPPDM DQDAQSHASVSWVDPKPKDSKRSL TN	None	None	None	None	None	None	None		