

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8VI51	SORC3_MOUSE	Mus musculus	VPS10 domain-containing receptor SorCS3	24.474581	NaN	NaN	33300544	MEAAGTERPAGWPGAPLARTGLLLL STWVLAGAEITWGATGGPGRLLVSPA SRPPVLPPLLPRAAENRWPEELASA RRAAAPRRRSRLEPLSQASRGEIRTE AAGMSPEGARWVPGIPSPSQAGSAR RTRRAQPPSPLERGDSDWATALADGA KGSRPHTKGSREEVRAIRTRGGASTE ELRLPSTSFALTDGSAHNQAMVHW SGHNSSVILILTCLYDFNLGVSSTESS LWRSVDYGATYEKLNKVKGLKTVLS YLYVNPNTNKRKIMLLSDPEMSESVLI SSDEGATYQKYRLTFYIQSLLFHPKQ EDWVLAYSLLDQKLYSSMDFGRRWQ LMHERITPNRFYWSVSGLDKEADLV HMEVRTADGYAHYLTCTRIQCAETT RSGPFARSIDISLWVQDEYIFIQVTIG GRASYYSYRREAFQAQIKLPKYSLPK DMHIISTDENOVFAAVQEWNQNDT YNLYISDTRGIYFTLAMENIKSSRGL MGNIIIELYEVAGIKGIFLANKRVDD QVKTYITYNKGRDWRLQAPDVLDR GSPVHCLLPFCSLHLHLQSENPPYS SGRISSKDTAPGLVVATGNIGSELSY TDIGVFISDGGNTWRQIFDEEYNV WFLDWGGALVAMKHTPLPVRHLWV SFDEGHSDWKYGFLLPLFVDGALV EAGVETHIMTVFGHFSLRSEWQLVK VDYKSIFSRRCTKEDFETWHLLNOG EPCVMGERKIFKRRKPGAQCALGRE YSGSVVSEPCVADWDFECYDYGER HGESCQVPAFWYNPASPSKDCSLG QSYLNSTGYRRIVSNCTDGLRDKY SAKTQLCPGKAPRGLHVTTDGRLV AEQGHNATFIILMEEGDLQRTNIQL DFDGVAVSYANFSPIEDGIRHVYKS AGIFQVTAYAENNLGSDTAFLFLHV VCPVEHVHLRVPFVAIRNKDVNISAV VWPSQLGTLTYFWWFNGSTKPLITL DSSISFTFLAEGTNTITVQVAAGNALI QDTKEIAVHEYFQSLLSFSPNLDY HNPDIPEWRQDIGNVIKRALIKVTSV PEDQILVAVFPGLPTSAELFILPPKNL TERRKGHEGDLEQIVETLFNALNQN LVQFELKPGVQVIVYVTLTLAPLVD SSAGHSSSAMLMLLSVVFVGLAVFL IYKFKRKIPWINIYAQVQHDKEQEMI GSVSQSENAPKITLSDFTEPELLDK ELDTRVIGSIATIASSESTKEIPNCTS V	False	True	2.029	1.359	1.633	1.209	1.707	4.686	1.465