

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	extracell region
Q3UVC0	KSR2_MOUSE	Mus musculus	Kinase suppressor of Ras 2	46.058797	T212;S213;T215	T273;T277;S475;T498	22645316;40885482	MDEENMTKSEEQQPLSLQKALQCC ELVQNMIDLSISNLEGLRTKCAASN DLTQKEIRTLKSLVKYFSRQLSCKK KVALQERNAELDGFPPQLRHWFRIVD VRKEVLEEISPDQLSLEDLLEMTDE QVCETVEKYGANQEECARLNASLSC LRNVHKS GGNLSKQDWIIQWPTTEP GOESNPVCPPEPSPWIRTHLSQSPR VOTKCPQHFCPTSPTPGTPVYTQVD RLTVDAYPNLCPPPPPLESGHRSLLP SPRQRHVVRTPPRTPNIVTTVTPPGT PPMRRNKLNKPPGTPPPSSRKLHLI PGFTALHRSKSHFQLGNRVDEANT PKAKKSKPLNLKIHSVGVGSCENIPA QORSPLLSERSLRSFFVGHGPFPLPST PPVHTEANFSANTLSVPRWSPQIPR RDLGNSIKHRFSTKYWMSQTCTVCG KGMLFGLKCKNCKLKCHNKCTKEA PPCHLLIHRGDPARLVRTESVPCDI NNPVRKPARYSDLHISQTLPKTNKIN KDHIPVPYQPDSSSNPSTTSTPSS PAPPLPPSATPPSPLHPSFQCPRQKK NFNLPASHYYKYKQFIFPDVVPVPE TPTRAPQVILHPVTSNTILEGNPLLQI EVEPTSENEESHNEAEESEDEFEEM NLSLLSARSFPRKASQTSIFLQEWDI PFEQLEIGELIGKGRFGQVYHGRWH GEVAIRLIDIERDNEQLKAFKREVM AYRQTRHENVVLFMGACMSPPHILAI ITSLCKGRTLYSVVRDAKIVLDVNKT RQIAQEIIVKGMGYLHAKGILHKDLK SKNVFYDNGKVITDFGLFSISGLQ AGRRDDKLRIONGWLCHLAPEIIRQ LSPDTEEDKLPFSKHSDFALGTIWI ELHAREWPFKTQPAEAIWQMG TG MKPNLSQIGMGKEISDILLFCWAFE QEERPTFTKLMMDMLEKLPKRNRRLS HPGHFWKSAEL	True	True	5.0	2.505	1.2	0.513	False	5.0	0.874

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