

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
095980	RECK_HUMAN	Homo sapiens	Reversion-inducing cysteine-rich protein with Kazal motifs	17.826004	NaN	NaN	38665916	MATVRASLRGALLLLAVAGVAEVA GGLAPGSAGALCCNHSKDNQMCRD VCEQIFSSKSESRLKHLQRAPDYCP ETMVEIWNCMNSSLPGVFKKSDGW VGLGCCELAIALCROACKQASSKN DISKVCRCKEYENALFSCISRNMESG VCCSYAGHHTNCREYCAIFRTDSS PGPSQIKAVENYCASISPOLIHCVNN YTQSYPMRNPTDSLCCDRAEDHAC QNACKRILMSKKTMEIVDGLIEGC KTQPLPQDPLWQCFLESSQSVHPGV TVHPPSTGLDGAKLHCCSKANTST CRELCTKLYSMWGTQSWQEFDR FCEYNPVEVSMILTCLADVREPCQLG CRNLTYCTNFNNRPTLFRSCNAQS DQGAMNDMKLWEKGSIKMPFINIP VLDIKKCQPEMWKAIACSLQIKPCH SKSRGSIICKSDCVELKKCGDQNKF PEDHTAESICELLSPTDDLKNCIPLD TYLRPSTLGNIVEEVTHPCNPNPCPA NELCEVNRKGCPSGDPCLPYFCVQG CKLGEASDFIVRQGTLIQVPSAGEV GCYKICSCGQSGLENMEMHCIDL QKSCIVGGKRKSHGTSFSDCNVCS CFAGNLVCSTRCLSEHSSEDDRRRT FTGLPCNCADQFVPCGQNGRTPS ACIARCVGLQDHFQFEFGSCMSKDPC NPNPCQKNQRCIPKQVCLTTFDKF GCSQYECVPRQLACDQVQDPVCDTD HMEHNNLCTLYQRGKSLSYKGPCQ PFCRATEPVCGHNGEYSSVCAAYS DRVAVDYGGDCQAVGLSEHSSVAE CASVKCPSLLAAGCKPIPPGACPL CAGMLRVLFDKEKLDTIKVTNKKPI TVLEILQKIRMHVSVPQCDVFGYFSI ESEIVLIIPVDHYPKALQIEACNKEA EKIESLINSDSPTLASHVPLSALIISQ VQVSSVPSAGVRRARPSCHSLLPLS LGLALHLLWTYN	False	True	1.545	2.249	1.734	1.658	1.184	4.669	4.676