	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q04693	RSE1_YEAST	Saccharomyces cerevisiae	Pre-mRNA-splicing factor RSE1	24.475439	NaN	NaN	33229814	MWGGGKMAVVSLSPHTAKMRKLF GQASTTMAYDGLKREAERRTRSDH NITMVAKDDELYLYHLTLKKQTNFV HSCIGHFVDLEAGSKREQSQLCVAT ETHLELYDTADGELKLIAKFQNLFAT ITSMKSLDLPHSGSRAKASNWPTFL ALTSDSGNLSIVQIIMHAGALRLKTL VNQPLTRTTLRRVSPISYMEIDPNGR CIILSSVEQNKLCFLVDYAQKLRISSP LEIIRPHMVTLDMAVVDVNFNNPCF VTLEIDNAATQLSVHLIFYVLELGLN HIVKKADYLVNPSANFVLSLPDLSRY NITTSLSDNNYDADYDTLFNPFVVIG FENHILVKDMNGFFSLKVEIPKRSIT NSRHKNVTIISGIVQKLKNDFFVLLQ SNHGDLFKLTVSPDTNDRNRPLVQL SYFDTIQNSHQLHIFKNGYLFALSE MNNNFLFQFEKLGVEKNDFSNVLT SKDPNKSLVFEPSIKLQNLSILSQQL NLNPSIKSQIVSDSPLSIATKHFTNN KIITLTNAVNYSNLISTSLPPNATKL WLIPDPATTGDNNTLLFITFPKKTMI LQIDNESMEELTPDEATRSAFKLSQ DTTIHTCLMGSHSIIQVCTAELRHIV PTGKSRYSNKLTWVPPAGIRIVCATS SKTQLIISLSNYELVYFKIDVSSDSLIE LTTHPELDTMPSKVAIVQDTQHADL LAIADNEGMIKIMSLKDQKEDFLTVI SLQLVSEKISDMIMVRDSSIGQLNL HVGLENGVYMKFHIGDVDGSFTDIK RRFLGLKPVSLSYLREISVSLNNEEE EEEEEDDDDEKEEEEINSSGAKWM SCVVCHSSSTWVSYTWKNVWTIRQL KDQNMLSCSKFVNADVAINGVCSIS SSGRLNIGRVSNFPTLDNWFHVHES SVNKQENGGGDESNEEEEDEMEEE MEMLQISTFRPRTILSFPNNPKSILFI DNHSGKKQCRISLQIDGECLKFGSS DHLYKILDDIDCVSAAIIDFTRQADH LIICAGDKRLLTYKILVNKDKLSFDIE LLHQTEIISPIHAMLKFKNFLLTAMG STIVLYGLGKKQLLRRSVTQTPVSITK IVSMHQWNYERLAVGDIHESVTLFI WDPAGNVFIPYVDDSVKRHVTVLKF LDEATVIGADRYGNAWTLRSPPECE KIMSNHDPSELSNGAIKYPLDVITLQ QKLPNTYDCKFKFQLLNHFFVNDIIT DFHILDSLSNSDRPGCIYMGLQGTV GCFIPLLSKGNVFMMGNIENIMAEA DDTFYLDYESRKKNNNMRKEDDEE ESGSVVLQGRHGIEDEIICEGSCSIL GRDHQEYRSYYAPVRKVIDGDLCEN FLRLSLNEQEFLAKNLKSVQVEDIIQ TINEVRTNYM	None	None	None	None	None	None	None	None	None