

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
B2RR83	YTDC2_MOUSE	Mus musculus	3'-5' RNA helicase YTHDC2	25.685855	NaN	S1104;S1105;S1107;S1278;S1282;S1296	39627609	MSRPSSVSPRPPAPSGGGTGGGGGG SGGGGGGGGGPASCPCGGGGRAK GLKDIRIDEEVKIADVNIALERFRYGD QREMEFPSSLTSTERAFIHRLSQSLG LVSKSKGKGANRYLTVKKGKDGSETA HAMMTCNLTHNTKHAVRSLIQRFP VTNKERTELLPKTERGNVFAVEAEN REMSKTSGRLLNNGIPIQVPVVRGESE FDSFRQSLPVFEKQEEIVKIKENKV VLIVGETGSGKTTQIPQFLDDCFKN GIPCRIFCTQPRRLAAIAVAERVAER RERIGQTIGYQIRLESRVSPKLLTFC TNGVLLRRLMAGDSTLSTVTHVIIVD EVHERDRFSDFLTKLRDLLQKHPT LKLILSSAALDVNLFIRYFGSCPVIYI QGRPFVVKEMFLEDILRTTGYTNKE MLKYKKEKQREEKQQTTLTEWYSA QENTFKPESQRQRAVASVSEEDLL DDGGDAVFSQLTEKDVNACLEPWLK EMDAACLSDIWLHKDVFDAFAQVFHLI LTENVSDYRHSSETSATALMVAAGR GFTSQVEQLISMGANVHASKASNGW MALDWAKHFGQTEIVDLLESYSASL EFGNLDSESSLVQTNNGNLSAEDREL LKAYHHSFDDKVDLDLIMHLLYNI CHSCDAGAILIFLPGYDEIVGLRDRIL FDDKRFADNTHRYQVFMHLSNMOT SDQKKVLKNPPAGVRKIILSTNIAET SITVNDVVFVIDSGKVKESFDALNF VTMLKMVWISKASAIQRKGRAGRGR PGICFRLFSRLRFQNMLEFQTPPELL RMPLQELCLHTKLLAPVNCTIADFL MKAPEPPALIVRNAVQMLKTIDAM DAWEDLTELGYLADLPVEPHLGK MVLCAVVLKCLDPILTIACLTAYRDP FVLPQTQASQKRAAMLRCRKRFTAGTF SDHMALLRAFQAWQKARSDGWER FCEKNFLSQATMEIIGMRTQLLGQL RASGFVRARGGDIRDVNTNSENW AVVKAALVAGMYPNLVHVDRENVIL TGPKEKKVRFHPTSVLSQPQYKIP ANGQAAIQALPTDWLIYDEMTRAH RIANIRCCSAVTPVTVLVFCGPARLA SNALQEPSSFRADGIPNDSSDSEME DRTTANLAALKLDEWLNFKLEPEAA SLLLQLRQKWHSLFLRRMRAPSKP WSQVDEATIRAHAVLSTEEQSAGLQ QPSGIGQRPRPMSSEELPLASSWRS NNSRKSTADTEFADGSTTGERVLMK SPSPALHPPQKYKDRGILHPRSTDD RSDQSSVKSTDSSSYSPCASPSPPS SGKGSKSPSRPNMIRYFIMKSSNL RNLEISQQGIWSTTPSNERKLNRA FWESSMVYLVFSVQSGHFQGFGR MSSEIGREKSDWGSAGLGGVFKV EWIRKESLPFQFAHLLNPWVNDNK KVQISRQGELEPQVGEQLLQLWER LPLGEKTTSD	False	False	2.359	3.745	2.078	2.682	False	1.559	1.745