

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
Q8BYA0	TBCD_MOUSE	Mus musculus	Tubulin-specific chaperone D	25.685855	NaN	NaN	39627609	MVLSNEPAASAAEEVEDDALVRAS ALEAFGESAE TRALLRSLPAVHRERA SREVAEERFRVIMDKYQEQPHLLDP HLEWMMNSLLDLVQDETSPLDLVH LAFKFLYIITKVRGYKVFRLRFPHEVA NVQPVLDMFTGQNPKDHWETRY MLLLWLSVTCLIPDFSRDLGNLST QTGETRVPTMDRILQIAESYLVVSDK ARDAAAVLVSKFITRPDVKQRKMAS FLDWSLCTLAHSSFQTIEGVITMDG MLQALAQIFKHGKREDCLPYANTVL QCLDGCRLPESSHTSLRKLGVKLVQ RLGLTFLKPKVATWRYQRGCRSLAA NLKLCAPGKSDQKLLSDSLTSDGDE DYDVPEGVETVIEQLLVGLKDKDTVV RWSAAKGIGRMAGRLPRELADDVVG SVLDCFSFQETDKAWHGGCLALAE GRRGLLPSRLEVVTVILKALTYDE KRGACSVGANVRDAACYVCWAFAR AYEPQELTPFVTAISSALVIAAVFDRN VNCRRASAAAFQENVGRQGTFFPHGI DILTTADYFAVGNISNCFIISVFIAG FQEYTKPMIDHLVSMKINHWDGAI ELSAKALHNLTPQVPEYIAMHVFP LLLMTQSPDLHTRHGAILACAEV ALYKLATQSNRLVTDYLDEKAVQSL KQIHQQLCDRHLRYRGLGGELMRQA VCILIEKLSLRMPFKGDATVEGWQ WLINDTLRSLHLVSSHRSRQIQEVA VSALTALCSEYYVKEPGEAGSSIAKE LIPQYLAELQSPEEMARCGFSSALGA LPGFLLRGHLQQVLSGLRRVTCISPN DVSFAEARRDGLKAIIRICQTVGVNT RGPPDEVICKENISEVYAALLGCM SDYTDSRVDVGAWVREAAAMTSLMDL MLLLARTEPVLIEAHICERVMCCVA QQASEKIDRFRAHAARVFLTLHFD SPPIPHVPHRQELSLFPRSDVATVN WNAPSQAFLITQLLGLPTRYHVLL GLAVSVGGTTESTVRHSTQSLFEYM KGIQKDAQVLQSFSETLLKVFEDNLL NDRVSVSLKMLDQLLANGCFDIFT AEENHPFCVKLLTLCKEIKKSKDIQ KLRSSIAVLGCMVQFNGDVRKKILL QLFLLGHFPFVIRKSTASQVYEMVL TYSDLVDAEVLDEVMSVLSDTAWDA ELPVVREQRNRLCDLLGVPRPQLVP KPIPGS	False	True	2.293	1.147	0.982	0.628	1.081	4.642	0.87