

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
B2RXC1	TPC11_MOUSE	Mus musculus	Trafficking protein particle complex subunit 11	25.161187	NaN	NaN	34887587	MSPTQWDFPVELCCRPMAFVTLTG LDVVYNVHRAVWDAFCANRRADR VPISFKVLPGDHEYPKCRSKRTSYEW YIPKGVLTGWMNKHLNLVLPALVVV FYELDWDEPQWKEKQSECATRVEIV RQSLQGRNTKVAVVLIQKKTPLPPGE DVIASERAAAALCNVCELSGKSLFVLP HTDHLVGYIIRLENAFYEAHQTYYYT EIRRVKSHKEFLNKTTHQLLFVRHQ FKIAFFSELKQDTQNALKNYRIAYNL VHELRAHETNILEIKTMAGFINYKIC RLCFQHNTPLDAIAQFRKHIDLCKK KIGSAELAFEHAAWMAKQFQAFGDL FDEAIKLGTLAIQTQNPGFYQQAAY YAQERKQHAKALCNHDAAVMYPNP DPLETQSGVLDYFGQRPWRQGILSF DLSDPKEKAGILAIQLKERSVVHSE IIALLSNAVAQFKKYKCPRMKSHLM VQMGEYYYYAKDYTKALKLLDYVMC DYRSEAWWLLTSILTALKCSYLM AQLKDYITYSLELLGRASTLKDEQKS RIEKNLMNVLNMNESPDPEPDCDVL AVKTAQKLWADRVSAGSNVFIQV QDFVFPVQCKAKFHAPSFHVDVPVE FDVFLKADCPHPIRFSKLCVSNQ VYNQFCVLEEASKASEVLENLTQGK MCLVPGKTRKLSFKFVAKTEDVGKK IEITSVDLFLGSESGRCVLSWQGGG GDAASSQEALQAARSFKRRPKLAED EIHWDSVHQASTMIISRVPNISVHLR HEPPALMNEMYCLVVTVQSHEKSPI RDVKLTAGLKPQDANLTQKTHVTL HGAELCDESPALLTDIPVGD LHPGE QLEKTVYVRCGTVGSRMFLVYVSYLI NTTVEGKEIICKCHKDDTVTIETVFP FDVAVRFVSTKFEHLERVYADIPFLL MTDVLSPWALTVSSELQLAPSM TAMDHLESQIDKVVLTGESASECF CLRCPSAGNIEGGVATGHYIISWKRA SVVESIPAVSTVITLPHVIAENIPLHV NADLPSFGRVRESLPVRYHLQNKTD LVQDVEISVEPSDAFMFSGLKKQIRLR ILPGTKQEMLYNFYPLMAGYQQLPS LNINLLRFPNFTNQLLRRFIPTSIFVK PQGRLEDTSIAAA	None	None	None	None	None	None	None		

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