

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8BFY9	TNPO1_MOUSE	Mus musculus	Transportin-1	25.685855	NaN	NaN	39627609	MVWDRQTKMEYEWKPDDEQGLQOIL QLLKEQSPDITTIQRTVQQKLEQLN QYPDFNNYLIFVLTCLKSEDEPTRSL SGLILKNNVKAHFQNFNPGVTDFFIK SECLNNIGDSSPLIRATVGLITTIASK GELQNWPDLLPKLCSLLDSEYNTC EGAFGALQKICEDSAEILSDVLDLDRP LNIMIPKFLQFFKHSSPKIRSHAVAC VNQFIISRTQALMLHIDSFIEMLFAL AGDEEAQVHRKNCVCRALVMLLEVRM DRLLPHMHNIVEYMLQRTQDQDEN VALEACEFWLTLAEQPICKDVLVRH LPKLIPLVNGMKYSDIDHLLKGDVE EDETIPDSEQDIRPRFHRRTVAQQ HEEDGIEEEDDDDEIDDDDTISDW NLRKCSAAALDVLANVYRDELLPHIL PLLKELLFHHEWVVKESGILVLGAIA EGCMQGMIPYLPPELPHLIQCLSDKK ALVRSITCWTLSRYAHWVVSQPPDT YLKPLMTELLKRILDSNKRQVQEAACS AFATLEEEACTELVPYLAYILDVLFVA FSKYQHKNLLILYDAIGTLADSVGHH LNKPEYIQLMPLLIQKWNMLKDE DKDLFPLECLSSVATALQSGFLPYC EPVYQRCVNLVQKTLAQAMLNNAQ PEQYEAQDKDFMIVALDLLSGLAEG LGGNIEQLVARSNLTLMYQCMQDK MPEVROSSFALLGDLTKACFQHVVP CIADFMPILGTLNLPFISVCNNAT WAIGEISIQMGIEMQPYIPMVLHQLV EIIINRPNTPKTLENTAITIGRLGYVC PQEVAPMLQQFIRPWCTSLRNIRDN EEKDSAFRGICTMISVNPQVQDFI FFCDAVASWINPKDDLDRDMFCILH GFKNQVGDENWRRFSDQFPLPLKE RLAAFYGV	True	False	3.597	4.248	1.763	1.46	1.274	1.589	1.34