

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q9UPV9	TRAK1_HUMAN	Homo sapiens	Trafficking kinesin-binding protein 1	47.251593	S447;S448;S452;S560;S563;S565;S566;S680;S719;S829;S830;S903;T935;S938	S537;S719;S919	34019948;28657654;39302247;24995978;30620550;40136647;12724313	MALVVFQFGQPVRAQPLPGLCHGKLI RTNACDVCNSTDLPEVEIISLLEEQL PHYKLRADTIYGYDHDWLHTPLISP DANIDLITTEQIEETLKYFLLCAERVG QMTKTYNDIDAVTRLLEEKERDLEL AARIGQSLLKKNKTLTERNELLEEQ VEHIREEVSQLRHELKSMKDELLOFY TSAAESEPEPESVCSTPLKRNESSSV QNYFHLDLQKLDLEENVVLRS EASQLKTETITTYEEKEQLVNDVCVK ELRDANVQIASISEELAKKTEDAARQ QEEITHLLSQIVDLQKKAKACAVENE ELVQHLGAAKDAQQLTAELELED KYAECMEMLHEAQEELKNLRNKTM PNTTSRRYHSLGLFPMDSLAAEIEGT MRKELQLEEAESPDIHQKRVFETV RNINQVVKQRSLTPSPMNIPGSNQS SAMNSLLSSCVSTPRSSFYGSDIGNV VLDNKTNSIILETEAADLGNDRSKK PGTPGTPGSHDLETALRRLSLRREN YLSERRFFEEEQERKLQELAEKQEL RSGSLTPTESIMSLGTHSRFSEFTGF SGMSFSSRSYLPEKQIVKPLEGSAT LHHWQQLAOPHLGGILDPRPGVVTK GFRTLVDLDEVYCLNDFEEDDTGD HISLPRLATSTPVQHPETSAHHPGKC MSQTNSTFTFTTCRILHPSDELTRVT PSLNSAPTACGSTSHLKSTPVATPC TPRRLSLAESFTNTRESTTTMSTSLG LVWLLKERGISAAVYDPQSWDRAGR GSLLSYTPKMAVIPSTPPNSPMQT PTSSPPSFEFKCTSPPYDNFLASKPA SSILREVREKNVRSSESQTDVSVSNL NLVDKVRRFVAKVNSGRAHVPTL TEEQGPLLCGPPGAPALVPRGLVPE GLPLRCPTVTSAGGLQLNSGIRNR SFPTMVGSSMQMKAPVTLTSGILMG AKLSKQTSLR	True	False	4.48	4.263	5.0	False