

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9CRT8	XPOT_MOUSE	Mus musculus	Exportin-T	25.685855	NaN	NaN	39627609	MDEQALLGLNPNADSDFRQRALAY FEQLKISPDAWQVCAEALAQKTYSD DHVKFFCFQVLEHQVKYKSELSTA QQQLIRETLLSWLQAQMNPQPEK TFIRNKAQV FALLFVTEYLTKWPKF FFDILSVVDLNPGRVDLYLRILMAID SELVDRD VVHTSEASGLENTLIKDT MREQCIPNLVESWYQILHNYQYTNS EVLCOCLEVVGAYVSWIDLSLIANDR FINMLLGHMSVEVLREEACDCLFEI VNKGM DPV DKMKLVESL CQVLQTA GFFSIDQEEDLDFVARF SKLVNMG QSLIVSWTKLIKNGAVKNAQEALAI ETKVPLMLQLLVHEDDDISSNIIGFC YDYLHILKQLPVLSDQKQANVEAIML AVMKKLT YDEEYNFENEGEDEAMF VEYRKQLKLLDRLAQVSP ELVLASV RRVFSATLQNWQTTRFMEVEVAVR LLYMLAEALPVSHGAHFGSDVSKAS ALQDMMRTLVTSGVSSYQHTSVTLE FFETVVRYEKFFTVEPQHIPCVM LAF LDHRGLWHSSAKVRSRTAYLFSRFV KSLNKQMNPIEIEILNRIQDLLALSP PENGYQSLSSDDQLFIYETAGALIV NSEYPAENKQALMKDLLTPLMERF KVLLEKLMMAQDEERQASLADSLN HAVGFASRTSKAFSNKQTVKQCGCS QVYLDCLQTFLPALSCPLQKDV LRS GVRTFLHRMIICLEEEVLPFIPSASE HMLKDCEAKDLQEFIPLINQITAKFK MQVSPFLQQMFMP LLHAIFEVLLRP AEDNDQSAALEKQMLRRSYFAFLQT VTGSGMSEVIANQGAENVEQVLVTII QGAVDYPDPIAQKTCFIILSKLVELW GGKDG PVGFADFVYKHIVPACFLAPL KQTFDLADAQTVLALSEC AVTLKTIH LKRGPCEVQYLQEQEYLP SLQVAP EII QEFQALQ QPDAKVFKNYLVFFQR AKP	True	False	3.563	4.095	1.244	0.954	False	0.858	0.535