

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	extrac region
043592	XPOT_HUMAN	Homo sapiens	Exportin-T	8.925602	NaN	NaN	33214551;23301498;34019948;34725712;28510447;38665916	MDEQALLGLNPNADSDFRQRALAY FEQLKISPDAWQVCAEALAQRTYS DHVKFFCFQVLEHQVKYKSELTTV QQQLIRETLISWLQAQMLNPQPEKT FIRNKAQVFFALLFVTEYLTWKWPKFF FDILSVVDLNPARGVDLYLRILMAIDS ELVDRDVVHTSEEARNTLIKDTMR EQCIPNLVESWYQILQNYQFTNSEV TCQCLEVVGAYVSWIDLSLIANDRFI NMLLGHMSIEVLREEACDCLFEVV NKGMDPVDKMKLVESLCQVLQSAG FFSIDQEEVDVFLARFSAKLVNMGMO SLIVSWSKLIKNGDIKNAQALQAIIE TKVALMLQLLIHEDDDISSNIIGFCY DYLHILKQLTVLSDQQKANVEAIML AVMKKLYDEEYNFENEGEDEAMF VEYRKQLKLLDRLAQVPELLASV RRVFSSTLQNWQTTRFMEVEVAIRL LYMLAEALPVSHGAHFSGDVS KASA LQDMMRTLVTSGVSSYQHTSVTLEF FETVVRYEKFFTVPEQHIPCVLMAFL DHRGLRHSSAKVRSRTAYLFSRFVK SLNKQMPFIEDILNRIQDLELSPP ENGHQSLSSDDQLFIYETAGVLIVN SEYPAERKQALMRNLLTPLMEKFKI LLEKLMQAQDEERQASLADCLNHAV GFASRTSKAFSNKQTVKQCGCSEVY LDCLQTFPALSCPLQKDILRSGVRT FLHRMIICLEEEVLPFIPSASEHMLK DCEAKDLQEFILINQITAKFKIQVSP FLQQMFPLLHAI FVLLRPAEEND QSAALEKQMLRRSYFAFLQTVTGSG MSEVIANQGAENVERVLVTVIQGAV EYDPPIAQKTCFIILSKLVELWGGKD GPVGFADVYKHIVPACFLAPLKQTF DLADAQTVLALSECAVTLKTIHLKRG PECVQYLQQEYLPQLQVAPEIIQEF QALQPDARVFNKYLKVVFFQRAKP	True	False	4.683	5.0	1.358	1.025	0.51	1.638	0.752