

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	ext
Q96QU8	XPO6_HUMAN	Homo sapiens	Exportin-6	24.685143	NaN	S199;T201;T204;S208;S224	40914422;34019948;37217939	MASEEASLRALLESMTTEFFHDCTTN ERKREIEELLNFAQQIGAWRFCLY FLSSTRNDYVMMYSLTVFENLINKM WLGVPSPQDKMEIRSLPKLLLAHHK TLPYFIRNKLCKVIVDIGRQDWP MFY HDFFTNQLIQSPVTTPLGLIMLKTT SEELACPREDLVSARKEELRKL LLD QVQTVLGLLTGILETVWDKHSVTAA TPPPSPTSGESGDLLSNLQSPSSAK LLNQPIILDVESEYICSLALECLAHL FSWIPLSASITP SLLTTFHFARFGCD IRARKMASVNGSSQNCVSGQERGL GVLAMSCINELMSKNCVPMEFEEY LLRMFQQTFYLLQKITKDNNAHTVK SRLEELDESYIEKFTDFLRLFVSVHL RRIESYSQFPVVEFLTL LFKYTFHQ THEGYFSCLDIWTFLFDYLT SKIKSR LGDKEAVLNRYEDALVLLLEVLNRI QFRYNAQOLEELDDDETLDDDOQTE WQRYLRQSLEVVAKVMELLPTHAFS TLFPVLQDNLEVYGLQQFIVTSGSG HRLNITAENDCRR LHCSRDLSSLL QAVGRLAEYFIGDVFAARFNDALT VERLVKVTLYGSQIKLYNIETAVPSVL KPDLDIVHAQSLAALQAYSHWLAQY CSEVHRQNTQQFVTLISTTMDAITPL ISTKVQDKLLSACHLLVSLATTVRP VFLISIPAVQKVFNRITDASALRLVDK AQVLVCRALSNILLPWP NLPENEQ QWPVRSINHASLISALSRDYRNLKPS AVAPQRKMP LDDTKLIHQTLSVLED IVENISGESTKSRQICYQLQESVQV SLALFPAFIHQSDVTDEMLSFFLTLF RGLRVQMGVPFTEQIIQTFLNMFTR EQLAESILHEGSTGCRVVEKFLKIQ VVVQEPGQVFKPFLPSIALCMEQVY PIAERPSPDVKAELFELLFRTLHHN WRYFFKSTVLSVQRGIAEEQME NE PQFSAIMQAFGQSFLQPDIIHLFKQN LFYLETLN TKQKLYHKKIFRTAMLFO FVNVLLQVLVHKSHDLLQEEIGIAIY NMA SVDFDGFFAAFLEPFLTSCDGV DANQKSVLGRNFKMDRDLPSFTQN VHRLVNDLRYRLCNDLSLPPGTVKL	True	True	4.603	4.667	1.027	0.882	0.997	4.464	0.8