

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
Q8VI75	IPO4_MOUSE	Mus musculus	Importin-4	28.4037	S99;T106	NaN	39627609;36852467;33300544	MEPAGLEQILKELLPLDTERIRRATE QLQTLIRDPALPALFDLLATATDSQI RQFAAVLTRRRLLNRRWRLAPEQR ESLKSLLVLTALQKETVHVSVSLAQL SATIFRKEGLQGWPFMNLLOHST HSSHSPKEKVGLLLLSVVSSQPEAF HAHQHELLQLNETLSDVSPFQVLF YSLRTLTAIARYVRPDDVSLARMLVP KVVTALRTLPLDEVKACEALEALDE MLETELPIINPHLSEVLTFCLEAVKN VALGEPLRVRVLCCLTFLVKVSKAL LKNRVPPLLHALFPLMAAEPMPGQ LDPEDQSDDDDDLEIGLMGETPKHF AVQVVDMLALHLPPEKLCPHVMPM LEALRSEDYPYQRKAGFLVAVLSD GAGDHIRORLLYPLLQIVCKGLDDPS QIVRNAALFALGQFSENLPQHISSYS EEVMPLLLSTYKSVPMGNTHHLAKA CYALENFVENLGPVQPYLPELMEC MLQPLKNPSKARTKELAVSAIGAIAT AAQDSSLPYFPTIMDLLREFLTGHG DFHLVQIQSLETGLVLRALGESMK PLAEECCQLGLGLCIHIDDPDVRCT YSLFAALSGLMGEGLPYLPQITLTM LLSLRSTEGIVPQYDGISSFLFDSDS EAESEELMDEDMEEEGDDSEISGY SVENAFFDEKEDTCTALGEISMNTC VAFLPFMDATFDEVYKLLCEPHMNV RKSAYEALGQFCALHKASORSSSD PSSSPVLQTSLARVMFAYMQAVKVE RERPVMVAVLESLTGLVTRTCGSLAL QPPGRLELCNVLKAVLQKKTACQD AEEDDEDDQAEYDAMLLEHAGE AIPVLAATAGGHAFAPFFATFLPLLL CKTKQSQCTVAEKSFVGTLEAESIQL GTASAQFVSRFPVLLNNAREADPE VRSNAIFGLGVLAHGGCPAQDHFP KLLGLLLPLLARERHDRVRDNCGAL ARVLMASPVGKTEPQVLATLLRALPL KEDMEEWLTIGHLFSFLHQNNPEQ VVDVASELLRICSLILPDNRIPPTKA ALLLLLTLAKOHTDSFHTALGSLPN DKAQELQAMMGLT	True	False	2.511	4.078	1.395	0.771	False	0.726	0.619