

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
Q7TMY7	IPO8_MOUSE	Mus musculus	Importin-8	25.685855	NaN	S902;S903	39627609	MDLNRIIQALKGTIDPKLRIAAETEL NQSYKIINFAPSLRLRIIVSDHVEFPVR QAAAAYLKNMVTQYWPDRPPPPGEV IFPFNIHENDRQQIRDNIVEGHIRSPD LVRVQLTMCLRVIIHRHDFPGHWPAV VDKIDYLLQSPNSGSLGSLCLYQL VKTYEYKKAEREPLLAAMQIFLPRI QQQILQLLPDASHYSVLLQKQILKIF YALVQYALPLQLVNHQMTTWTWMEIF RTIIDRTVPPETLQIDEDDRPELVVW KCKKVALHIVARLFERYGSPGNVTK EYFEFSEFFLKYAVGIQQVLLKILD QYRQKEYIAPRVLQAFNYLNQGVV HAVTWKQMKPHIQNISEDVIFSVMC YKDEDEELWQEDPYEYIRMKFDIFE DYASPTTAAQTLLYTAAKKRKEVLPK MMAFCYQILTDPNFDPKRDGALH VIGSLAEILLKSLFKDQIELFLQNH VFPLLMSNLGYLRARSCWVLFHAFSS LKFHNELNLRNAVELAKKSLIEDEE MPVKVEAALALQSLISNQAQAKEHM KPYVRFIMQELLHIVRETENDDDVTN VIQKLICEYSQDVASIAVDTTQHLAEI FGKVLQSDYEYEEIEDKTMAMGILH TIDTILTVEDHPHIIQLENICLRIID LVLQKHVIEFYEEILSLAYNLTCHTIS PQMWQLLGILYEVFQDQCFEYFTD MMPLLHNYVTVDTNALLSNPKHLE VLFTMCRKVLGCEAGEDAECYAAKL LEVHILQCKGRGIDQCIPLFIQLVLER LTRGVKTSSELRTMCLQVAIAALYSP ELLFHTLEQVQLPHNPGPVTSSQFIN QWMNDTDYFLGHHRKMCIGLSV LLELQNRPPAVDAVAQILPSILFLFL GLKQVQCATRQTVNRENHKAQKVDI EENEISSEEEEEETSVSQAQAMQSQI GRSEEDDDWDEEVLEETALEGFS TPLDLDNSVDEYQFFTQALLTVQNR DAAWYQLLVAPLSEDDQKRKLQEVYT LAEHRRTLAAGQFHI	True	False	2.884	4.15	1.853	1.238	0.877	4.298	1.49