

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	ex
Q9Y5L0	TNPO3_HUMAN	Homo sapiens	Transportin-3	22.926716	NaN	S74;T896	33214551;34725712;40914422;35132862;34019948;28510447	MEGAKPTLQLVYQAVQALYHDPDPS GKERASFWLGELOQRSVHAWAISDQL LQIQDVESECYFAAQTMMKIQTSF YELPTDSHASLRDSSLTHIQNLKDL PVIVTQLALAIADLALQMPMSWKGCV QTLVEKYSNDVTSPLPFLLEILTVLPE EVHSRSLRIGANRRTEIIEDLAFYSST VVSLLMTCVEKAGTDEKMLMKVFR CLGSWFNLGVLDSNFMANNKLLAL LFEVLQDDKTSSNLHEAASDCVCSA LYAIENVETNPLAMQLFQGVLTLET AYHMAVAREDLKVLNYCRIFTELC ETFLEKIVCTPGQGLGDLRTRLELLI CAGHPQYEVVEISFNFWYRLGEHLI KTNDEVIHGIFKAYIQRLLHALARHC QLEPDHEGVPEETDDFGEFRMRVS DLVKDLIFLIGSMCEFAQLYSTLKEG NPPWEVTEAVLFIMAAIAKSVDPEN NPTLVEVLEGVVRLPETVHTAVRYTS IELVGEMSEVVDNRNPQFLDPVLGYL MKGLCEKPLASAAAKAIHNICSVCR DHMAQHFNGLLEIARSLSFLLSPE AAVGLLKGTAIVLRLPLDKITECLS ELCSVQVMALKKLLSQEPSNGISSD PTVFLDRLAVIFRHTNPIVENGTQHP CQKVIQEIWPVLSLTKHRADNRIV ERCCRCLRFVRCVCGKSAALLQPL VTQMNVVYVHVHQSFLYLGSLVD EYGMEEGCRQGLDMLQALCIPTFQ LLEQQNGLQNHPTVDDLFRLATRF IQRSPVTLRSQVVIPLQWAIASSTL DHRDANC SVMRFLRDLIHTGVAND HEEDFELRKEIGQVMNQLGQQLVS QLLHTCCFCLPPYTLDPVAEVLWEI MQVDRPTFCRWLENSLKGPKETTV GAVTVTHKQLTDFHKQVTSAECKQ VCWALRDFTRLFR	True	False	4.518	5.0	1.649	0.971	0.604	1.536	1.5