

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasm mem
Q76FK4	NOL8_HUMAN	Homo sapiens	Nucleolar protein 8	26.689553	T1065	S268;S298;T302;S304;S331;S365;T376;S378;T381;S432;S723;T795;S801;S837;S838;S843;S845;T888;S890;S1036;S1082;S1083;S1084;S1099	28411811;23301498;29351928	MKVNRETKRLYVGGLSQDISEADLQ NQFSRFGVEVSDVEIITRKDDQGNPQ KVFAYINISVAEADLKKCMSVLNKT WKGGLTQIQLAKEFLHRLAQEREA AKAKKEESTTGNANLLEKTTGGVDFH MKAVPGTEVPGHKNWVVSKFGRVL PVLHLKNQHKRRIKYDPSKYCHNL KKIGEDFSNTIPISSLTWELEGNDP MSKRRRGEFSDFHGPPKIIKVKQD ESSTGSLAMSTRPRRVIERPPLTQQQ AAQKRTCDSITPSKSSPVPVSDTQKL KNLFPFKTSGLEAKKRNSISDDDDTD SEDELRRMMIAKEENLQRTTQPSINE SESDPFVVRDDFKSGVHKLHSLIG LGIKNRVSDHSDDDIMRNDREYDS GDTDEIHAMKKNVAKVKNSTEFSSQM EKSTKKTSTFKNRENCESLSDHCIKLQ KRKSNVESALSHGLKSLNRKSPSHS SSSEDADSAELADSEGGEYNYAM MKNCLRVNLTADLEQLAGSDLKVP NEDTKSDGPETTTQCKFDRGSKSPK TPTGLRRGRQIRPAEIVASLLEGEE NTCGKQPKENNLKPKFQAFKGVG CLYEKESMKSLLKDSVASNNKQDN SMKHEDPSIISMEDGSPYVNGSLGE VTPCQHAKKANGPNYIQPKRQTTT ESQDRKAVSPSSSEKRSKNPISRPLE GKKSLSLSAKTHNIGFDKDSCHSTT KTEASQEERSDSSGLTSLKSPKVSS KDTREIKTDFSLISNSSDVSAKDKH AEDNEKRLAALAEARQAKEVQKLV HNALANLDGHPEDKPTHIIIFGSDSE CETEETSTQEQSHPGEEVVKESMG KTSGLKFDSSDDESDSEDDSNRFX IKPQFEGRAGQKLMDLQSHFGTDDR FRMDSRFLTDSEEEQEEVNEKTA EEEELAEEKKKALNVVQSVLQINLS NSTNRGSAAKKFKDIIHYDPTKQD HATYERKRDDKPKESKAKRKKKREE AEKLEPVSKEMYNIAMDLKEIFQTT KYTSEKEEGTPWNEDCGKEKPEIQ DPAALTSDAEQPSGFTSFSDSDTK DIKEETYRVETVKPGKIVWQEDPRLQ DSSSEEDVTEEDHRNSSPGEASL LEKETTRFFFFSKNDRLQGSDFW RGVGSNMSRNSWEARTTNLRMDC RKKHKDAKRMKPK	None	None	None	None	None	None	None	

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