

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	extra regio
Q92499	DDX1_HUMAN	Homo sapiens	ATP-dependent RNA helicase DDX1	18.746291	NaN	S481	34229054;30397120;23301498;37217939;33214551;16408927;38665916;21740066;34725712;32119511;40914422;34019948;35138101;28510447	MAAFSEMGMPEIAQAVEEMDWLLPTDIAESIPLILGGGDVLMMAETGS GKTGAFSIPVIQIVYETLKDQEGKK GKTTIKTGASVLNKGWQMNPDYRGS FAIGSDGLCCQSREVKWEGCRATK GLMKGKHYYEVSCHDQGLCRVWGS TMQASLDLGTDFGFGGGTGKKS HNKQFDNYGEEFTMHDITIGCYLDID KGVKFSKNGKDLGLAFEIPPHMK NQALFPACVLKNAELKFNFGEEEFK FPPKDGVALSKAPDGYIVKSQHS NAQVTQTKFLPNAPKALIVEPSRELA EQLNNIKQFKKYIDNPKLRELLIG GVAARDQLSVLENGVDIVVGTGRL DDLVTGKLNLSQVRFVLDEADGL LSQGYSDFINRMHNQIPQVTS DGKR LQVIVCSATLHSDVKKLSEKIMHFP TWVDLKGEDSVPTVHHVVVPVNP KTDRLWERLGS SHIRTDDVHAKDNT RPGANSPBMWSEAIKILKGEYAVRAI KEHKMDQAIIFCRTKIDCDNLEQYFI QGGGPDKKGHQFSCVCLHGD RKP HERKQNLERFKKGDVRFICTDVAA RGIDIHGVPYVINVTLPDEKQNYVHR IGRVGRAERMGLAISLVATEKEKVV YHVCSSRGKGCYNTRLKEDGGCTIW YNEMQLLSEIEEHLNCTISQVEPDIK VPVDEFDGVYGGKRAAGGGSYKG HVDILAPTQVELAALEKEAQTSLHL GYLPNQLFRTF	True	False	4.764	5.0	4.349	1.581	1.183	1.506	1.458