

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extr regi
Q7Z478	DHX29_HUMAN	Homo sapiens	ATP-dependent RNA helicase DHX29	13.560748	T1159	S71;S192;S200	28510447;34229054;38665916;30379171;39302247	MGGKNNKHKAPAAAVVRAAVSASR AKSAEAGIAGEAQSCKPVSRRPATAAA AAAGSREPRVKQGPKIYFNFSTNDS SGPANLDKSLKVVINNKLQRIIGVI NEHKKQNNDKGMISGRLLTAKKLQD LYMALQAFSFKTKDIEDAMTNTLLY GGDLHSALDWLCLNLSDDALPEGFS QEFEEQPKSRPKFQSPQIQATISPP LQPKTKTYEEDPKSKPKKEEKNMEV NMKEWILRYAEQQNEEKNENSKS LEEEEFDPNERYLHLAAKLLDAKE QAATFKLEKNKQGQKEAQEKIRKFKQ REMETLEDHPVFNPMKISHQQNE RKKPPVATEGESALNFNLFKESAAA TEEEKDKKKEPHDVRNFDYTARSWT GKSPKQFLIDWVRKNLPKSPNPSFE KVPVGRYWKCRVVRVKSIEDDLVVC PTILTEDGMQAQHLGATLALYRLVK GQSVHQLLPPTYRDVWLEWSDAEK KREELNKMETNKRDLFIKLLNKL KQQQQQQQHSENKRENSEDPEES WENLVSDEDFSALESANVEDLEP VRNLFRLQSTPKYQKLLKERQQLP VFKHRDSIVETLKRHRVVVAGETG SGKSTQVPHFLEDDLLEWEASKC NIVCTQRRISAVSLANRVCDELGCE NGPGGRNSLCGYQIRMESRACESTR LLYCTTGVLRLKLOEDGLLSNVSHVI VDEVHERSVQSDFLLIILKEILQKRS DLHLILMSATVDSEKFTYFTHCPIL RISGRSYPVEVFHLEDIIEETGFVLEK DSEYQCQFLEEEETINVTSKAGGI KKYQEIYVQTAHADLNPFYQKYSS RTQHAILYMNPBKINLDLILELLAYL DKSPQFRNIEGAVLIFLPLGLAHIQQL YDLLSNDRRFYSERYKVIALHSILST QDQAAAFTLPPPQVVRKIVLATNIAET GITIPDVVVIDTGRTKENKYHESQ MSSLVETVFSKASALQRQGRAGRVR DGFCEFRMYTRERFEGFMDYSVPEIL RVPLEELCLHIMKCNLGSPEDFLSK ALDPPQLQVISNAMNLLRKIGACEL NEPKLTPLGQHLAALPVNVKIGKMLI FGAIFGCLDPVATLAAVMTEKSPFTT PIGRKDEADLAKSALAMADSDHLTIY NAYLGWKKARQEGGYRSEITYCRRN FLNRTSLLTLEDVKQELIKLVKAAGF SSSTTSWEGNRASQTLFSQEIALL KAVLVAGLYDNVGIYTKSVVDVTEK LACIVETAQGKAQVHPSSVNRDLQT HGWLlyQEIKIRYARVYLRETTLITPF PVLLFGGDIEVQHRERLLSIDGWYIF QAPVKIAVIFKQLRVLIDSVLRKLE NPKMSLENDKILQITELIKTENN	True	False	4.561	3.506	1.068	0.877	False	0.911	0.73