

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q92841	DDX17_HUMAN	Homo sapiens	Probable ATP-dependent RNA helicase DDX17	22.729939	S52;T55;S64;S663;T665;T666;S667;T668;S671;S672;S674;S675;S676	S64;T523	35083852;28657654;35132862;35254053;37217939;34725712;32119511;32870666;34019948;30059200;30379171;38665916;21740066;22967762;31373491;28510447;30397120;29351928;32522876;39531497;31492838;33214551	MPTGFFVAPILCVLLPSPPTREAAATVAS ATGDSASERESAAPAAAPTAEAPPPS VVTRPEPQALPSPAIRAPLPDLYPFGT MRGGGFGDRDRDRDRGGFGARGG GGLPPKFGNPGERLRKKKWLSELE PKFEKNFYVEHPEVARLTPYEVEDEL RRKKEITVRGGDVCPKPVFAFHAN FPQYVMDVLMQHFTEPTIQCCQGF PLALSGRDMVGIQGTGSGKTLAYLLP AIVHINHQPYLERGDGPICLVLAPTR ELAQQVQQVADDYKCSRLKSTCIY GGAPKGPQIRDLEGRVEICIAITPGRLI DFLESGKTNLRRCTYLVLDEADRML DMGFEPQIRKIVDQIRPDRQTLMWS ATWPKEVRQLAEDFLRDYTOINVGN LELSANHNILQIVDVCMESEKDHKL IQLMEEIMAEKENKTIHFVETKRRC DLTRRMRRDGWPAMCIHGDKSQPE RDWVWLNFRSGKAPILLIATDVASRGL DVEDVKFVINYDYPNSEDYVHRIGR TARSTNKGTYTFFTPGNLQARELI KVLEEANQAINPKLMQLVDHRGGG GGGGGRSRYRTSSANNPNLMYQD ECDRRLRGVKDGGRRDSASYRDRSE TDRAGYANGSGYSPNSAFGAQAG QYTYGQGTGAAAYGTSSYTAQEY AGTYGASSTTSTGRSSQSSSQFSGI GRSGQQPQLMSQQAQPPGATNM IGYMGQTAYQYPPPPPPPPSRK	True	False	4.09	5.0	1.71	1.7