

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q86XP3	DDX42_HUMAN	Homo sapiens	ATP-dependent RNA helicase DDX42	17.332114	S35;S37;T42;S43;S45;S709;S715;T778	S58;S96;S104;S109;S111;S185;S754	34019948;34725712;32119511;28510447;31492838;37217939;35254053;31373491;29237092;33214551;30059200;35289036;34846842;29485866;35132862	MNWNKGGPGTKRFGFGGFAISAG KKEPKLPQQSHSAFGATSSSSGFG KSAPPQLPSFYKIGSKRANFDEENAY FEDEEDSSNVLDLPYIPAENSPTRQ QFHSKPVDSDSDDDPLEAFMAEVE DQAARDMKRLEEKDKERKNVKGIR DDIEEEDDQEAYFRYMAENPTAGVV QEEEEENLEYSDGNIAPTCKIIDP LPPIDHSEIDYPPFEKNFYNEHEEIT NLTPQQLIDLRHKLNLRVSGAAPPR PGSSFAHFGFDEQLMHQIRKSEYTO PTPIQCQGVVALSGRDMIGIAKTGS GKTAAFIWPMLIHIMDQKELEPGDG PIAVIVCPTRELCQQIHAECKRFGKA YNLRVAVYGGGSMWEQAKALQEG AEIVVCTPGRLLIDHVKKKATNLQRV YLVFDEADRMFDMGFYQVRSIASH VRPDRQTLFSAFRKKIEKLARDILI DPIRVVQGDIGEANEDVTQIVEILHS GPSKWNWLTRRLVEFTSSGSVLLFV TKKANAEELANLKKQEGHNLGLLH GDMDQSERNKVISDFKKKDIPVLA TDVAARGLDIPSIKTVINYDVARIDIT HTRHRIGHTGRAGEKGVAYTLLPKD SNFAGDLVRNLEGANQHVSKELLDL AMQNAWFRKSRFKGGKGLNIGG GGLGYRERPLGSENMDRGNMNV MSNYEAYKPSTGAMGDRLTAMKAA FQSQYKSHFVAASLSNQKAGSSAAG ASGWTSAAGSLNSVPTNSAQQGHNS PDSPVTSAAKGIPGFGNTGNISGAPV TYP SAGA QGVNNTASGNN SREGTG GSNGKRERYTENRGSSRHSGETG NRHSDSPRHGDGGRHGDGYRHPES SSRHTDGHHRHGENRHGG SAGRHGE NRGANDGRNGESRKEAFNRESKME PKMEPKVDSKMDKVDSKTDKTAD GFAVPEPPKRKKS RWD S

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