

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6P2Q9	PRP8_HUMAN	Homo sapiens	Pre-mRNA-processing-splicing factor 8	16.977753	S1251	S859;S1358	34229054;35083852;34019948;34725712;32119511;28510447;37217939;26853435;33214551;33465208;30444036;22121020	MAGVFPYRGPNGNPVPGPLAPLPDYM SEEKLOEKARKWQQLQAKRYAEKR KFGFVDAQEKEDMPPEHVRKIIRDHG DMTNRKFRHDKRVYLGALKYMPHA VLKLEENMPMPWEQIRDPVPLYHIT GAISFVNEIPWVIEPVYISQWGSMWI MMRREKRDRRHFKRMRFPFDDEE PPLDYADNILDVEPLEAIQLELDPEE DAPVLDWFYDHQPLRDSRKVNGST YQRWQFTLPMMSSTLYRLANQLLTD LVDDNYFYFLFDLKAFFTSKALNMAIP GGPKFEPLVRDINLQDEDWNEFNDI NKIIIRQPIRTEYKIAFPYLYNNLPHH VHLTWYHTPNVVFIKTEDPDLPAFY FDPLINPISHRHSVKSQEPLPDDDEE FELPEFVEPFLKDTPLYTDNTANGIA LLWAPRPFNLRSGRTRRALDIPLVK NWYREHCPAGQPVKVRVSYQKLLKY YVLNALKHRPPKAQKKRYLFRSFA TKFFQSTKLDWVEVGLQVCRQGYN MLNLLIHRKNLNYLHLDYNFNLPV KTLTTKERKKSFRGNFHLCREVLR LTKLVVDSHVQYRLGNVDAFQLADG LQYIFAHVGQLTGMRYKYKLMRQI RMCKDLKHLIYRFNTGPVKGKPGC GFWAAGWRVWVFFMRGITPLLERW LGNLLARQFEGRHSKGVAKTVTKQR VESHFDELELRAAVMHDILDMMPEGI KQNKARTILQHLSEAWRCWKANIP WKVPGLPTPIENMILRYVKAKADWW TNTAHYNRERIRRGATVDKTVCKKN LGRLTRLYLKAEQERQHNYLKDGPY ITAEAVAVYTTTTVHWLESRRFSPIP FPPLSYKHDTKLLILALERLKEAYSV KSRLNQSQREELGLIEQAYDNPHEA LSRIKRHLLTQRAFKEVGFIEFMDLYS HLVPVYDVEPLEKITDAYLDQYLWYE ADKRRLFPPWIKPADTEPPPLLKYKW CQGINNLQDVWETSEGECEVMLES RFEKMYEKIDLTLNRLRLRLIVDHNI ADYMTAKNNVVINYKDMNHTNSYG IIRGLQFASFIVQYYGLVMDLLVLGL HRASEMAGPPQMPNDFLSFQDIATE AAHPIRLFCRYIDRIHIFRFTADEAR DLIQRYLTEHPDPNNENIVGYNNKK CWPRDARMRLMKHDVNLGRAVFW DIKNRLPRSVTTVQWENSFVSVYSK DNPNFLFMCGFECRILPKCRTSYE EFTHKDGVVNQLQNEVTKERTAQCF

LRVDDDESMQRFHNRVRQILMASGS
TTFTKIVNKWNTALIGLMTYFREAVV
NTQELLDLLVKCENKIQTRIKIGLNS
KMPSRFPPVVFYTPKELGGLGMLSM
GHVLIPQSDLRWSKQTDVGIHFERS
GMSHEEDQLIPNLYRYIQPWESEFID
SQRVWAEYALKRQEAIQNRRLTLE
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THQRHDGKLWNLNNYRTDMIQALG
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KASGFEE SMKWKLTNAQRSGLNQ
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QLDLTGIFMHGKIPTLKI SLIQIFRAH
LWQKIHESIVMDLCQVFDQELDALE
IETVQKETIHPKSYKMNSSCADILL
FASYKWNVSRPSLLADSKDVMSTT
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MKANPALYVLRERIRKGLQLYSSEPT
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KKNNVNVAASLTQSEIRDIILGMEISA
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DIKETGYTYILPKNVLKKFICISDLRA
QIAGYLYGVSPDPNPQVKEIRCIVMV
PQWGTHTQTVHLPQQLPQHEYLKEM
EPLGWIHTQPNESPQLSPQDVTTHA
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TLTAYKLTPSGYEWGRQNTDKGNP
KGYLPSHYERVQMLLSDRFLGFFMV
PAQSSWYNFMGVRHDPNMKYEL
QLANPKEFYHEVHRPSHFLNFALLQ
EGEVYSADREDLYA