	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membra
0	Q14191	WRN_HUMAN	Homo sapiens	Bifunctional 3'-5' exonuclease/ATP-dependent helicase WRN	24.40031	NaN	S426;S440;S453;S467;S478;S1133;S1400	28411811;23301498	MSEKKLETTAQQRKCPEWMNVQN KRCAVEERKACVRKSVFEDDLPFLE FTGSIVYSYDASDCSFLSEDISMSLS DGDVVGFDMEWPPLYNRGKLGKVA LIQLCVSESKCYLFHVSSMSVFPQGL KMLLENKAVKKAGVGIEGDQWKLL RDFDIKLKNFVELTDVANKKLKCTE TWSLNSLVKHLLGKQLLKDKSIRCS NWSKFPLTEDQKLYAATDAYAGFIIY RNLEILDDTVQRFAINKEEEILLSDM NKQLTSISEEVMDLAKHLPHAFSKL ENPRRVSILLKDISENLYSLRRMIIGS TNIETELRPSNNLNLLSFEDSTTGGV QQKQIREHEVLIHVEDETWDPTLDH LAKHDGEDVLGNKVERKEDGFEDG VEDNKLKENMERACLMSLDITEHE LQILEQQSQEEYLSDIAYKSTEHLSP NDNENDTSYVIESDEDLEMEMLKH LSPNDNENDTSYVIESDEDLEMEML KSLENLNSGTVEPTHSKCLKMERNL GLPTKEEEEDDENEANEGEEDDDK DFLWPAPNEEQVTCLKMYFGHSSF KPVQWKVIHSVLEERRDNVAVMAT GYGKSLCFQYPPVYVGKIGLVISPLIS LMEDQVLQLKMSNIPACFLGSAQSE NVLTDIKLGKYRIVYVTPEYCSGNMG LLQQLEADIGITLIAVDEAHCISEWG HDFRDSFRKLGSLKTALPMVPIVALT ATASSSIREDIVRCLNLRNPQITCTG FDRPNLYLEVRRKTGNILQDLQPFLV KTSSHWEFEGPTIIYCPSRKMTQQV TGELRKLNLSCGTYHAGMSFSTRKD IHHRFVRDEIQCVIATIAFGMGINKA DIRQVIHYGAPKDMESYYQEIGRAGR DGLQSSCHVLWAPADINLNRHLLTE IRNEKFRLYKLKMMAKMEKYLHSS RCRRQIILSHFEDKQVQKASLGIMGT EKCCDNCRSRLDHCYSMDDSEDTS WDFGPQAFKLLSAVDILGEKFGIGLP ILFLRGSNSQRLADQYRRHSLFGTG KDQTESWWKAFSRQLITEGFLVEVS RYNKFMKICALTKKGRNWLHKANT ESQSLILQANEELCPKKLLLPSSKTV SSGTKEHCYNQVPVELSTEKKSNLE KLYSYKPCDKISSGSNISKKSIMVQS PEKAYSSSQPVISAQEQETQIVLYGK LVEARQKHANKMDVPPAILATNKIL VDMAKMRPTTVENVKRIDGVSEGKA AMLAPLLEVIKHFCQTNSVQTDLFS STKPQEEQKTSLVAKNKICTLSQSM AITYSLFQEKKMPLKSIAESRILPLMT IGMHLSQAVKAGCPLDLERAGLTPE VQKIIADVIRNPPVNSDMSKISLIRM LVPENIDTYLIHMAIEILKHGPDSGL QPSCDVNKRRCFPGSEEICSSSKRS KEEVGINTETSSAERKRRLPVWFAK GSDTSKKLMDKTKRGGLFS	True	False	2.926	5.0	2.152	1.443	0.885	1.595