

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6VNB8	WDFY3_MOUSE	Mus musculus	WD repeat and FYVE domain-containing protein 3	22.083095	S3283;S3290	S1942;S2277;S2474;S3317;S3321	22645316;33300544;35822049;30016717	MNMVVKRIMGRPRQEECSPODNALGLMHLRRLFTLCHPPRHMTQKEQE EKLYMMLPVFNRFVGNAPPNTMTE KFSDLLQFTTQVSRLMVTEIRRRAS NKSTEAASRAIVQFLEINQSEEASRG WMLLTTINLLASSGQKTVDGMTM SVPSTLVKCLYLFFDLPHVPEAGGGA QNELPLAERRGLLQKAFVQILVKLCS FVSPAEEELAQKDDLQLLFSAITSWCP PYNLPWRKSAGEVLMTISRHGLSVN VVKYIHEKECLSTCVQNMQQSDDL PLEIVEMFAGLSCFLKSSDVSQTL DDFRIWQGYNFLCDLLRLEQGKEA ECRDALKDLVSLVSLTTYGVSELKP AGVTTGAPFLLPGFAVPQAGKGHS VRNIQAFAVLQNAFLKAKTNFLAQII LDAITNIYMADNANYFILESQHTLSQ FAEKISKLPVQNKYFEMLEFVVFSL NYIPCKELISVSILLKSSSSYHCSIAM KTLLKFTRHDYIFKDVFREVGLEEV MVNLLHKYAALLKDPALNEQGD SRNNSVEDQKHLALLVMEALTVLL QGSNTNAGIFREFGGARCAHNIVKY PQCRQHALMTIQQLVLSPNGEDDM GTLGLMHSAPTELQKTDILRALL SVLRESHRSTVFRKVGGFVYITSL VAMERSLSSPPKNGWEKVSQSQVL ELLHTVFCTLTALRYEPANSHFFKT EIQYEKLADAVRFLGCFSDLRKISAV NVFPSNTQPFQRLLEEGAVSVDSVS PTLRHCSKLFYLYKVATDSFDSHAE QIPPCLTSESSLPSPWGTPLSRKRH AFHCVSTPPVYPAKNVTDLKLQVTSS PLQSSDAVHHPGAMLAMDLLASV GSVTQPEHALDLQLAVANILQSLVH TERNQQVMCEAGLHARLLQRCGAA LADEDHSLHPPLQRMFERLASQALE PMVLREFRLASPLNCGAWDKLLK QYRVHKPSSLSFEPERMSSVITSLEG LGSDNVFSSHEDNHYRISKSLVKS EGSTVPLTRVKCLVSMTPHDIRLH GSSVTPAFVEFDTSLEGFGCLFLPSL APHNAPTNTVTTGLTDGAVVSGM GSGERFFPPPSGLSYSCWFCIEHFSS PPNNHPVRLTVVRRANSSEQHYVC LAIVLSAKDRSLIVSTKEELLQNYVD DFSESSFYEILPCCARFRCGELVVE GQWHHLALLMSRGMLKNSTAALYL DGQLVSTVKLHYVHSTPGGSGSANP PVLSTVYAYVGTTPAQRQIASLVWRL GPTHFLEEVLPPSSVTIYELGPNYV GSFQAVCVCKDAKSEGVTSPVSL VAEEKVSFGLYALSVSSLTVARIRKV YNKLDKAIKQLGISSHENATPVKL VHNAAGHLNGPARTIGAALIGYLG

RTFVPKPVATTLOYIGGAAAILGLVA  
MASDVEGLYAAVKALVCVVKSNPLA  
SKEMERIKGYQLLAMLLKKRSLN  
SHLHLTFSLVGTVDSGHETSIIPNST  
AFQDLLCDFEVLHAPYELHLSLFE  
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FGLAEARMTTWHIMIPSDIEPDGG  
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KKQVLEELFKVSLPVNDRGHVDIAL  
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AVVPSLTDSSSVSGQRPNTSVEQG  
SGLLSTLVGEKSVTQRWERGEISNF  
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WILSDYDSEEVDLTNPKTFRNLAKP  
MGAQTDERLAQYKKRYKD WEDPNG  
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PFTQIFLRLQGGHFDLADRMFHSVR  
EAWYSASKHNMADV KELIPEFFYLP  
EFLFNSNNFDLGCKQNGTKLGDVIL

PPWAKGDPREFIRVHREALECDYVS  
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VFHHLFYEGQVDIYNINDPLKETATI  
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LEMQEGCPEAQIGQQAQDDSSDS  
ETEEPSVSQDPKDTSSQPSSTSHRP  
RAASCRAATWCTDSGSDSRRWS  
DQLSLDEKDGFIHVNYSEGQTRAHL  
QGPLAHPHPNPIEARSYSRLKPGYR  
WERQLVFRSKLTMHTAFDRKDNTH  
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CSGCSVRFSLTERRHHCRNCGQLF  
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