

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
E9Q8I9	FRY_MOUSE	Mus musculus	Protein furry homolog	9.108089	NaN	T213;S1382;S1383;S1936;S1940;S2427;S2428;S2495;T2516;S2815	33300544	MASQQDSGFFEISIKYLLKSWSNAS PVGNGYIKPPVPPASGTHREKGP MLPINVDPDSKPGYVLKSLFVNFTT QAERKIRIIMAEPLEKPLTKSLQRGE DPQFDQVISSMSSLSEYCLPSILRTL FDWYKRQNGIEDESHEYRPTS KSDEQQRDYLMEERRDLAIDFIFSLVL IEVLKQIPLHPVIDSLIHDIIINLAFKHF KYKEGYLGPNTGNMHIVADLYAEVI GVLAQAKFPAVKKKFMALKELRHK EQSPYVVSIIISLIMGKFFRIKMYP VEDFEASLQFMQECAHYFLEVKDK DIKHALAGLFVEILVPVAAAVKNEVN VPCLRNPFVESLYDTTLELSSRKKHSL ALYPLVTCLLCVSQKQLFLNRWHVF LNNCLSNLKNKDPKMARVALESY RLLWVYMIRIKCESNTATQSRITITT TLFPKGSRGVPRDMLNIFVKIIQFI AQERLDFAMKEIIFDFLCVGP SLNPERMNIGLRAFLVIADSLQK GEPMPVTGAVLPSGNTLRVKKTYL SKTLTEEEAKMIGMSLYYSQVRKAV GNILRHLDKVGRCMMLTNVQMLN KEPEDMITGERKPKIDLFRTCVA RLLPDGMSKLELIDLLARLSIHMD ELRHIAQNSLQGLLVDFSDWREDVL FGFTNFLREVNDMHHTLLDSSLKL LLQLLTQWKLVIQTQGRAYEQANKI RNSSELIPNGSSHRMQSERGPHCSVL HAVEGFALVLLCSFQVATRKL KEIRALFLALGQPEDDDRPMIDVMD QLSSSILEFSIHVAVSDSATLPPTHN VDLQWLVEWNAVLVNSHYDVKSPS HVWIFAQSVKDPWVLCFLSFLRQEN LPKHCPTALSYAWPYAFTRLQSVMP LVDPNSPVNAKKTSTASSGDNYVTL WRNYLILCFGVAKPSIMSPGHLRAS TPEIMATTPDGTVSYDNKAIGTPSVG VLLKQLVPLMRLESIEITESLVLGFG RTNSLVFRELVEELHPLMKEALERR PENKKRRRERRDLLRLQLLRFELLAD AGVISDSTNGALERDTLALGALFLEY VDLTRMLLEAENDKEVEILKDIRAH FSAMVANLIQCVPVHRRFLFPQQS LRHHLFILFSQWAGPFSIMFTPLDRY SDRNHQITRYQYCALKAMSAVLCCG PVFDNVGLSPDGYLYKWL DLRVHQLGCEVVMLLELNPQINL FNWAI DRCYTGSYQLASGCFKAIATV

CGNRNYPFDIVTLLNLVLFKASDTN
REIYEVSMQLMQILEAKLFVHSHKV
AEQRPGSILYGTGHPPLYSVSLAL
LSCELARMYPELTLPLFSEVSQRFP
THPNGRQIMLTYPWLNHNLVDS
RLLPGSSPSSPEDEVKDREGEVTS
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TAKYGDEVGAEMENAWNALANNE
KWSNNLRVTLQFLISLCGVSSDTILL
PYIKKVATYLCRNNTIQTMEELLFEL
QQTEPVNPIVQHCDNPPFYRFTASS
KASAAASGTTSSSNTVVAGQDSFPD
PEESKILKESDDRFSNVIRAHTRLES
RYSNSSGGSYDEDKNDPISPYTGWL
LSITEAKQPQPLMPCSGGCWAPLV
DYLPEITPRGPHRCNIAVIFMTEM
VVDHSVREDWALHPLLLHAVFLGL
DHYRPEVFEHSHKLLLHLLIALSCNS
NFHAIASVLLQTTREMGAEKTLTMQP
AYQPEYLYTGGDFLREDQSSPVPDS
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EVEDVEAATETDEKASKLIEFLTTRA
FGPLWCHEDITPKNONSKEAQLSN
FLRHVVSVFKDSRSGFHLEQHLSEV
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PRSATLDRIQACTQQGLSSKTRSNSS
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ALMESDFEFEYLMALRLLNRLLAH
MPLEKAENREKLEKLQAQLKWADF
PGLQQLLLKGFTSLTTTDLTLQLFSL
LTSVSKVPMVDSSQAIGFPLNVLCLL
PQLIQHFENPNQFCKDIAERIAQVCL
EEKNPKLSNLAHVMTLYKTHSYTRD
CATWVNVVCRYLHEAYADITLNMVT
YLAELLEKGLPSMQPLLQVIYSLLS
YMDLSVVPVKQFNMEVLKTIKYVQ
SIHWREALNILKLVVSRASLVLP
QHSKLSKIELHRVWTSASKELPGKT
LDFHFDISETPIIGRRYDELQNSSGR
DGKPRAMAVTRSASSTSSGSNSNVL
VPVSWKRPQYSQKRTKEKLVHVL
CGQEVGLSKNPSVIFSSCGDLDLPE
HQTSLVSSSEDGPREQENMDDTNSE
QQFRVFRDFDFLDVELEDGEGESM
DNFNWGVRRRSLDSLKCDMQILE
ERQLSRSTPSLNKMSHEDSDESSEE
DLTASQILEHSDLMNLSPSEANP

							MELLTSACDSAPADPHSFNTRMAN FEASLPDINNLOISEGSKAEAVPEEE DTTVHEDDLSSSINELPAAFECSDSF SLDMTEAEKGNRGLDQYTLASFGE GDRGVSPPPSPFFSAILAAFQPAACD DAEEAWRSHINQLMCDSDGSCAVY TFHVFSSLFKNIQRFCFLTCDAAASY LGDNLRGIGSKFVSSSQMLTSCSEC PTLFVDAETLLSCGLLDKLFVLEL QEYLDTYNNRKEATLSWLANCKATF AGGSRDGVITCQPGDSEEKQLELCQ RLYKLFQLLLLLYQSYCKLIGQVHEV SSPELLNMSRELSDLKRNLKEATA AIATDPLYIEGAWSEPTFTSTEAAIQS MLECLKNNELGKALRQIKECRSLWP NDIFGSSDDEVQTLNLIYFRHQTLG QTGTALVGSNHSLTEICTKLMELN MEIRDMIRRAQNYRVLTAFLPDSSV SGTSL
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