

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P02751	FINC_HUMAN	Homo sapiens	Fibronectin	10.484972	T1001	T2454;S2475	33214551;37340703;34019948;34725712	MLRGPGPGLLLLAVQCLGTAVPSTG ASKSKRQAQQMVQPQSPVAVSQSKP GCYDNGKHYQINQQWERTYLGNAL VCTCYGGSRGFNCESKPEAEETCFD KYTGNTYRVGDTYERPKDSMIWDCT CIGAGRGRISCTIANRCHEGGQSYKI GDTWRRPHETGGYMLECVCCLGNGK GEWTCKPIAEKCFDHAAGTSYVVGE TWEKPYQGWMMVDCTCLGEGSGRI TCTSRNRCNDQDTRTSYRIGDTWSK KDNRGNLLQCICTGNRGEWK CER HTSVQTTSSSGPFTDVRAAVYQPQ PHPQPPPYGHCVTDSGVVYSVGMQ WLKTQGNKQMLCTCLGNGVSCQET AVTQTYGGNSNGEPCVLPFTYNGRT FYSCTTEGRQDGHLCSTTSNYEQ DQKYSFCTDHTVLVQTRGGNSNGAL CHFPLYNNHNYTDCTSEGRRDNM KWC GTTQNYDADQKFGFCPMAAHE EICTTNEGVMYRIGDQWDKQHDMG HMMRCTCVGNRGEWTCAYSQLR DQCIVDDITYNVNDTFHKRHEEGH MLNCTCFGQGRGRWKCDPVDQCQ DSETGTFYQIGDSWEKYVHGVRYQC YCYGRGIGEWHCQPLQTYPSSSGPV EVFITETPSQPN SHPIQWNAPQPSHI SKYILRWRPKNSVGRWKEATIPGHL NSYTIKGLKPGVVYEGQLISIQQYGH QEVTRFDFTTTSTSTPVTSTNTVTGET TPFSPLVATSESVTEITASSFVVS WV SASDTVSGFRVEYELSEEGDEPQYL DLPSTATSVNIPDLLPGRKYIVNVYQI SEdgeQSLILSTSQTAPDAPPD TT V DQVDDTSIVVRWSRPQAPITGYRIVY SPSVEGSSTELNLPETANSVTLSDLQ PGVQYNITIYAVEENQESTPVVIQQE TTGT PRSDTVPSRDLQFVEVTDVKV TIMWTPPESAVTGYRVDVIPVNL PGE HGQRLPISRNTFAEVTGLSPGVTY YF KVFAVSHGRESKPLTAQQTTKLDAP TNLQFVNETDSTVLVRWTPPRAQIT GYRLTVGLTRRGQPRQYNVGPSVSK YPLRNLQPASEYTVSLVAIKGNQESP KATGVFTTLQPGSSIPPYNTEVTETTI VITWTPAPRIGFKLGVRPSQGGEAPR EVTSDSGSIVVSGLTPGVEYVYTIQVL RDGQERDAPIVNKVVTP LSPPTNLH LEANPDTGVLTVSWERSTTPDITGYR ITTTPTNGQQGNSLEEVVHADQSSC

TFDNLSPGLEYNVSVYTVKDDKESV
PISDTIPEVPQLTDLSFVDITDSSIGL
RWTPLNSSTIIGYRITVVAAGEGIPIF
EDFVDSSVGYTGTGLEPGIDYDISVI
TLINGGESAPTTLTQQTAVPPPTDLR
FTNIGPDTMRVTWAPPPSIDLTNFLV
RYSVKNEEDVAELSISPSDNAVLT
NLLPGTEYVSVSSVYEQHESTPLRG
RQKTGLDSPTGIDFSDITANSFTVH
WIAPRATITGYRIRHHPEHFSGRPRE
DRVPHSRNSITLTNLTPGTEYVVSIV
ALNGREESPLLIGQQSTVSDVPRDLE
VVAATPTSLLISWDAPAVTVRYRITY
GETGGNSPVQEFTVPGSKSTATISGL
KPGVDYTTVYAVTGRGDSPASSKPIS
INYTEIDKPSQMQVTDVQDNSISVK
WLPSSSPVTGYRVTTTPKNGPGPTKT
KTAGPDQTEMTIEGLQPTVEYVSVY
AQNPSGESQPLVQTAVTNIDRPKGL
AFTDVDVDSIKIAWESPQQQVSRYRV
TYSSPEDGIHELFPAPDGEEDTAELO
GLRPGSEYTVSVVALHDDMESQPLI
GTQSTAIPAPTDLKFTQVTPTSLSAQ
WTPPNVQLTGYRVRVTPKEKTGPMK
EINLAPDSSSVVSGLMVATKYEVS
YALKDTLTSRPAQGVVTTLENVSPPR
RARVTDATETTITISWRKTETITGFQ
VDAVPANGQTPIQRTIKPDVRSYTT
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PRPGVTEATITGLEPGTEYTIYVIALK
NNQKSEPLIGRKKTDDELQPLVTLPH
PNLHGPEILDVPSTVQKTPFVTHPGY
DTGNGIQLPGTSGQQPSVGQQMIFE
EHGFRRTTPPTTATPIRHRPRPYPPN
VGEEIQIGHIPREDVDYHLYPHGPG
NPNASTGQEALSQTTISWAPFQDTS
EYIISCHPVGTDDEEPLQFRVPGTSTS
ATLTGLTRGATYNVIVEALKDQQRH
KVREEVTVGNSVNEGLNQPTDDSC
FDPYTVSHYAVGDEWERMSESGFKL
LCQCLGFGSGHFRCDSSRWCHDNG
VNYKIGEKWDRQGENGQMMMSCTCL
GNGKGEFKCDPHEATCYDDGKTYH
VGEQWQKEYLGAICSCTCFGGQRG
WRCDNCRPPGGEPSPEGTTGQSYN
QYSQRYHQRTNTNVNCPICFMPLD
VQADREDSRE