

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P35556	FBN2_HUMAN	Homo sapiens	Fibrillin-2	4.49472	NaN	NaN	34725712;34019948	MGRRRRLCLQLYFLWLGCVVWLAQ GTAGQPQPPPKPPRPQPPQVRS ATAGSEGGFLAPEYREEGAASRV RRRGQQDVLGRPNVCGSRFHSYCC PGWKTLPGGNQCVIPICRNSCGDGF CSRPNMCTCSSGQISSTCGSKSIQQ CSVRCMNGGTCADDHCQCQKGYIG TYCGQPVCENGCGNGGRCIGPNRC ACVYGFTGPQCERDYRTGPCFTQVN NQMCQGQLTGIVCTKTLCCATIGRA WGHPCEMCPAQPCRRGFIPNIRT GACQDVDECQAIPGICQGGNCINTV GSFECRPCAGHKQSETTQKCEDIDE CSIIPGICETGECSTVGSYFCVCPR GYVTSTDGSRCIDQRTGMCFSGLVN GRCAQELPGRMTKMCCCEPGRG WGIGTIPEACPVRGSEYRRLCMDG LPMGGIPGSAGSRPGGTGGNGFAPS GNGNGYGGGTGFIPPGGNGFSPG VGGAGVGAGGQGPITGLTILNQTIDI CKHHANLCLNGRCIPTVSSYRCECN MGYKQDANGDCIDVDECTSNPCTN GDCVNTPGSYCKCHAGFQRTPTKQ ACIDIDECIQNGVLCKNGRCVNTDG SFQCICNAGFELTTDGKNCVDHDEC TTTNMCLNGMCINEDGSFKCICKPG FVLAPNGRYCTDVDECQTPGICMNG HCINSEGSFRDCPPGLAVGMDGRV CVDTHMRSTCYGGIKKGVCVRPFPG AVTKSECCANPDYGFGEPCQPCPA KNSAEFHGLCSSGVGITVDGRDINE CALDPDICANGICENLRGSYRCNCN SGYEPDASGRNCIDIDECLVNRLLCD NGLCRNTPGSYSCTCPPGYVFRTE ETCEDINECESNPCVNGACRNNLGS FNCECSPGKLSSTGLICIDSLKGTC WLNIQDSRCEVNINGATLKSECCAT LGAAWGSPCERCELDTACPRGLARI KGVTCEDVNECEVFPGVCPNGRCV NSKGSFHCECPEGLTLDGTGRVCLD IRMEQCYLKWDEDECIHPVPGKFR MDACCAVGAAWGTECECECPKPGT KEYETLCPRGAGFANRGDVLTRPF YKDINECKAFPGMCTYGKCRNTIGS FKCRCNSGFALDMEERNCTDIDECR ISPDLCGSGICVNTPGSFECECFEGY ESGFMMMKNCMDIDECERNPLLC RGGTCVNTGEGSFQDCPLGHELSPS REDCVDINECSLSDNLCRNGKCVN

MIGTYQCSCNPGYQATPDRQGCTDI
DECMIMNGGCDTQCTNSEGSYECS
CSEGYALMPDGRSCADIDECENNP
ICDGGQCTNIPGEYRCLCYDGFMAS
MDMKT CIDVNECDLNSNICMFGEC
ENTKGSFICHQCLGYSVKKGTTGCT
DVDECEIGAHNCDMHASCLNIPGSF
KCSCREGWINGIKCIDLDECSNGT
HQCSINAQCVNTPGSYRCACSEGFT
GDGFTCSVDDECAENINLCENGQCL
NVPGAYRCECEMGFTPASDSRSCQ
DIDEC SFQNICVFGTCNNLPGMFHC
ICDDGYELDRTGGNCTDIDECADPIN
CVNGLCVNTPGRYECNCPDFQLNP
TGVGCVDNRVGNCYLKFGPRGDGS
LSCNTEIGVGVSRSSCCSLGKAWG
NPCETCPPVNSTEYITLCPGGEGFR
PNPITHILEDIDECQELPGLCQGGNCI
NTFGSFQCECPQGYLSEDTRICEDI
DECFAHPGVCGPGTCYNTLGNYTCI
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RSYNGTTCENELPFNVTKRMCCCTY
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GNIPGFTFDIHTGKAVIDECKEIPGI
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LLLVCEIDIDEC SNGDNLQORNADCI
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DGSFRCECPMGYNLDYTGVRVDT
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NEGFEPGPMMNCE DINECAQNPLL
CAFRCMNTFGSYECTCPIGYALRED
QKMCKDLDECAEGLHDCESRGMM
CKNLIGTFMCICPPGMARRPDGEGC
VDENECRTKPGICENGRVNIIGSYR
CECNEGFQSSSSGTECLDNROGLCF
AEVLQITICMASSSRNLVTKSECCC
DGGRGWGHQCELCPLPGTAQYKKIC
PHGPGYTTDGRDIDECKVMPNLCTN
GQCINTMGSFRFCFKVGYTTDISGT
SCIDLDECSQSPKPCNYICKNTEGSY

