

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5T1H1	EYS_HUMAN	Homo sapiens	Protein eyes shut homolog	6.287639	NaN	NaN	34019948	MTDKSIVILSLMVFHSSFINGKTCRR QLVEEWHQPSSYVVNWTLTENICL DFYRDCWFLGVNTKIDTSGNQAVPQ ICPLQIQIGDILVISSEPSLQFPEINL MNVSETSFVGCVQNTTTEDQLLFGC RLKGMHTVNSKWLSVGYHYFITVM ASGPSPCPLGLRLNVTVKQQFCQES LSSEFCSGHGKCLSEAWSKTYSCHC QPPFSGKYCQELDACSFKPCKNNGS CINKRENWDEQAYECVCHPPFTGK NCSEIIGQCQPHVCFHGNCSNITSN SFICECDEQFSGPFCEVSAKPCVLL FWKRGICPNSSSAYTYECPKGSSSQ NGETDVSEFSLVPCQNGTDCIKISN DVMCICSPIFTDLLCKSIQTSCEFPPL RNNATCKKCEKDYPSCISGFTEKN CEKAIDHCKLLSINCLNEEWCFNIIIG RFKYVCIPGCTKNPCWFLKNVYLIIH QHLCYCGVTFHGICQDKGPAQFEYV WQLGFAGSEGEKCGVIDAYFFLAA NCTEDATYVNDPEDNNSCWFPHE GTKEICANGCSCLSEEDSQEYRYLCF LRWAGNMYLENTTDDQENECQHE AVCKDEINRPRCSCSLSYIGRLCVVN VDYCLGNHSISVHGLCLALSHNCNC SGLQRYERNICEIDTEDCKSASCKN GTTSTHLRGYFFRKCVPVGFKGTQCEI DIDECASHPCCKNGATCIDQPGNYFC QCVPPFKVVDGFSLCNPVGYVIRG EQDIDDCILNACEHNSTCKDLHLSY QCVCLSDWEGNFCEQESNECKMN PCKNNSTCTDLYKSYRCECTSGWTG QNCSEEINECDSDPCMNGGLCHES TIPGQFVCLCPPLYTGQFCHQRYNL CDLLHNPCRNNSTCLALVDANQHCI CREEFEGKNCIDVKDCLFLSCQDY GDCEDMVNNFRICRPGFSGSLCEI EINECSSEPCKNNGTCVDLTNRFFC NCEPEYHGPFCELDVKCKISPCLD EENCVYRTDGYNCLCAPGYTGINCEI NLDECLSEPCLHDGVCIDGINHYTC DCKSGFFGTHCETNANDCLSNPCL HGRCTELINEYPCSCDADGTSTQCKI KINDCTSIPCMNEGFCQKSAHGFTC ICPRGYTGAYCEKSIDNCAEPELNSV ICLNGGICVDGPGHTFDCRCLPGFS GQFCEININECSSPCLHGADCEDH INGYVCKCPGWSGHHCENELECIP NSCVHELCEMENEPEGSTCLCTPGFM

TCSIGLLCGDEIRRITCLTPIFQRTDPI
STQTYTIPPSETLVSSFPSIKATRIPAI
MDTYPVDQGPQGTGIVKHDILPTTGL
ATLRISTPLESYLLQELIVTRELSAKH
SLLSSADVSSSRFLNFGIRDPAQIVQ
DKTSVSHMPIRTSAATLGFFFFPDRRA
RTPFIMSSLMSDFIFPTQSLLFENCQ
TVALSATPTTSVIRSIPGADIELNRQS
LLSRGFLIAASISATPVVSRGAQEDI
EEYSADSLISRREHWRLLSPSMSPIF
PAKVIISKQVTILNSSALHRFSTKAFN
PSEYQAITEASSNQRLTNIKSQAADS
LRELSQTCATCSMTEIKSSREFSDQV
LHSKQSHFYETFWMNSAILASWYAL
MGAQTITSGHSFSSATEITPSVAFTE
VPSLFPSKKSARKTILSSSLEESITLS
SNLDVNLCLDKTCLSIIVPSQTISSDL
MNSDLTSKMTTDELSVSENILKLLKI
RQYGITMGPTVNLNQUESLLDMEKSK
GSHTLFLKHPDSSSLDFELNLQIYPD
VTLKTYSEITHANDFKNNLPPLTGSV
PDFSEVTTNVAFYTVSATPALSIQTS
SSMSVIRPDWPYFTDYMTSLKKEVK
TSSEWSKWELQPSVQYQEFPTASRH
LPFTRSLTLSSLESILAPQRLMISDFS
CVRYYGDSYLEFQNVALNPQNNISL
EFQTFSSYGLLLYVKQDSNLVDGFFI
QLFIENGLKYHFYCPGEAKFKSINT
TVRVDNGQKYTLLIRQELDPCNAEL
TILGRNTQICESINHVLGKPLPKSGS
VFIGGFDDLHGKIQMPVPVKNFTGCI
EVIEINNWRSFIPSKAVKNYHINNCR
SQGFMLSPTASFVDASDVTQGVDT
MWTSVSPSVAAPSVQCQDVCHNGG
TCHAIFLSSGIVSFQDCPLHFTGRF
CEKDAGLFFPSFNGNSYLELPFLKF
VLEKEHNRTVTIYLTIKTNSLNGTILY
SNGNCGKQFLHLFLVEGRPSVKY
GCGNSQNILTVSANYSINTNAFTPITI
RYTTPVGSPPGVVCMIEMTADGKPPV
QKKDTEISHASQAYFESMFLGHIPA
NVQIHKKAGPVYGFRCILDQVNN
KEFFIIDEARHGKNIENCHVPWCAH
HLCRNNGTCISDNENLFCECPRLYS
GKLCQFASCENNPCGNGATCVPKS
GTDIVCLCPYGRSGPLCTDAINITQP
RFSGTDAFGYTSFLAYSRLDISFHYE
FHLKFQLANNHSALQNNLIFFTQK
GHGLNGDDFLAVGLLNGSVVYSYNL
GSGIASIRSEPLNLSLGVHTVHLGKF
FQEGWLKVDDHKNKSIIAPGRLVGL
NVFSQFYVGGYSEYTPDLLPNGADF

							KNGFQGCIFTLQVRTEKDGHFRLG NPEGHPNAGRSVGQCHASPCSLMK CGNGGTCIESGTSVYCNCCTTGWKGS FCTETVSTCDPEHDPHHC SRGATC ISLPHGYTCFCPLGTTGIYCEQALILI VILEKPKPAERKVKKEALSISDPSFRS NELSWMSFASFHVRKKTHIQLQFQP LAADGILFYAAQHLKAQSGDFLCISL VNSSVQLRYNLGDRTHILETLQKVTI NGSTWHIIKAGRVGAEGYLDLDGIN VTEKASTKMSSLD TNTDFYIGGVSSL NLVNPMAIENEPVGFQGCIRQVIIN NQELQLTEFGAKGGSNVGDCDGTA CGYNTCRNGGECTVNGTTFSCRCLP DWAGNTCNQSVSCLNNLCLHQSLC IPDQSFSYSCCLTLGWVGRYCNKT SFSTAKFMGNSYIKYIDPNYRMRNL QFTTISLNFSTTKTEGLIVWMGIAQN EENDFLAIGLHNQTLKIAVNLGERIS VPMSYNNGTFCCKWHHVVIQNQ TLIKAYINNSLILSEDIDPHKNFVALN YDGICYLGGFEYGRKVNIVTQEIFKT NFVGKIKDVVFFQEPKNIELIKLEGY NVYDGDEQNEVT
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