

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8VDJ3	VIGLN_MOUSE	Mus musculus	Vigilin	14.099706	NaN	T8;S11;S31;T295;T296;S317;T437;S645;S1247;S1252	28528544;34887587;36288343;29187734	MSSVAVLTQESFAEHRSGLVPPQIQIK VATLNSEEENDPPTYKDAFPPLPEKA ACLESAQEPAGAWSNKIRPIKASVIT QVFHVPLEERKYKDMNQFGEQEQA KICLEIMQRTGAHLELSLAKDQGLSI MVSGKLDVAVMKARKDIVARLQTQAS ATVPIPKHEHHRFVIGKNGEKLQDLEL KTATKIQIPRPDDPSNQIKITGTKEGI EKARHEVLLISAEQDKRAVERLEVE KAFHPFIAGPYNRLVGEIMQETGTRI NIPPPSVNRTEIVFTGEKEQLAQAVA RIKKIYEKKKKTTTTIAVEVKKSQHKY VIGPKGNSLQEILERTGVSVEIPPSDS ISETVILRGEPEKLGQALTEVYAKAN SFTVSSVSAPSWLHRFIIIGKKGQNLA KITQQMPKVHIEFTEGEDKITLEGPT EDVNVAQEQIEGMVKDLINRMDYV EINIDHKFHRHLIGKSGANINRIKDQ YKVSVRIPPDSEKSNLIRIEGDPQGV QOAKRELLELASRMENERTKDLIIE QRFHRTIIGQKGERIREIRDKFPVEVII NFPDPAQKSDIVQLRGPKNEVEKCT KYMQKMVADLVENSYSISVPIFKQF HKNIIGKGGANIKKIREESNTKIDLPA ENSNSETIIITGKRANCEAARSILSI QKDLANIAEVEVSIPAKLHNSLIGTK GRLIRSIMEECCGVHIFPVEGSGSD TVVIRGPSSDVEKAKKQLLHLAEKQ TKSFTVDIRAKPEYHKFLIGKGGGKI RKVRDSTGARIIFPAAEDKDQDLITII GKEDAVREAQKELEALIQLNLENVVE DYMLVDPKHHRHFVIRRGQVLREIA EEYGGVMVSFPRSGTQSDKVTLKGA KDCVEAAKKRIQEIIEDLEAQVTVEC AIPQKFHR SVMGPKGSRIQQITRDYN VQIKFPDREENPVHSVEPSIQENGD EAGEGREAKETDPGSPRRCDIIISGR KEKCEAAKEALEALVPVTIEVEVPFD LHRYIIGQKSGGIRKMMDEFVNIH VPAPELQSDTIAITGLAANLDRKAG LLDRVKELQAEQEDRALRSFKLSVT VDPKYHPKIIIGRKGAVITQIRLEHEV NIQFPDKDDGNQPQDQITITGYEKN TEAARDAILKIVGELEQMVSEDEVPLD HRVHARIIGARGKAIRKIMDEFKVDI RFPOSGAPDPCVTVTGLPENVEEAI DHILNLEEEYLADVVDSEALQVYMK PPAHEESRAPSKGFVVRDAPWTSNS SEKAPDMSSEEFPSFGAQVAPKTL PWGPKR