

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8VDJ3	VIGLN_MOUSE	Mus musculus	Vigilin	13.077453	NaN	T8;S11;S31;T295;T296;S317;T437;S645;S1247;S1252	28528544;29187734	MSSVAVLTQESFAEHRSGLVPOQIK VATLNSEENDPPTYKDAFPPLPEKA ACLESAQEPAGAWSNKIRPIKASVIT QVFHVPLEERKYKDMNQFGEQEQA KICLEIMQRTGAHLELSLAKDQGLSI MVSGKLDAVMKARKDIVARLQTQAS ATVPIPKHHRFVIGKNGEKLQDLEL KTATKIQIPRPDDPSNQIKITGTKEGI EKARHEVLLISAEQDKRAVERLEVE KAFHPFIAGPYNRLVGEIMQETGTRI NIPPPSVNRTEIVFTGEKEQLAQAVA RIKKIYEEKKKTTTIAVEVKKSQHKY VIGPKGNSLQEIERTGVSVEIPPSDS ISETVILRGEPEKLGQALTEVYAKAN SFTVSSVSAPSWLHRFIIIGKKGQNLA KITQQMPKVHIEFTEGEDKITLEGPT EDVNVAQEQIEGMVKDLINRMDYV EINIDHKFHRHLIGKSGANINRIKQD YKVSVRIPPDSEKSNLIRIEGDPQGV QQAKRELLELASRMENERTKDLIE QRFHRTIIGQKGERIREIRDKFPEVII NFPDPAQKSDIVQLRGPKNVEVEKCT KYMQKMVADLVENSYSISVPIFKQF HKNIIGKGGANIKKIREESNTKIDLPA ENSNSETIIITGKRANCEAARSRLSI QKDLANIAEVEVSIPAKLHNSLIGTK GRLIRSIMEECCGGVHIHFVPEGSGSD TVVIRGPSSDVEKAKKQLLHLAEEKQ TKSFTVDIRAKPEYHKFLIGKGGGKI RKVRDSTGARIIFPAEDKDQDLITII GKEDAVREAQKELEALIQNLENVVE DYMLVDPKHHRHFVIRRGQVLREIA EEYGGVMVSFPRSGTQSDKVTLKGA KDCVEAAKKRIQEIIEDLEAQVTVEC AIPQKFHRSVMGPKGSRIQQITRDYN VQIKFPDREENPVHSVEPSIQENGD EAGEGREAKETDPGSPRRCIIIISGR KEKCEAAKEALEALVPVTIEVEVPFD LHRYIIGQKSGIRKMMDEFEVNIH VPAPQLQSDTIAITGLAANLDRKAG LLDRVKELQAEQEDRALRSFKLSVT VDPKYHPKIIIGRKGAVITQIRLEHEV NIQFPDKDDGNQPQDQITITGYEKN TEAARDAILKIVGELEQMVEDVPLD HRVHARIIGARGKAIRKIMDEFKVDI RFPQSGAPDPCVTVTGLPENVEEAI DHILNLEEEYLADVVDSEALQVYMK PPAHEESRAPSKGFVVRDAPWTSNS SEKAPDMSSEEFPSFGAQVAPKTL

