

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9NT68	TEN2_HUMAN	Homo sapiens	Teneurin-2	24.917859	NaN	S90;S124;T155;S157	28411811	MDVKKRRHRSRLTRGRGCKECRYTS SSLDSEDCRVPTQKSYSSSETLKAYD HDSRMHMYGNRVTDLIHRESDEFPR QGTNFTLAEELGICEPSPHRSGYCS MGILHQGYSLSTGSDADSDTEGGMS PEHAIRLWGRGIKSRSSGLSSREN SALTLTSDNENKSDDENGRPIPPT SSPPLLPSAQLPSSHNPPVSCOMP LLDSNTSHQIMDTNPDEEFSPNSYL LRACSGPQQASSSGPPNHHQSSTLR PPLPPHNHTLSHHSSANSLNRN SLTNRRSQIHAPAPAPNDLATTPEV QLQDSWVLSNSVPLETRHFLFKTSS GSTPLFSSSSPGYPLTSGTVYTPPPRL LPRNTFSRKAFLKPKSKYCSWKCA ALSAIAAALLAILLAYFIAMHLLGLN WQLQPADGHTFNNGIRTGLPGNDD VATMPSGGKVPWSLKNSSIDSGEAE VGRRTVQEVPPGVFWRSQIHISQPO FLKFNISLGKDALFGVYIRRGLPSSH AQYDFMERLDGKEKWSVVEPRER RSIQTLVQNEAVFVQYLDVGLWHLA FYNDGKDKEMVSFNTVVLDVQDC PRNCHGNCECVSGVCHCFPGFLGA DCAKAACPVLCSGNGQYSKGTQCY SGWKGAECDVPMNQCIDPSCGGHG SCIDGNCVCSAGYKGEHCEEVDCLD PTCSSHGVCVNGECLCSPGWGGLN CELARVQCPDQCSGHGTYLPDTGLC SCDPNWMGPDCSVEVCSVDCGTHG VCIGGACRCEEGWTGAACDQRVCH PRCIEHGTCKDGKCECREGWNGEH CTIGROTACTETDGCPLCNGNGRC TLGQNSWQCVCQTGWRGPGCNVA METSACDNKDNKDGDLVDCLDPDC CLQSACQNSLLCRGSRDPLDIIQQG QTDWPAVKSFYDRIKLAGDSTHII PGENPFNSSLVSLIRGQVTTDGTPL VGVNVSFVKYKPYGYTITRQDGTFDL IANGGASLTLHFERAPFMSQERTVW LPWNSFYAMDTLVMKTEENSIPSCD LSGFVRPDPHIISSPLSTFFSAAPGQN PIVPETQVLHEEIELPGSNVKLRYLS SRTAGYKSLKITMTQSTVPLNLRV HLMVAVEGHFLFQKSFQASPNLAYTF IWDKTDAYGQRVYGLSDAVVSVGFE YETCPSLILWEKRTALLQGFELDPSN LGGWVSLDKHHILNVKSGILHKGTE NQFLTQQAIIITSMGNRRRSISCP SCNGLAEGNKLLAPVALAVGIDGSLY VGDFNYIRRFPSRNVTSILELRNKE FKHSNPAHKYLLAVDPVSGSLYVS DTNSRRIRYVKSLSGTKDLGNSEV VAGTGEQCLPFDEARCGDGGKADA TLMSPRGAIVDKNGLMYFVDATMIR KVDQNGIISTLLGSNDLTAVRPLSCD SSMDVAQVRLEWPTDLAVNPMDNS LYVLENNVILRITENHQVSIAGRPM HCQVPGIDYSLSKLAHSALESASAIA ISHTGVLYITETDEKINRLROVTTN GEICLAGAASDCCKNDVNCNCYS GDDAYATDAILNSPSSLAVAPDGTIYI ADLGNIRIRAVSKNKPVLNAFNQYE AASPGEQELYVFNADGIHQYTVSLVT GEYLYNFTYSTDNDVTELDIDNNGNS LKIRRDSSGMPRHLLMPDNQIITLV GTNGLLVSTQNLLEGLMITYDGN TGLLATKSDETGWTTFYDYDHEGRL TNVTRPTGVVTSLHREMEKSITIDIE NSNRDDVTVITNLSVEASYTVVQ DQVRNSYQLCNGTLRVMYANGM GISFHSEPHVLAGTITPTIGRCNISLP MENGLNSIEWRLRKEQIKGKVTFIG RKLRVHGRNLLSIDYDRNIRTEKIYD DHRKFTLRHYYDQVGRPFLWLPSSGL AAVNVSYFFNGRLAGLQRGAMSSERT	None	None	None	None	None	None	None		

DIDKQGRIVSRMFADGKVVWSYSYLD  
KSMVLLQSQRQYIFEYDSSDRLLAV  
TMPFSVARHSMSTHTSIGYIRNIYNPP  
ESNASVIFDYSDDGRILKTSFLGTGR  
QVFYKYGKLSKLSSEIVYDSTAVTFGY  
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GPLVDKQIYRFSEEGMVNARFDYTY  
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MYWMTVQYDSMGRVIKRELKLGPY  
ANNTTKYTYDYGDLQSVAVNDRP  
TWRYSYDLNGLHLLNPGNSVRLM  
PLRYDLRDRITRLGDVQYKIDDDGYL  
CQRGSDIFEYNSKGLLTRAYNKASG  
WSVQYRYDGVGRRASYKTNLGHHL  
QYFYSDLHNPTRITHVYNHNSSEITS  
LYYDLQGHLFAMESSGEEYVVASD  
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VHFTQRDYDVLGRWTSPTYTMWK  
NVGKEPAPFNLYMFKSNNPLSSELD  
LKNYVTDVKSWLVMFGFQLSNIIPG  
FPRAKMYFVPPPYELSESQASENGQ  
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KKLHASIREKAGHWFATTTPIIGKGI  
MFAIKEGRVTTGVSSIAESDSRKVAS  
VLNNAYYLDKMHSIEGKDTHYFVK  
IGSADGDLVTLGTTIGRKLVESEGVNV  
TVSQPTLLVNGRTRRRFTNIEFYSTL  
LLSIRYGLTPDLDDEEKARVLDQARQ  
RALGTAWAKEQQKARDGRECSRLW  
TEGEKQQLLSTGRVQGYEGYYVLPV  
EQYPELADSSSNIQFLRQNEGKGR