

*C. gigas* 1  
*C. elegans*  
*H. sapiens*  
*M. musculus*  
*D. rerio*

10	20	30	40	50	60	70		
MYDMFOALLQPPSYHLFFEALLIWI <del>F</del> KLLFFSKAYAP	---	ESVLT EKEKEELIA	EWQP EPLA PEIFDD					
MGFLFD <del>SWH</del> FYIETLLVALLAYVVMRNRSKRQOEKL <del>S</del> KKLT	E	EROKDELIA	DWTP EPLVPETFQD					
MATAT EQWVLVEMVOALYEAPAYHLLILEGIL <del>L</del> ILWI <del>R</del>	-L	LF SKTYKLOE	-- RSDLTVKEKEELIE	EWQP EPLVPPVPKD				
MATVA EQWVLVEMVOALYEAPAYHLLILEGIL <del>L</del> ILWI <del>R</del>	-L	VF SKTYKLOE	-- RSDLTAKEKEELIE	EWQP EPLVPPVSKN				
MASG -OOWVLVEMVOAFYEA <del>P</del> A <del>Y</del> HLLILEGFL <del>L</del> ILWI <del>R</del>	-L	LF SKTYKLOE	-- RSDLTEKEKEELIE	EWQP EPLVSPVSKD				
100	110	120	130	140	150	160	170	
KPGKYVTINGKSCVN <del>M</del> ATLNFLGMAGNPSEAEAIKTLK <del>K</del> YGVGSCGPRGFYGT	M	DVHLELEDKI <del>A</del> KFMNC	EEAIIYLAF					
KMTKDVSIDGEKYLNMASTNFLSF <del>I</del> GVKRIEDRAKOTIF <del>K</del> YGVGSCGPRGFYGT	V	DVHLDLEKELAKFMG	C	EEAVLYSY				
PPSHKTVVNGKECINFASFNFLGLLDNP <del>R</del> VKA <del>A</del> ALA <del>S</del> LKKYGVGT	CGPRGFYGT	FDVHLDLEDRLAKFM	K	TEEAIIY	SY			
PPTHNIVVNGKECVNFASFNFLGLLDNP <del>R</del> VKA <del>T</del> AF <del>S</del> SLKKYGVGT	CGPRGFYGT	FDVHLDLEERLAKFM	K	TEEAIIY	SY			
PPSHKIIIVNGKECINFASFNFLGLLDNERVKLKALASLKKYGVGT	CGPRGFYGT	FDVHLEERLAKFM	R	TEEAIIY	SY			
190	200	210	220	230	240	250	260	
RGDVIFADEGVCFAIQKGLVASRSKIKWF <del>H</del> NMD <del>E</del> LERLLI	E	QAKEDDKNPKAKV	TRRF	LVVEGLYIN	Y	GDL	CPLPK	
KGDVIFVD <del>E</del> GVNFAIQKGLQASRSRVEYFKHNDME <del>H</del> LERLLI	E	EQEQRDKDPKAK	SVRR	FIVV	EGLYVN	Y	ADL	CPLPK
RGDIVFVDRAACFAIQKGLQASRS <del>D</del> IKLF <del>H</del> NMD <del>A</del> LERLLK	E	QE <del>E</del> EDOKNPRKAR	V	TRRF	FIVV	EGLYMN	TGT	ICPLPE
RGDIIFVD <del>S</del> AACFAIQKGLQASRS <del>D</del> IKLF <del>H</del> NDA <del>V</del> ADERLLK	E	QE <del>E</del> EDOKNPRKAR	V	TRRF	FIVV	EGLYMN	TGT	ICPLPE
RGDIIFVD <del>E</del> AACFSIQKGLQASRSFIKYFKHNDM <del>E</del> LERLLK	E	QE <del>E</del> EDOKNPRKAR	V	IRR	FIL	VEGLYINT	TAD	CPLPE
280	290	300	310	320	330	340	350	
EESLSFGILGSNGKVT <del>E</del> HYNI <del>S</del> PDDIDLIAASLENAIGST	GGFCCGKKY	I	VDHQRLSGL	GYCF	SASLPPMLATAAIE			
DESW <del>S</del> FGVIGKTGRGVTEHFNVPMEDVDMVMASLENALAST	GGFCVGR	SYVVGH	HQRLSGL	GYCF	SASLPPPLLATAASEA			
EESLSFGVLGEHGRGVTEHYG <del>I</del> SIDIDLISANMENALAS <del>I</del>	GGFCCGR	SFVID	HQRLSG	GYCF	SASLPPLLAAAI	EA		
EESLSFGVLGEHGRGVTEHYG <del>I</del> SIDIDLISANMENALASVGGFCCGRSFVVDHQRLSG	GGFCCGR	SFVID	HQRLSG	GYCF	SASLPPLLAAAI	EA		
EESMSFGVLGEHGRGVTEHF <del>G</del> VNIIDIDLISANMENALAS <del>I</del>	GGFCCGR	SFVID	HQRLSG	GYCF	SASLPPMLAAAI	EA		
380	390	400	410	420	430	440	450	
RENCEKITHSKLSDING	--	-TVIVGEPI <del>S</del> PVKHIRLAEPSTD <del>R</del> DFDVOTLOK <del>I</del> ADL <del>S</del> RDN	-KVAVT	LARYLEEEEHK	LPL			
TEMAINGQKKL <del>O</del> DALSGSKF <del>S</del> LQGCPESP <del>M</del> KHIYYNG	--	ED <del>E</del> EKQLDTF <del>V</del> ETVFTKNHLLL <del>T</del> RARYL <del>D</del> KDEL <del>F</del> KIR						
KEKCGQIHKALOGI <del>S</del> G	--	LKVVGESLSPA <del>H</del> LQLEESTG <del>S</del> REQDVRL <del>L</del> OEV <del>I</del> VDQCMNR	-SIALTO	QARYLEKEEK	CLPP			
KKKQNIHK <del>S</del> LOGVSG	--	LKVVGESLSPA <del>H</del> LQLEESTG <del>S</del> REKDV <del>K</del> LLOA <del>I</del> VDQCMDK	-GIALTO	QARYLDKEEK	CLPP			
REKCRNVHKALOGTAG	--	LKVVGESFAPAL <del>H</del> LQLEN <del>S</del> TG <del>S</del> RVNDL <del>K</del> LLRA <del>I</del> VDYSLDR	-QVALT	LARYLDKEER	F	CLPP		
470	480	484						
EIDTVFTTLSEAFQK <del>I</del> TH								
EIQRAVDAIRVVAHKF								
ELERA <del>A</del> STIKEVAQAVLL								
ELQRA <del>A</del> STIREAAAQAVLL								
EIEKAAQCIREAALHV <del>L</del> K								