

ScAtg11	-----MADADEYSTAPQQEITPLQTTATIINAI	SGECITTNVDF---FV	42																				
PfAtg11	YTKNKSDFNNYNLNESLNDSYKDKTFEELKSQ	LIKQKNLNSCLSLKL	SIHKLNNFL	60																			
ScAtg11	SLDKFKQFIARKWKIPPDQLLILLPYGNKL	KPSMFELLINRSFTLN	---	DFYVYDRRL	98																		
PfAtg11	KKNEFEKTINN	KEIELQFLIQNVTHEN	KGIPISKELEQEKH	LQNSHENVAI	YDTHE	120																	
ScAtg11	FSLVSKPTPTNLLTSKDSNPMNSPNSNDLTETLEY	LIKNSHISQYQGSDTIMIKPMPSP	L	158																			
PfAtg11	IENNDKKKLYTNFHNEEKDHLC	KLLEEYSKTL	EIYKMGKIQLEFELKCC	KEKLNEEIEKN	180																		
ScAtg11	EDADVDSL--RLNYHSVTSLLTTNLGWLSALEIDVHYFKS	-----LIPDIIAHIK	206																				
PfAtg11	NNYNNKMKSYEIHDVVKNENCKN	LEELNDLKLQLEKT	KSENNQNYVKNKILN	DEKNNLD	240																		
ScAtg11	RIFDGLTVCSQYLKLYCFDVES	-----LYNSNVQFLNQLVDNGMTS	247																				
PfAtg11	KINNDLKIKIKNF	TLLNDAQNKEYILNNFTQKIFNIITYLKN	NDEHNFLNDKIHNKLDI	300																			
ScAtg11	KWEKCFNDTLSKLTALEGDSLQKF	INIESLLEN-----EKSVKILNHSINGKLN	KIREI	302																			
PfAtg11	NQDQIYVQTELYIDIIS	-SSIRNLINFKKTLEERNVEL	EKVTHEM	KELKELILKKNYE	359																		
ScAtg11	DENASFRDIITVNIDRRLQMFTPNESK	--FELLEDQMAESFEVLVSEMRTR	SRNVLDKEE	359																			
PfAtg11	ELRLKLNHLECVERDSVKINSEKEGEKVIY	EIKEKLDNDEKI	INDLKKNSYQVYKMD	419																			
ScAtg11	E EF NSQEF FL SMN	--VMLEKD-----KKESVKT	LFTISQALYSQIGELIDLKKS	406																			
PfAtg11	YEKRENNLINE	IN	KLKL	FIEENKMTVERGNEMNNKLEEMKQKNEL	INNLNDISDELKN	479																	
ScAtg11	LQKHAVAILGNI	IAFTQMEILGIKRLL	NNECNKDLELYKKYEVEFA	QVEDLPLIYGLYLIE	466																		
PfAtg11	CIEQVNSVSRN	MANVEKEKENI	INELQILRMK	NNDTMRKRISKFVE	QEKVLFKLYTLNND	539																	
ScAtg11	KYRRLSWFQQILSFISNF	NQDLELFKQNELRTRNKWV	KNFGSIATVF	CEDLLSSDF	KRL	526																	
PfAtg11	IFSKNEKLNDMQKKLNDV	NEYK	KNIVECLNNYKTEHKE	QIEKKIERINTL	KQNYYYLKKE	599																	
ScAtg11	NEYHSHTSPPNEDEEDE	DENENSIANYRQDLV	KVSQA	IDNYMTQIKETDV	SEPII	DLLSKTL	586																
PfAtg11	YDLKNKELEK	NIEHGKK	LEHEL	SHCYEENQ	KLNEE	IKRRNSFI	KNKDRKIDLL	TNIENE	L	659													
ScAtg11	FETKRFHI	IYSNFKN	--NNNNSSNGNSISPEGSIAL	KSDVV	KGYKT	RIKKLESLLHEF	643																
PfAtg11	PKLKE	KEINNI	KLMEK	QNVIKNN	EQLL	KDIKDEN	EKMNEHVNL	QNLQNELIK	RELQNCISKDI	719													
ScAtg11	QYS-----DIGHWPQGVLN	--THLKPF	GRGSATSINK	---	KKFLGASVL	LEPANISEVN	691																
PfAtg11	EFCKKE	EDK	IKNLED	DLLEKK	KCIEN	LKDELIN	IKKKM	MEDK	MHMTNEMD	LLSNKVE	ELN	779											
ScAtg11	-IDS	VSQANH	QI	KELES	NVDD	LLHQLQL	LKEE	NNR	KSMQI	SEM	GKK	ISDLEV	EKTAYRE	750									
PfAtg11	RINKTYE	KNIV	ELN	NEL	NDV	IKKKL	LNDEF	LKEEE	KKKN	IDMVY	KI	KEYE	IQIKE	EKEN	839								
ScAtg11	TLTNL	QELAR	LTNE	QSHRTE	IFTLN	ASF	KKQ	LN	NDI	IISQDNE	KIEK	---	LTG	DYDDVS	806								
PfAtg11	SLKKN	EQNLH	V	IK	NEEL	NEKEI	ILK	N	--	KYD	KE	I	NM	II	EQ	YNKKI	QEE	KDM	LN	NN	IK	KSMD	897
ScAtg11	KSRERL	LQMDL	DE	SNKK	HEQEVN	-----	LLK	A	DIE	RLGKQ	I	V	T	SEKSYA	E	T	NSSS	855					
PfAtg11	QTHKN	QIE	M	EE	NKK	EL	KRL	KN	VCD	MNL	QSQI	L	I	K	NE	KHM	QE	KV	EEY	KN	LL	KQDQEL	957
ScAtg11	MEKGEKFETI	PLAEDPG	REN	QI	SAYT	QT	L	QDR	IFD	---	I	I	STN	I	FILE	N	IG	LLL	TFD	NN	911		
PfAtg11	KNII	Q	HEY	DERIE	I	QN	KEME	DIV	NDCE	EK	L	KQAK	I	NN	KKL	TT	A	TN	MANN	NN	MLMDEN	KEK	1017

Figure S1. Alignment of the yeast Atg11 (ScAtg11) and its putative *P. falciparum* homolog (PfAtg11) was performed using the Clustal W2 program. Shown are conserved residues (in red).