

NADP-specific Glutamate Dehydrogenase

Heat Shock Protein 70

LSU Ribosomal Protein L10P

1	10	20	30	40	50	6
-----	MPKAKRDKLITLT	KTKRSTKEQKNLVEEIRNAIEFNN	-----	IYI		
-----	MAKS KRVKKVLM	T KDLKK-KRKDKSEIIENVHEYIGKF	-----	FIYV		
-----	MAKS KRVKKVLM	T KDLKK-KRKDKSEIIENVHEYIGKF	-----	FIYV		
-----	MAISKR-RNPLHTSICK	-DGKKKIKLVENIRETLENFIKKNNDNSTFVYL	-----			
-----	MAVS KR-RNP	AVTSSCKK-DSKKVKLVENIRETLDKFIKNNDNSTFVYL	-----	VYT		
-----	MPKSKRAKIVPLT	KTQAK-TREDKD K LIERIREALEDYS	-----	VYT		
-----	MPKSKRAKIVPLT	KTQAK-TREDKD K LIERIREALEDYS	-----	VYT		
-----	MPKSKRAKIVPLT	KTRAK-TREDKD K LINRIRKA LDYTD	-----	VYS		
-----	MPKSKRATIVPLT	KTRSK-TREEKDELINKIRDALDDYTD	-----	VYT		
MIYFLKCNLLM	PRSKRSKTVVLAKVEKK-TREAKQEIIKQIREAFDTYDT	-----	VYV			
MIYFLKCNLLM	PRSKRSKTVVLAKVEKK-TREAKQEIIKQIREAFDTYDT	-----	VYV			
-----	MPPRSKRSKTVVL	TKVEKK-TREAKQEIIKQIREAFDTYDT	-----	VYV		
-----	MPPRSKRSKTVVL	TKVEKK-TREAKQEIIKQIREAFDTYDT	-----	VYV		
-----	MAKL SKQ--QKKQMYIEKLSSLIQQYSK	-----	ILI			
-----	MAKL SKA--QKKQIYMDKLSSLIQQYTK	-----	ILI			
-----	MPFEKSAHQ--NKDDYFKILHDLFHNYNK	-----	VAV			
LDLSNCRTNHV	KEIRRDFNESKFFMTKNKVMRVALGTSPENEVENDLN	HASKFL T A	R			
VKLKNQRNAALKQLRVR	LEPGK-LLVGKNKL	LQVAFGADSDSESAKNAHKISSFLRG	ERG			
VKLKNQRNAALKQLRVR	LEPGK-LLVGKNKL	LQVAFGADSDSESAKNAHKISSFLRG	ERG			
ISLNNQRNSPLKTLRSLLL	PGR-VFYGKNKVMRIAGFTKPEDEIHNI	H KISKNINGETA				
IALNNQRNSPLKTLRSILL	PGR-VFYGKNKVMRIAGFTKPEDEIHNI	H KISKNINGETA				
FQLHNIRTNILQQIREER	AGDSRIFLGNNKIMMIAIGRDEESAQ	KPNLHKLSPFLTGLCG				
FQLHNIRTNILQQIREER	AGDSRIFLGNNKIMMIAIGRDEESAQ	RQNLHKLSPFLTGLCG				
FTLSNVRTNILQQIREER	KGDCLFLGNKLIMIALGRDEESSQRP	NLHKLCFLVGSCG				
FELSNIRTNILQQIREER	RND SRLFLGNKLIMIALGRDDSSQRP	NLHKLSKYLTGSCG				
IDSHNMNTSSWQQLRTSM	KGYARIFMGKNQLMRYALGKTEESYRT	KTWQLGRLLKGMTG				
IDSHNMNTSSWQQLRTSM	KGYARIFMGKNQLMRYALGKTEESYRT	KTWQLGRLLKGMTG				
IDSHNMNSSSWQQLRTAM	KDYARIFMGKNQLMRYALGKTEESYRSK	TWQLGRLLKGMTG				
VHVDNVGSNQMASVR	KS LRKGATILMGKNTRIRTA	LALKN--LQAVPQIEKLLPLV	KLNMG			
VHVDNVGSNQMATVR	QSLRGKAIILMGKNTRIRTA	LALKN--LQTPQIEKLLPLV	KLNMG			
VTSQQVKS	NQLMKIRTDI	SSFAEIVFCNSLMRAVDQLK--REIPSITELE	KYL TNGAG			
L FITDRPKA	EVAKYFANFKRS	DYAKVGFATKTITLEA	G PVD--WMPFSLEER	LLNLGLP		
LIFTDLAPS	NLK VLEES	STMEFGREGSLSDITCV	VEPN-TELECL	YKNAEFYMRKQ-FP		
LIFTDLAPS	NLK VLEES	STMEFGREGSLSDITCV	VEPN-TELECL	YKNAEFYMRKQ-FP		
VLITSEDPEVV	VIKKVKEYKVRDFSKAGNIATDT	IVLKVVDGEFDEIPGSME	PQFRQLGLMP			
VLITSENPEVV	VNKVKG YKVRDFSKAGNIATDT	IVLKVVDGEFDEIPGSME	PQFRQLGLP			
LLFTNL	SKKEVKEYFATVGAPVYARTG	QTATESLVLKGGLP--QFP	PHSMFDHLAKLGLP			
LLFTNL	SKKEVKEYFATVGAPVYARTG	QTATESLVLKGGLP--QFP	PHSMFDHLAKLGLP			
LLFTNL	PKKEVKSYFASVGAQVYARTG	QTATSSLVLRAGGLP--QFT	PHSMFDHL	SKLGLP		
LLFTNL	PHQDVKEYFKGVSADV	FARTGQVATCP	LLLRTGPLA--QFP	PHSMFDHLSRLGLP		
LLFTSAPEEKV	QVQ SALASVARP	CLARGDVATKTIVIPQ	GPLDRDKYSFALEPEL	RKLGLP		
LLFTSAPEEKV	QVQ SALASVARP	CLARGDVATKTIVIPQ	GPLDRDKYSFALEPEL	RKLGLP		
LLFTSAPEEKV	QVQ SALASVARP	CLARGDVATKTIVIPQ	GPLDRDKYSFALEPEL	RKLGLP		
FVFC	KDDLSEIRNIILDNKSP	A PARLGVIAPIDVF	IPPGPTG--MDPSHTSFF	QSLGIS		
FVFC	KDDLTEVRNIILQNKS	PAPARLGVIAPIDVF	IPPGPTG--MDPSHTSFF	QSLGIS		
LIFTNGSF	KAIKEVIDANC	CRYSAAKAGSIAPCDV	I K R Q L T S--IPYYEYNL	FFDLQIP		
IEVK	NKVINILRPFNL	C TFGEPIAQ	QAKLLKHFGHELTEFN	ASIVCHYDKKTH	LEKFV	
QLKPTL	FGSDQGKVI	CEKGNPLNKYQY	LLLKHLEIPSVK	FEIKPIAC	HNEELTHFENA	
QLKPTL	LGSDQGKVI	CEKGNPLNKYQY	LLLKHLEIPSVK	FEIKPIAC	HNEELTHFENA	
TALNMGK	IILMGDFTL	CEKDPLTPNQ	THLLKLFGIRMSL	FEANV	LGFWNDGHYKQLA	
TALNMGK	IILMGDFTL	CEKDPLTPNQ	THLLKLFGIRMSL	FEANV	VGFWNDGHYKQLA	
IKLDRGV	VIVLLQDT	TVCEPGDTL	SAAQ	LLKLFGVQSAE	KMDLTAHWTNG--VAKK	
IKLDRGV	VIVLLQDT	TVCEPGDTL	SAAQ	LLKLFGVQSAE	KMDLTAHWTSG--VAKK	
LKLDKG	VIVLLQDT	CEANDTL	SAAQ	LLKLFGIQSAE	FRIDLTAHWANG--AASRI	
IKLDKG	VIVLLRDT	TVCEVGDTL	TA	AAQ	LLKLFGIQSAE	FKLET
IKLDKG	VIVLLRDT	TVCEVGDTL	TA	AAQ	LLKLFGIQSAE	FKLET
TSLQNTV	IHVLC	DHVLCKEGDVL	TSQA	QARLLKHF	DVL	
TSLQNTV	IHVLC	DHVLCKEGDVL	TSQA	QARLLKHF	DVL	
TSLQNTV	IHVLC	DHVLCKEGDVL	TSQA	QARLLKHF	DVL	
TSLQNTV	IHVLC	DHVLCKEGDVL	TSQA	QARLLKHF	DVL	
TSLQNTV	IHVLC	DHVLCKEGDVL	TSQA	QARLLKHF	DVL	
TKIVKGQIEI	QEHVHL	IKQGEVTASSAT	LLQFNMKPF	SYGV	DVVRTVYDDGVIYDAK	
TKIVKGQIEI	QEVHHL	IKQGEVTASSAT	LLQFNMKPF	SYGV	DVVRTVYDDGVIYDAK	
CKIFKG	TIEVAGKQL	VWKQKIRASEARV	LEMLGI	KPF	KHTLKIEVLYDNGT	CDPEAC
EEPTGDEEEGDDDEMEY	--	--	--	--	--	
DME	--	--	--	--	--	
DME	--	--	--	--	--	
SN-SASKRSGK	--	--	--	--	--	
SS-SASKRSEK	--	--	--	--	--	
VGRNEENV	TGKA-	--	--	--	--	
SKKKKEE	EN--	--	--	--	--	
PNCLYLLD	TAPAEEDEQSD	--	--	--	--	
PNCLYLLD	TAPAEEDEQSD	--	--	--	--	
ADCLYLLD	TAPAEEDEQSD	--	--	--	--	
ANCLYLLD	TAPAEEDEQSD	--	--	--	--	
DITDEDILE	KFSKG	VNSVA	ALSRATGV	IT EASY	PHVF	EA
DITEEDIL	A KFSKG	VANVA	ALSRSIGV	IT EASY	PHVF	EA
GLGNEIDEH	QEEEEEEPM	ILDDSNY	FDFDF	--	--	

Protein Translation Elongation Factor 2 (EF-2)

Sec61 Alpha Subunit

Ribosomal Protein L14p/L23e

	1	10	20	30	40	50	60	
Naegleria gruberi strain NEG-M	-----	MASKRATGGLGNKF	RISTALPTAAVMNCAD	NSGAKNLYVIAVNQTQ	GRLNRLP			
Giardia lamblia GS_B	-----	MASKGRSG-GIGNK	FRISCGLPVGAVINC	CADNSGAKSLNMIAVKNW	KGHLNRFP			
Giardia lamblia WB_A1	-----	MASKGRSG-GIGNK	FRISCGLPVGAVINC	CADNSGAKSLNMIAVKNW	KGHLNRFP			
Giardia lamblia DH_A2	-----	MASKGRSG-GIGNK	FRISCGLPVGAVINC	CADNSGAKSLNMIAVKNW	KGHLNRFP			
Giardia lamblia Pig_E	-----	MASKGRSG-GIGNK	FRISCGLPVGAVINC	CADNSGAKSLNMIAVKNW	KGHLNRFP			
Trichomonas vaginalis G3	-----	MSKRRTG-QQGT	KFAMTAGLPVGAVINCC	CDNSGAKNMFIISVRGH	KGRLNRLP			
Cryptosporidium hominis TU502	-----	MKRGRA-AGGAKM	RITLGLNVGALINCCD	NSGGKNLYIIAVKG	TGSCLNRLP			
Cryptosporidium parvum Iowa II	YSLYLEKMKRGRGA-	AGGAKM	RITLGLNVGALINCCD	NSGGKNLYIIAVKG	TGSCLNRLP			
Theileria annulata strain Ankara	-----	MKRGRRGG-SGGNKL	RVTLGLPVGALINCCD	NSGGKNLYLIAVKG	TGACLNRLP			
Theileria parva strain Muguga	-----	MKRGRRGG-SGGNKL	RVTLGLPVGALINCCD	NSGGKNLYLIAVKG	TGACLNRLP			
Plasmodium falciparum 3D7	-----	MKRGAG-TLKNKM	RITLSPVGALINCCD	NSGGKNLYIIAVQGF	GSCLNRLP			
Plasmodium knowlesi strain H	-----	MKRGAG-TLKNKM	RITLSPVGALINCCD	NSGGKNLYIIAVQGF	GSCLNRLP			
Leishmania infantum JPCM5	-----	MGKDQAN-VKGCR	FRVSVALPVGAVVNC	ADNTGAKNLYVISVK	GYHGRLNRLP			
Leishmania major strain Friedlin	-----	MGKDQAN-VKGCR	FRVSVALPVGAVVNC	ADNTGAKNLYVISVK	GYHGRLNRLP			
Trypanosoma brucei gambiense DAL972	-----	MGDKAN-VKGCR	FRVSLALPVGAVVNC	ADNTGAKNLYIISVK	GYHGRLNRLP			
Trypanosoma cruzi strain CL Brener	-----	MGDKAN-VKGCR	FRVSLALPVGAVVNC	ADNTGAKNLYIISVK	GYHGRLNRLP			
Naegleria gruberi strain NEG-M	AAGVGDMVMATVKK-	GKPELRKKVMPAVVIR	QRPWRRDGTFIYFEDN	AGVIVNNK	GEM			
Giardia lamblia GS_B	KASVGDMVLITVGRDG	KPELRRKVTPAVVV	RQRKVWRRKDGV	WIYFQDNAG	VIINPK	GEL		
Giardia lamblia WB_A1	KASVGDMVLITVGRDG	KPELRRKVTPAVVV	RQRKVWRRKDGV	WIYFQDNAG	VIINPK	GEL		
Giardia lamblia DH_A2	KASVGDMVLITVGRDG	KPELRRKVTPAVVV	RQRKIWRRKDGV	WIYFQDNAG	VIINPK	GEL		
Giardia lamblia Pig_E	KASVGDMVLITVGRDG	KPELRRKVTPAVVV	RQRKIWRRKDGV	WIYFQDNAG	VIINPK	GEL		
Trichomonas vaginalis G3	AASVSDLIVVTC	KK-GKPALRK	KVSMGVVV	RQRAIWR	RKDGV	VIGFQDNAG	VIINDK	GEM
Cryptosporidium hominis TU502	SASIGDMVLATVKK-	GKPELRKKVWP	AVIVRQRKAFRR	PEGTF	FLYFEDN	AGVIVNP	KGEM	
Cryptosporidium parvum Iowa II	SASIGDMVLATVKK-	GKPELRKKVWP	AVIVRQRKAFRR	PEGTF	FLYFEDN	AGVIVNP	KGEM	
Theileria annulata strain Ankara	SASVGDMVLATVKK-	GRPDLRK	KVLP	AVIVRQRKAWRR	REGYFIYFEDN	AGVIVNP	KGEM	
Theileria parva strain Muguga	SASVGDMVLATVKK-	GRPDLRK	KVLP	AVIVRQRKAWRR	REGYFIYFEDN	AGVIVNP	KGEM	
Plasmodium falciparum 3D7	SASVGDMVLATVKK-	GRPDLRK	KVLP	AVIVRQRKAWRR	REGYFIYFEDN	AGVIVNP	KGEM	
Plasmodium knowlesi strain H	AASLGDMVLATVKK-	GKPDLRK	KVLN	AIICRQSKAWRR	HEGYYIYFEDN	AGVIVNP	KGEM	
Leishmania infantum JPCM5	AASLGDMVLATVKK-	GKPDLRK	KVLN	AIICRQSKAWRR	HEGYYIYFEDN	AGVIVNP	KGEM	
Leishmania major strain Friedlin	SAALGDMVMCSVKK-	GKPELRKKV	LNAVI	IRQRKS	WRRKDGT	VIYFEDN	AGVIVNP	KGEM
Trypanosoma brucei gambiense DAL972	SAALGDMVMCSVKK-	GKPELRKKV	LNAVI	IRQRKS	WRRKDGT	VIYFEDN	AGVIVNP	KGEM
Trypanosoma cruzi strain CL Brener	AAALGDMVMASVKK-	GKPELRKKV	LNAVI	IRQRKS	WRRKDGT	VIYFEDN	AGVIVNP	KGEM
Naegleria gruberi strain NEG-M	AAALGDIVMASVKK-	GKPELRKKV	LNAVI	IRQRKS	WRRKDGT	VIYFEDN	AGVIVNP	KGEM
Giardia lamblia GS_B	KGSAVAGPVAKECAD	MWPRIASSASC	VV					
Giardia lamblia WB_A1	KGSSITGPVAKEAAD	RWPKISTTAEC	CVH					
Giardia lamblia DH_A2	KGSSITGPVAKEAAD	RWPKISTTAEC	CVH					
Giardia lamblia Pig_E	KGSSITGPVAKEAAD	RWPKISTTAEC	CVH					
Trichomonas vaginalis G3	KGSAITGPVAKEAAE	LWPKVASVAPAV	F					
Cryptosporidium hominis TU502	KGSAITGPVGKECA	EELWPKVSAAPSIV						
Cryptosporidium parvum Iowa II	KGSAITGPVGKECA	EELWPKVSAAPSIV						
Theileria annulata strain Ankara	KGSAINGPVAKECA	EELWPKISAAAPSIV						
Theileria parva strain Muguga	KGSAINGPVAKECA	EELWPKISAAAPSIV						
Plasmodium falciparum 3D7	KGSAITGPVARECA	EELWPKLSSAASAIV						
Plasmodium knowlesi strain H	KGSAITGPVARECA	EELWPKLSSAASAIV						
Leishmania infantum JPCM5	KGSGIAGPVAKEAAE	LWPKISTHAPAIV						
Leishmania major strain Friedlin	KGSGIAGPVAKEAAE	LWPKISTHAPAIV						
Trypanosoma brucei gambiense DAL972	KGSGIAGPVAKEAAE	LWPKISTHAPAIV						
Trypanosoma cruzi strain CL Brener	KGSGIAGPVAKEAAE	LWPKISSHAPAIV						

SSU Ribosomal Protein_S2P

	1	10	20	30	40	50	60
Naegleria gruberi strain NEG-M	-MSLTNAKSNI	DLTPTDVQQMLACECHGKTNLE	TDMQKVYHGRSQKDGT	HVINVAKT			
Cryptosporidium hominis TU502	-----MPPAETRK	LDDIAKMIVCKTHIGTKNVED	KMLSYVYKRT	-HEGIFLNLAKT			
Cryptosporidium parvum Iowa II	----LYSNMPPAETRK	LDDIAKMIVCKTHIGTKNVED	KMLSYVYKRT	-HEGIFLNLAKT			
Plasmodium falciparum 3D7	----MS-NKKGQSP	KEESIAKMLICKVHIGTKNLEN	ENKMKRYVYTRA	-KDGVIHILNLAKT			
Plasmodium knowlesi strain H	----MSTNKKVQSP	KEESIAKMLICKVHIGTKNLEN	ENKMKRYVYTRA	-KDGVIHILNLAKT			
Theileria annulata strain Ankara	----MAAPTSKL	TLPDEDSIRMMTAKVHIGTKN	VENKMRKVYVSRT	-QEGVHLINLAHT			
Theileria parva strain Muguga	----MAAPTSKL	TLPDEDSIRMMTAKVHIGTKN	VENKMRKVYVSRT	-QEGVHLINLAHT			
Leishmania infantum JPCM5	--MTAVESGSKVL	RMKESDAQKLLAMRCHIGTRN	QSSAMKKYIYGRT	-AEGSHIIDLHMM			
Leishmania major strain Friedlin	--MSAVEGS	SKVLRMKESDAQKLLAMRCHIGTRN	QSSAMKKYIYGRT	-AEGSHIIDLHMM			
Trypanosoma brucei gambiense DAL972	--MTSVESSGAKV	LRMKESDVQKLAMHCHLG	TKNSSSAMKKYIHGRT	-SDGTNIIDIHMT			
Trypanosoma cruzi strain CL Brener	--MTSVESGAKV	LRMKEGDVQKLVAMHCHLG	TKNRSNAMKKYIHSRT	-KEGTNIIDLHMT			
Trichomonas vaginalis G3	-----MSTA	AAAADREAVIKMMLAAKVYL	GSTNL	FGMKPVFTCR	-KDGTHVINLAMT		
Giardia lamblia DH_A2	MSTEKT	SQASKEYQLKEADVKKML	VATTHLGVRNIDRRM	QFYIFDRQ	-KDGTFVFNLQKV		
Giardia lamblia WB_A1	MSTEKT	SQASKEYQLKEADVKKML	VATTHLGVRNIDRRM	QFYIFDRQ	-KDGTFVFNLQKV		
Giardia lamblia Pig_E	MSTEKT	SQASKEYQLKEADVKKML	VATTHLGVRNIDRRM	QFYIFDRQ	-KDGTFVFNLQKV		
Giardia lamblia GS_B	MSTEKTSHAS	EHQLKEADVKRML	VATTHLGVRNIDRRM	QFYIFDRQ	-KDGTFVFNLQKV		
Naegleria gruberi strain NEG-M	HEKLLAARVLVTIENPKDICA	ISSKQYGHRAVLKFGSYTG	ANYLAGKFTP	PGTFTNQVQA			
Cryptosporidium hominis TU502	WEKIQIAARI	IATIDNLADV	VVSQRPYGSR	PALKFAQHTGAH	AMVGRFT	PGTLTNQITQ	
Cryptosporidium parvum Iowa II	WEKIQIAARI	IATIDNLADV	VVSQRPYGSR	PALKFAQHTGAH	AMVGRFT	PGTLTNQITQ	
Plasmodium falciparum 3D7	YEKLQLAARI	IVAI	SNPADV	VVSARPGF	SRAVLKFAQYTGA	QAIAGRWT	PGMLTNQIIQ
Plasmodium knowlesi strain H	YEKLQLAARI	IVAINN	NPADV	VVSARPGF	SRAVLKFAQYTGA	QAIAGRWT	PGMLTNQIIQ
Theileria annulata strain Ankara	LEKLKVAARA	I	IAITVSNPEE	VVSARPGF	SRAVLKFSH	YVGSHPIAGR	WIPGTLTNQITQ
Theileria parva strain Muguga	LEKLKVAARA	I	IAITVSNPEE	VVSARPGF	SRAVLKFSH	YVGSHPIAGR	WIPGTLTNQITQ
Leishmania infantum JPCM5	WEKLILAAR	VIAAVENPKDVC	CVCSRLY	GTRAIYKF	SQHVGT	FHGGRF	IPGTTFNQIJK
Leishmania major strain Friedlin	WEKLILAAR	VIAAVENPKDVC	CVCSRLY	GTRAIYKF	SQHVGT	FHGGRF	IPGTTFNQIJK
Trypanosoma brucei gambiense DAL972	WEKLILAAR	VIAAVENPQDVT	CSTR	FGQRAIFKF	SQLVGT	FLGGRF	IPGTTFNQIJK
Trypanosoma cruzi strain CL Brener	WEKLILAAR	VIAAVENPQDVT	CSTR	FGQRAIFKF	SQLVGT	FLAGRF	IPGTTFNQIJK
Trichomonas vaginalis G3	YEKIKLAARM	YAVEEP	KDVL	AISTKSF	TRAVHKFAEFL	GANYVEQR	FTPGLFTNYSIK
Giardia lamblia DH_A2	WAKIVFAARI	LVTIDDPA	EIAVVANRPDA	QRSILK	FCKYTHATAFPGR	FIPGNFT	NRMNP
Giardia lamblia WB_A1	WAKIVFAARI	LVTIDDPA	EIAVVANRPDA	QRAILK	FCKYTHATAFPGR	FIPGNFT	NRMNP
Giardia lamblia Pig_E	WAKIVFAARI	LVTVD	DPAEI	AVANRPDA	QRAILK	FCKYTHATAFPGR	FIPGNFT
Giardia lamblia GS_B	WAKIVFAARI	LVTIDDPA	EIAVVANRPDA	QRAILK	FCKYTHATAFPGR	FIPGNFT	NRMNP
Naegleria gruberi strain NEG-M	KFMEPRVLIVSDP	RSDYQAIKEASK	KANI	PVIAFAD	TDSPLKFVDIAIPVNN	KNKNSIALM	
Cryptosporidium hominis TU502	KFMEPRLLIVTD	PRVDSQAVI	ESSYANIP	VIALCDT	DSPLQYV	DVAIPCN	NKGKESIALM
Cryptosporidium parvum Iowa II	KFMEPRLLIVTD	PRVDSQAVI	ESSYANIP	VIALCDT	DSPLQYV	DVAIPCN	NKGKESIALM
Plasmodium falciparum 3D7	KFTEPRLLIVTD	PRVDSQAVI	ESAYANIP	VIALCSD	DSPL	EHDIAIP	CNNKGKESIALM
Plasmodium knowlesi strain H	KFTEPRLLIVTD	PRVDSQAVI	ESAYANIP	VIALCSD	DSPL	EHDIAIP	CNNKGKESIALM
Theileria annulata strain Ankara	KFIEPRLLVATD	PRVDSQAVI	ESAYANIP	VIALCSD	DSPL	EHDIAIP	CNNKGKESIALM
Theileria parva strain Muguga	KFIEPRLLVATD	PRVDSQAVI	ESAYANIP	VIALCSD	DSPL	EHDIAIP	CNNKGKESIALM
Leishmania infantum JPCM5	KFVQPRVLVVT	DPRTDHQAIRE	ASLVNIP	VIALCDT	DAPLEYV	DIAIPC	NNRGIKSIGMM
Leishmania major strain Friedlin	KFVQPRVLVVT	DPRTDHQAIRE	ASLVNIP	VIALCDT	DAPLEYV	DIAIPC	NNRGIKSIGMM
Trypanosoma brucei gambiense DAL972	KFMQPRVLVVT	DPRTDHQAIRE	ASLVNIP	VIAFC	CDT	DAPLEYV	DIAIPC
Trypanosoma cruzi strain CL Brener	KFMQPRVLVVT	DPRTDHQAIRE	ASLVNIP	VIAFC	CDT	DAPLEYV	DIAIPC
Trichomonas vaginalis G3	NFCEPRLMIVCDP	NTDSQAV	HEAAYANIP	CIALCDT	DAHLDYVDC	VIPCN	TKNNSMLV
Giardia lamblia DH_A2	NYCEPRLLLVNDP	VVDRQAILE	ASYVNIP	TC	TFN	YD	TKPMSIGLI
Giardia lamblia WB_A1	NYCEPRLLLVNDP	VVDRQAILE	ASYVNIP	TC	TFN	YD	TKPMSIGLI
Giardia lamblia Pig_E	NYCEPRLLLVNDP	VVDRQAILE	ASYVNIP	TC	TFN	YD	TKPMSIGLI
Giardia lamblia GS_B	NYCEPRLLLVNDP	VVDRQAILE	ASYVNIP	TC	TFN	YD	TKPMSIGLI
Naegleria gruberi strain NEG-M	YWMLAREV	RLKGV	VSRK	KEWDV	KPD	FLH	RDLDE
Cryptosporidium hominis TU502	YYLLAREVN	FLRGK	TDK	---	-----	KKEEREAKESQ	--ETKVQAV
Cryptosporidium parvum Iowa II	YYLLAREVN	FLRGK	TDK	---	-----	ETKVQAV	
Plasmodium falciparum 3D7	YWLLAQEV	VLYLKG	VP	RE	-----	-----	
Plasmodium knowlesi strain H	YWLLAQEV	VLYLKG	VP	RE	-----	-----	
Theileria annulata strain Ankara	YWLLAQEV	VLYLKG	VP	RE	-----	-----	
Theileria parva strain Muguga	YWLLAQEV	VLYLKG	VP	RE	-----	-----	
Leishmania infantum JPCM5	YWLLAQEV	VLYLKG	VP	RE	-----	-----	
Leishmania major strain Friedlin	MWLLTREV	RLRGAL	TRTQEWSVLP	DFFYRDAAD	--EAKIAE	EAEGEETV	SEEGAI
Trypanosoma brucei gambiense DAL972	YWLLAREV	RLKG	SISRT	EEWDVKP	D	DEESED	-----FYD
Trypanosoma cruzi strain CL Brener	YWLLAREV	RLKG	SISRT	EEWDVKP	D	DEESED	-----FYD
Trichomonas vaginalis G3	YWLLAREV	RLKG	SISRT	EEWDVKP	D	DEESED	-----FYD
Giardia lamblia DH_A2	YWLLAREV	RLKG	SISRT	EEWDVKP	D	DEESED	-----FYD
Giardia lamblia WB_A1	YWLLAREV	RLKG	SISRT	EEWDVKP	D	DEESED	-----FYD
Giardia lamblia Pig_E	YWLLAREV	RLKG	SISRT	EEWDVKP	D	DEESED	-----FYD
Giardia lamblia GS_B	YWLLAREV	RLKG	SISRT	EEWDVKP	D	DEESED	-----FYD
Naegleria gruberi strain NEG-M	VAAETTE	EEAEYE	--GEGGQ	TAEWGMDQ	-----		
Cryptosporidium hominis TU502	DSAAAAN	EWGGV	SGNW	GGSAADEWRNAP	-----		
Cryptosporidium parvum Iowa II	DSAAAAN	EWGGV	SGNW	GGSAADEWRNAP	-----		
Plasmodium falciparum 3D7	AAEAPYD	EWE	TKKE	-EWNDNTNEDWK	KNPIA-AEEW	-----	
Plasmodium knowlesi strain H	ATEAPYEE	WNKKE	-EWNDN	NEEW	WK	-----	
Theileria annulata strain Ankara	VPLHPL	PDW	STVD	-PSHANS	-SDWKVMAAGHEEW	GSFVD	TRAQWQ
Theileria parva strain Muguga	VPLHPL	PDW	STVD	-PAHANS	-SDWKVMAAGHEEW	GSFVD	TRAQWQ
Leishmania infantum JPCM5	PAAEAE	EGFGW	VE	---	RNDDNAWEA	-----	
Leishmania major strain Friedlin	PAAEAE	EGFGW	VE	---	RNDDNAWEA	-----	
Trypanosoma brucei gambiense DAL972	VHEEAD	E	EGY	GW	VE	-----	
Trypanosoma cruzi strain CL Brener	PVAEVDE	EGFGW	VE	---	RDNN-AWEQ	-----	
Trichomonas vaginalis G3	NAAPKAEI	KAGAT	NEEW	EN	WKQEWDA	-----	
Giardia lamblia DH_A2	DDEEEDE	EFSAG	---	GNLF	DEY	-----	
Giardia lamblia WB_A1	DDEEEDE	EFSAG	---	GNLF	DEY	-----	
Giardia lamblia Pig_E	DDEEEDE	EFSAG	---	GNLF	DEY	-----	
Giardia lamblia GS_B	DDDEEDE	EFS	STGN	---	GNLF	DE	-----

SSU Ribosomal Protein S15

	1	10	20	30	40	50	60
Naegleria gruberi strain NEG-M						MAMENDELKNLFTSALRRRGK-G	
Cryptosporidium hominis TU502						MADTEQKKRTFRTYSYRGVLDLKLLTMKLDEVVELLPARKRRKIARGC	
Cryptosporidium parvum Iowa II						TSTREKKMADTEQKKRTFRTYSYRGVLDLKLLTMKLDEVVELLPARKRRKIARGC	
Leishmania infantum JPCM5						MASNITAERYEQLKKERTFHFKFTYRGLEIDPLLALSEEFRALVHARARRNMNRH-	
Leishmania major strain Friedlin						MASNITAERYEQLKKERTFHFKFTYRGLEIDPLLALSEEFRALVHARARRNMNRH-	
Trypanosoma brucei gambiense DAL972						MASNITQERYDQLKKERTFHFKFTYRGHEIDPLLALTEEKFALVHARARRNMNRH-	
Trypanosoma cruzi strain CL Brener						MTSNITQERYEQLKKERTFHFKFTYRGLEIDPLLALSEEFKTLVHARARRHMNRH-	
Plasmodium falciparum 3D7						MEDANKPKK-RTFRTFQYRGVLDKLLDLSQDELIKLFKARQRRKFQRG-	
Plasmodium knowlesi strain H						MEDANKPKK-RTFRTFHYRGVELDKLLDLSQEELVKLFRARQRRKFKG-	
Theileria annulata strain Ankara						MDGANRDRRSILKK-RTFRTFKYRGYELEKLLEMPIKLTLLPARQRRFSRG-	
Theileria parva strain Muguga						MVEEQGVQILKK-RTFRTFKYRGYELEKLLEMPIKLTLLPARQRRFSRG-	
Giardia lamblia DH_A2						MADQVKTRRRGLAQPNFRGVEVEKLATMKITDILPKLNARCRRIGKHG	
Giardia lamblia WB_A1						MTDQVKTRRRGLAQPNFRGVEVEKLATMKITDILPKLNARCRRIGKHG	
Giardia lamblia Pig_E						MADQVKTRRRGLVQPNFRGVEVEKLATMKITDILPKLNARCRRIGKHG	
Giardia lamblia GS_B						MDSYISFLEGTMADQTKTRRRGLAQPNFRGVEVEKLVTMKITDILPKLNARCRRIGKHG	
Trichomonas vaginalis G3						MSASGKFAQNYTYRGKTLDDLAMELKDFVELLPTKRRRFISRS-	
Naegleria gruberi strain NEG-M						ISKKYKTLIEKAQKKRD--TAPYGEKPDAVKTHLRDMIILPSMIGCVIGVYQGKDYVSVE	
Cryptosporidium hominis TU502						LNRRTAAFIAKLRKSKA--ECPMGEKPVAVRTHLRNMVILPEMVGSVAGVYNGKTYVTVE	
Cryptosporidium parvum Iowa II						LNRRTAAFIAKLRKSKA--ECPMGEKPVAVRTHLRNMVILPEMVGSVAGVYNGKTYVTVE	
Leishmania infantum JPCM5						VDRRPPVLLKRLREAKK--HVKGEGPKAVKTHLRDVVITPEMVGSVVAIYNGHQFNAVE	
Leishmania major strain Friedlin						ADRRRAPVLLKRLREAKK--HVKGEGPKAVKTHLRDVVITPEMVGSVVAIYNGHQFNAVE	
Trypanosoma brucei gambiense DAL972						ADRRRAPVLLKRLREAKK--NVKAGEKPKAVKTHLRDVVITPEMVGSVVAIYNGHQFNAVE	
Trypanosoma cruzi strain CL Brener						ANRRRAPVLLKRLREAKK--DVKAGEKPRAVKTHLRDVVITPEMVGSVVAIYNGHQFNAVE	
Plasmodium falciparum 3D7						ISKKAKSLLKKIRKSKK--NCEPGEKPNPVPTHLRNMTIIPPEMVGSIVAVHNGKQYTNVE	
Plasmodium knowlesi strain H						ISKKEKSLLKKLRKAKK--ECEVGEKPRAIPTHLRNMTIIPPEMVGSIVAVHNGKQYNNVE	
Theileria annulata strain Ankara						VKRQSLTLLNKLRAAKK--DLPYGQKPEPVKTHLRNMVVIPEMIGSIVGVHNGKQYINVE	
Theileria parva strain Muguga						VKRQSLTLLKKLRAAKK--DLPYGQKPEPVKTHLRNMVVIPEMIGSIVGVHNGKQYINVE	
Giardia lamblia DH_A2						LSMKHLRFVNKLRLRRA--AQPS-QKPKIVRTHLRDMIIFPEMIGMTVSYNGRQFIPVE	
Giardia lamblia WB_A1						LSMKHLRFVNKLRLRRA--AQPS-QKPKIVRTHLRDMIIFPEMIGMTVSYNGRQFIPVE	
Giardia lamblia Pig_E						LSMKHLRFVNKLRLRRT--AQPS-QKPKIVRTHLRDMIIFPEMIGMTVSYNGRQFIPVE	
Giardia lamblia GS_B						LSMKHLRFVNKLRLRRA--AQPS-QKPKIVRTHLRDMIIFPEMVGMTVSYNGRQFIPVE	
Trichomonas vaginalis G3						FDLKHVHLIEKLKKARQAVEGQAGVRPPVKTHLRHMVILPEFVDNVIGIYDGHVFFEVQ	
Naegleria gruberi strain NEG-M						IKPEMVGMLSDFLSITYRPVKHGKPGIGATSSSKFVPLK	
Cryptosporidium hominis TU502						IKPEMIGMYLGEFSITYKPVRHGKPGVGSTSSSRIFIPLK	
Cryptosporidium parvum Iowa II						IKPEMIGMYLGEFSITYKPVRHGKPGVGSTSSSRIFIPLK	
Leishmania infantum JPCM5						IKGEMIGHYLGEFSMSYRPVLHGRPGVGATHSSRFIPIK	
Leishmania major strain Friedlin						IKGEMIGHYLGEFSMSYRPVLHGRPGVGATHSSRFIPIK	
Trypanosoma brucei gambiense DAL972						IKGEMIGHYLGEFSITYKPVAHGRPGFGATHSSRFIPLK	
Trypanosoma cruzi strain CL Brener						IKGEMIGHYLGEFSMTYKPVIIHGRPGVGATHSSRFIPLK	
Plasmodium falciparum 3D7						IKPEMIGYYLGEFSITYKHTRHGKPGIGATHSSRFIPLK	
Plasmodium knowlesi strain H						IKPEMIGYYLGEFSITYKPVRHGKPGIGATHSSRFIPLK	
Theileria annulata strain Ankara						IKPEMIGYYLGEFSITYKPVRHGKPGIGATHSSRFIPLK	
Theileria parva strain Muguga						IKPPMVGRALREYSMSYKIVSHGKVGIGATRSSKFVPLR	
Giardia lamblia DH_A2						IKPPMVGRALREYSMSYKIVSHGKVGIGATRSSKFVPLR	
Giardia lamblia WB_A1						IKPPMVGRALREYSMSYKIVSHGKVGIGATRSSKFVPLR	
Giardia lamblia Pig_E						IKPPMVGRALREYSMSYKIVSHGKVGIGATRSSKFVPLR	
Giardia lamblia GS_B						IKPEMIGHVLADFAPSKKVVKHSKAGVGATRSSSATSLK	
Trichomonas vaginalis G3							

SSU Ribosomal Protein S15P

	1	10	20	30	40	50	60
Naegleria gruberi strain NEG-M							
Cryptosporidium hominis TU502	---MGRMHGAGNGKSQS	VIPYRRSPP	TWLKVTAEEVEDKIVKL	AKGYTPSQIGAILRDS			
Cryptosporidium parvum Iowa II	RGKMG	MGRVYGRGKG	GISSSAKPYRRRPTWIKL	KPSEAEDLITKLARKGLTPSQIGVTLRDS			
Plasmodium falciparum 3D7	---MGRMYGKGKG	GISSSTLPYKRQPSW	LQKPSEIEDAI	IKLAKKGQTPSQIGATLRDN			
Plasmodium knowlesi strain H	---MGRMYGKGKG	GISSSTIPYKRQPSW	LQKPSEIEDAI	IKLAKKGQTPSQIGATLRDN			
Theileria annulata strain Ankara	---MGRMYGKGKG	GISSSSIPYGRKPPS	WLTKPFEVEE	QIAKLAKKGQTPSQIGVSLRDS			
Theileria parva strain Muguga	---MGRMYGKGKG	GISSSSIPYGRKPPS	WLTKPFEVEE	QIAKLAKKGQTPSQIGVSLRDS			
Trichomonas vaginalis G3	---MGRLYHSQGKG	GISSSTCPFRRSAP	HWLQLTAEKCVQM	ICDYAKKGVRSQIGALLRDR			
Giardia lamblia DH_A2	---MGRMHSKGKG	ISRSAIPYTRAPAP	WNKYTPEDVCEQICK	LAKKGTPPSRIGLILRDQ			
Giardia lamblia WB_A1	---MGRMHSKGKG	ISRSAIPYTRAPAP	WNKYTAEDVCEQICK	LARKGTPPSRIGLILRDQ			
Giardia lamblia GS_B	---MGRMHSKGKG	ISRSAIPYTRAPAP	WNKYTPEDVCEQICK	LARKGTPPSRIGLILRDQ			
Giardia lamblia Pig_E	---MVRMHGNRGRK	ASSALPYRRTPAW	LKIASRN	VVKMVCKSSRKGMMP	SQIGMELRDS		
Leishmania infantum JPCM5	---MVRMHGNRGRK	ASSALPYRRTPAW	LKIASRN	VVKMVCKSSRKGMMP	SQIGMELRDS		
Leishmania major strain Friedlin	---MVRMHGNHGK	AAASALPYRRTPTW	LKSSRDVIDAV	CKLA	KGLSPSRIGMQLRDS		
Trypanosoma brucei gambiense DAL972	---MVRMQGVGHGK	SASSLPFCRTPPS	WLKIASRDVIDAV	CKLA	KGLTPSRIGMQLRDS		
Trypanosoma cruzi strain CL Brener	DGIGRVKSVT	GKKILRMLKS	NGLAPEI	PEDLYQLIKKAVGIR	KHLEKTRKDRKAKYQLIL		
Naegleria gruberi strain NEG-M	HGIPMVQSLTG	SKILRILKKNGLAPD	IPE	DLYFLIKRAVSIRK	HLEKFRKDTAAKYRLIL		
Cryptosporidium hominis TU502	HGIPMVQSLTG	SKILRILKKNGLAPD	IPE	DLYFLIKRAVSIRK	HLEKFRKDTAAKYRLIL		
Cryptosporidium parvum Iowa II	HGIPMVQSLTG	SKILRILKKNGLAPD	IPE	DLYFLIKRAVSIRK	HLEKFRKDTAAKYRLIL		
Plasmodium falciparum 3D7	YGIPQVKSV	TGNKILRILRAQGI	ATTI	PEDLYFLIKKAVSMRK	HLEKNNKKDKCKFRLIL		
Plasmodium knowlesi strain H	YGIPQVKAV	TGNKILRILRAQGI	ATTI	PEDLYFLIKKAVSMRK	HLEKNNKKDKCKFRLIL		
Theileria annulata strain Ankara	MAIPQVKAV	TNNKILRILKAQGLA	PEI	PEDLYFLIKKAVSMRK	HMEQNLNNDKSKFRLIL		
Theileria parva strain Muguga	MAIPQVKAV	TNNKILRILKAQGLA	PEI	PEDLYFLIKKAVSMRK	HMEHNLNNDKSKFRLIL		
Trichomonas vaginalis G3	MAIPQVKAV	TNNKILRILKAQGLA	PEI	PEDLYFLIKKAVSMRK	HMEHNLNNDKSKFRLIL		
Giardia lamblia DH_A2	NGVMVR	AVTGVKVLRILKANG	LAPEI	PEDLYFLIKKAVSMRK	HMEQNLNNDKSKFRLIL		
Giardia lamblia WB_A1	FAIQQVK	NVTGSKIVRILKAKG	LAPEI	PEDLYFLIKKAVSMRK	HMEQNLNNDKSKFRLIL		
Giardia lamblia GS_B	FAIQQVK	NVTGSKIVRILKAKG	LAPEI	PEDLYFLIKKAVSMRK	HMEHNLNNDKSKFRLIL		
Giardia lamblia Pig_E	FAIQQVK	NVTGSKIVRILKAKG	LAPEI	PEDLYFLIKKAVSMRK	HMEHNLNNDKSKFRLIL		
Leishmania infantum JPCM5	MGIAQVK	NVTGRKILRILKHSG	LAPEI	PEDLYFLVKRATQMRK	HLERHTTDRDTKYRLIL		
Leishmania major strain Friedlin	MGIAQVK	NVTGRKILRILKHNG	LAPEI	PEDLYFLVKRATQMRK	HLERHTTDRDTKYRLIL		
Trypanosoma brucei gambiense DAL972	MGIAQVK	NVTGRKILRILKHKG	MAPEI	PEDLYCLIKRATQMRK	HLERNTKDKDTKFRLIL		
Trypanosoma cruzi strain CL Brener	MGIAQVK	NVTGRKILRILKHKG	MAPEI	PEDLYCLIKRATQMRK	HLERNTKDKDTKFRLIL		
Naegleria gruberi strain NEG-M	CESKVHRLS	RYYKSTKVL	APNWKYESATAEALINA	--			
Cryptosporidium hominis TU502	VESRIHRL	ARYFKRSKQL	PANWKYQAATASTLVA	--			
Cryptosporidium parvum Iowa II	VESRIHRL	ARYFKRSKQL	PANWKYQAATASTLVA	--			
Plasmodium falciparum 3D7	TESKIHR	ISRYYKRKKLLPSN	WKYQSSTASALIA	--			
Plasmodium knowlesi strain H	TESKIHR	ISRYYKRKKLLPSN	WKYQSSTASALIA	--			
Theileria annulata strain Ankara	VESRIHRL	ARYYKKKRQL	PATWKYQASTAGTLVA	--			
Theileria parva strain Muguga	VESRIHRL	ARYYKKKRQL	PATWKYQASTAGTLVA	--			
Trichomonas vaginalis G3	IESRIHRL	TRYKTRRV	LAPNWKYNAQTASALVA	--			
Giardia lamblia DH_A2	VESRIHRL	ARYRQARRLPPT	FRYKSDQAAALLATYA				
Giardia lamblia WB_A1	VESRIHRL	ARYRQARRLPPT	FRYKSDQAAALLATYA				
Giardia lamblia GS_B	VESRIHRL	ARYRQARRLPPT	FRYKSDQAAALLATYA				
Giardia lamblia Pig_E	VESRIHRL	ARYRQARRLPPT	FRYKSDQAAALLATYA				
Leishmania infantum JPCM5	VESRIHRL	ARYKRVKQLP	PPTWKYESSTASAMVA	--			
Leishmania major strain Friedlin	VESRIHRL	ARYKRVKQLP	PPTWKYESSTASAMVA	--			
Trypanosoma brucei gambiense DAL972	VESRIHRL	ARYRVRVKQLP	PPTWKYESSTASAMVA	--			
Trypanosoma cruzi strain CL Brener	VESRIHRL	ARYKRVKQLP	PPTWKYESSTASAMVA	--			

S-Adenosylmethionine Synthetase

Trios-Phosphate Isomerase

1 10 20 30 40 50 60
| | | | | | |

-----MSN-----
MEGNESISSLMSKLVIGVNHHLNTSMRSNHIFIHWLNYLINQSYPYENWIIYGSSNNVG

-----MRTPYIGGNWKCN-GTVESIKQLVQDLNQGPSTEGKVE
-----MARKYFVAANWKCN-GTLESIKSLTNSFNNLDFDPSKL
-----MTRKYFVSANWKCN-GTQESIKALAASFNELEFDPAKLD
-----MSRKYFVGGNFKCN-GTKESLKTLDTSFKQVESSNS--E
-----MSRKYFVGGNFKCN-GTKESLKTLDTSFKQVESSNS--E
-----RRKWLGGNWKCN-GSKQSISDLLLAHFNNLKSCHKNNLD
LTLQRKLLRCLIKRFIIFYLKYLKIRKWLGGNWKCN-GTKQSISDLLLHFNRHQTKNNNLD
-----MSKPQPIAAANWKCN-GSQQSLSELIDLNFN-STSINHDVQ
-----MASKPQPIAAANWKCN-GSESLLVPLIETLN-AATFDHDVQ
-----MSSKPQPIAAANWKCN-GTTASIEKLVQVLN-EHNISHDVQ
-----MSAKPQPIAAANWKCN-GTTASIEKLVQVLN-EHNISHDVQ
-----MPARRPFIGGNFKCN-GSLDFIKSHVAAIA-AHKIPDSVD
-----MPARRPFIGGNFKCN-GSLDFIKSHVAAIA-AHKIPDSVD
-----MPARRPFIGGNFKCN-GSLDFIKSHVAAIA-AHKIPDSID
-----MPARRPFIGGNFKCN-GSLDFIKSHVASIA-SYKIPESVD
-----MRTFFVGGNWKANPKTVEEAEKLIEMLN-GAKVEGNVE

VIVAPSSIHIGLVQSLNLRK---DWKISAQNIVQEKGNGAFTGEISASMLKDLGIDSTLL
VVVFVPSVHYDHTRKLLQS---KFSTGIQNVSKFGNG-SYTGEVSAEIAKDLNIEYVII
VVVFVPSVHYELAKSLLKP---KFHTGIQNVSKYNG-SYTGEISAEIAKDLNIEYVII
VVVFPTSLHISLVKEFFGNDHPGVFKIGSQNISCTNG-AFTGEVSCEMLKDMDDVDCSLV
VVVFPTSLHISLVKEFFGNDHPGVFKIGSQNISCTNG-AFTGEVSCEMLKDMDDIDCSLV
VVLFPPSLYVEHTRNVLKT---GVFELGVQNVSQSKSG-AFTGELSPLMFSDFGLKWSLV
VVLFPPSLYVEHTRNELKS---EVFELGVQNVSQSKSG-AFTGELSMTDFGLKWSLV
CVVASTFVHLAMTKERLSP---KFVIAAQNAIAKS-G-AFTGEVSLPILKDFGVNWIVL
CVVAPTFLHIPMTKARLTNP---KFQIAAQNAITRS-G-AFTGEVSLQILKDGYIKWVVL
CVVAPTFVHIPLVQAKLRNP---KYAVSAENAIAKS-G-AFTGEVSMPLKDLGINWVIL
CVVAPTFVHIPLVQAKLRNP---KYVVAENAIAKS-G-AFTGEVSMPLKDLGINWVIL
VVIAPSAVHLSTAIAANTS---QLRIAQQNVYLENG-AWTGETSVEMLQDMGLKHVIV
VVIAPSAVHLSTAIAANTS---QLRIAQQNVYLENG-AWTGETSVEMLQDMGLKHVIV
VVVAPSAVHLSTAIAANTS---QLKIAAQNVYLENG-AWTGETSVEMLQDMGLEYVII
VVVAPSFVHLSTAIAANTS---CLKIAAQNVYLENG-AWTGETSVEMLLDMGLSHVII
VVVAAPFIFLPTLQQK-LRK---DWKVSAENVFTKPNG-AFTGEVTVPMIKSFGEWTL

GHSERRHVFGESNTVIAKKVKVSEEVGLTIIGCVGETLEERERGETNQVITSQLKDFN---
GHFERRKYFHETDEDVREKLQASLKNNLKAVVCFGESLEQREQNKTIETVKQVKAFVDSL
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----VTNWDFKVIAYEPVWAIGTGKTAASPQQAEAHKVLRDWLADNVNSEVAQKVRIIYGD
ID----NFDNVILAYEPLWAIGTGKTATPEQAQLVHKEIRKIVKDTGEGKQANQIRILYGD
IN----NFDNVILAYEPLWAIGTGKTATPEQAQLVHKEIRNIVKEKCQGEKNAQQIRILYGD
VS----DLSNLVIAYEPIWAIGTVVATPGQQAQEAHAFIREYVTRMNPQVSSNLRIIYGD
VS----DLSNLVIAYEPIWAIGTVVATPEQAQEAHAFIREYVTKMYNSQVSSNLRIIYGD
VK----DWDKVVLAYEPMWAIGTGKVATVEQVNQAHKFVREHVRGLAG-AVADKVRVYVG
VK----DWDKVVLAYEPMWAIGTGKVATVDQVKEAHKFVRDYVRGLVG-DVADKVRVYVG
LK--KADWAKVVIAYEPVWAIGTGKVATPQQAEHALIRSWSVSSKIGADVAGELRILYGD
LS--KEAWAHVVIAYEPVWAIGTGKVATPQQAEVHELLRRWRSKLGADIAQLRILYGD
LP--KEAWDQIVLAYEPMWAIGTGKVATPEQAQEVHALLRKWVSEKIGADVAARLRILYGD
LP--KEAWDQIVLAYEPMWAIGTGKVATPEQAQEVHALLRKWVSEKIGTDVAAKLRILYGD
LGESKMLWKEVVIAYEPVWSIGTVVATPEQAEEHVGLRKWFAEKVCAEGAQHIRIIYGD
LGESKMLWKEVVIAYEPVWSIGTVVATPEQAEEHVGLRKWFAEKVCAEGAQHIRIIYGD
LGESKLLWKKVVIAYEPVWSIGTVVATPEQAEEHVGLRKWFAEKVCAEGAQHIRIIYGD
IGESKKLWENVVIAYEPVWSIGTVVATPEQAEEHVGLRKWFAEKVCAEGAQHIRIIYGD
IP--AGKWDDVVIAYEPIWAIGTGKAVASTQDAQEMCKVIRDILAAKVGADIANKVRILYGD

GSVNSANSDELIKQPDIDGFLVGGASLKSDFLSIVNSGLKSVE
GSVNTENCSSLIQQEDIDGFLVGNASLKE-SFVDIICKSAM----
GSVNTENCASLIKQEDIDGFLVGNASLKP-SFVEIIKSAM----
GSVTPDNCNELIKCADIDGFLVGGASLKP-TFAKIIESAQ----
GSVTPDNCNELIKCADIDGFLVGGASLKP-TFAKIIESAQ----
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GSANGSNCEKLQCPNIDGFLVGGASLKP-EFMTMIDLAKTRT
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GSANGSNCEKLQCPNIDGFLVGGASLKP-EFTTMIDLAKTRA

Alpha Tubulin