		Sh F	
Estrame Aano XP 009033281.1	 . <u>M</u>	NM F	DRFK
Ealveol Vbra CEL99851.1	 .M	N <mark>FI</mark>	D <mark>R</mark> AF
Eviridi Vcar XP 002949072.1	 . <mark>M</mark>	N <mark>L</mark> F	S <mark>RV</mark> F8
Bspiroc Tcal AEJ18398.1	 . M	G <mark>IF</mark>	S <mark>RL</mark> K8
Bgemmat Gpho AMW04363.1	 . M	G <mark>IF</mark>	D <mark>RL</mark> S 8
Bthermo Tter ACZ42165.1			
Bacidob Gmal AEU35097.1	 . <mark>M</mark>	S <mark>L</mark> M	E <mark>RV</mark> A8
Bdeinoc Tthe BAD69831.1			
Bchloro Caur ABY35517.1			
Bproteo Daut ACN14859.1			
Bchloro Ctha ACF15037.1			
Bbacter Hsp AIZ65498.1			
Baquifi Aaeo AACO7332.1			
Bactino Cacn AFU40489.1			
Athauma CNit AFU59479.1	 . <u>M</u>	GL M	N <mark>RF</mark> S 8
Aeuryar Nmag ADD05844.1	 · M · · · · · · · · · · · · · · · · · ·	G I L	SRTS8
Bproteo Afer ACK79767.1	 	SI F	K H A A
Bthermo Mpri AFK05994.1			
Bfusoba Lbuc ACV38698.1			
Emetazo Hsap NP 001231573.1	 		
Emetazo Bmor XP 004922821.1			
Eviridi Atha NP 001332586.1			
Bnitros Lfer BAM07649.1			
Bproteo Alip CBS85649.1 Bproteo Shal A0066257.1			
Bproteo Bcep ALK18953.1	 		
Bplanct Psp AMV20944.1			
Bproteo Maes ANG64508.1	 M	ς Τ	KRI Q Q
Barmati Fgin AIE88136.1			
Aasgard CLok OLS16333.1			
Bcyanob Hsp AFZ44603.1			
Bfirmic Gthe AMV09772.1			
Eamoebo Ddis XP 637438.1	 M	GILLI	GREK
Eamoebo Einv XP 004256843.1			RQ
Eeuglen Tbru XP 011780226.1			
Eparaba Tvag XP 001308534.1	 . <u>M</u>	W <u>L</u> W	G <mark>K</mark> KK
NAarchae a13 OLB69573.1	 . <u>M</u>	S <mark>I</mark> T	D <mark>RL</mark> R 8

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	<u>Pelvi</u>						
Estrame Aano XP 009033281.1		<mark>QK</mark> DLV.TI	<mark>R</mark> Q.SY.A	E VM A	TQKR.	MQRQKDQADQLADEWYKR	49
Ealveol Vbra CEL99851.1		QSDLG . RI	<mark>K</mark> Q.SY.A	<u>E</u> VT	TQRR.	GERQREALLEQVQEWERR	49
Eviridi Vcar XP 002949072.1						MAKQKSLQGSADQWLSR	49
Bspiroc Tcal AEJ18398.1						LERETLNQQAQAQEWERK	49
Bgemmat Gpho AMW04363.1		<mark>RN</mark> QLA . <mark>K</mark> A	<mark>K</mark> Q . Q <mark>V</mark> . A	AAI <mark>A</mark>	DEKR.	LKDQADAEFKLADDWEKR	49
Bthermo Tter ACZ42165.1		ASA <mark>I</mark> A . <mark>E</mark> A	<mark>R</mark> A . Q <mark>V</mark> . A	<u>E</u> MI	QEKI.	IEANLERNRSLANEWRQK	49
Bacidob Gmal AEU35097.1	L <mark>DM</mark>	<mark>EN</mark> Q <mark>L</mark> L . <mark>Q</mark> L	<mark>K</mark> T . Q <mark>V</mark> . A	<u>I</u> AI	DKHL.	LEKKSKEQSQTAEDWRLK	49
Bdeinoc Tthe BAD69831.1		<mark>KQ</mark> A <mark>L</mark> R . <mark>E</mark> A	<mark>R</mark> E . Q <mark>V</mark> . A	<u>E</u> AL	EQKR.	LEREVESHEREAALWEEK	49
Bchloro Caur ABY35517.1			<mark>R</mark> T . G <mark>V</mark> . A	Q A M A	DETR.	LEQRRIAAEGEAEQWQIK	49
Bproteo Daut ACN14859.1		<mark>KK</mark> D <mark>L</mark> T . AA	MT.S <mark>L</mark> .A	E V K G	VAIR.	TRKEADKKKNMAQDYERK	49
Bchloro Ctha ACF15037.1		<mark>KN</mark> D <mark>L</mark> S . <mark>Q</mark> A	MQ . G <mark>L</mark> . A	Q V K A	SAIR.	KKDADDQNSLAADYERK	49
Bbacter Hsp AIZ65498.1		<mark>RQ</mark> D <mark>L</mark> D . <mark>K</mark> A	LH . A <mark>L</mark> . A	<mark>E</mark> V K A	LTIR.	SRNDAETERSRALDYENK	49
Baquifi Aaeo AACO7332.1						FEREARNEKERAKALVKK	49
Bactino Cacn AFU40489.1	Q <mark>R</mark> Q	L <mark>D</mark> M <mark>L</mark> Q . <mark>K</mark> V	<mark>R</mark> R . G <mark>V</mark> . A	D V A T	SRKR.	VELQMNQLQTDVDKMQNA	49
Athauma CNit AFU59479.1						LEMQKSKLWDNIRLLDEQ	49
Aeuryar Nmag ADD05844.1	E Q M	<u>RDQLQ . <mark>E</mark>V</u> .	<mark>K</mark> R . G <mark>I</mark> . A	DLT <mark>T</mark>	QKKR.	LEMQKRRLEENVEKHNEQ	49
Bproteo Afer ACK79767.1	E <mark>K</mark> M	IANLQ . DT	KR . H A	<u>D</u> VV <u>T</u>	ERIS.	LENQMAQAQKAADKADAD	49
Bthermo Mpri AFK05994.1		SEDLA . EV	KR . GT . A	GVI A	EETR.	TKRLVDENAAEVAKYADF	49
Bfusoba Lbuc ACV38698.1		ERDLN.SV	KS.ET.A	A V M A	QESA.	AKRKVIECEDEIKKMDTY	49
Emetazo Hsap NP 001231573.1					EL	E	4
Emetazo Bmor XP 004922821.1	TP	DEMLR . KN		KAM R	RDLD	.R.ERMKMEQUEKKVIAD	45
Eviridi Atha NP 001332586.1						IEREITSLQLEEKRLVAE	44
Bnitros Lfer BAM07649.1		DDKLQ . KA		K A V A	EIKL.	LEQKIRDDESQVALYQER	49
Bproteo Alip CBS85649.1						IERRVADLHRERDEWDRK	49 49
Bproteo Shal A0066257.1		DDGEG DA	KL.EL.G	<u>K</u> DVV	KUKL.	IKERIDSYNAEHNKLLAQ QRSKRDAADDKVKKYEDG	49 49
Bproteo Bcep ALK18953.1						TELEIGEQTQQVQQWLGR	49 49
Bplanct Psp AMV20944.1 Bproteo Maes ANG64508.1	DEM	DUZWA OA	Nr. SS . K	I A A	INTOK.	LQSRIAALHEDEQLWAER	49 49
	DDM	OCATA CN	RA . R N		LEEAU.	LQMMLDDERKKAAQLESQ	49 46
Barmati Fgin AIE88136.1 Aasgard CLok OLS16333.1			VE VI A	UAL I	UNN	IELKSRQFRQKANVMRVN	41
Bcyanob Hsp AFZ44603.1	EM	OMCWE OI	TO AV T	ELLH	INAU U	AEERYSAKVQEFKDLETK	41
Bfirmic Gthe AMV09772.1	DOC		ז ע . איי . ז	VQVV	INIQQ.	FAREHREAE QMVEKRSKQ	
Eamoebo Ddis XP 637438.1		EDKIG KT		NTV T	LUBU	K.LKNYQTQVLEIAI	49
Eamoebo Einv XP 004256843.1		EDRED . KI	KK AV A	KTU E	RIK F	LKLKIDHDTQIISEELKL	33
Eeuglen Tbru XP 011780226.1			NS SL F	FI.O S	KIE I	LEKREGVLTKKVEDELAK	33
Eparaba Tvag XP 001308534.1		KEPTV NΔ	OT AG F	KNV E	TTE M	MNKKIEQLDRQAENMAKE	46
NAarchae a13 OLB69573.1	FΔΠ	KPTPV KR	BT OF A	KI.R. T	OTN R	LEKTINQMEHRDRILHDK	49
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Estrame Aano XP 009033281.1		.AQL	AL	E K G .	DDELAI	R.EA	. S. RKTQQTDASAGDEQIAVQTDS 89
Ealveol Vbra CEL99851.1		.AQL	<mark>A</mark> V	SK <mark>G</mark> .	RDDMAI	RE <mark>A</mark>	.LQ.RRVVLTTKVQGLDEQLRIQKEA 89
Eviridi Vcar XP 002949072.1							.LK.RRKVLQEDADRMNEQVALQTNA 89
Bspiroc Tcal AEJ18398.1		.AML	<mark>A</mark> V	RAG.	QDDLAI	KE <mark>A</mark>	.LL. <mark>rk</mark> qeye <mark>nn</mark> yveykkqweaq <mark>k</mark> as 89
Bgemmat Gpho AMW04363.1		.AML	<mark>A</mark> V	<mark>Q</mark> E <mark>G</mark> .	RDDLAI	KQ <mark>A</mark>	LM.RGQEHLEHGQQLAATWEAHKME 89
Bthermo Tter ACZ42165.1		.AEL	AV	QKG.	RDDLAI	REA	.LR.RYNDYL <mark>EN</mark> ASVYEQQLDAQ R KI 89
Bacidob Gmal AEU35097.1		.AEL	<mark>A</mark> V	<u>T</u> K <mark>G</mark> .	KDDLAI	RA <mark>A</mark>	.LD.RALTHEKMAGDLLHQVEEQTEE 89
Bdeinoc Tthe BAD69831.1							. <mark>L</mark> R. <mark>RK</mark> KRAL <mark>D</mark> LAEG <mark>F</mark> KAQLSEH <mark>K</mark> AL 89
Bchloro Caur ABY35517.1		.AER	<mark>A</mark> L	<mark>R</mark> A <mark>G</mark> .	DEALAI	KA <mark>A</mark>	. <mark>L</mark> A. <mark>RK</mark> VQAQ <mark>K</mark> LAEQŸRAQEKAQ <mark>E</mark> EQ 89
Bproteo Daut ACN14859.1		.AML	L <mark>L</mark>	<mark>Q</mark> KM .	QERLA	ΓΕ <mark>Α</mark>	.LN.LKEQSLA <mark>E</mark> SVR <mark>L</mark> GTEATAH <mark>E</mark> QR 89
Bchloro Ctha ACF15037.1		. <mark>A</mark> ML	L <mark>L</mark>	<mark>Q</mark> K <mark>G</mark> .	EERLA	ΓΕ <mark>Α</mark>	.LS.RKEEASQRSKTLMSEHQGQQEM 89
Bbacter Hsp AIZ65498.1		. <mark>A</mark> VL	L <mark>L</mark>	<mark>Q</mark> K <mark>A</mark> .	HDRLA	ΓΕ <mark>Α</mark>	.LL.KKNQHETQSSRAAQDQEKFETS 89
Baquifi Aaeo AACO7332.1		.AES	L <mark>L</mark>	<mark>Q</mark> E <mark>G</mark> .	K EKLA	<u>A</u> EI	.IQ. <mark>K</mark> AELHE <mark>KN</mark> ARR <mark>L</mark> EÈEAKRQ <mark>R</mark> EM 89
Bactino Cacn AFU40489.1							.LT.RKATLQAQLND <mark>L</mark> QTQHATL <mark>Q</mark> AE 89
Athauma CNit AFU59479.1		. <mark>A</mark> SR	<mark>A</mark> L	<mark>D</mark> A <mark>G</mark> .	REDLAI	RL <mark>A</mark>	.LE.RKNTNLLQIQDLDKQIAELQEE 89
Aeuryar Nmag ADD05844.1		.ART	<mark>A</mark> V	<mark>Q</mark> QD .	REDLAI	RR <mark>A</mark>	.LE. <mark>KK</mark> KTKM <mark>NQ</mark> IED <mark>L</mark> ERQISEL <mark>Q</mark> NQ 89
Bproteo Afer ACK79767.1							.LA. <mark>QK</mark> QSEL <mark>QK</mark> IAS <mark>L</mark> KEAHDTV <mark>A</mark> AQ 89
Bthermo Mpri AFK05994.1		.AKR	AL	<u>L</u> AK .	N DGDAI	KVF	. <mark>I</mark> A. <mark>KK</mark> QELE <mark>N</mark> VGAGLMTAYAAA <mark>H</mark> EN 89
Bfusoba Lbuc ACV38698.1		.ARK	<u>A</u> L	KS <mark>G</mark> .	N ESDAI	KLF	.LE.KKATLQTKLESLNNDKEVAAQN 89
Emetazo Hsap NP 001231573.1		.IKK	MA	KI <mark>G</mark> .	N KEACI	KVI	.AK.QLVHLR <mark>KQ</mark> KTRTFAVSSKVTSM 44
Emetazo Bmor XP 004922821.1		.IKK	LA	KE <mark>G</mark> .	Q MD A VI	KIM	.AK. <mark>D</mark> LVRTR <mark>RY</mark> VRK <mark>E</mark> MLMKANI Q AV 85
Eviridi Atha NP 001332586.1							.AR. <mark>Q</mark> LVRLR <mark>QQ</mark> ITNLQGSRAQI <mark>R</mark> GV 84
Bnitros Lfer BAM07649.1							.LV.EKSRLSADLSEMTRQRDEQQKV 89
Bproteo Alip CBS85649.1		. AEV	AL	TRD .	REDLAI	KGA	.LM.AKSKVAEQADALAEQLVQVEAA 89
Bproteo Shal A0066257.1		. VKI	AL	KED .	R EDT	E A G	.LA.EQISIEEKIKANQVQMDALQKE 89
Bproteo Bcep ALK18953.1		AKR	AL	A G .	DEALAI	REA	.LG.AQSNAEAERDALAAELSTLEPS 89
Bplanct Psp AMV20944.1		ARE	AL	AAE.	DEENAL	RQA	LL.RKREHESLLAGLEQQLGAARST 89
Bproteo Maes ANG64508.1							.IE.RRQQCRSQAEHLAQGLISYRQS 88
Barmati Fgin AIE88136.1		. ATM	AL	<u>K</u> Q <mark>G</mark> .	DRELA	IQF	.MR.EKMNHDATIATLQQSYEQAVAT 86
Aasgard CLok OLS16333.1		. AKK	AL	<u>K</u> V <mark>G</mark> .	DKEGAI	RNL	LV. RWKKYKTNLDRYQNIVGRIERH 81
Bcyanob Hsp AFZ44603.1		. AKT	. <u>L</u> K	A Q G .	N TEAAI	К <u>.</u> Г	.MI.QAIQLEKILPQLKENVENAEKF 80
Bfirmic Gthe AMV09772.1		. ALV	AA	A G .	EMELAI	L F A	.RR.EQAQYAERAARLKQLLEQASRE 89
Eamoebo Ddis XP 637438.1		AKE	<u>C</u> S	KAG.	KKNUAI	LL <mark>A</mark>	.LK.KKKYQEKMLDESFANLQNIEEL 81
Eamoebo Einv XP 004256843.1							.LK.KKKMYEKAVSETQNFIINLESQ 73
Eeuglen Tbru XP 011780226.1		AKE	rï	SKK.	N KSMAI	L W	.MK.RKKMAEDELTKTACQKQNLDTL 73 .MK.RKKMYENQSTKESQMRDNLEAQ 86
Eparaba Tvag XP 001308534.1 NAarchae a13 0LB69573.1	• • • • • • • • • • • • • • • • • • • •	ANA	AIM	AKK.	TOGAT	LĽ <mark>(</mark> r tr	.MK.RKMYENQSIKESQMRDNLEAQ 86 .AN.ECVQIRKLIKTSLSSQISLEQV 89
MAAICHAE AIS ULBO95/3.1	• • • • • • • • • • • • • • • • • • • •	v v	<mark>н</mark> г	<u>=</u> v R .	<u>u</u>	1Г	.An. <mark>E</mark> CVQIR <mark>K</mark> LIKTSLSSQISL <mark>E</mark> QV 89

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Estrame Aano XP 009033281.1	LAKLYDSMT
Ealveol Vbra CEL99851.1	LLQLRSSMR
Eviridi Vcar XP 002949072.1	LEQLQANIRV 9
Bspiroc Tcal AEJ18398.1	VE <mark>KLK</mark> ESLK
Bgemmat Gpho AMW04363.1	TE <mark>KLK</mark> QSLRD
Bthermo Tter ACZ42165.1	VD <mark>K</mark> L <mark>K</mark> NELQ
Bacidob Gmal AEU35097.1	VE <mark>G</mark> M <mark>R</mark> MTYN
Bdeinoc Tthe BAD69831.1	TE <mark>R</mark> LMTQLK
Bchloro Caur ABY35517.1	VEA <mark>LQ</mark> QALI
Bproteo Daut ACN14859.1	AAG <mark>LQ</mark> TQVS <mark>K</mark> 9
Bchloro Ctha ACF15037.1	AD <mark>QLQ</mark> AKVQ <mark>K</mark> 9
Bbacter Hsp AIZ65498.1	AQ <mark>QL</mark> SQNVQ <mark>Q</mark> 9
Baquifi Aaeo AAC07332.1	AV <mark>KLQ</mark> AKID <mark>E</mark> 9
Bactino Cacn AFU40489.1	EE <mark>KL</mark> Ī
Athauma CNit AFU59479.1	AR <mark>KME</mark> EAEK <mark>R</mark> 9
Aeuryar Nmag ADD05844.1	QD <mark>QL</mark> TDQKD <mark>E</mark> 9
Bproteo Afer ACK79767.1	ÂQ <mark>KL</mark> TDYES
Bthermo Mpri AFK05994.1	AV <mark>KMR</mark> QMHD <mark>K</mark> 9
Bfusoba Lbuc ACV38698.1	AL <mark>KMR</mark>
Emetazo Hsap NP 001231573.1	STQTKVMNS
Emetazo Bmor XP 004922821.1	SL <mark>KTQ</mark> TLKS
Eviridi Atha NP 001332586.1	TTHTQALYA
Bnitros Lfer BAM07649.1	VSETE
Bproteo Alip CBS85649.1	LS <mark>KAN</mark> EDIGR9
Bproteo Shal A0066257.1	ITTYO EYLD. K. 9
Bproteo Bcep ALK18953.1	VDKIK
Bplanct Psp AMV20944.1	RDHMT CTLH A 9
Bproteo Maes ANG64508.1	AARLS
Barmati Fgin AIE88136.1	VDQVK TAIK 9
Aasgard CLok OLS16333.1	LEALENVMGK9
Bcyanob Hsp AFZ44603.1	VQAAKEQLT
Bfirmic Gthe AMV09772.1	LFETEAKYEE9
Eamoebo Ddis XP 637438.1	IANVEQAEI
Eamoebo Einv XP 004256843.1	ILETEN
Eeuglen Tbru XP 011780226.1	IFTIONQTMN.
Eparaba Tvag XP 001308534.1	VITIO
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Estrame Aano XP 009033281.1	DE SKUSEAKSMK				TKVN.DML	131
Ealveol Vbra CEL99851.1		EQ∏IA	RARTA	<mark>KT</mark> . A	QQIQ.DML	131
Eviridi Vcar XP 002949072.1	LEGKMTEARGKK	ETLK A	RAAS <mark>A</mark>	<mark>KT</mark> . S	LQ.EMI	131
Bspiroc Tcal AEJ18398.1	LQNKIEEAQRKK	NLLIA	RAKR <mark>A</mark>	<mark>EA</mark> . Q	QKIQ.NTI	131
Bgemmat Gpho AMW04363.1	LNDKIEEAKRKK	NLLLA	RQRR <mark>A</mark>	<mark>EA</mark> . Q̀	IS.QTM	131
Bthermo Tter ACZ42165.1	<mark>LE</mark> S <mark>K</mark> YEDA <mark>R</mark> RNR	EVLIA	RHKR <mark>A</mark>	<mark>ET</mark> . Q		131
Bacidob Gmal AEU35097.1	<mark>LQ</mark> G <mark>KL</mark> KETQAHC	VLLIA	QHRR <mark>A</mark>	<mark>Q</mark> M . A	KA <mark>I</mark> T.ARE	131
Bdeinoc Tthe BAD69831.1	LEAKIDEAEARK	KLLLA	RKKGV	<mark>E A</mark> . A	ETVR.RME	131
Bchloro Caur ABY35517.1	<mark>LE</mark> T <mark>RI</mark> AEV <mark>K</mark> ARK	ELIIA	KKNR <mark>A</mark>	<mark>QT</mark> . Q	LQ.RTA	131
Bproteo Daut ACN14859.1					IN.AQL	
Bchloro Ctha ACF15037.1	LKQTITKYENEL				IN.KKIN.KQL	
Bbacter Hsp AIZ65498.1	<mark>LK</mark> QT <mark>I</mark> SQW <mark>D</mark> N <mark>E</mark> L	KTLKA	<mark>(R</mark> VTV <mark>S</mark>	<mark>T </mark>	KT <mark>I</mark> N. <mark>Q</mark> QL	131
Baquifi Aaeo AACO7332.1					KK <mark>V</mark> N. <mark>K</mark> QI	
Bactino Cacn AFU40489.1	<mark>LQ</mark> A <mark>RV</mark> DAF <mark>R</mark> TKK	ETIKA	<mark>I</mark> TYSA <mark>A</mark>	<mark>E A</mark> . Q	IRIN.EAF	131
Athauma CNit AFU59479.1	MAVKVEEFKSKK	EVIKA	<mark>(R</mark> YSA <mark>A</mark>	<mark>E A</mark> . Q	IK. <mark>E</mark> SV	131
Aeuryar Nmag ADD05844.1	LQNRIEEFRTKK	ETMKA	RYEA <mark>A</mark>	<mark>EA</mark> . <u>S</u>	VS. <mark>E</mark> AM	131
Bproteo Afer ACK79767.1	LQDHIEQFRTQK	EVTKS	EMAA <mark>A</mark>	<mark>QA</mark> . <u>E</u>	VKVs. <mark>v</mark> s. <mark>e</mark> sl	131
Bthermo Mpri AFK05994.1	LVSDIESLKNRR	A M I K A	KVAV <mark>A</mark>	<mark>KT</mark> . Q	IN.EVS	131
Bfusoba Lbuc ACV38698.1	A SDIQKLSAKR	SEIKA	KMKMA	<mark>KT</mark> . S	ERIS.SLS	131
Emetazo Hsap NP 001231573.1	MK MA	G A	MSTT <mark>A</mark>	<mark>KT</mark> . M	QAVN.KKM	75
Emetazo Bmor XP 004922821.1	STMA	Q	MKGVT	<mark>R.A</mark> . M	ATMN.RQL	116
Eviridi Atha NP 001332586.1	TS <u>I</u> S	S	MKGAT	<mark>K A</mark> . M		115
Bnitros Lfer BAM07649.1	LRDMHDEFARKQ	SMLSL	REER <mark>A</mark>	<mark></mark> . S	QEVN.SIR	131
Bproteo Alip CBS85649.1	AKLTDAKTRE	KALV A	RTQT <mark>A</mark>	SN.R		131
Bproteo Shal A0066257.1	SAKKDMDSEL	KALS S	KPNA		ID.SLA	125
Bproteo Bcep ALK18953.1	MRQRKNDLNARS	N I 🖪 Q A	K.M. T. F.T.W.	<mark>Q A</mark> . K	DVĀA.SAL	131
Bplanct Psp AMV20944.1	LEARLADALRRK		DO GOT		ADLE TA.DSE	115
Bproteo Maes ANG64508.1	SEIKLHSLNNKH	SIMKA	KUSGI	A <mark>A</mark> . K		130 128
Barmati Fgin AIE88136.1	QE EEVKKKIAEA	LAMKA	ME TOE	<mark>d</mark> T . M		128 123
Aasgard CLok OLS16333.1	SAKILEKISINI	SPEKA	MEISE	G <mark>A</mark> . E		123 122
Bcyanob Hsp AFZ44603.1 Bfirmic Gthe AMV09772.1	ME HELLINGEL	ANMQ A	T L M V V			122
Eamoebo Ddis XP 637438.1	MADE ESTROCH	EGI V	rncNA <mark>A</mark>	<mark>п.а</mark> . п		131
Eamoebo Einv XP 004256843.1	LR		V		KKVS.DSM	90
Eeuglen Tbru XP 004256645.1						
Eparaba Tvag XP 001308534.1					DKID.EIR	
NAarchae a13 OLB69573.1						
marchae are ordered.						110

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Estrame Aano XP 009033281.1		V G		AF	ER M 144
Ealveol Vbra CEL99851.1		LG	TS <mark>L</mark> Q	(A <mark>F</mark>	ER M 144
Eviridi Vcar XP 002949072.1		LR	TSWA	.A <mark>F</mark>	EK 144
Bspiroc Tcal AEJ18398.1					
Bgemmat Gpho AMW04363.1					
Bthermo Tter ACZ42165.1	AQ				
Bacidob Gmal AEU35097.1	EQ				
Bdeinoc Tthe BAD69831.1					
Bchloro Caur ABY35517.1		I G	SA <mark>L</mark> D	K <mark>L</mark>	
Bproteo Daut ACN14859.1	AN				
Bchloro Ctha ACF15037.1	ST				
Bbacter Hsp AIZ65498.1		<u> </u>	GT <mark>V</mark> T	<u> </u>	ER M 144
Baquifi Aaeo AACO7332.1		<u>VD</u>	DT <u>I</u> A	.ML	ER M 144
Bactino Cacn AFU40489.1		<u> </u>	D G M	[A]	TR A 144
Athauma CNit AFU59479.1		<u> </u>	D <mark>V</mark> G M	AM	NR
Aeuryar Nmag ADD05844.1	EE				
Bproteo Afer ACK79767.1		<u>M</u> N	DAG D	AM	RR A 144
Bthermo Mpri AFK05994.1					
Bfusoba Lbuc ACV38698.1					
Emetazo Hsap NP 001231573.1 Emetazo Bmor XP 004922821.1					
Ewiridi Atha NP 001332586.1		<u> </u>	Kullari	MT	EK QS 130
Bnitros Lfer BAM07649.1	AQ				
Bproteo Alip CBS85649.1 Bproteo Shal A0066257.1					
Bproteo Bcep ALK18953.1		TC	ANACE Na dem		OKL 144
Bplanct Psp AMV20944.1					
Bproteo Maes ANG64508.1			 D <mark>∏</mark> F T	' q	ER W 143
Barmati Fgin AIE88136.1					
Aasgard CLok OLS16333.1					
Bcyanob Hsp AFZ44603.1					
Bfirmic Gthe AMV09772.1		A G	PAT A	AF	ADA 144
Eamoebo Ddis XP 637438.1	ED				
Eamoebo Einv XP 004256843.1	KD	LN		RL	
Eeuglen Tbru XP 011780226.1		MD	E <mark>L</mark> L E	EM	
Eparaba Tvag XP 001308534.1	EE	NE	E <mark>V</mark> TD	Q <u>L</u>	
NAarchae a13 OLB69573.1			.	. .	

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Bgemmat Gpho AMW04363.1			N			156
Bthermo Tter ACZ42165.1			E			156
Bacidob Gmal AEU35097.1			Ā			156
Bdeinoc Tthe BAD69831.1			M			156
Bchloro Caur ABY35517.1			R			156
Bproteo Daut ACN14859.1			D			156
Bchloro Ctha ACF15037.1			E			156
Bbacter Hsp AIZ65498.1			E			156
Baguifi Aaeo AACO7332.1			E			156
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Emetazo Bmor XP 004922821.1						136
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Eamoebo Einv XP 004256843.1			$\texttt{M}\dots$			111
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Bbacter Hsp AIZ654						
Baquifi Aaeo AACO7						
Bactino Cacn AFU40						
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Bplanct Psp AMV209						
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Aasgard CLok OLS16 Bcyanob Hsp AFZ446						
Bfirmic Gthe AMV09						
Eamoebo Ddis XP 63						
Eamoebo Einv XP 00						
Eeuglen Tbru XP 01						
Eparaba Tvag XP 00						
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Bchloro Caur ABY35517.1	S <mark>ie</mark> A	
Bproteo Daut ACN14859.1	· · · · · · · · · · · · · · · · · · ·	.E 168
Bchloro Ctha ACF15037.1		.Q 168
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Bnitros Lfer BAM07649.1	D <mark>LD</mark> T	
Bproteo Alip CBS85649.1	S <mark>I</mark> TE	
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Bproteo Bcep ALK18953.1	A <mark>ID</mark> D	. <mark>K</mark> 168
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Bspiroc Tcal AEJ18398.1	<mark>F</mark> AA	71
Bgemmat Gpho AMW04363.1	<mark>F</mark> KQ	
Bthermo Tter ACZ42165.1	=	71
Bacidob Gmal AEU35097.1	<mark>F</mark> AK	
Bdeinoc Tthe BAD69831.1	<mark>F</mark> QA	
Bchloro Caur ABY35517.1	<mark>F</mark> RD	. –
Bproteo Daut ACN14859.1	<mark>I</mark> DN	71
Bchloro Ctha ACF15037.1	<mark>I</mark> NK	71
Bbacter Hsp AIZ65498.1	<mark>I</mark> DK	. –
Baquifi Aaeo AACO7332.1	<u>I</u> KE 17	. –
Bactino Cacn AFU40489.1	<mark>L</mark> DS	. –
Athauma CNit AFU59479.1	<u>L</u> EK 17	
Aeuryar Nmag ADD05844.1	<u>L</u> ED 17	
Bproteo Afer ACK79767.1	<u>I</u> DE 17	
Bthermo Mpri AFK05994.1	<u>A</u> KA	
Bfusoba Lbuc ACV38698.1	<mark>M</mark> KK	
Emetazo Hsap NP 001231573.1		05
Emetazo Bmor XP 004922821.1		50
Eviridi Atha NP 001332586.1		33
Bnitros Lfer BAM07649.1		. –
Bproteo Alip CBS85649.1	17 AE 17	. –
Bproteo Shal A0066257.1		53
Bproteo Bcep ALK18953.1	<mark>L</mark> AA	
Bplanct Psp AMV20944.1		25
Bproteo Maes ANG64508.1		69
Barmati Fgin AIE88136.1		68
Aasgard CLok OLS16333.1		56
Bcyanob Hsp AFZ44603.1		62
Bfirmic Gthe AMV09772.1		59
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Bgemmat Gpho AMW04363.1	LER		
Bthermo Tter ACZ42165.1	I.E.G.		E 179
Bacidob Gmal AEU35097.1	I.E.R.	D . E K V	E
Bdeinoc Tthe BAD69831.1		VE.KEV	
Bchloro Caur ABY35517.1		E . T E	
Bproteo Daut ACN14859.1			
Bchloro Ctha ACF15037.1			S
Bbacter Hsp AIZ65498.1	ALH		
Baguifi Aaeo AAC07332.1			
Bactino Cacn AFU40489.1		D . H Å	
Athauma CNit AFU59479.1	<mark>I</mark> TV		
Aeuryar Nmag ADD05844.1			
Bproteo Afer ACK79767.1			
Bthermo Mpri AFK05994.1			
Bfusoba Lbuc ACV38698.1			
Emetazo Hsap NP 001231573.1	SDD		
Emetazo Bmor XP 004922821.1			
Eviridi Atha NP 001332586.1			
Bnitros Lfer BAM07649.1			
Bproteo Alip CBS85649.1	LEA	to the control of the	The state of the s
Bproteo Shal A0066257.1			
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Bproteo Maes ANG64508.1		D . A R L	
Barmati Fgin AIE88136.1			<mark>E</mark>
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Bcyanob Hsp AFZ44603.1			
Bfirmic Gthe AMV09772.1		<mark>I</mark>	D
Eamoebo Ddis XP 637438.1	<mark>E</mark> E		
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Ealveol Eviridi	Aano XP 009033281.1 Vbra CEL99851.1 Vcar XP 002949072.1 Tcal AEJ18398.1	DE EE AE	.SA. .LR. .QA. .ME.	MKR LKQ LRQ LKE	. Q . L . . Q . Q . . S . S .	.TA .GI .SS .KA	191 191 191 191
Bthermo Bacidob	Gpho AMW04363.1 Tter ACZ42165.1 Gmal AEU35097.1 Tthe BAD69831.1	V Q A Q RL ED	.QE.	LKR LKE	. <mark>Q</mark> . <mark>L</mark> . . K . Q .	. <mark>T</mark> A . <mark>A</mark> H	191 191 191 191
Bchloro Bproteo Bchloro	Caur ABY35517.1 Daut ACN14859.1 Ctha ACF15037.1	S <mark>E</mark> T <mark>D</mark> DS	.AE. .LE. .LA.	L <mark>KR</mark> LKK LKK	. <mark>K . L .</mark> . R . M . . K . M .	. <mark>G</mark> L .GI .GM	191 191 191
Baquifi Bactino	Hsp AIZ65498.1 Aaeo AAC07332.1 Cacn AFU40489.1 CNit AFU59479.1	ED AE EE	.EE.	L <mark>KK</mark> LKA	. <mark>R</mark> . <mark>L</mark> . . Q . L .	. <mark>G</mark> L . <mark>G</mark> T	191 187 191 191
Aeuryar Bproteo	Nmag ADD05844.1 Afer ACK79767.1 Mpri AFK05994.1	A <mark>E</mark> DD EE	.ET. .AR. .EQ.	L <mark>K</mark> S LKA MKK	. <mark>D</mark> . M . . E . M . . E . L .	. <mark>G</mark> K . AK . GI	191 191 191
Emetazo Emetazo	Lbuc ACV38698.1 Hsap NP 001231573.1 Bmor XP 004922821.1 Atha NP 001332586.1	A <mark>E</mark> DI AV	V.NQ. V.SQ.	V <mark>LD</mark> ILD	. E . I . . E . L .	. <mark>G</mark> I	191 125 170 135
Bnitros Bproteo	Lfer BAM07649.1 Alip CBS85649.1 Shal A0066257.1	SE ED ER	.SN.	LKK LKA	.T.L. .R.I.	. <mark>A</mark> A	191 191 173
Bplanct Bproteo	Bcep ALK18953.1 Psp AMV20944.1 Maes ANG64508.1	DR AE AE	.AA. .EA. .NA.	L <mark>KK</mark> LKR LLQ	. Q . L . . E . V . . E . D .	.DT . <mark>S</mark> G .AK	191 143 189
Aasgard Bcyanob	Fgin AIE88136.1 CLok OLS16333.1 Hsp AFZ44603.1 Gthe AMV09772.1	SE KE RR AR	M.EK. L.E	LKK	. <mark>E</mark> . <u>L</u> .	. ES . <mark>G</mark> F	188 176 176 175
Eamoebo Eamoebo Eeuglen	Ddis XP 637438.1 Einv XP 004256843.1 Tbru XP 011780226.1	NE	.DE. .QE. .AE.	MEK LEK LTE	. <mark>Q</mark> . <mark>L</mark> . . Q . I . . E . M .	.NA .DE .NE	158 139 147
	Tvag XP 001308534.1 e a13 OLB69573.1	D <mark>E</mark>	.NQ.	LME LTT	. E . G . . E . I .	. DE . <mark>G</mark> E	163 137

X external

X ambivalent
X internal