

	10	20	30	40	50	60	70	80	90	100
Enterococcus	MA	LSK	LDNLYRQVILDHSSHPHH	HGTLDA	SSQTIELNN	PTCGDVIELDVAIE	DGV	IKDIAFQGS	GCSISTASASMMT	DAV
Carnobacterium	MA	LTK	LDQLYRQVILDHSSHPHH	HGKLEQ	STTQIELLN	PTCGDVLQLQLVVK	DQV	IKDIRFDGSGCTISQASASMMT	DAV	
Lactococcus	MA	LSK	LDNLYRAVILDHSSNPRH	AGELQT	G-CMVDLNN	PTCGDVIRLTVFEFE	NDV	ISNIAFSGHGCTISTASASMMTVAV		
Streptococcus	MA	LSK	LNHLMAVVADHSKRPHH	HGQLDG	V-EAVQLNN	PTCGDVISLTVKFD	EDK	IEDIAFAGNGCTISTASSMMT	DAV	
Lactobacillus	MG	LSK	LNGLYREVILDYADHPHN	KGELAT	TTNAMTLHN	PTCGDTINLQVEVE	DNK	IKNIAFTGDCGTISQASASMMT	DAV	
Leuconostoc	MS	LHN	LDLLYRQTVMEYAQHPHH	YKPMGLT	ETYHVRKYN	PTCGDIIDLAFEMT	DDK	VTDIYFYGDGCAISKASASMMT	DLV	
Lysinibacillus	MS	FNN	LDQLYRSVIMDHYKNPRN	KGSLEG	EAVTIDMNN	PTCGDRIHLTLKVT	DGV	VEDAKFEGEGCSISMSSASMMT	QLI	
Geobacillus	MS	SNHP	LDQLYRQVIMDHYKNPRN	RGVLEG	TNVDVMNN	PTCGDRIHLTMKVE	DGK	VVDVKFEGEGCSISMSSASMMT	QAI	
Bacillus	MS	FNN	LDQLYRQVIMDHYKNPRN	KGSLED	GSLTVDMNN	PTCGDRIHLTMKVE	DGK	VTDAKFDGEGCSISMASASMMT	QIV	
Exiguobacterium	MD	FNN	LDHLYRQVIMDHYKTPRN	RGALDG	G-VTIDMNN	PTCGDTIRLQLAIE	DDI	VKDAKFDGEGCSISMASASMMT	QAV	
Staphylococcus	MN	FNN	LDQLYRSVIMDHYKNPRN	KGVLDN	GSMTVDMNN	PTCGDRIRLTFDIE	DGI	IKDAKFEFEGEGCSISMASASMMT	QAV	
Paenibacillus	MQ	LDD	LYRRVIMDHYKNPRN	RGTLNT	DAVTINLNN	PTCGDRISLQLQVE	DGK	VIDAKFSGEGCSISLSSASMMTEAV		
Symbiobacterium	MN	LSA	LYQQVILDHYKKPRN	KGAVER	AVLKKHLHN	PTCGDDIEVQVSLGEDGR	IDD	VKWNRGCSISMASASMMSVAL		
Acholeplasma	MD	LKT	LYRSVIMDHYKNPKN	KGLINDS	SYLTVHLNN	PTCGDDLIVQLLIK	DSK	IDLKQQKGCISCCASASVASELL		
Oenococcus	MG	LEG	LNQLYRQVILDAVDQRY	RGQLLN	PTAVQLAHN	PNCGDVLELQIKVS	DRH	ITAIAFQGGCTISQASASILADLA		
Pediococcus	MSN	IQ	ELYQKIIIVNHAKYPIG	VGVLPQ	YNYHAKLKN	PDCGDDIEVYLKVV	DQH	LEQISFTGQGCIIISQASASMMVDAL		
Peptostreptococcus	MD	IR	EIYTEIITEESRNTKN	KRYLEH	PTHEEKGN	PSCGDEITLQLDIE	DGI	IKDASYVGVGCAISQASTSLMIDL	LI	
Finegoldia	ME	LG	SLYTELILEKSRDKSN	RRELEH	PTHCELGN	PSCGDEITLQLKVQ	DDT	IEDIAYTGMGCAISQASTSIMCDVI		
Clostridium	MD	LN	AIYTELIMEHSTSKHN	KRNLDN	PDIKEKGN	PSCGDEITLKLKN	NGV	IEDLAFTGQGCIAISQASTSIMIDL	LI	
Thermoanaerobacter	MSD	LN	QLYSEVIMEHYENSPH	RRELKD	ATHKERGN	PLCGDDITLYLKMN	GDI	IEDASFTGHGCAISQASTSMIDL	LI	
Coprococcus	MQN	NIQ	NRNFYNEILTEHMRPEF	KYDLPA	ANVVEGVN	PNCGDDIWLKLVKE	NGV	IEDGSFVGDGCAISQASADIMLGM	I	
Nitrococcus	MSD		LRDLYQEVILDHNKHPRN	FRSVEP	HSHQADGYN	PLCGDRVTIQLSVEDG		MIASIGFQGDGCAISTASASIMTEVL		
Nitrosococcus	MSE		LLDLYQEVILDHRKRPRN	FHSMEN	ADFQADGHN	PLCGDRVTVFLKMHNG		IIEVVSFQEGGCAISIASASLMTESL		
Polaromonas	MSD		LRDLYQELIVDHYRHPHN	FGPLAG	ANRRAEGFN	PLCGDRLTLYLQVVDG		VIENARFEGSGCAISTASASLMTDAL		
Coxiella	MSD		LNDLYHQLIIDHGRNPRH	FGRLM	PTHHEGYN	PLCGDRLTVFFQEKNG		VITDARFEGSGCAISMASASLMMEAL		
Myxococcus	MSASD		D-LKDLYQEVVLEHSHKRPRN	YRVVEG	ATAEAAGHN	PLCGDQLVVTLKVEGG		VIKDVAFQGGCAISKASASLMTGAV		
Stigmatella	MSS		E-LKDLYQEVVLDHGKRPRN	FRAVEG	ANHRAEGFN	PLCGDQLSVALKVEDG		VIRDIGFQGGCAISRASASLMTGAV		
Anaeromyxobacter	MSE		LTDLYQEVVLDHGKRPRN	YGLEG	ATHRAEGLN	PLCGDRITVAARLEGG		VVRDVRFEFGSGCAISKASASVMTGVA		
Thauera	MSGMQ		DSLRELYQEVIFDHNRP	YRLLPA	ANHHADGHN	PLCGDRLTVYLQVEDG		IVRDAFVGHGCAISTASASLMTGAV		
Rickettsiella	MMS		HLRELYQEMILDHGRNPRH	HHTMPL	ANRLAHGFN	QICGDRLTLYLKDIDHG		KIKAISFQSGCAIAIASASLMSEIL		
Methylococcus			MED		ANRTVEGFN	PLCGDRITLYLVIDDAG		VIRDVSFQGGCAISTASASLMTGAV		
Legionella	MSM		ELRELYQEIIDHNRP	HHAMED	ATTEAKGFN	PLCGDKLTVYLKIQD		LIRDVSFQGGCAISQASASLMTDAL		
Dichelobacter	MND		LSDLYQELILDHKNPRN	FHALTP	CTHSATGHN	PLCGDNLKVTVRLNEG		VIADLSFVGDGCAISKASASMMTELA		
Mariprofundus	MFE		LRDLYQQVIVDHNKSPRN	FGKLAS	FNHEADGYN	PLCGDKLHIYLVNSDDG		IIEVVSFQEGGCAISVASASLMTDAL		
Psychromonas	MND		ELRALYQEVIIDHGRPRN	SKKLEH	PSCTQEGYN	PLCGDRLTLYLRIIDN		RIIDASFQEGGCAISMAASSLMTERI		
Nitrospira	MTT		KSLYQEVILDHNRP	YGKLDK	ASHHAUGHN	PLCGDHLIDIDLNVEGK		HIEGIAFHGESCAICKASASMMTTVV		
Rhodoferrax	MSADP		KALYQEVILDHNRP	YGELDH	PSHHAEGVN	PLCGDHIHVALDLKGD		SVERIAFHGESCAICKASASMMTVAV		
Nitrosomonas	MSL		KSIYQEVILDHNRP	YGALRS	PTHHATGHN	PLCGDRIELDINMLDG		HIEEIAFQGESCAICKASSMMTNAV		
Acidovorax	MTGFTGNA		ENELYQEVVLEHKRAPRN	FGHLPQ	PTHQAQGN	PSCGDRIAVELQLQGG		RVQDIRFTGQGCIAICMASTSMTEAV		
Burkholderia	MNPAASHTVPDDA		QSTLYQELVVEHKRAPRN	FGRLAE	PTHEARGHN	QCGDLDLQVLRIEGG		RIGDIRFDHGCAICIASASMMTEAV		
Acidiphilium	MSGAAD		VGDLYQRLIMERARAPLH	AGRPAR	FDAEAEGDN	PMCGDRVHLRLSC		AGG-AIGEVDHETRGCAICVASADLMADAV		
Gluconacetobacter	MD		QDGLYQROVIERARAPVH	AGPLDG	ATHQEGTN	PMCGDRVRLGVTLDAA		RVVMVRHQTRGCAICVASADMMADLA		
Thermosinus			MYTEKVMDFHTNPRN	VGEIEDAN		GIGEVGNACG DIMRIYLKIE	NDI	IKDVKFKTFGCGAAIATSSMVTEMV		
Desulfotobacterium			MYTEKVMDFHTNPRN	VGEIEDAN		GVGEVGNACG DIMRIYLDVE	GDI	IKDVKFKTFGCGAAVATSSMVTEMV		
Heliobacterium			MYTDKVMDFHTNPRN	VGEIENAS		GVGEVGNASC DIMRIYLDVE	DNI	IKDVKFKTFGCGAAIATSSMVTELI		
Desulfotomaculum			MYTEKVMDFHTNPRN	VGEIENAD		GIGQVGNPSCG DIMKITLKVE	DNI	IKDIKFKTFGCGAAVATSSMVTEMA		
Moorella			MYSEKVMDFHTNPRN	VGEIENAD		GVGQVGNPVCG DIMRLYIKVE	DGI	IKDVKFKTFGCGAAIATSSMVTEMV		
Carboxydotherrus			MMYSEKVMDFHTNPRN	VGEIPDAD		GVGEVGNPSCG DIMRIYIKVD	GDK	ITDVKFKTFGCGAAIATSSMVTEMV		
Halothermothrix			MYSEKVMDFHTNPRN	VGEIKDAD		AVGEVGNPVCG DIKLYLIKIK	DDT	IEDIKFKTFGCGAAVATSSMVTELV		
Pelotomaculum			MYSEKVIHFHTNPRN	VGEIPDAD		GVGEIGNQVCGDITKIFIKVE	DNV	IKDIKFKTFGCGAAIASGSMTEMA		
Natranaerobius			MYNEKVMDFHTNPRN	VGEVESP		GVGEVGNVTCG DIMRISIKVN	NNEEIED	IKFKTFGCGAAIATSSIVTEMA		
Anaerofustis			MYNDIVLDHFHTNPRN	VGVIEDAT		VVAKEASPCGDTTEFFLKDID	DNDVITD	IKFRTFGCAAIAIASASMSTELI		
Desulfuromonas			MYTDKVMDFHTNPRN	VGQIENPN		VVVKVGDPGCGDAVLIFLKDID	DN	VITDVKYKVGCGAAIATSSMASTMV		
Pelobacter			MSFAHYSPKVFDFHTNPRN	NGVLEDAN		GIGEIFDPECGDHLKVYVKIE	DD	IVKDIKFKQIKGCPAAIACASAMTELV		
Geobacter			MAEIYSAKVVDHVRNPRN	VGSLEDAN		VVVQAGDPTCGDAVLYFLRIE	ED	IVRDIKFKLIKCGAAIATSSVATELV		
Desulfococcus			MIYSKVTMDHFRNPRN	VGVLENAA		GVGEVGNP ICGDMMTIYLDIQ	DD	RIADIKFQTFGCGSAIAVSSMLTELA		
Anaerotruncus			MLYSKVDVLEHFANPHN	VGEIENAD		GIGEVGNACG DIMKMYLKI		KDGVIEDAKFKTFGCGAAIATSSMATDMI		
Faecalibacterium			MASMYSAKVMHFANPHN	VGELPDAN		GVGEVGNPKCG DIMRMYLKI		ENNVIVDVKFLTFGCGAAIATSSMATDLI		
Anaerostipes			MYSEKVMDFHEHPRN	VGEIEGAS		GVGTVGNACG DIMRIYFDID	DNQIIQD	VKFKTFGCGAAVATSSMATELV		

Sequence alignment table showing protein sequences for various bacterial species, aligned by position (10-100). The sequences are color-coded by amino acid type. Species listed include Dorea, Ruminococcus, Alkaliphilus, Ehrlichia, Cowdria, Anaplasma, Wolbachia, Plesiocystis, Sorangium, Oceanospirillum, marine_proteobac., Pseudomonas, Azotobacter, Colwellia, Moritella, Shewanella, Aeromonas, Pseudoalteromonas, Alteromonadales, alpha_proteobac., Haemophilus, Pasteurella, Actinobacillus, Mannheimia, Neisseria, Vibrio_cholerae, Vibrionales, Photobacterium, Shigella, Escherichia_coli, Salmonella, Enterobacter, Klebsiella, Photorhabdus, Xenorhabdus, Serratia, Yersinia, Erwinia, Providencia, Buchnera, Janthinobacterium, Herminiimonas, Azoarcus, Thiobacillus, Ralstonia, Dechloromonas, Chromobacterium, Limnobacter, Bordetella, Polynucleobacter, Methylibium, Leptothrix, Delftia, Comamonas, Verminephrobacter, Methylobacillus, Methylophilales, and Acinetobacter.

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      10      20      30      40      50      60      70      80      90     100
.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
Psychrobacter-----MAYSNQVIDHYENPRN-VGNLDKDAKNVGTGMVGAPACGDVMRLQIQVG-DDGIIEDARFKTYGCGSAIASSSLVTEWL
Candidatus-----MAYGEKVMDHYENPRN-VGVLDKDAKNVGTGMVGAPACGDVMRLQIQVD-DNGVIEDAKFKTYGCGSAIASSSLTEWV
Endoriftia-----MAYSDKVLDHYENPRN-VGAFDNEEKSVGTGMVGAPACGDVMRLQIKVN-DAGVIEDARFKTYGCGSAIASSSLTEWV
Magnetococcus-----MAYNEKVLDHYEKPRN-VGSMDKDADDVGTGMVGAPACGDVMKLQIKVD-DNGVIEDAKFKTFGCGSAIASSSLVTEWI
Magnetospirillum-----MAYSDKVIDHYEHPRN-VGALDKDDSAVGTGLVGAPACGDVMKLQIKVS-AEGIIEDAKFKTFGCGSAIASSSLVTEWV
Rickettsia-----MTMAYSKKVIDHYENPRN-VGSLDKENKNVGTGLVGAPACGDVMKLQIAVD-DDGIITDAKFKTFGCGSAIASSSLVTEWV
Neorickettsia-----MAYSDSVLNHYNNPRN-VGTIDKNKNVGTGLVGAPACGDVMRLQIEVD-NNGCIVDAKFKTFGCGSAIASSSLATEYL
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	110	120	130	140	150	160	170	180	190	200
Enterococcus	LGKTIAEATALAEDFSQLVQGN	---	EVAED	-EKLGD	AAMLSGVA	-KFPARIK	CATLAWKALEQAVA	-----	NNGQGEAGHLHCEK	-----
Carnobacterium	MGKTVSEALALADQFSLLVQGK	---	DAPKL	-EELGDA	ALLNGVA	-KFPARIK	CATLSWKALEKALV	-----	EK	-----
Lactococcus	LGKTKEEAKELATIFSAMVTGE	---	TDERQ	-EKLGDA	QFLAGVS	-KFPARVK	CSTLAWNALKKAID	-----	VGEAQET	-VIHGE
Streptococcus	IGKSKEEALALADIFSEMVQGG	---	ENPAQ	-KELGDA	EELLGVA	-KFPQRIK	CSTLAWNALKEAIK	-----	RSANAQHLTDQNVKEGKNV	-----
Lactobacillus	KGKTTEEALAMAKTFSDMAIGKE	-HSEADL	-DQLGD	ARILTSIM	-EFPARIK	CATLSWWALQALL	-----	KDSEEEENNE	-----	
Leuconostoc	LGQRREAVATLLEEFskLTRG	---	EVADT	-KLLGEA	QILAGVT	-KFPTRIK	CATLAWHALDELIS	-----	VEK	-----
Lysinibacillus	KGKKVDEALELADIFSKMMGE	---	EYSDK	-YLEDVE	ALQGV	-QFPARIK	CATLAWKAMEKG	-----	VK	-----
Geobacillus	KGKTVEEALRLAHIFSDMIQGK	---	EYDSS	-VDLGD	IEALQGV	-KFPARIK	CATLAWKALEKG	-----	LHNHEHEGR	-----
Bacillus	KGKDVESALKLSEVFSNMVQGG	---	EYDED	-IDLGD	IEALQGV	-KFPARIK	CATLAWKAMEKG	-----	VKEGQ	-----
Exiguobacterium	KGKTVEEALQLANVFSEMVQGG	---	DYDEK	FDLGD	EALSGVT	-KFPARIK	CATLAWKALERG	-----	VEEGK	-----
Staphylococcus	KGHSLGEAMQMSQEFTKMMLGE	---	DYVIT	-EEMGD	IEALQGV	-QFPARIK	CATLAWKALEKGTV	-----	AKEGKAEGTTEEE	-----
Paenibacillus	KGKTYEEALGLAERFSGLMKGE	---	DVDFE	-ENED	IEALSGVN	-KFPARIK	CATLAWNALRKG	-----	IEQKQV	-----
Symbiobacterium	KGKTLAEAQELMQSFYRMIQGE	---	QGNKY	-ALGEI	QALSGVS	-KFPVRIK	CATLAWHCLIEGIR	-----	EYEGGDANG	-----
Acholeplasma	KNKDIFEAKELIQTFYDMLTGE	---	EIKDKS	-VLEDAL	AFEGVG	-QFPARIK	CATLAWKAYEKGLN	-----	PLEGEQNE	-----
Oenococcus	LHQSLQSMQTRISQFQKMITGK	---	ESVEI	-DQLGD	ASVFSKIS	-QFPTRVK	CAGLVWDALEQMLK	-----	NN	-QFD
Pediococcus	SGKRILEAQQLMNFQNLILG	---	KDYDE	-SQLGD	LIAFATLN	-QFPTRVK	CGMLAWHAMADGLN	-----	QGGISFGK	-----
Peptostreptococcus	KGKTLDEAKELCEFTFLAMIT	-E	-GLEGDEL	-KKLKDA	IALQNIS	-TMPARVK	CAVLAWHTLKNILE	-----	AN	-----
Fingoldia	KGKSVNEAIDLCKNFISMIKGE	-ITDRKEL	-KVLKDA	AVCFQNIS	-TLPARVK	CAVLAWHTLKDML	-----	NDKSEGSFIPD	-----	
Clostridium	KGKNIEEALKLTETFIGMIKRE	-IKDDEEL	-YALEDA	MAFKNIS	-NMPARVK	CAVLAWHTLKEALE	-----	K	-----	
Thermoanaerobacter	KGKDKKEALRLVQEFIDMIH	---	KKDVNL	-DELGD	AQVLHGVS	-DFPARVK	CALLAWKTLQEIL	-----		-----
Coprococcus	IGKTKEEALKLGLFLKMIQGG	---	EATDEEI	-DQLEE	ASALKDIA	-HMPARVK	CAVLGWHITLLEALK	-----	NIS	-----
Nitrococcus	EGKTVDEAQAQFDQFHDLVTRDEACPEA	---	ALGKIA	VLAGVR	-DYPMRVK	CATLAWHTLHAALT	-----	EKDEVTE	-----	
Nitrosococcus	KGKSSQEAEGFLGKFDLVTDESTHREEG	---	LSLGL	GLVLAGVK	-AFPMPRVK	CATLAWHTLHAALA	-----	HENKMATTE	-----	
Polaromonas	KGKTEETEVENLFASFHALVAGGSDSPS	-T	-VALGK	LEVLAVGR	-EFPVVRVK	CATLAWHTLRAALR	-----	EVSQPVSTE	-----	
Coxiella	KGKSIQAAEILFSQFHDLVTG	---	TKRE	-T	-AQLGK	LAVLAGVA	-EYPARVK	CATLAWHTLRAALH	-----	HNTTTLVKTE
Myxococcus	KDRTRAEEADLFERVHKLVTGEPESVD	-V	-DALGK	LAVLSGVS	-EFPARVK	CASLAWHTLRAALE	-----	GRGEAVSTE	-----	
Stigmatella	KDKTCEEAEMLFARVHQLVTEGPAEVD	-T	-EALGK	LTVLSGVS	-EFPARVK	CASLAWHTLRAALH	-----	EEPEPVSTE	-----	
Anaeromyxobacter	KGRTPAEIDAVFERFHRVTEGPGAVD	-A	-GQLGK	LAVFGGVH	-EYPTRVK	CASLAWHALRQALR	-----	GGGEAVSTE	-----	
Rhauera	KGKPVAAEVEALFRDVHALLTEGREATDPA	---	RDFGK	LEVLAVGR	-EFPARVK	CATLAWHTLHNLV	-----	GEHETAHTE	-----	
Rickettsiella	SGKTCKEAETCFERFHQLLTRELTTAEQL	---	QDLEK	LAVFAGVK	-AYPARVK	CATLAWHTLLAALK	-----	QNSILVSTESSLLSYDHTLSAE	-----	
Methylococcus	RNMHESEAHALFETFHRMATG	---	DDAVNL	---	EELGK	LAVLAGVR	-AYPARVK	CATLAWHSLAAL	-----	NQETTVTTE
Legionella	KGKSIKEAHELFHRVHRMLTQE	---	EEDSL	---	VSMDK	LTVLAGVK	-AFPMPRVK	CATLAWHTLEAALN	-----	KETEIVVKTE
Dichelobacter	MGKTLLEEFQKLYDLFHYIATTONPIQGE	---	VGKLQ	ALAGVR	-QYPARVK	CATLAWHTLDAALH	-----	HKNLVKTE	-----	
Mariprofundus	KGTELAAFQQRFEQFQHMVTADIDEEDT	---	DVLGK	LAVLSGVR	-EFPSPRIK	CASLAWHTMKAAL	-----	DSGKAATE	-----	
Psychromonas	KNMTVTEARQLFTAFQNLVTQ	---	AEPDN	IREYLG	KLSVLGGVR	-DFPVRVK	CATLAWHTLHAALN	-----	NTKQIVSTE	-----
Nitrosospira	KGKSLGDAEMLIKEFRDMATGALDLAH	-P	-HHLGR	LTVFAGVR	-DLPTRVK	CAILPWHITLHAALN	-----	SVSTTSTEAEDDPMHAAIGDA	-----	
Rhodoferrax	KGKSAIEAKTLIHEFVDMATGRIAAKD	-S	-PHIGR	LAVFSGIS	-ELPMRVK	CAILPWHITLQAAFN	-----	AEPSASTEAEADPMHTPLGDA	-----	
Nitrosomonas	KGKSHQEAELIQEFREMLVSGEDKS	---	F	-DHLGR	LKVLAVGR	-DLPTRVK	CAILPWHITLHAAMN	-----	STDSATTEADHASKLVANNH	-----
Acidovorax	KGKEVAAAQALQQHFR	---	AVLTG	QAVHDE	---	APLGK	LVSLAVGR	-QYPSRIK	CALLGWHALHAIA	-----
Burkholderia	IGRDVEAARELQQRF	---	AVLTG	QAVHDE	---	ASLGK	LESLAVQ	-RYPRIK	CALLGWHALHAIA	-----
Acidiphilium	AGRSRAAAEELADAFEMVATG	---	AVPDRED	---	FSELRA	LSGVH	-EYRSRHR	CATLAWHTLQALRAALT	-----	KTMETGHGG
Gluconacetobacter	PGRSVAELGVLSTRAFDMRLRTGGDAPN	---	PELAT	FAGLH	-RHRSRIR	CATLAWHTLQALRAALT	-----	ESKEG	-----	
Thermosinus	KGKTIDEALKIS	-----	NQAVA	EALGGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Desulfotobacterium	KGKTIEEALKIT	-----	NAAVA	EALDGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Heliobacterium	KGKTIDEALGLT	-----	NRAVA	DALGGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Desulfotomaculum	MGKTIDEALTIT	-----	NKAVA	EALGGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Moorella	KGKTVEEALKIS	-----	NAAVA	EALDGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Carboxydotherrhus	KGKTIEEAMKIT	-----	NKAVA	DALDGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Halothermothrix	KGKKIDEALKVS	-----	KEITVA	EALDGL	PSNKMHCNLAADALHEAIKDYLNKKGK	-----			-----	
Pelotomaculum	KGKTIEEALQIT	-----	NKHVA	DMLGGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Natranaerobius	KGKHINEARDIK	-----	NKDVA	DELDEL	GLPKNKLHCNLAADALHEAIKDYLNKKGK	-----			-----	
Anaerofustis	KGKTVEEARKIT	-----	NKMVV	DSLGL	PAPAKMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Desulfuromonas	EGKTLLEAEVLT	-----	DEKVA	EALGGL	PDSPMHCSNLAADALHEAIKDYLNKKGK	-----			-----	
Pelobacter	MGKPVGEAMLVS	-----	DDQII	EYIDGL	PEFKVHCSALGASGFRVAVMDYSIKSKLFGATEG	-----			-----	
Geobacter	MGKGLDEVMLGS	-----	DQIIA	QALDGL	PEEKMHCNMAASALHAAVEQY	-RATVAGETKPL	-----		-----	
Desulfococcus	KGKTLLEAKKIT	-----	NKDVA	EALGGL	PKNKLHCNLAADALHEAIKDYLNKKGK	-----			-----	
Anaerotruncus	KGSTIEDALKLS	-----	NKAVV	EALDGL	PASKIHCNLAADALHEAIKDYLNKKGK	-----			-----	

	110	120	130	140	150	160	170	180	190	200	
										
Acinetobacter	KGKTLDEAQA	IK	-----		NIDIA	TELA	LPPVKVHCSVLA	EDA	IKAAIEDYRSKKS	KA	-----
Psychrobacter	KGKTLDQASE	EIK	-----		NKHIA	EELA	LPPVKVHCSVLA	EDA	IKAAIEDYRAKSGTVEA		-----
Candidatus	KGKTLNEAAE	EIK	-----		NSDIA	EELS	LPPVKIHCSVLA	EDA	IKAAINDIKS	KA	-----
Endoriftia	KGKTLDEAQQ	IK	-----		NSEIA	EELA	LPPVKVHCSVLA	EDA	IKAAIKDYTDKNAG		-----
Magnetococcus	KGKTIDEAL	TIK	-----		NKDVA	EELA	LPPVKVHCSVLA	EDA	IKAAVKDYREKQOKKGA		-----
Magnetospirillum	KGKTLDEAAS	IK	-----		NTDIA	QELA	LPPVKIHCSVLA	EDA	IKAAIADYKKKSG		-----
Rickettsia	KGRSVEDAET	IK	-----		NTEIA	KELS	LPPVKLHCSLLA	EDA	IKAAIADYKQKESKKS	D	-----
Neorickettsia	IGKSIEEA	EKIK	-----		NTEIA	ASTLC	LPPIKMHCSMLA	EDA	IKAAIKDFREKQVTSSTEEAGNENTENKS		-----