

Laterally Transferred Genes between Chlamydiales and Actinobacteria (MurA)
 E. Griffiths and R.S. Gupta (2006) J Mol Evol. 63(2):283-296.

		327		374
Proteobacteria	<i>E. coli</i>	4972324	ETVFENRFMHVPELSRMGAHAEI	ESNTVICHGV EKLSGAQMVDLRA
	<i>Pas. multocida</i>	P57821	--I-----I---I---GK---	-G--A-----DH---E-----
	<i>Pse. aeruginosa</i>	NP_350124	-----Y-MN----QILV	-G--A-VT--P--K--P-----
	<i>Bru. suis</i>	NP_697288	--I-----Q--A-L--KISL	SGQ-ATVE--R-K-----
	<i>Ri. prowazekii</i>	3861126	-NI-----C-----DIVV	RG-KAVVR--M-K-E-----S
	<i>Ca. crescentus</i>	0AAK24321	--I-----A---M-L--DISV	SGGEARVR--DQ-E-----
	<i>Ral. solanacearum</i>	NP_521074	--I-----Q--N-L--NIIT	-G--AAVT--Q-----T-----
	<i>Nei. meningitidis</i>	AL162759	-I-----N-----NITT	-G--AFVQ--R-----V-K-----
	<i>Des. vulgaris</i>	YP_012468	-NI-----L--V-----DIR-	SGRSVVR--KR-T--P--S-----
	<i>Bde. bacteriovorus</i>	NP_967097	-----T-----L--DITP	KTRVAVVR-CP G--T--P-----
	<i>Wol. succinogenes</i>	NP_906592	-RL-----S--Q-L--GIHL	RG--ATVN-G TE--LC-D-----
	<i>Camp. jejuni</i>	CAB73123	-RL-----S--L--DIKL	NGHIATIV-G KE-NA-D-----
	<i>Hel. pylori</i>	26695	--L-----AS--Q-L--NISL	KT-VATIS-S TE-T-SD-----
	<i>Chl. trachomatis</i>	4887172	---H---LGYLKG-VK---CDL	FHECLSAKSCRYSTGN HPHSAVI--P TP-QATDLVIP---
	Chlamydiales	<i>Chl. muridarum</i>	Q9PJT7	---H---LGYLSG-AK---CDL
<i>Chlam. caviae</i>		NP_829044	---H---LGYLRG-QQ---SC-L	FYQCLSSKACRYATGN FPHSAVI--P TP-RSSHLVIP---
<i>Chlam. pneumoniae</i>		4376864	---H---LGYLHG-QH---ECQL	FHQCLSTKACRYAIGN FPHSAVI--A TP-WASHLVIP---
<i>Chlam. abortus</i>		AY038586	---H---LGYLRG-QK---NC-L	FYQCLSSKACRYATGN FPHSA-I--- TP-KASHLVIP---
<i>Chlam. psittaci</i>		AY038585	---H---LGYLRG-QQ---NC-L	FYQCLSSKACRYATGN FPHSAVI--A TP-KASELVIP---
<i>Chlam. felis</i>		AY038587	---H---LGYLRG-QK---NC-L	FYQCLSSKACRYATGN FPHSA-I--- TP-KAS-LVIP---
<i>Proto. amoebophila</i>		YP_007228	---Y---GYTDT-KE---DITP	FRQCLGGKSCRFASQS F-HSA-IV-A TP-V-KEIRIP---
<i>Sim. negevensis</i>		AY845411	---H---GYIKT-KE---DV-L	FTQCLGGRPCRFGAQN YEHS-L-VK-P TPFV-KEIEVP---
<i>Wad. chondrophila</i>		AF468694	---Y---GYTDT-KE---EITL	FRQCLGGKRCRFSSQA F-HSL-VK--- SP-T-REINIP---
<i>Str. lividans</i> (1)		BAA85335	---Y-S-LGFTSA-NQ---IQL	YRECLGGSDCRFGRQN FLHSAVVS-P T--E--DLVIP---
<i>Str. coelicolor</i> (1)		CAB72195	---Y-S-LGFTSA-NQ---IQL	YRECLGGSDCRFGRQN FLHSAVVS-P T--E--DLVIP---
<i>Tro. whipplei</i>		NP789193	---Y-S--GFTEA-N---DISV	HENGFQGTLLRRVPHRA IEQAAVIN-P TP-T-QDIDVP---
<i>Bif. longum</i>		ZP_00121228	---Y---GFTDA-I---TIQL	YRECLGSLPCRFRQRN YKHSVIF-P TP-T-RDIDVP---
<i>Act. naeslundii</i>		TIGR_240017	---YE--GFTGA-CK---TIQV	YRECLGGTECRFGQRN FYHSAVIS-P TP-T--DIVVP---
Insert-Containing Actinobacteria		<i>Brev. linens</i>	NZ_AAGP01000007	---Y---GFTHA-GE---QIQL
	<i>Art. sp.</i>	EAL96147	---Y---GFTDA-I---NIQV	HRECLGSDVPCRFRQRN FLHSAVIS-S TP-K-TDIDVP---
	<i>Leif. xyli</i>	AAT88840	---Y-Q--GF-DA-MD---KIQV	HKECLGGHACRFGRQN FKHSAVIM-P VD-K--DVIEIP---
	<i>Aqu. aeolicus</i>	2983705	-NI--H--H-AQ--N-L--NITV	RG--AYVE--R-Y-SE-YS---
	<i>Cb. tepidum</i>	NP_661455	DRIYLE--N-I--N-L--I--	RD-WALV--P QE-T-TK--S-----
	<i>Por. gingivalis</i>	NP_905531	QKM--S-LFF-DK-ID---QIILC	DPHRATII-LDKRVP-RA-TMVSP-I--
	<i>Cyto. hutchinsonii</i>	ZP_00308053	QKM--S-LFF-DK-ID---QIILC	DPHRATVI-LGKQVA-R-ISMSPP-I--
	<i>Bact. fragilis</i>	YP_100979	QKM--S-LFF-DK-ID---QIILC	DPHRAVVI-HNHGFT-R-GNMTSP-I--
	<i>Tre. pallidum</i>	2780178	-KM--S-MFF-DK-IT---RIILC	DPHRALVS-P SA-H-SDLVSP-V--
	<i>Bor. burgdorferi</i>	2688372	-KM--S-MFF-DK-IK---RIVLC	DPHR-VVT-K SS-K-NVLSSP-V--
	<i>Lep. interrogans</i>	YP_003158	-KM--S-LFF-DNIIA---QIILC	DPHRA-VI-H SR-Y-QK-ASP-I--
	<i>Gloe. violaceus</i>	NP_926071	-----L-----N-L--DIRV	R-GHA-VR--L-----P-V-----
	<i>Nostoc. sp.</i>	NP_484218	-S-----LR-AS--N-L--DIRV	KG--AFVR--PL----P-IG-----
	<i>Therm. elongatus</i>	NP_681569	-----G--A--N---DIRV	KG-HAAIR--PH----P-TG-----
	<i>The. thermophilus</i>	YP_143406	DRIYPD--T--G--A---ELYL	RDRILTVQ--RR-H---K-L-I--
Other Gram(-)ve Bacteria	<i>D. radiodurans</i>	AAF10696	DP-YPD-LT--A--H---TITV	SGY-QVIQ--GT-HA-P-K-A---
	<i>T. maritima</i>	4980599	-N--KT--L--D--K---DI-V	SG-VA-VK-------P-EG-----
	<i>Bac. halodurans</i> (1)	BAB07503	D-IYNA--K-ID--R---DIKV	-GRSA-IN-K T--Q--K-R-S---
	<i>Bac. halodurans</i> (2)	BAB07468	-----E-FR--NGNIK-	-GRSA-IS-P CQ-Q--E-T---
	<i>Clo. acetabutylicum</i> (1)	NP_349465	-I-----K-F--NIK-	DGRSAVIV--KE-T-CSAR-----
	<i>Clo. acetabutylicum</i> (2)	NP_350124	-SIW-S-LK--D--KK---NIKV	-GTVA-ID--T--N-K---
	<i>Sta. aureus</i> N315 (1)	BAB43186	-----A-FK--N-NINV	-GRSAKLE-K SQ-Q---K-----
	<i>Sta. aureus</i> (2)	BAB43210	D-IYPE--K--E--K---NI-V	DEG-ATIKP ST-H--E-Y-S---
	<i>Lis. innocua</i> (1)	CAC97896	-----E-MR--N-DMK-	-GHS--IS-P A--Q--E---
	<i>Lis. innocua</i> (2)	CAC97923	D-IYPS--K-IA-IE---GKFKL	-GRSAVVS-P V--Q-SK-T-----
	<i>Strep. mutans</i> (1)	NP_721869	-----Q-LE-MR---LTS--	LRD-AMI--G -Q-Q-----SS---
	<i>Strep. mutans</i> (2)	NP_721911	D-IY-K-VN--A--A---KISV	LGGQI-YE-P NE---P-K---
	<i>Cor. diphtheriae</i>	NP_940224	-NI--A--RF-D--V-L--D-TV	DGHH-VMR--STP-WSS-I---
	<i>Myc. tuberculosis</i>	I361416	-N--A--RF-E-MI-L--D-RT	DGHHAVVR-L PQ--S-P-WCS-I--
	<i>Noc. farcinica</i>	YP_117278	-NI--A--RF-E-MI-L--D-RT	DGHHAVVR-I PQ--S-P-WSS-I--
<i>Sym. thermophilum</i>	YP_077129	-LY-D--RF---V---QIRV	-GR-A-VK--CR-Y--P-E---I--	
<i>Rub. xylanophilus</i>	ZP_0018649	-N-Y---LQVAE--N---GIDL	FGGHRALVR-P RR---TT-Q-P---	
<i>Thermo. fusca</i>	ZP_00293281	-AI-DG--ALAD--NK---KI-V	-G-RA-V--P SE-H-TK-T-H---	
<i>Kin. radiotolerans</i>	AAEF02000010	---YD--AIYLTE-NRL--RV-L	LDPHR-LVE-P TRW---E-L---	
<i>Str. coelicolor</i> (2)	NP_733681	---YD--AIYLTD-N-L-GRLQL	LDPHR-LVE-P TRWRA-EM-CPPA--	
<i>Str. avermatis</i> (2)	NP_823436	---YD--AIYLTD-N-L-GRLQL	LDPHR-LVE-P TRWRA-EM-CPPA--	
<i>Pro. acnes</i>	YP_055933	WCY---AI-LLD-GKL--DVQL	LDPHRL-VR-P TRWR-RELI	

Figure 5. Partial sequence alignment of MurA proteins from different species showing a 16aa conserved insert that is commonly shared by various *Chlamydiales* and a subset of Actinobacteria. Additional abbreviations in species names are: *Act.*, *Actinomyces*; *Art.*, *Arthrobacter*; *Therm.*, *Thermosynechococcus*.