



Cloning and Expression of Novel Aminoglycoside Phosphotransferase Genes from *Campylobacter* and Their Role in the Resistance to Six Aminoglycosides

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ABSTRACT Nine *aph* genes, including *aph*(2")-*lb*, *aph*(2")-*lc*, *aph*(2")-*lg*, *aph*(2")-*lf*, *aph*(2")-*lf*1, *aph*(2")-*lf*3, *aph*(2")-*lh*, *aac*(6')-*le*–*aph*(2")-*la*, and *aac*(6')-*le*–*aph*(2")-*lf*2, were previously identified in *Campylobacter*. To measure the contribution of these alleles to aminoglycoside resistance, we cloned nine genes into the pBluescript and expressed them in *Escherichia coli* DH5 α . The nine *aph* expressed in *E. coli* showed various levels of resistance to gentamicin, kanamycin, and tobramycin. Three genes, *aac*(6")-*le*–*aph*(2")-*la*, *aph*2"-*lf*1, and *aph*2"-*lg*, showed increased MICs to amikacin, and five *aph* genes were transferrable.

KEYWORDS Campylobacter, gentamicin resistance, NARMS, cloning and expression

The aminoglycoside 2"-phosphotransferase [APH(2")] family is a major contributor to gentamicin resistance (Gen^r) in *Campylobacter*. Nine variants of *aph* genes were previously identified in *Campylobacter* isolated from human and retail chickens, including seven monofunctional *aph*(2") genes, *aph*(2")-*lb*, *aph*(2")-*lc*, *aph*(2")-*lg*, *aph*(2")-*lf*, *aph*(2")-*lf*1, *aph*(2")-*lf*3, and *aph*(2")-*lh*, and two bifunctional *aph*(2") genes, *aac*(6')-*le*/*aph*(2")-*la* and *aac*(6')-*le*-*aph*(2")-*lf*2. Five of them, *aph*(2")-*lg*, *aph*(2")-*lf*1, *aph*(2")-*lf*3, *aph*(2")-*lh*, and *aac*(6')-*le*-*aph*(2")-*lf*2, were novel genes (1).

Toth et al. showed that aph(2'')-If confers resistance to several aminoglycosides (2). Since the APH(2'') family is genetically diverse and the percentage of amino acid identity between subfamilies can be as low as 25.9% (1–3), it is important to investigate transferability and the attribution of variants of *aph* gene resistance to other aminoglycosides, in addition to gentamicin.

Eleven *Campylobacter* strains, five *C. jejuni* and six *C. coli*, obtained from the National Antimicrobial Resistance Monitoring System (NARMS) program were used in this study (Table 1). All isolates were previously sequenced using the Illumina MiSeq (Illumina, San Diego, CA, USA) (1, 4). The coding sequences of nine *aph* genes were synthesized and cloned in a pBluescript expression vector by GenScript (Piscataway, NJ). For two bifunctional *aph*(2") genes, only the *aph*(2")-*la* and *aph*(2")-*lf2* genes were cloned into the pBluescript vector. The pBluescript::*aph*(2") recombinant plasmids were transformed into competent *Escherichia coli* DH5 α , according to the protocol provided by Thermo Fisher Scientific (Invitrogen, Carlsbad, CA). The clones were grown on LB agar plates containing 50 μ g/ml ampicillin and 8 μ g/ml gentamicin.

Expression of the cloned *aph* genes in *E. coli* DH5 α was first determined by MICs of gentamicin using broth microdilution (CMV3AGNF; Thermo Fisher Scientific, Trek Diagnostics, Cleveland, OH), following standard protocols (5). *E. coli* DH5 α carrying the pBluescript vector without the *aph* genes was used as a control. To measure the contribution of different *aph* genes to additional aminoglycoside resistance, agar dilution was performed, and the MICs of six aminoglycosides, including gentamicin,

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TABLE 1	Campylobacter	strains	used in	this	study
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				Resistance			
Strain ID ^a	Species	Source	Yr	phenotype ^b	Resistance genes ^c	Mutation	Purpose
N29710 ^d	C. coli	Chicken breast	2011	GEN TET	aadE aad9 aph(2")-Ig aph(3')-Illa	None	Cloning donor
					bla _{OXA-61} sat4 tetO		
N20344 ^d	C. coli	Chicken breast	2009	GEN TET	aph(2")-Ic aph(3')-Illa bla _{OXA-61} tetO	None	Cloning donor
41912 ^d	C. jejuni	Human	2008	CIP GEN NAL TET	aadE aac(6')-le– aph(2")-lf2 bla _{OXA-61} tetO	GyrA86l	Cloning donor
41921 ^d	C. jejuni	Human	2008	CIP GEN NAL TET	aadE aad9 aac(6')-Im aph(2")-Ib tetO	GyrA86l	Cloning donor
41945 ^d	C. coli	Human	2010	CIP AZI CLI ERY TEL	aadE aad9 aac(6')-le– aph(2")-la aph(2")-	GyrA86I 23S rRNA	Cloning donor
				GEN NAL TET	lf1 aph(3')-Illa bla _{OXA-61} catA tetO	2075G	
41898 ^e	C. coli	Human	2003	CIP GEN NAL TET	aadE aac(6')-le–aph(2")-la bla _{OXA-61} tetO	GyrA86l	Donor
41902 ^e	C. jejuni	Human	2005	CIP GEN NAL TET	aph(2")-If tetO bla _{OXA-61}	GyrA86l	Cloning donor
41904 ^e	C. coli	Human	2006	CIP AZI CLI ERY TEL	aadE aad9 aph(2")-If3 InuC aph(3')-IIIa	GyrA86I 23S rRNA	Cloning donor
				GEN NAL TET	sat4 tetO	2075G	
41905 ^e	C. jejuni	Human	2007	CIP AZI CLI ERY TEL	aadE aad9 aph(2")-Ih sat4 aph(3')-Illa	GyrA86I 23S rRNA	Cloning donor
				GEN NAL TET	bla _{OXA-61} tetO	2075G	
N18880	C. jejuni	Chicken breast	2008	AZT ERY TEL	None	23S rRNA 2075G	Recipient
N46788F	C. coli	Cattle	2014	CIP CLI FEN NAL TET	bla _{OXA-61} aph(3')-Illa tetO	GyrA86l	Recipient

^aID, identifier.

^bGEN, gentamicin; TET, tetracycline; CIP, ciprofloxacin; NAL, nalidixic acid; AZI, azithromycin; CLI, clindamycin; ERY, erythromycin; TEL, telithromycin; FFN, florfenicol. 'The genes in bold are aminoglycoside 2"-phosphotransferase [aph/2"]] genes.

^dThe aph(2") genes from these isolates were successfully transferred to Gen^s recipient strains.

^eThe aph(2") genes from these isolates were not successfully transferred to Gen^s recipient strains.

kanamycin, streptomycin, neomycin, tobramycin, and amikacin, were determined. Agar dilution plates were prepared with concentrations ranging from 0.125 μ g/ml to 1,024 μ g/ml. The MIC was recorded as the lowest concentration of antimicrobial agent that completely inhibited the growth of an organism after incubation at 35°C for 16 to 20 h, according to guidelines of the Clinical and Laboratory Standards Institute (CLSI) (6)

For conjugation, nine Gen^r *Campylobacter* strains that carried variants of *aph(2")* genes were used as donor strains, including four *C. jejuni* (41912, 41921, 41902, and 41905) and five *C. coli* (41945, 41898, 41904, N29710, and N20344) strains. Two gentamicin-susceptible (Gen^s) strains, *C. jejuni* N18880 and *C. coli* N46788F, were used as recipient strains (Table 1). Conjugation was performed as described by Chen et al. (7). Successful transconjugants were then confirmed by antimicrobial susceptibility testing (AST) and whole-genome sequence (WGS) analysis (4, 8).

AST showed that the expression of the nine *aph* genes in *E. coli* resulted in various levels of resistance to gentamicin, kanamycin, and tobramycin. The MICs from different clones ranged from 16 to 64 μ g/ml for gentamicin, 64 to 512 μ g/ml for kanamycin, and 8 to 128 μ g/ml for tobramycin. Comparing the MICs of *E. coli* that carry pBluescript:: *aph*(2") to the control strain, there were 32- to 128-, 64- to 512-, and 16- to 256-fold increases in the MICs to gentamicin, kanamycin, and tobramycin, respectively (Table 2). The MIC differences from nine clones to the same drug may explain the diversity of this family. Furthermore, three clones carrying *aph*(2")-*Ia*, *aph*2"-*If*3, and *aph*2"-*Ig* showed 8-

TABLE 2 MICs of six aminoglycosides for E. coli DH5 α expressing the nine aph(2") genes

		MIC (µg/ml)						
Bacterial strain name	aph genes in pBluescript	Gentamicin	Kanamycin	Tobramycin	Amikacin	Neomycin	Streptomycin	
DH5a	pBluescript	0.5	1	0.5	1	1	2	
E. coli-la	aac(6')-le–aph(2")-laª	64	512	128	8	0.5	2	
E. coli-lb	aph(2")Ib	16	64	16	1	1	2	
E. coli-lc	aph(2")-Ic	32	64	16	0.5	0.5	2	
E. coli-lf	aph(2")If	32	128	16	2	1	4	
E. coli-lf1	aph(2")-If1	32	128	16	2	1	2	
E. coli-lf2	aac(6')-le–aph(2")-lf2 ^a	32	128	32	2	1	2	
E. coli-lf3	aph(2")-If3	32	128	8	8	0.5	2	
E. coli-lg	aph(2")-Ig	32	128	16	16	1	2	
E. coli-lh	aph(2")-lh	64	128	16	1	0.5	2	

^aFor bifunctional aminoglycoside resistance genes, only aminoglycoside phosphotransferase genes [aph(2")-la and aph(2")-lf2] were cloned to the pBluescript vector.



FIG 1 The chromosome-carried Gen^r island integrated into the chromosome of Gen^s isolate through conjugation.

to 16-fold increases in their MICs to amikacin compared to those of the *E. coli* control strain. None of the nine *aph* gene clones showed resistance to streptomycin or neomycin (Table 2).

A similar study conducted by Toth et al. showed that when aph(2'')-Ia and aph(2'')-Ia were cloned in pET22b(+) vector and expressed in *E. coli* BL21, the MICs increased 32-to 128-fold to kanamycin, tobramycin, and gentamicin. There was only a 1- to 2-fold increase in the MICs to neomycin and amikacin compared with control strain *E. coli* JM83 (2). Our results agreed with their findings for these two antimicrobial agents, except that pET22b::aph(2'')-Ia expressed in *E. coli* JM83 had a 4-fold lower MIC to tobramycin (32 μ g/ml) than to pBluescript::aph(2'')-Ia expressed in *E. coli* DH5 α (128 μ g/ml). The difference in MICs to tobramycin between these two experiments could be due to the use of different cloning vectors and expression strains of *E. coli*, as well as the use of different AST methods.

The conjugation study showed that aph(2'')-*Ib* (41921), aph(2'')-*Ic* (N20344), aph(2'')-*If* (41945), aph(2'')-*If* (41912), and aph(2'')-*Ig* (N29710) were successfully transferred to Gen^s strains, either in *C. jejuni* N18880 or *C. coli* N46788. Strain 41945 carried two aph(2'') genes, bifunctional aac(6')-*Ie*-aph(2'')-*Ia* and monofunctional aph(2'')-*If* 1, and only aph(2'')-*If* 1 was transferred based on the comparative genomic analysis. The other four aph(2'') genes, including three monofunctional aph(2'')-*If* (41902), aph(2'')-*If* (41904), aph(2'')-*Ih* (41905) genes and one bifunctional aac(6')-*Ie*-aph(2'')-*Ia* (41898) gene, were not transferred to Gen^s strains (Table 1).

WGS analysis of donors, recipients, and transconjugants showed that transferred *aph(2")* genes, including *aph(2")-lb*, *aph(2")-lc*, and *aph(2")-lg*, and *aac(6')-le–aph(2")-lf2*, are located on plasmids. However, the *aph(2")-lf1* gene from strain 41945, which is located on a chromosome, also was transferred. Comparative genomic analysis of donor (41945), recipient (N46788F), and transconjugant (41945-TC) strains showed that the aminoglycoside resistance island from the donor chromosome was integrated into the chromosome of recipient cells (41945-TC) through a recombination event (Fig. 1). WGS data showed that *aadE*, *aad9*, *sat4*, and *aphA-3*, one of the *aph(2")* genes, and the *tetO* genes were often clustered together, forming a resistance island located either on the chromosome or the pTet plasmid (4, 7). Similar aminoglycoside resistance genomic islands were found either on the plasmid pCG8245 of *C. jejuni* or in the chromosome of *C. coli* SX81 (9, 10).

In summary, all nine variants of $aph(2^n)$ genes are responsible for resistance to

gentamicin, kanamycin, and tobramycin but not to neomycin or streptomycin. Three variants, including *aph*(2")-*la*, *aph*(2")-*lf3*, and *aph*(2")-*lg*, showed decreased susceptibility to amikacin. Both the plasmid- and chromosome-carried *aph*(2") gene can be transferred by conjugation. This study highlights the need for continuous monitoring of emergent resistance genes in foodborne pathogens.

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