Plasmid-encoded trimethoprim resistance in salmonellas isolated in Britain between 1970 and 1981

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SUMMARY

Trimethoprim resistance was plasmid-encoded in all trimethoprim-resistant Salmonella typhimurium and in the majority of trimethoprim-resistant salmonellas of other serotypes isolated since 1970 from humans and food animals in Britain. In S. typhimurium, non-autotransferring plasmids of compatibility group 3 and autotransferring plasmids of group H₂ predominated. The predominance of these plasmid types has resulted from the spread of clones of trimethoprim-resistant strains of phage types 18, 170 and 204c. In other salmonellas, a variety of plasmid compatibility groups have been identified. Almost all plasmids which conferred resistance to trimethoprim also coded for sulphonamide resistance.

INTRODUCTION

An earlier report described the incidence of trimethoprim resistance in salmonellas which had been isolated in Britain from humans and food animals since 1970 and which had been referred to the Division of Enteric Pathogens (Ward, Rowe & Threlfall, 1982). Resistance to trimethoprim was particularly evident in Salmonella typhimurium and by 1981, 6.7% of strains from humans and 14.0% of strains from cattle were trimethoprim-resistant. In other salmonellas, 0.5% of strains isolated in 1981 from humans were trimethoprim-resistant. This paper provides an account of the transferability of trimethoprim resistance in these strains and of the plasmids which specified resistance to trimethoprim.

MATERIALS AND METHODS

Bacterial strains

The source and identification of the trimethoprim-resistant strains have been described previously (Ward et al. 1982).

Resistance transfer and plasmid characterization

Strains were tested for the ability to transfer trimethoprim resistance, either directly or by mobilization, to an *Escherichia coli* K12 F⁻ lac⁺ nalidixic acidresistant recipient strain. The methods were those of Anderson & Threlfall (1974).

		No. tested	Resista	Total with transferable trimethoprim	
Serotype	No.*		Directly	Mobilization	resistance
S. typhi S. typhimurium	3	3	1	0	1
Human	693	693	243 (35.1)	450 (64.9)	693 (100.0)
Bovine	718	718	647 (90-1)	71 (9.9)	718 (100·0)
Others†	37	37	17 (45.9)	20 (54·1)	37 (100.0)
Other serotypes	101	67	29 (43.3)	19 (28.3)	48 (71.6)

Table 1. Transfer of trimethoprim resistance

In parentheses: strains with autotransferring or mobilizable trimethoprim resistance as percentage of number tested.

- * Number of trimethoprim-resistant strains referred to the Division of Enteric Pathogens.
- † Ovine, porcine, avian.

Plasmids which conferred trimethoprim resistance were classified by their incompatibility with autotransferring and non-autotransferring plasmids of the compatibility groups listed by Willshaw et al. (1980), and with other non-autotransferring plasmids described by Smith, Humphreys & Anderson, (1974).

RESULTS

Transfer of trimethoprim resistance

Trimethoprim resistance was transferable either directly or by mobilization in one of three resistant strains of S. typhi, in all strains of S. typhimurium (100%) and in 48 of 67 strains of other salmonella serotypes that were tested (71.6%) (Table 1).

S. typhi

Trimethoprim resistance was directly transferable in one strain of R-type TmCSSu (Tm = trimethoprim, C = chloramphenicol, S = streptomycin, Su = sulphonamides). The plasmid which specified resistance to trimethoprim was of the I complex (Datta, Richards & Datta, 1981); the strain carried a second autotransferring plasmid of the H_2 compatibility group which coded for CSSu. In the two other trimethoprim-resistant strains, trimethoprim resistance was neither transferable nor mobilizable and the levels of resistance were low (8 μ g/ml). In contrast, in the strain with plasmid-encoded trimethoprim resistance, the level of resistance was 1000 μ g/ml.

S. typhimurium

The plasmids which conferred trimethoprim resistance in strains from humans are shown in Table 2 and those in strains from food animals in Table 3.

Autotransferring trimethoprim resistance plasmids were of the H_2 , I_1 and I_2 groups; one plasmid which coded for TmC was not identified. Non-autotransferring plasmids were of two types. Those of one type specified resistance to up to eight drugs including trimethoprim, and were of the F_1 me compatibility group. Plasmids of the second type coded for resistance to TmSu or Tm alone. These plasmids were

Table 2. Trimethoprim resistance plasmids in S. typhimurium of human origin

Autotransferring			Non-autotransferring			
Group	Resistances specified	No.	Gŗoup	Resistances specified	No.	`
H_2	TmCSSuT	170	3*	TmSu	382	
	TmSSuT	22		Tm	3	
	TmC	12		Total	387	
	TmACSSuT	1				
	Total	205	$\mathbf{F_{I}}me$	TmACGKSSuT	46	
			•	TmACKSSuT	7	
$\mathbf{I_1}$	TmKS	13		TmACSSuT	5	
-1	TmS	8		TmACGSSuT	2	
	TmK	6		TmAGKSSuT	1	
	TmSu	3		TmAKSSuT	1	
	TmT	1		\mathbf{TmT}	1	
	Tm	1		Total	63	
	Total	32				
I,	TmS	5				
ÜC†	TmC	1				
1	Grand total	243		Grand total		450

Resistance symbols: A, ampicillin; C, chloramphenicol; G, gentamicin; K, neomycin-kanamycin; S, streptomycin; Su, sulphonamides; T, tetracyclines; Tm, trimethoprim;

Table 3. Trimethoprim resistance plasmids in S. typhimurium from food animals

		Autotransferring			Non-autotransferring			
Host	Group	Resistances specified	No.	Group	Resistances specified	No.		
Cattle	H ₂	TmCSSuT TmSSuT TmC Total	476 74 11 561	3	TmSu Tm Total	70 1 71		
	I, I,	TmKS TmS TmSu TmK Tm Total TmS	31 21 2 1 1 56 30					
		Grand Total	647	•	Grand Total	71		
Pigs, sheep, poultry	H ₂	TmCSSuT TmC Total	15 1 16	3	TmSu Tm Total	18 2 20		
	I_1	TmS Grand Total	1 17		Grand Total	20		

^{*} In accordance with scheme of Smith (1975) for the classification of non-autotransferring plasmids.

[†] Not classified.

Table 4. Trimethoprim resistance plasmids in other serotypes

			Autotransferring		Non-autotransferring			
Serotype*		Group	Resistances specified	No.	Group	Resistances specified	No.	
S. agona	(2)	_				TmT†	1	
•						Tmt	1	
S. anatum	(1)	I,	TmS	1				
S. bareilly	(1)	Č	TmACKSu	1				
S. brandenburg	(1)	Ĭ,	TmS	1	_		_	
S. chicago	(2)	H,	TmASSuT	2				
S. derby	(1)	H ₂	TmSuT	1	-			
S. enteritidis	(1)	I ₂	TmS	1			_	
S. hadar	(1)		_		3	TmSu	1	
S. heidelberg	(8)	H ₂	TmCSSuT	2	$\mathbf{H_2}$	TmCSSuT	1	
		H,	TmACKSSuT	1	_	-		
		H,	TmACGKSSuT	1	_		_	
		آ،	TmKS	1	_		_	
		T,	TmS	1			_	
		I,	Tm	1	_		_	
S. infantis	(1)	н,	TmCSSuT	1	_			
S. mbandaka	(1)	Η,	TmSSuT	1	_	_	_	
S. meleagridis	(5)	N.	TmASSuT	5				
S. montevideo	(3)	C	TmCSuT	1	_		_	
		C	TmACKSuT	1		-		
		I ₁	TmKS	1	_		_	
S. muenchen	(10)	<u> </u>				TmACKSSuT	9	
					$\mathbf{F_{I}}me$	TmAGKSSu	1	
S. oranienburg	(2)	$\mathbf{F_1}me$	TmACGKSSuT	1	-			
		I_1	TmSu	1	_	_		
S. panama	(1)	Н,	TmCSSuT	1		_	_	
S. saintpaul	(2)	H ₂	TmCSSuT	1	H ₂	TmCSSuT	1	
S. virchow	(1)	N	TmASSuT	1			-	
S. wien	(4)		_	<u> </u>		TmACKSSuT	1	
				_		TmACSSuT	2	
		_	-		$\mathbf{F_{i}}me$	TmACGKSSu	1	

^{*} Number of strains in parentheses.

incompatible with the tetracycline resistance determinant NTP 5 (Smith $et\,al.$ 1974) and were therefore assigned to group 3 of non-autotransferring plasmids in accordance with the scheme of Smith (1975). Non-autotransferring TmSu plasmids were identified in strains from humans and food animals, whereas $F_{I}me$ plasmids were found only in strains from humans.

Other serotypes

Sixty-seven trimethoprim-resistant strains of scrotypes other than S. typhi and S. typhimurium were tested. Trimethoprim resistance was directly transferable in 29 strains of 15 scrotypes (43.3%) and was mobilizable in a further 19 strains of 6 scrotypes (28.4%) (Table 1).

The plasmids which coded for trimethoprim resistance were characterized and the results are summarized in Table 4. Autotransferring plasmids were of the C, F_1me , H_2 , I_1 , I_2 and N groups. Non-autotransferring F_1me plasmids were identified in strains of S. muenchen and S. wien, a group 3 resistance determinant which coded

[†] Compatible with standard non-autotransferring plasmids.

for TmSu was identified in a strain of S. hadar, and strains of S. saint paul and S. heidelberg carried transfer-defective H₂ plasmids which coded for TmCSSuT. Two non-autotransferring plasmids in strains of S. agona were not classified.

DISCUSSION

These studies have shown that trimethoprim resistance was plasmid-encoded in all trimethoprim-resistant S. typhimurium and in the majority of trimethoprim-resistant strains of other salmonella serotypes that have been isolated in Britain since 1970. This constrasts with previous findings with E. coli, Klebsiella aerogenes, Proteus mirabilis and Enterobacter spp isolated from humans (Brumfitt, Hamilton-Miller & Grey, 1977; Towner et al. 1978), although an increase in plasmid-specified resistance in these organisms has recently been reported (Towner et al. 1980). Furthermore, plasmids which coded for trimethoprim resistance in salmonellas were, in general of different compatibility groups than those in coliforms, where plasmids of the W, F_{II}, I₂ and B groups have predominated. (Grey, Hamilton-Miller & Brumfitt, 1979; Towner, 1979).

In S. typhi, the one strain with high-level transferable trimethoprim resistance had acquired resistance whilst the patient was being treated with co-trimoxazole (Datta et al. 1981). The low level of resistance in the two other trimethoprim-resistant strains was probably chromosomal.

In both human and bovine trimethoprim-resistant $S.\ typhimurium$, the predominance of non-autotransferring plasmids of group 3 and autotransferring plasmids of group H_2 has resulted from the spread of trimethoprim-resistant clones of phage types 18, 170 and 204c (Ward et al. 1982). Strains of types 18 and 170 carried non-autotransferring plasmids of group 3 which coded for TmSu whilst type 204c strains carried H_2 plasmids which transferred TmCSSuT (T = tetracyclines) or TmSSuT (Threlfall et al. 1980). Group I_1 plasmids were distributed amongst various phage types but I_2 plasmids were found only in related strains of phage types 49, 204, 204a and 193 (Threlfall, Ward & Rowe 1978; Threlfall, 1982). Non-autotansferring F_1me plasmids were found exclusively in strains from humans. These strains were of phage types 208, 66/122 or untypable and were isolated from persons infected in the Middle East or South East Asia, or from persons of Asian origin (Rowe et al. 1980; Frost et al. 1982).

The incompatibility between the tetracycline resistance determinant NTP 5 and the non-autotransferring TmSu plasmids was particularly interesting. NTP 5 was identified in a strain of S. typhimurium phage type 49 isolated in 1969 and, until these studies, has proved to be compatible with all other non-autotransferring plasmids we have tested. NTP5 is a multiple-copy plasmid of molecular weight (MW) 6.5×10^6 (Smith et al. 1974). Studies are in progress to determine the MW of the non-autotransferring TmSu plasmids and the number of copies of these plasmids per chromosome.

A variety of plasmid types were encountered in trimethoprim-resistant salmonellas of other scrotypes. Non-autotransferring $F_{1}me$ plasmids were found in strains of S. muenchen and S. wien and in several cases the patient had been infected in areas where $F_{1}me$ plasmids are widely distributed in number of salmonella scrotypes (Rowe et al. 1980). Of the other plasmid types, I_{1} , I_{2} , H_{2} and group 3 plasmids have been observed in trimethoprim-resistant S. typhimurium

and group C plasmids which coded for trimethoprim resistance have been identified in S. bareilly from outbreaks in India (DEP, unpublished observations).

Until recently, trimethoprim has been available only in trimethoprim-sulphonamides combinations. A consequence of the use of such combinations has been that the majority of plasmids which conferred resistance to trimethoprim also coded for sulphonamide resistance. Trimethoprim-containing products have now been released for use without the sulphonamide component and it is possible that the use of this drug alone will result in a proliferation of plasmids which confer resistance to trimethoprim but not to sulphonamides. This may be avoided by restricting the use of trimethoprim alone for prophylaxis and, in the treatment of salmonella infections, by using this drug only when the strains spread extra-intestinally.

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