

Present significance of resistance to trimethoprim and sulphonamides in coliforms, *Staphylococcus aureus*, and *Streptococcus faecalis*

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SYNOPSIS The incidence of trimethoprim resistance in coliforms and multiresistant strains of *Staphylococcus aureus* isolated in Bristol from 1970 to 1972 is low—2·3 and 1·0% respectively. The resistance is probably intrinsic; there is no evidence that it is R-factor or plasmid mediated. A single mechanism that confers resistance to both trimethoprim and sulphamethoxazole has not been detected. Normal growing one-step mutants of *S. aureus* and *Escherichia coli* resistant to trimethoprim could not be isolated *in vitro*. For these reasons cotrimoxazole should retain its usefulness against these bacteria for some years. However, cotrimoxazole was found not to be bactericidal against many coliforms.

The usefulness of cotrimoxazole against *Streptococcus faecalis* seems limited because mutants resistant to trimethoprim occurred at high frequency in one step.

Since the combination of trimethoprim and sulphamethoxazole (cotrimoxazole) was introduced about four years ago, many reports have shown it to be effective against a variety of bacteria, except *Pseudomonas aeruginosa*. The initial low numbers of bacteria resistant to both its components gave considerable hope for a long therapeutic future for cotrimoxazole, particularly as resistance to sulphamethoxazole alone does not necessarily impair the synergism between the two drugs *in vitro* (Bushby, 1969). However, in the last year there have been some reports of organisms resistant to both components of cotrimoxazole. Amongst coliforms, Lacey, Gillespie, Bruten, and Lewis (1972) found that about 2·5% were resistant to trimethoprim, most also being resistant to sulphamethoxazole. In *Streptococcus faecalis*, resistance to trimethoprim appears to develop after exposure of the organism to cotrimoxazole *in vivo* (Chattopadhyay, 1972). In strains of *Staphylococcus aureus* isolated from hospital sources about 1·6% are now resistant to both trimethoprim and sulphonamides, with 18·5% resistant to sulphonamides alone (Nakhla, 1972). Previously no trimethoprim resistance has been reported in this organism. The overall picture, therefore, is of a small but significant increase in bacteria resistant to cotrimoxazole.

In 1971 R factors producing resistance to both

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trimethoprim and sulphonamides were detected in coliforms isolated from a few London hospitals (Fleming, Datta, and Grüneberg, 1972); Datta and Hedges (1972) have anticipated a dissemination of this R factor to other hospitals and throughout the coliform population generally. We have been monitoring coliforms isolated in Bristol for the last 18 months for the appearance of such an R factor and have found none. We also report some laboratory studies on the resistance to trimethoprim and sulphamethoxazole which might relate to the usefulness of this combination in the future.

Materials and Methods

ORGANISMS

Single colony isolates of coliforms (Gram-negative bacilli; lactose-positive on MacConkey agar), *Streptococcus faecalis*, and *Staphylococcus aureus* were obtained during 1970-1972 from clinical specimens initially plated on antibiotic-free medium. The cultures were stored at room temperature on nutrient agar slopes. Strain J53 (Su, Tp) which is a derivative of *E. coli* K12 harbouring an R factor conferring resistance to sulphonamides and trimethoprim was supplied by Dr N. Datta.

MEDIA

Blood agar was Oxoid blood base no. 2 containing

5% (v/v) lysed horse blood. Antibiotic sensitivity was determined on Oxoid diagnostic sensitivity test (DST) agar containing 5% (v/v) lysed blood and stored overnight at 4°C before inoculation. Oxoid nutrient broth (no. 2) containing 5% (v/v) lysed blood (Harper Cawston broth) was used for estimating the minimum bactericidal concentrations (MBCs) of the drugs. In addition, double-strength Difco nutrient broth and agar were used for the cultivation of *S. aureus*.

ANTIBIOTIC SENSITIVITY

Preliminary testing was done by the use of disks (Lacey *et al.*, 1972). On strains giving absent or reduced zones, minimum inhibitory concentrations (MICs) were determined by the plate dilution method. Inocula were standardized to give about 20 colonies in 0.01 ml broth per test organism. Known sensitive and resistant strains were included in each batch. The endpoint was taken as that concentration of either a single drug or combination that prevented any visible growth after 20 hours' incubation at 37°C. For the determination of MBCs, trimethoprim and sulphamethoxazole were added to Harper Cawston broth the day before inoculation and stored at 4°C. Ten ml volumes of the broth were then inoculated with about 10⁶ organisms suspended in 0.1 ml broth. After incubation for 24 hr at 37°C the broths were subcultured into solid medium. The MBC was defined as the concentration of cotrimoxazole that destroyed > 99% of the inoculum. In estimating MICs the MBCs of mixtures of trimethoprim and sulphamethoxazole, the ratio of trimethoprim/sulphamethoxazole was 1:20, and generally increased in twofold concentrations.

ATTEMPTS AT TRANSFER OF ANTIBIOTIC RESISTANCE IN COLIFORMS

The method was essentially that of Moorhouse and McKay (1968) using mutants of *E. coli* K12 resistant to either rifampicin (100 µg/ml) or nalidixic acid (50 µg/ml) as recipient. Controls included a strain known to transfer tetracycline resistance to this and other recipients. Concentrations of trimethoprim used to detect possible transfer of resistance were 0.5, 1.0, and 5.0 µg/ml. For several organisms all three levels were used. In some experiments, other recipients were used; these were rifampicin-resistant mutants of strains recently isolated from different sources.

ATTEMPTS TO DEMONSTRATE LOSS OF TRIMETHOPRIM RESISTANCE IN *S. AUREUS* AND MUTAGENESIS OF CULTURES

Cultures were grown overnight at 42°C, diluted, and inoculated into nutrient agar plates to give discrete

colonies. These were then replicated onto nutrient agar containing 5% (v/v) lysed blood and 5 µg trimethoprim per ml. The method of mutagenesis of cultures of *S. aureus* and *E. coli* J53 (Su, Tp) has been previously described (Lacey, 1971).

TRANSDUCTION OF TRIMETHOPRIM

RESISTANCE IN *S. AUREUS* STRAIN NO. TPR-A
Cultures of the trimethoprim-resistant culture no. TPR-A were induced with mitomycin C. The resultant lysates were checked for sterility, and then irradiated with ultraviolet (UV) light for intervals of up to 5 min and added to the recipient strain 6936 (for further details see Lacey, 1971). The recipient was then plated on nutrient agar containing 5% (v/v) lysed blood and 5 µg trimethoprim/ml. After incubation, trimethoprim-resistant colonies were examined for sulphonamide resistance by replica plating onto media containing 500 µg sulphamethoxazole/ml.

DEVELOPMENT OF RESISTANCE TO TRIMETHOPRIM BY STRAINS OF *E. COLI*, *STREP. FAECALIS*, AND *S. AUREUS* IN VITRO

After assessing the numbers of bacteria by the Miles and Misra method, the test organisms, either from centrifuged broth cultures or directly from plate cultures, were inoculated onto DST agar containing 5% lysed blood and various levels of trimethoprim. The inoculated plates were incubated for 48 hours at 37°C for *E. coli* and *S. aureus* and 96 hours at 37°C for *Strep. faecalis*.

Trimethoprim lactate and sulphamethoxazole were obtained from Burroughs Wellcome Ltd, Beckenham, Kent.

Results

TRIMETHOPRIM RESISTANCE IN COLIFORMS

From November 1971 to October 1972, a total of 711 strains of coliforms from hospital and non-hospital patients were tested for resistance to trimethoprim by the disk method. All coliform strains were tested during three one-month periods; at other times strains were tested when it was clinically relevant. Of these strains, 16 (2.3%) were resistant to between 2 and 100 µg trimethoprim/ml. Further identification of these organisms showed that three were atypical *Klebsiella*, four *Enterobacter* species, and nine *Escherichia*. Since no property investigated was associated with a particular species, this group of organisms is referred to as coliforms. A similar incidence (2.5%) was found in the strains isolated earlier in 1971 (Lacey *et al.*, 1972). Of the 16 strains isolated since November 1971, 13 were also resistant to sulphamethoxazole (MIC > 4000 µg/ml). The

sensitivity to cotrimoxazole varied from an MIC of 0.2 µg trimethoprim and 4 µg sulphamethoxazole/ml to >25.6 µg trimethoprim and 512 µg sulphamethoxazole/ml. Most strains had MICs near the higher figures.

TRANSFERABILITY OF TRIMETHOPRIM RESISTANCE

Each trimethoprim-resistant isolate was tested for its ability to transfer this resistance to *E. coli* K12. Six of the isolates were also tested for transfer of resistance to 13 other coliforms. No transfer occurred. These findings are similar to those of Datta and Hedges (1972) who found that transfer of trimethoprim resistance only occurred when the MIC of trimethoprim was >1000 µg/ml.

IS COTRIMOXAZOLE BACTERICIDAL TO COLIFORMS AND *S. AUREUS*?

The correlation of the MIC with the MBC of cotrimoxazole was investigated in 30 coliforms and 24 strains of *S. aureus*. Although there was a close association of the MIC and MBC in most of the strains of *S. aureus*, a very poor association between these existed for coliforms. The presence of sulphamide resistance in the latter had little effect on the MBC (table I). In many coliform strains the MBC was >500 times that of their minimum inhibitory concentration. Such a disparity between the MIC and MBC indicates that cotrimoxazole cannot be considered bactericidal, at least under these conditions. Ratios of trimethoprim to sulphamethoxa-

zole other than 1:20 did not produce better cell destruction.

MUTATION TO TRIMETHOPRIM RESISTANCE IN VITRO IN STRAINS OF *E. COLI*, *S. AUREUS*, AND *STREP. FAECALIS*

Darrell, Garrod, and Waterworth (1968) and Bushby (1969) have found that serial transfer of bacteria on media containing trimethoprim can cause the appearance of organisms resistant to high levels of trimethoprim. However, there is no evidence that bacteria acquire resistance by such a process under natural conditions. We have found that repeated subculture of strains of *S. aureus* and *E. coli* in the presence of trimethoprim yields colonies, that although trimethoprim-resistant, have much reduced growth rates and some other defects eg, loss of coagulase production in *S. aureus*. But bacteria develop high-level resistance to streptomycin after a short exposure under natural conditions (eg, Garrod, 1950). Cultures were therefore tested for their ability to give rise to one-step mutants resistant to high levels of trimethoprim. Strains of sulphamide-sensitive and resistant *E. coli* and *S. aureus* were plated on media containing levels of trimethoprim 10, 40, 100, and 250 times that of the corresponding MIC of the strain. Results of these experiments are summarized in table II. Although some colonies grew on each medium, they were slow growing after subculture to antibiotic-free agar, and unstable with only a slight increase in resistance. But from strains of *Strep. faecalis*, mutants resistant

Organism	Resistance to Sulphamethoxazole (No. of Isolates)	Range of MIC (fig.) of Trimethoprim (T) and Sulphamethoxazole (S) (No. of Isolates)	MBC (fig.) Levels of Trimethoprim (T) and Sulphamethoxazole (S) (No. of Isolates)
Coliforms	1 Sensitive (MIC > 32 µg/ml) (9)	0.025T+0.50S-0.1T+2.0S	(9) 0.8T + 16S (3) >12.8T + 256S (6)
	<i>S. aureus</i> (3)	0.025T+0.50S-0.1T+2.0S	(3) 0.8T + 16S (3)
Coliforms	2 Moderately resistant (MIC 64 µg/ml-256 µg/ml) (4)	0.1T+2.0S-0.4T+8.0S	(4) 0.8T + 16S (1) 3.2T + 64S (1) 12.8T + 256S (2)
	<i>S. aureus</i> (21)	0.1T+2.0S-0.4T+8.0S	(21) 0.8T + 16S (8) 1.6T + 32S (5) 3.2T + 64S (6) 6.4T + 128S (1) 25.6 + 512S (1)
Coliforms	3 Highly resistant (MIC 512 µg/ml- > 4000 µg/ml) (17)	0.1T+2.0S-0.4T+8.0S	(15) 0.1T + 2.0S (1) 0.8T + 16S (1)
		1.6T+32S	(2) 1.6T + 32S (1) 3.2T + 64S (1) >12.8T + 256S (13)

Table I Minimum inhibitory concentration (MIC) and minimum bactericidal concentration (MBC) to cotrimoxazole of 30 coliforms and 24 *Staph. aureus* strains sensitive to trimethoprim but of variable resistance to sulphamides

Organism	No. of Strains Tested	MIC of Sulphamethoxazole ($\mu\text{g/ml}$)	MIC of Trimethoprim ($\mu\text{g/ml}$)	Proportion of Inoculum Producing Colonies	No. of Colonies Examined Further	Properties of Colonies	
						Growth Rate	MIC Compared to Wild Strain (number)
<i>E. coli</i>	12	< 8	<0.4	10^{-10} - $<10^{-12}$	34	SC, U ¹	$\times 1$ - $\times 4$
	2	> 256	<0.4	10^{-10} - $<10^{-12}$	21	SC, U	$\times 1$ - $\times 4$
<i>S. aureus</i>	2	< 8	<0.4	10^{-10} - $<10^{-12}$	16	SC, U	$\times 1$ - $\times 2$
	2	> 256	<0.4	10^{-11} - $<10^{-12}$	14	SC, U	$\times 1$ - $\times 4$
<i>Strep. faecalis</i>	10	> 4000	<0.5	10^{-5} - $<10^{-9}$	55	Normal growth	$\times 1$ - 4 (31) $\times 4$ - 16 (11) $\times 500$ (13)

Table II Frequency of mutation to trimethoprim resistance after a single plating on medium containing levels of trimethoprim 10-250 times that of the MIC for each strain

¹SC, U denoted small colony growth usually unstable with large, normal growing and fully sensitive colonies appearing at high frequency.

to high levels of trimethoprim were isolated from each medium. The growth rates and cultural characteristics of these mutants resembled those of the wild strains; the resistance was stable over several subcultures.

TRIMETHOPRIM RESISTANCE IN STAPHYLOCOCCUS AUREUS

Between June 1970 and October 1972, 309 multi-resistant strains of *S. aureus* from both hospital and non-hospital sources were tested by the disk method for resistance to trimethoprim. Three (1.0%) were resistant. Minimum inhibitory concentrations to the drug of two strains (TPR-A and TPR-B) was 6.4 $\mu\text{g/ml}$ and 25.6 $\mu\text{g/ml}$ for strain TPR-C. Each was resistant to sulphamethoxazole (MIC > 500 $\mu\text{g/ml}$ and cotrimoxazole (MIC 3.2 μg trimethoprim/ml and 64 μg sulphamethoxazole/ml) for strain nos. TPR-A and TPR-B; that for TPR-C was twice these levels). Each isolate was also resistant to penicillin (penicillinase), erythromycin, neomycin, streptomycin, and tetracycline, but sensitive to methicillin, fusidic acid, and gentamicin. The phage patterns were 84 in two and 84/85 in one. In properties, they resemble closely trimethoprim-resistant staphylococci isolated from other hospitals (Nakhla, 1972).

DOES ONE MECHANISM CONFER RESISTANCE TO BOTH TRIMETHOPRIM AND SULPHAMETHOXAZOLE?

Since both trimethoprim and sulphamethoxazole act on the same biosynthetic pathway (Hitchings, 1969), it is theoretically possible that a single biochemical mechanism can produce resistance to both drugs simultaneously, in which case the same gene(s) will determine both resistances. We have found that sulphonamide resistance in some coliforms may be transferred to *E. coli* K12 without trimethoprim resistance (Lacey *et al.*, 1972). In these organisms the genes determining each resistance

must be distinct. The question whether these genes were also different was investigated in a strain of *E. coli* harbouring an R factor conferring resistance to both drugs (J53 (Su, Tp)) and in two cultures of *S. aureus*. Cultures were treated with a mutagen (nitrosoguanidine) and examined for loss of trimethoprim resistance by replica plating. About 0.1% colonies of each strain became trimethoprim sensitive (table III). Although some of these were also sulphonamide sensitive, at least one colony fully sensitive to trimethoprim (MIC 0.1 $\mu\text{g/ml}$) and resistant to sulphamethoxazole (MIC > 500 $\mu\text{g/ml}$) was obtained from each strain. The mutants resembled the wild strains in other properties. Thus the gene(s) coding for resistance to the two drugs are distinct in the three strains. It is interesting to note that whenever trimethoprim resistance was lost, the MIC of the culture to cotrimoxazole became sensitive regardless of the level of resistance to sulphonamide (table III).

CHROMOSOMAL LOCUS OF GENES DETERMINING TRIMETHOPRIM RESISTANCE IN *S. AUREUS*

Because staphylococcal plasmids may be lost after growth at 43°C (eg, May, Houghton, and Perrett, 1964), strain nos. TPR-A, TPR-B, and TPR-C were grown overnight at 43° and then examined for loss of trimethoprim resistance by replica plating. No loss of resistance was found amongst 22000 colonies of strain no. TPR-A, 5000 of TPR-B, and 9500 of TPR-C. The stability of the resistance after growth at 43°C suggested a chromosomal locus for the genes in question. This was confirmed in strain no. TPR-A by transduction. Cultures of strain no. TPR-A were induced with mitomycin C and the resultant lysate irradiated with ultraviolet light for various intervals and then added to the recipient (strain 6936). The transduction frequency from the unirradiated lysate was low (2×10^{-9}) but after

Strain	MIC to Trimethoprim ($\mu\text{g/ml}$)	MIC to Sulphamethoxazole ($\mu\text{g/ml}$)	MIC to Co-Trimoxazole ($\mu\text{g/ml}$)
<i>E. coli</i> J53 (Su Tp) wild	>1000	>4000	6.4T + 128S
Mutant 1	0.4	128	0.1T + 2S
Mutant 2	0.4	128	0.1T + 2S
Mutant 3	0.1	>4000	0.025T + 0.5S
Mutant 4	0.4	2	0.025T + 0.5S
Mutant 5	0.4	2	0.025T + 0.5S
Mutant 6	0.4	>4000	0.1T + 2.0S
Mutant 7	0.4	64	0.025T + 0.5S
<i>S. aureus</i> TPR-A wild	25.6	> 512	6.4T + 128S
Mutant 1	1.6	128	0.4T + 8S
Mutant 2	0.4	128	0.2T + 4S
Mutant 3	0.4	4	0.2T + 4S
Mutant 4	0.1	256	0.2T + 4S
Mutant 5	1.6	> 512	0.8T + 16S
Mutant 6	3.2	128	1.6T + 32S
Mutant 7	0.1	32	0.1T + 2S
<i>S. aureus</i> TPR-B wild	6.4	>4000	3.2T + 64S
Mutant 1	0.1	>4000	0.1T + 2S

Table III Properties of trimethoprim-sensitive mutants of *E. coli* J53 (SU, TP) and two strains of *Staph. aureus* (TPR-A and TPR-B)

ultraviolet irradiation the frequency rose to a maximum of 3.2×10^{-6} (table IV). Such stimulation of the frequency is indicative of chromosomal genes (Arber, 1960; Asheshov, 1966).

Duration of Exposure of Lysate to Ultraviolet Light (Min)	Transduction Frequency ¹
0	2.0×10^{-9}
0.5	4.5×10^{-7}
1	3.2×10^{-6}
2	4.0×10^{-7}
5	6.8×10^{-8}

Table IV Increase in the frequency of transduction of trimethoprim resistance from *S. aureus* strain TPR-A by ultraviolet light

¹ Expressed as the number of colonies per phage particles (titred against strain 6936) in the unirradiated lysate.

About 1.2% of the trimethoprim-resistant transductants were resistant to $> 500 \mu\text{g}$ sulphamethoxazole; the remainder were as sensitive as the recipient. This suggests that although the genes determining resistance to sulphonamides and trimethoprim, although distinct, are linked. Similar conclusions may be drawn from the results following nitrosoguanidine treatment (table III) when several of the trimethoprim-sensitive mutants were sensitive to sulphamethoxazole (nitrosoguanidine causes high rates of comutation of adjacent genes: Guerola, Ingraham, and Cerdá-Olmedo, 1971).

Discussion

In trimethoprim-sensitive strains of *E. coli* and *S. aureus* moderate or even high levels of resistance

to sulphonamides have little influence on either the MIC or MBC of cotrimoxazole compared with those of sulphonamide-sensitive cultures. But resistance of an organism to moderate levels of trimethoprim (MIC about $6 \mu\text{g/ml}$) and high levels of sulphamethoxazole produces resistance to the combination (eg, MIC about $12 \mu\text{g}$ trimethoprim/ml and $240 \mu\text{g}$ sulphamethoxazole/ml). Since many coliforms and staphylococci are resistant to sulphonamides, the usefulness of cotrimoxazole against these organisms depends on the absence of trimethoprim resistance in sulphonamide-resistant cells.

There is no evidence for a single mechanism that confers resistance to both drugs simultaneously; even in a strain of *E. coli* in which an R factor gives high-level resistance to both drugs, the genes are distinct. Mutation to trimethoprim resistance in sulphonamide-resistant strains of *E. coli* and *S. aureus* is unlikely to be important under natural conditions. We have been unable to obtain one-step mutants resistant to moderate levels of trimethoprim which are not grossly defective. Although it is possible for resistance to develop by serial transfer in the presence of the drug, there is no evidence that this process occurs *in vivo*.

None of the coliform isolates could transfer trimethoprim resistance *in vitro*. Information on the 'epidemiology' of R factors under natural conditions is scanty, although much is known about their properties *in vitro*. The possibility that R factors determining high-level resistance to trimethoprim and sulphonamides will spread through the enterobacteriaceae as suggested by Datta and Hedges (1972) remains a threat, but has not happened yet, at least in Bristol. The resistance to trimethoprim in these

coliforms and staphylococci is probably intrinsic, present in a minority of strains. The incidence of these resistant isolates may be expected to increase—probably slowly—with the use of cotrimoxazole. The three trimethoprim-resistant strains of *S. aureus* described here are very similar in properties to those described by Nakhla (1972). All could have evolved from a single clone. The gene(s) determining trimethoprim resistance is chromosomal in strain no. TPR-A and probably also in strains TPR-B and TPR-C. If the genetic locus for trimethoprim resistance in all staphylococci is chromosomal, this might explain why trimethoprim resistance has not 'spread' to other staphylococci. Chromosomal genes are transferred between cultures of *S. aureus* at much lower frequencies than plasmid genes (Lacey, 1972a). Fortunately trimethoprim-resistant strains are sensitive to methicillin; methicillin resistance is unlikely to be transferred to other strains (Lacey, 1972b). Based on these findings *in vitro*, strains of *S. aureus* should be expected to retain their sensitivity to either cotrimoxazole or methicillin for some time to come.

In contrast, the value of cotrimoxazole in treating infections due to *Strep. faecalis* seems limited. Chattopadhyay (1972) showed that strains of *Strep. faecalis* developed resistance to cotrimoxazole during therapy with this drug; the reason for this would seem to be the frequent occurrence of mutants of *Strep. faecalis* which possess resistance to trimethoprim.

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