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## Cloning and Sequencing of an Ampicillin Resistance Gene of a Multiple Antibiotic Resistance Transposon from Salmonella typhi

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#### ABSTRACT

Four Salmonella typhi isolates (S2, S6, S8, and S9), each harbouring a large conjugative R plasmid that confers resistances to ampicillin (Ac'), chloramphenicol (Cm'), cotrimoxazole (Ctm'), streptomycin (Sm'), and tetracycline (Tc'), were investigated for antibiotic resistance transposons. Transposition experiments were performed with pUB307, a conjugative R plasmid conferring resistances to kanamycin (Km') and Tc', as the recipient replicon. Two major antibiotic resistant transconjugants were obtained, Ac'Ctm'Km'Sm'Tc' and Ac'Cm'Ctm'Km'Sm'Tc'. The results showed that the transposons isolated from the four S. typhi isolates were similar despite their different origins.

Results of the transposition, transformation, and secondary transposition experiments indicated the existence of two multiple antibiotic resistance transposons. One encoded Ac'Ctm'Sm' and the other, Cm' and probably also Tc'. On the R plasmid of S. typhi, another copy of the Ac'Ctm'Sm' transposon may be associated with the Cm' transposon and it could transpose independently of the Cm' transposon into pUB307. The insertion of the Cm' transposon on pUB307 was unstable and the Cm' transposon often transposed again onto the Escherichia coli chromosome by a conservative or 'cut and paste' mode of transposition.

An Ac'Ctm'Sm' transposon derived from S. ηphi S8 was inserted into a recombinant pUB307, pCL8, and it appeared to be identical to Tn5410, previously detected in S8 by Kadambeswaran (1993). In a shotgun cloning experiment, DNA fragments of Sall-digested pCL8 were cloned into plasmid vector pKan in E. coli DH5α. Recombinant pKan that carried 2.5 kb Ac' and 5.5 kb Sm' inserts were designated as pCLA25 and pCLS55, respectively. A Ctm' transformant that harboured a recombinant pUB307, pCLC600, was also obtained. pCLC600 resulted from recircularization of Sall-digested pCL8. DNA fragments of Sau3AI partially-digested pCLC600 were cloned into BamHI-digested pKan in E. coli DH5α. Five Ctm' transformants were obtained. One of them harboured a recombinant pKan, pCLC2, that carried a 4.3 kb insert.

Southern hybridization confirmed that the 2.5 kb Ac' Sall-insert in pCLA25 originated from the R plasmid pST8 of S. nyphi S8. The 2.5 kb Ac' insert was subcloned into a bacteriophage vector M13mp19 in E. coli DH5 $\alpha$ F'. An Ac' transfectant, harbouring M13mp19-Ac'(2.5), was obtained and its RF DNA was purified for nested deletions by using a nested deletion kit. After unidirectional nested deletions and transfection of the nested deleted M13mp19-Ac'(2.5) DNA into E. coli DH5 $\alpha$ F', a library comprising a series of progressively deleted clones was obtained. The smallest recombinant M13mp19 RF DNA isolated from this library that still conferred Ac' carried a 1.8 kb insert and was designated as M13mp19-Ac'(1.8). The 1.8 kb insert of M13mp19-Ac'(1.8) was subcloned into bacteriophage vector M13mp18 in E. coli DH5 $\alpha$ F'. An Ac' transfectant, harbouring M13mp18-Ac'(1.8), was obtained and its RF DNA was purified for construction of a second nested deletion library.

Deleted M13 derivatives from both nested deletion libraries were used to determine the nucleotide sequences of both strands by manual and automated dideoxyribonucleotide chain termination sequencing. Analysis of the 1814 nucleotides sequenced showed the presence of three distinct segments. One segment of 1444 nucleotides was identical to the right section of transposon Tn2, extending from the middle of the mpR resolvase gene to the right inverted repeat (IR) end, including the Ac' TEM-1 \(\beta\)-lactamase gene. The mpR resolvase gene was disrupted by a second segment comprising a 351 nucleotide sequence common to the IS26R element and its variants. The IS26-like segment extended from its left IR end to its unique Sall site. A third segment of 19 nucleotides flanked the IR end of the Tn2-like segment and the 19 nucleotides were identical to that of those flanking the sullI strA strB (Su'Sm') gene cluster on plasmid RSF1010.

Results from this study and circumstantial evidence from other workers suggest that Tn5410 was a Tn5086-like element from the Tn21 family and might contain three inserted copies of IS26-like elements. A genetic structure of Tn5410 was proposed.

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#### ABBREVIATIONS

Most of the abbreviations used are standard. However, attention is drawn to the following:

> Ac ampicillin

bp base pair

BPB: bromophenol blue

 $^{\circ}C$ degree Celcius

ccc covalently closed circular

chloramphenicol

Cm

conc. : concentration Ctm co-trimoxazole

DHFR: dihydrofolate reductase

DHPS: dihydropteroate synthetase

DNase: deoxyribonuclease

DNA: deoxyribonucleic acid

e.g. for example EtBr : ethidium bromide

Fig. figure

g gramme

Gen gentamicin

hr hour

i.e.

Inc Incompatibility

that is

kb kilobase pair

Km : kanamycin

Lac<sup>+</sup> lactose fermenting

Lac\* non-lactose fermenting

LB Luria-Bertani medium

M molar

milligramme mg

min minute ml millilitre

mM millimolar

microgramme μg

μl microlitre

MW molecular weight

ng nanogramme

No. number

Nx nalidixic acid

% percentage

p.s.i. pound per square inch

RF replicative form RNase: ribonuclease

revolutions per minute rpm

SDS sodium dodecyl sulphate

sec second

SSC

Sm streptomycin

Smz sulfamethoxazole

sodium chloride-sodium citrate Su sulfonamides

TBE : Tris-borate-EDTA

Tc tetracycline

TE Tris-borate

Tris tris(hydroxymethyl)methylamine

U unit

uv ultra-violet

volt

Vol. : volume

(v/v) : volume per volume (w/v) : weight per volume

> : greater than

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