SHORT PAPERS

The orientation of transfer of the plasmid RP4

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(Received 26 November 1980 and in revised form 30 March 1981)

The probably identical broad host range plasmids RP4, RP1 and RK2 (Datta *et al.* 1971; Grinsted *et al.* 1972; Olsen & Shipley, 1973; Meyer *et al.* 1975; Beringer, 1974; Towner & Vivian, 1976) have been extensively studied as vectors for *in vitro* recombination and mediators of conjugative transfer of DNA between species (Olsen & Gonzalez, 1974; Jacob *et al.* 1976; Dixon *et al.* 1976; Stepanov *et al.* 1976; Meyer *et al.* 1977; Nagahari *et al.* 1977). Studies of the conjugation system have led to the identification of transfer (*tra*) genes and have mapped the drug resistance determinants (Barth & Grinter, 1977; Grinsted *et al.* 1977; Thomas *et al.* 1979). We present here a deletion analysis which shows in which direction the plasmid is transferred during conjugation.

A short (5.7 kb) λ fragment (Szybalski & Szybalski, 1979; Daniels *et al.* 1980) containing *att* and *int* was inserted into the single *Eco*RI site of RP4 (Jacob & Grinter, 1975; Pastrana, 1976; Pastrana & Brammar, 1979). The recombinant plasmid can integrate into *att*_{λ} on the *Escherichia coli* chromosome to form a stable Hfr strain (Watson & Scaife, 1978). It transfers the chromosome in the orientation: O-*lac-leu-thr...trp.* We have recently reported (Watson & Scaife, 1980) that plasmid excision specifically depends on the *xis* function of λ , confirming that Hfr formation is mediated by *int*-directed, site-specific recombination and, by implication, uses *att* in the same orientation as the parent phage. Chromosome transfer by the Hfr thus allows us to establish the orientation of the λatt fragment relative to the chromosome. Here, we present deletion studies with an RP4 λatt derivative, pZD100 (Al-Doori and Scaife, in preparation) which establish the orientation of the phage fragment in the original plasmid.

The plasmid pZD100 carries $\lambda drif^{D}$ 18 inserted at *att* (Fig. 1*b*). This phage makes a dominant rifampicin-resistant (rif^{D}) RNA polymerase β subunit (Kirschbaum & Konrad, 1973) and a temperature-sensitive phage repressor (*c*1857) (Sussman & Jacob, 1962). Electron microscope studies on pZD100 DNA (date not shown) indicate that it occurs in several non-multimeric sizes. For this reason we have preferred to analyse deletion mutants of pZD100 which exist in a single form and were made as follows.

Bacteria carrying pZD100 cannot form colonies on rifampicin medium at 42 °C since the phage is induced and is either excised from the plasmid or interferes with its replication. However, rare mutants do grow on this medium. They have deletions extending through most of the prophage into the plasmid DNA (Fig. 1). Two such deletion plasmids, pZD23 and pZD44 have been analysed in detail (Plate 1*a* and *b*).

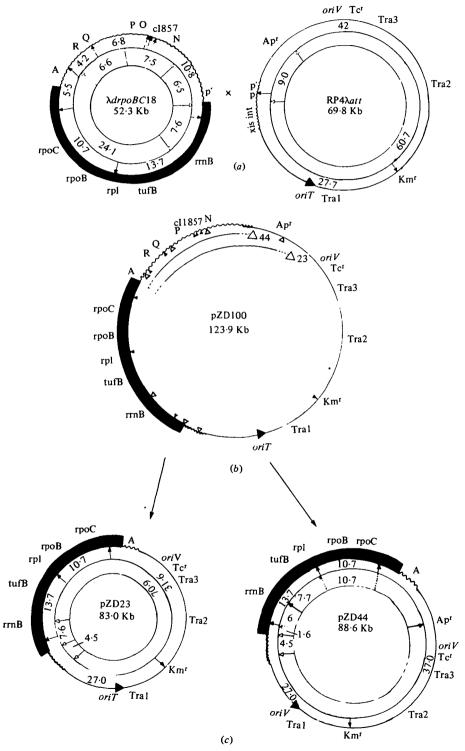


Fig. 1. For legend see opposite.

2. THE STRUCTURES OF pZD23 AND pZD44

Genetic tests show that both of these plasmids retain determinants for resistance to rifampicin (100 μ g/ml), tetracycline (10 μ g/ml) and kanamycin (25 μ g/ml). The plasmids differ in that pZD23 has lost and pZD44 retains Ap^r. Both plasmids have lost λ immunity (tested according to Miller, 1972) but still produce pill since they are still sensitive to PRR1 phage (Olsen & Shipley, 1973; Olsen & Thomas, 1973). However, only pZD23 is Tra⁺, a point which will be considered elsewhere (Al-Doori and Scaife, in preparation). Restriction patterns of the two, independently isolated plasmids can be simply interpreted as the result of single deletions arising in the parent plasmid.

Some of the DNA in pZD44 comes from $\lambda drif^{D}18$. It appears (Plate 1b) as the 10.7 and 13.7 kb HindIII fragments, the 7.6 kb BamHI fragment and the 10.7, 7.7, 5.2 and 1.6 kb fragments from HindIII/BamHI double digests. About half of the $\lambda drif^{D}18$ DNA present in pZD100 (data not shown) is absent from pZD23 and pZD44 (Plate 1a). The 10.8, 6.8, 5.5 and 4.2 kb HindIII fragments are missing from both plasmids. In confirmation of our genetic results, the restriction analysis shows that the BamHI site located in the Ap^r gene is absent from pZD23 but present in pZD44 (Plate 1a, track 2).

The restriction results can be simply combined to give two mutually consistent plasmid maps (Fig. 1c), which can be derived by single deletions from a common parent plasmid, pZD100 (Fig. 1b). The structure inferred for pZD100 is precisely that predicted for the product of insertion of $\lambda drif^{D}$ 18 into RP4 λatt (Fig. 1a). It also shows us that the phage fragment is inserted in the orientation...Km^R(xis int att P'OP) Ap^r...

It will be recalled that the phage fragment in the RP4 λatt Hfr strain is transferred in the order – xis int att P' – (Watson & Scaife, 1978). This fact, together with the finding reported here leads us to conclude that RP4 transfers its markers anticlockwise (Fig. 1a). The origin of transfer, oriT, has recently been located on the RK2 map (Thomas et al. 1979). Assuming structural identity of these plasmids we conclude that during mating the plasmid transfer genes enter the recipient last. This property is also shown by the sex factor, F, of E. coli (Willetts, 1972; Guyer & Clark, 1977).

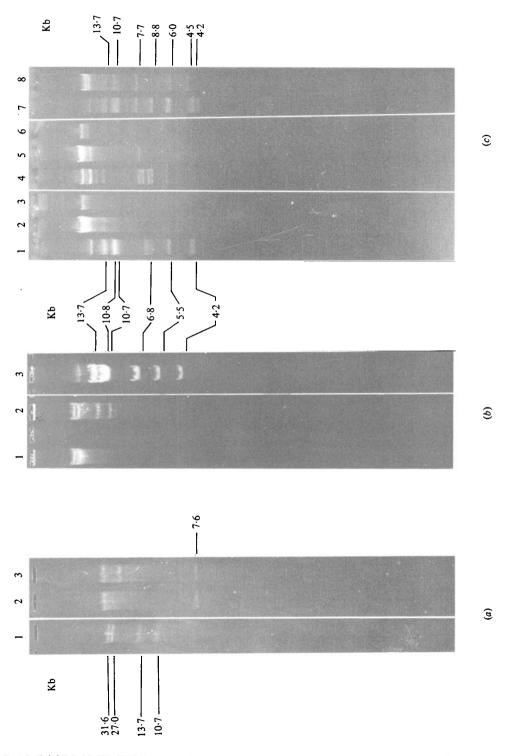
We thank Dr D. R. Helinski for information about oriT, the Medical Research Council and the Iraqi Ministry of Higher Education and Scientific Research for financial support and Dr N. S. Willetts for helpful comments on the manuscript. We thank Drs T. Linn and A. Newman for gifts of $\lambda drif^{D}18$ DNA.

Fig. 1. The lysogenisation of RP4 att with $\lambda drif^{D}18$. (a) Postulated mechanism confirmed by our results. (b) The inferred structure of the temperature-sensitive plasmids pZD100. $\Delta 23$ and $\Delta 44$ represent the two deletions in pZD23 and pZD44 respectively. (c) The two temperature-resistant plasmids pZD23 and pZD44, showing the *Hind*III (\triangle , outer circle) and *Bam*HI (\triangle , inner circle) sites established in this study. The thick, thin and wavy lines represent *E. coli*, RP4 and λ DNA respectively. *oriT* is sited according to its position in RK2 (Thomas *et al.* 1979).

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Z. AL-DOORI, M. WATSON AND J. SCAIFE

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PLATE 1

Restriction analysis of pZD23 and pZD44. Plasmid digests were electrophoresed in tris-acetate buffer on 0.7% agarose gels containing ethidium bromide $(1.5 \,\mu g/ml)$ (McDonell et al. 1977). Unlabelled fragments of $\lambda drif^{D}18$ are from the contaminating DNA of helper phage λ . (a) pZD23 DNA digestion with (1) HindIII and (2) BamHI. Track 3 shows pZD44 digested with BamHI. (b) A HindIII digest of pZD44 (2) compared with HindIII digests of RP4 λatt (1) and $\lambda drif^{D}18$ (3) (c) HindIII, BamHI double digest of pZD44. Controls – HindIII: (1) $\lambda drif^{D}18$, (2) pZD44. BamHI: (3) RP4 λatt (4) $\lambda drif^{D}18$. Double digest: (6) RP4 λatt , (7) $\lambda drif^{D}18$, (8) pZD44.