# Role and Specificity of Plasmid RP4-Encoded DNA Primase in Bacterial Conjugation

## ANDREW MERRYWEATHER,<sup>1</sup> PETER T. BARTH,<sup>2</sup> AND BRIAN M. WILKINS<sup>1\*</sup>

Department of Genetics, University of Leicester, Leicester LEI 7RH,<sup>1</sup> and ICI Corporate Bioscience Group, The Heath, Runcorn, Cheshire WA7 4QE,<sup>2</sup> England

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The role of the DNA primase of IncP plasmids wasexamined with <sup>a</sup> derivative of RP4 containing Tn7 in the primase gene (pri). The mutant was defective in mediating bacterial conjugation, with the deficiency varying according to the bacterial strains used as donors and recipients. Complementation tests involving recombinant plasmids carrying cloned fragments of RP4 indicated that the primase acts to promote some event in the recipient cell after DNA transfer and that this requirement can be satisfied by plasmid primase made in the donor cell. It is proposed that the enzyme or its products or both are transmitted to the recipient cell during conjugation, and the role of the enzyme in the conjugative processing of RP4 is discussed. Specificity of plasmid primases was assessed with derivatives of RP4 and the IncI<sub>1</sub> plasmid ColIb-P9, which is known to encode a DNA primase active in conjugation. When supplied in the donor cell, neither of the primases encoded by these plasmids substituted effectively in the nonhomologous conjugation system. Since Collb primase provided in the recipient cell acted weakly on transferred RP4 DNA, it is suggested that the specificity of these enzymes reflects their inability to be transmitted via the conjugation apparatus of the nonhomologous plasmid.

Plasmids belonging to the IncP-1 group, such as RP4, are remarkable for their ability to promote conjugation between, and to be maintained within, a broad range of gram-negative bacterial species. One of the best-characterized products specified by RP4 is DNA primase, an enzyme that synthesizes oligoribonucleotides for starting DNA chain growth. The *pri* locus has been mapped, and it is thought to be the penultimate gene in a group of coordinately transcribed genes situated at the end of the Tral region of the plasmid (17, 19, 20). The gene encodes two immunologically related polypeptides of 118 and 80 kilodaltons (kDa), and primase activity is specified by a domain located towards the Cterminal region of both products (17, 20). In vitro, the enzyme can generate RNA primers on <sup>a</sup> variety of singlestranded bacteriophage DNA templates, and it has the potential to substitute in vivo for the primase of Escherichia coli (dnaG protein) in discontinuous replication of the bacterial chromosome (17, 18, 20).

The physiological role of the primase specified by IncP-1 plasmids is unclear. It was inferred from genetic studies with RP4 pri mutants that it optimizes the priming of vegetative replication of the plasmid and that it promotes efficient transconjugant formation in 5 of 11 gram-negative bacterial species tested as recipient strains in conjugation (17). Of these, Salmonella typhimurium recipients, but not E. coli recipients, were found to be deficient in inheriting Priplasmids. A similar, recipient-specific variability was observed when primase-negative mutants of another IncP-1 plasmid, R18, were tested for conjugation proficiency (16). Thus, the primase apparently contributes to the broad host range of IncP-1 plasmids by facilitating the conjugative metabolism of these DNA elements.

The purpose of our studies was to investigate further the role of RP4-specified primase in bacterial conjugation and to determine whether the enzyme acts in the donor or the recipient cell. A genetic approach was taken, testing for complementation of a primase-negative RP4 mutant by re-

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combinant plasmids carrying <sup>a</sup> cloned pri gene. We also used  $Incl<sub>1</sub>$  plasmids to assess the specificity of genetically different plasmid primases in bacterial conjugation. The primase encoded by the sog gene of  $Incl<sub>1</sub>$  plasmids such as ColIb-P9 differs from the RP4 enzyme in molecular mass and antigenic specificity, but it is known to act in bacterial conjugation and can generate primers on a similar diversity of phage and bacterial DNA templates as can the RP4 pri product  $(8, 17, 24)$ . In view of these similarities, we examined whether sog and pri DNA primases are functionally interchangeable in bacterial conjugation.

#### MATERIALS AND METHODS

Bacterial strains and cultural conditions. Bacterial strains and plasmids are described in Tables <sup>1</sup> and 2. Spontaneous nalidixic acid-resistant mutants of E. coli C600 and S. typhimurium SL329 were used as recipients. E. coli BW103 was isolated as a Thy<sup>+</sup> recombinant of E. coli BW85, constructed in a 30-min mating with KL16-99 donors  $(thy^+$ recAl  $[10]$ ) at 30°C. The recombinant was chosen for its sensitivity to UV light, and the recombination-defective phenotype was confirmed in test matings with HfrH. Plasmid-containing strains were grown in medium containing antibiotics to maintain the presence of the plasmid. Kanamycin (100  $\mu$ g ml<sup>-1</sup>) was used for cells harboring pLG261 or derivatives of RP4 and ColIb, tetracycline  $(7.5 \mu g \text{ ml}^{-1})$  was used for cells harboring pLG215, chloramphenicol (25  $\mu$ g  $ml^{-1}$ ) was used for cells harboring pLG260, and ampicillin (200  $\mu$ g ml<sup>-1</sup>) was used for cells harboring pJF107 and its derivative. The permissive temperature for BW86 was 30°C, and thymine-requiring strains were routinely grown in media containing thymine (20  $\mu$ g ml<sup>-1</sup>).

Construction of pLG260 and pLG261. The constructions of pLG260 and pLG261 are summarized in Fig. 1. Plasmids pBR328 and R300B were purified by the Triton X-100 method (15), and pRP26 was purified by a procedure involving lysozyme and Brij <sup>58</sup> (23). DNAs were digested with appropriate restriction endonucleases (Bethesda Research Laboratories, Inc.) by the supplier's instructions, treated

<sup>\*</sup> Corresponding author.

with phenol, and precipitated with ethanol. pRP26 fragments and cleaved vector molecules were mixed in a ratio of 1:5 and ligated with T4 DNA ligase (New England BioLabs, Inc.). The resulting DNA was used to transform competent cells of BW86, and transformants were selected at 30°C on nutrient agar containing chloramphenicol for pLG260 and on nutrient agar containing kanamycin for pLG261. After replica plating, replicas were incubated at 40°C to identify recombinant plasmids that suppressed the *dnaG3*(Ts) mutation in BW86. The rapid method of Birnboim and Doly (5) was used for all other plasmid preparations required for mapping and strain construction.

Filter matings. Conjugation experiments involving pRP1 and pRP2-containing donor cells were carried out on solid medium. Cultures of donor and recipient strains, shaken overnight in nutrient broth, were diluted to an  $A_{600}$  of 0.35  $(-2.5 \times 10^8 \text{ cells m}^{-1})$  and mixed in a 1:5 (vol/vol) ratio. A 1-ml amount of mixture was filtered onto cellulose nitrate disks (Whatman; 25-mm diameter,  $0.45-\mu m$  pore size), which were placed on prewarmed nutrient agar plates and incubated at 37°C, unless stated otherwise. After 3 h, bacteria were washed from the filters, and the yield of transconjugants was determined by plating on nutrient agar containing nalidixic acid (50  $\mu$ g ml<sup>-1</sup>) and either kanamycin or tetracycline. Each result is the average value from at least three experiments.

Conjugative DNA synthesis. Details of conjugative DNA synthesis have been described previously (7). Cells were grown in a salts-glucose-Casamino Acids (Difco Laboratories) (SGC) medium. Recipients were irradiated with UV light (400 Jm<sup>-2</sup>), and matings were carried out at 41<sup>o</sup>C in the presence of rifampin (100  $\mu$ g ml<sup>-1</sup>), 2-deoxyguanosine (200  $\mu$ g ml<sup>-1</sup>), and [2-<sup>14</sup>C]thymine (63 mCi mmol<sup>-1</sup>, 1  $\mu$ g ml<sup>-1</sup>).

Colony formation by BW86. Suitable dilutions of overnight cultures, shaken in nutrient broth at 30°C, were plated on prewarmed nutrient agar plates and incubated at 30°C or 40°C. All media incubated at 30°C contained appropriate antibiotics.

DNA synthesis at 41°C. Overnight cultures were diluted in SGC medium supplemented with thymine  $(3 \mu g \text{ ml}^{-1})$  plus 2-deoxyguanosine (200  $\mu$ g ml<sup>-1</sup>) and were grown at 30°C for about three mass doublings to an  $A_{450}$  of ~0.35. At 5 min after a shift to 41°C, 1.5 ml of culture was added to an equal volume of the same medium containing  $[methyl-<sup>3</sup>H]thymine$ to yield 320 mCi mmol<sup>-1</sup>. The  $A_{450}$  was determined immediately and after 90 min at 41°C. Radioactivity incorporated into acid-precipitable material in this period is given as counts per minute  $\times$  10<sup>-3</sup> per unit increase of  $A_{450}$ .

TABLE 1. Bacterial strains used

Strain and strain no.	Genotype <sup>a</sup>	Source or reference	
$E.$ coli $K-12$			
W3110T <sup>-1</sup>	thyA36 deoC	17	
C600	thr-1 leu-6 thi-1 supE44 lacYl fhuA21	17	
<b>BW85</b>	leu thyA deoB rpsL cir	6	
<b>BW86</b>	leu thyA deoB rpsL cir $dnaG3$ $\Delta$ (chlA-uvrB)	ጸ	
<b>BW96</b>	leu deoA deoC tdk rpsL cir rpoB	8	
<b>BW103</b>	leu deoB rpsL cir recAl	This work	
S. typhimurium LT2			
<b>SL329</b>	trp	17	

<sup>a</sup> Genetic symbols are defined in the linkage map of Bachmann (2).

TABLE 2. Bacterial plasmids used

Plasmid	Description <sup>a</sup>	Source or reference
pLG215	pBR325 $\Omega(sog + EcoRI 8.0$ kb) Ap <sup>r</sup> Tc <sup>r</sup>	24
pLG221	Collb-P9drd-1 cib::Tn5 IncI, Km <sup>r</sup>	8
pLG250	pLG221 carrying sog-217	8
pLG260	$pBR328 \Omega$ (pri <sup>+</sup> HindIII 14.5 kb) $Apr$ Cm <sup>r</sup>	This work
pLG261	R300B $\Omega(pri^+ \; PstI \; 12.6 \; kb) \; IncQ$ Km <sup>r</sup>	This work
pLG270	Collb-P9drd-2 tra::Tn5 IncI <sub>1</sub> Km <sup>r</sup>	See text
pRP1	RP4 pri::Tn7 IncP-1 Ap <sup>r</sup> Tc <sup>r</sup> Km <sup>r</sup> Tp' Sm' Sp'	3
pRP2	RP4::Tn7 Tra+ IncP-1 Apr Tcr Km <sup>r</sup> Tp <sup>r</sup> Sm <sup>r</sup> Sp <sup>r</sup>	3
pRP26	RP4 tra::Tn7 IncP-1 Apr Tcr Kmr Tp' Sm' Sp'	3
pJF107	pBR325 $\Omega(pri^+$ HindIII 14.5 kb $\Delta$ 10.6 kb) Ap <sup>r</sup> Cm <sup>r</sup>	20
$pJF107\Delta32$	pBR325 $\Omega(pri^+ HindIII$ 14.5 kb $\Delta$ ~13.7 kb) Ap <sup>r</sup>	20

<sup>a</sup> Abbreviations for drug resistance phenotypes are: Ap, ampicillin; Tc, tetracycline; Km, kanamycin; Cm, chloramphenicol; Tp, trimethoprim; Sm, streptomycin; and Sp, spectinomycin.

#### RESULTS

Experimental plan and bacterial plasmids. We assessed the requirement for primase in conjugation from the yield of E. coli K-12 and S. typhimurium transconjugants generated in matings mediated by a primase-negative mutant of RP4. The mutant plasmid used was pRP1, which carries a Tn7 insertion preventing the synthesis of an active primase and of the 118-kDa pri polypeptide (17). The transposon insertion is about 1.5 kilobases (kb) from the HindIII site in RP4 (Fig. 1) and therefore is located in the <sup>3</sup>' region of pri, as shown on



FIG. 1. Construction of  $pri$ <sup>+</sup> recombinant plasmids pLG260 and pLG261. The central linear map shows the pri region of pRP26 (RP4::Tn7). The map coordinates of RP4, the location of pri, and the putative transcript (line with arrowhead) are taken from the data of Lanka et al. (19, 20). Details of the Tn7 region (thick line) are taken from the data of Gosti-Testu et al. (11). Plasmid pRP1 is an RP4 derivative containing a Tn7 insertion at about coordinate 40.5 (17, 19).

TABLE 3. Plasmid-mediated suppression of the E. coli dnaG3 mutation

Plasmid in <b>BW86</b>	Colony formation <sup>a</sup>	<b>DNA</b> synthesis <sup>b</sup>
pLG215	$1.5 \times 10^{-4}$	116
pLG221	$2.2 \times 10^{-1}$	233
pLG250	$< 10^{-7}$	10
pLG260	$5.9 \times 10^{-1}$	208
pLG261	$1.3 \times 10^{-2}$	137
pLG270	1.1	261
pRP1	${<}10^{-7}$	12
pRP2	$2.2 \times 10^{-6}$	82
pRP26	$3.0 \times 10^{-5}$	100
pJF107	$3.7 \times 10^{-3}$	96
$pJF107\Delta32$	$5.1 \times 10^{-5}$	85

<sup>a</sup> Colony formation at 40°C relative to that at 30°C. Value of  $\leq 10^{-7}$  for **BW86** 

 $<sup>b</sup>$  Value of 10 for BW86. Synthesis was at 41°C.</sup>

the 60-kb map of RP4 (19). The control plasmid was pRP2; this RP4 derivative has Tn7 inserted near the tetracycline resistance  $(Tc^r)$  gene in a region that is devoid of known transfer or maintenance genes (3).

Complementation studies involving  $pRP1$  and  $Pri<sup>+</sup>$  recombinant plasmids were undertaken to determine whether the primase acts to promote events in the donor or in the recipient cell, and Pri<sup>+</sup> recombinants based on R300B and pBR328 were constructed to this end. R300B is an 8.7-kb plasmid of the IncQ group (13), which is maintained in a broad range of bacterial species, including S. typhimurium, and it was used as the vector for the establishment of the cloned pri gene in recipient strains. However, R300B is an unsuitable vector for studying the effect of cloned genes in donor cells, because it is mobilized by RP4 to recipient cells during conjugation (4). Thus, the nonmobilizable plasmid  $pBR328$  (22) was used in the construction of a Pri<sup>+2</sup> recombinant for the necessary complementation tests in donor cells.

The source of DNA for cloning the *pri* gene was pRP26. This plasmid is an RP4 derivative containing Tn7 near the putative promoter of the transcriptional unit containing pri, and the mutant has merit in that it specifies overproduction of primase (17) and the inserted transposon provides useful restriction sites for the cloning of the adjacent DNA. The constructions of pLG260 and pLG261 are summarized in Fig. 1. The pri gene in each recombinant is presumably expressed from the normal RP4 promoter, but the promoter of the sulfonamide resistance-streptomycin resistance (Sur-Sm<sup>r</sup>) operon in the R300B vector (4) may also contribute to the expression of pri in pLG261. The ability of these plasmids to suppress the temperature-sensitive dnaG3 mutation in BW86, as judged by rescue of colony formation and DNA synthesis at high temperature, is shown in Table 3. These parameters provide a good estimate of plasmid primase activity (20, 24). Although pLG260 apparently specifies more primase than does pLG261, both recombinants directed the synthesis of greater amounts of primase than did pRP2, the control plasmid used in the conjugation experiments. Presumably, these differences in primase activity reflect the different copy numbers of the plasmids.

Requirement for plasmid primase in the recipient cell. Initial experiments involved W3110T<sup>-</sup> donor cells mated with E. coli C600 and S. typhimurium SL329 recipients, since these strains were used previously (17). About  $\bar{5} \times 10^8$  $pRP2$  transconjugants  $ml^{-1}$  were obtained from the E. coli matings, but the yield was approximately four orders of magnitude lower when S. typhimurium was the recipient species. The yields of pRP1 transconjugants relative to pRP2  $transconjugants$  in matings of  $W3110T^-$  donors and recipient strains C600, C600(pLG261), SL329, SL329(pLG261), and SL329(R300B) were 0.59, 0.84, 0.01, 0.37, and 0.03, respectively. Thus, formation of pRP1 transconjugants was slightly and reproducibly reduced relative to that of pRP2 transconjugants when C600 was the recipient strain, but a 100-fold reduction was detected in the S. typhimurium strain. These deficiencies were mostly overcome when the recipients contained the Pri<sup>+</sup> recombinant pLG261.

These results confirm previous findings (17) that the deficiency of pRP1 transconjugants depends on the strain used as the recipient. Furthermore, the complementation of pRP1 by pLG261 implies that the RP4 mutant fails to specify one or more gene products that promote some event in the recipient cell and that plasmid primase is one such product.

Contribution of plasmid primase in the donor cell. The apparent requirement for plasmid primase in the recipient cell might normally be satisfied by enzyme that is either supplied by the donor or synthesized in the recipient after expression of the transferred DNA. To distinguish between these possibilities, we determined whether the  $Pr<sup>+</sup>$  recombinant pLG260 can complement pRP1 in the donor cell. This pBR328-based recombinant was not mobilized by pRP1 or pRP2 (data not shown). W3110T- proved to be an unsuitable donor strain for these experiments because it was transformed inefficiently by pLG260 and because most transformants were found to contain deletion mutants of the plasmid. BW86 was used instead, since it allowed the presence of a  $Pri<sup>+</sup> plasmid to be monitored by the rapid method of  $dn aG$$ suppression.

In contrast to results obtained with W3110T<sup>-</sup>, BW86 donors of pRP1 were almost equally deficient (>94%) in contributing to transconjugants in E. coli C600 recipients as they were in S. typhimurium SL329 recipients (Table 4). The deficiency in BW86  $\times$  C600 matings was not due to mating at 30°C, the permissive temperature for BW86, as indicated by control matings with W3110T- donors at 30°C and 37°C (data not shown). Furthermore, the deficiency cannot be attributed to the *dnaG3* mutation in BW86 because a  $dna<sup>+</sup>$  donor strain (BW85) of the same pedigree gave similar results. The yield of pRP1 transconjugants generated by donors of a third K-12 pedigree (C600) was intermediate between yields obtained with W3110T<sup>-</sup> and BW86 strains, confirming that the genotype of the E. coli K-12 donor cells influences formation of Pri- plasmid transconjugants in K-12 recipients. Finally, Table 4 shows that the presence of the  $Pri<sup>+</sup>$  recombinant

BW86, BW85, and C600 donor cells

TABLE 4. Relative yield of pRP1 transconjugants generated by BW86, BW85, and C600 donor cells				
<b>Bacterial strains</b>		Temp	Yield of pRP1	
Donor <sup>a</sup>	Recipient	(C)	transconjugants relative to pRP2 matings <sup>b</sup>	
<b>BW86</b>	C600	30	0.016	
<b>BW86</b>	SL329	30	0.06	
<b>BW85</b>	C600	30	0.008	
C600	C600	37	0.16	
C600	<b>SL329</b>	37	0.032	
<b>BW86</b>	C600(pLG261)	30	0.28	
<b>BW86</b>	SL329(pLG261)	30	0.58	

<sup>a</sup> Each donor strain contained either pRP1 or pRP2 as the conjugative plasmid.

 $<sup>b</sup>$  Yield of pRP1 transconjugants expressed as a fraction of the yield of pRP2</sup> transconjugants.

plasmid (pLG261) in the recipient strains enhanced the yield of pRP1 transconjugants generated by BW86 donor cells. This indicates that BW86(pRP1) donor cells resemble W3110T<sup>-</sup> donors (see the previous section) in failing to specify some product that acts in the recipient cell.

BW103 containing the nonmobilizable,  $Pri<sup>+</sup>$  recombinant plasmid pLG260 and either pRP1 or pRP2 was used as the donor strain to test whether the requirement for plasmid primase in pRP1 transconjugant formation can be satisfied by enzyme supplied by the donor. BW103 is homogenic with BW86 and was chosen for its Rec<sup>-</sup> phenotype to limit recombination in the donor strain between the cloned RP4 DNA and the homologous region in pRP1. The presence of pLG260 in the donors rescued the deficiency of pRP1 transconjugants in E. coli and S. typhimurium, as shown in Table 5. This complementation was efficient, since little or no further rescue was detected if the recipients also contained functional primase specified by a  $Pr<sup>+</sup>$  recombinant plasmid. To ensure that primase, rather than the product of some other RP4 gene product, was responsible, the complementation tests were repeated with  $Pr<sup>+</sup>$  recombinant plasmids pJF107 and pJF107A32, constructed by Lanka et al. (20). pJF107 $\Delta$ 32 contains only the *pri* gene, whereas pJF107 carries this gene and the promoter-distal DNA up to the HindIII site at coordinate 39 in Fig. 1. These two recombinants complemented pRP1 in the donor with moderate but approximately equal efficiency, implying that the conjugation deficiency of pRP1 is not due to a polar effect of the transposon insertion on the expression of the gene downstream of pri. However, neither pJF plasmid complemented pRP1 with the efficiency of pLG260. Since pLG260 apparently specifies more primase activity than do the pJF plasmids (Table 3), presumably because of the fact that the pri genes in the two sets of recombinants are expressed from different promoters, the different complementation efficiencies may reflect the amount of primase in the donor cells.

It is interesting that neither pLG260 nor the pJF plasmids complemented pRP1 fully, despite the indication that they specify at least as much plasmid primase as the control RP4 derivative, pRP2 (Table 3) (20). Since the Tn7 insertion in pRP1 maps at the <sup>3</sup>' end of pri, the mutant may encode truncated polypeptides. Competition between such inactive pri polypeptides and proteins specified in trans may explain the inability of the recombinants to fully complement pRP1.

Complementation of pRP1 by plasmids carrying the sog gene of ColIb. Complementation of pRP1 was detected when the recipient strain harbored a  $\text{Sog}^+$  Incl<sub>1</sub> plasmid (pLG221),

TABLE 5. Complementation of pRP1 in the donor strains by nonmobilizable Pri<sup>+</sup> recombinant plasmids

<b>Bacterial strains</b>		Yield of pRP1	
Donor <sup>a</sup>	Recipient	transconjugants relative to pRP2 matings <sup>b</sup>	
<b>BW103</b>	C600	0.002	
BW103(pBR328)	C600	0.004	
BW103(pLG260)	C600	0.63	
<b>BW103(pLG260)</b>	C600(pLG260)	0.85	
BW103(pJF107)	C600	0.15	
BW103(pJF107432)	C600	0.19	
<b>BW103</b>	<b>SL329</b>	0.067	
BW103(pLG260)	<b>SL329</b>	1.6	
BW103(pLG260)	SL329(pLG261)	0.71	

<sup>a</sup> Each donor also carried either pRP1 or pRP2.

bYield of pRP1 transconjugants as a fraction of the yield of pRP2 transconjugants.





<sup>a</sup> Each donor also carried either pRP1 or pRP2.

<sup>b</sup> Yield of pRP1 transconjugants as a fraction of the yield of pRP2 transconjugants.

with the extent of the rescue depending on the strains used (Table 6). This complementation is attributed to the activity of sog primase, because no such effect was observed when the recipient cells carried a primase-negative mutant of pLG221, namely, pLG250. Interestingly, pRP1 was not complemented by a small recombinant plasmid (pLG215) containing a cloned sog gene, suggesting that other ColIbspecified products are required to allow sog primase to act on transferred RP4 DNA.

In contrast, no complementation of pRPl was observed when the donor strain harbored a  $Sog<sup>+</sup> Incl<sub>1</sub> plasmid,$ pLG270. This is a mutant of ColIbdrd-2 that contains a normal sog primase gene (Table 3), but it is more than 105-fold deficient in conjugation as a result of a TnS insertion (C. Rees and B. M. Wilkins, manuscript in preparation). The plasmid was chosen here for its conjugation deficiency to ensure that sog primase was only synthesized in the donor cell. Furthermore, it was used to ensure that, subject to effects of the TnS insertion, other ColIb-specified products that may cooperate with sog primase would be present in the donor cell.

Specificity of RP4 primase in conjugation. It was shown previously that sog primase of ColIb functions to promote the synthesis of DNA complementary to the transferred  $Incl<sub>1</sub> plasmid strand (8), and we examined whether RP4$ primase can substitute for the *sog* product in this process. The method involved mating donors of a ColIb plasmid with rifampin-treated, heavily UV-irradiated BW86 recipients at 41°C. In response to ColIb transfer, such recipients incorporate exogenously supplied, labeled thymine into plasmid DNA by a process dependent on sog primase, and this incorporation is readily detectable if uptake of label into DNA of donor cells is prevented by use of a tdk strain (7, 8).

Figure 2 shows  $[$ <sup>14</sup>C]thymine incorporation under such conditions when the donors harbored the control Sog+ plasmid pLG221, and it shows the deficiency when its primase-negative derivative pLG250 was used instead. Conjugative synthesis of pLG250 in recipient cells is complemented efficiently in this system by the presence of the pBR325-sog<sup>+</sup> recombinant, pLG215, in the donor strain (25), but to optimize detection of any effect caused by the pBR328-pri<sup>+</sup> recombinant (pLG260), the complementation tests were done with the recombinant plasmids present in both the donor and the recipient cells. The results (Fig. 2) indicate that, whereas pLG215 complemented pLG250 fully, pLG260 had no such effect. When pRP26, the RP4 derivative that specifies overproduction of primase, was introduced into the donor cells carrying pLG250, the deficiency of conjugative DNA synthesis in the recipients was only marginally rectified. It should be noted that, since pRP26 is



FIG. 2. Primase-dependent conjugative synthesis of ColIb DNA in recipient cells. BW96 and BW86 were used as donor and recipient strains, respectively. Plasmids present were:  $\circlearrowleft$ , pLG221 (Sog<sup>+</sup>) in donor strains;  $\bullet$ , pLG250 (Sog<sup>-</sup>) in donor strains; **A**, pLG250 and  $pLG215$  (Sog<sup>+</sup>) in donor strains with  $pLG215$  in recipient strains;  $\blacksquare$ , pLG250 and pLG26O (Pri+) in donor strains with pLG260 in recipient strains;  $\Box$ , pLG250 and pRP26 (Pri<sup>+</sup>) in donor strains. Values have been corrected by the subtraction of counts per minute incorporated  $(< 400$  cpm ml<sup>-1</sup>) by donor and recipient strains incubated separately.

conjugation defective and was not mobilized by the  $Incl<sub>1</sub>$ plasmids used here (data not shown), the amount of conjugative DNA synthesis due to the transfer of pRP26 is negligible in such an experiment.

### DISCUSSION

The results show that pRPI was consistently defective in generating transconjugant colonies, but the value varied according to the strains and genera participating in conjugation. A deficiency of  $\sim$ 20-fold or more was detected in E. coli K-12  $\times$  S. typhimurium matings, but, depending on the donor strain used in K-12  $\times$  K-12 matings, the deficiency ranged from 1.7 to in excess of 100. Since pRP1 was maintained in both K-12 and S. typhimurium with comparable and nearly normal frequencies (17), the primase-negative mutant must be defective in mediating some aspect, of conjugation rather than in subsequent vegetative replication required for formation of the transconjugant colony.

We investigated the nature of the conjugative defect of pRP1 through complementation tests. These indicated that the defect is caused by lack of RP4 DNA primase and that any possible polarity effect of the Tn7 insertion in pri on the expression of the gene downstream of the insert is not responsible. Although it was impossible to quantify pRP1 DNA transfer itself in these genetic experiments, it is evident that substantial transfer of the mutant plasmid occurred from donors lacking the plasmid primase and that the enzyme is required to promote establishment of the transferred DNA in the recipient cell. However, the requirement for plasmid primase in the recipient cell was satisfied by enzyme specified in the donor by a nontransferable *pri* gene. This leads to the important conclusion that plasmid primase or its RNA products or both are transmitted from the donor cell to promnote some aspect of the conjugative metabolism of RP4 in the recipient cell. Presumably, this reflects the normal mode of action of IncP plasmid primase.

The role of plasmid primase in the recipient cell will remain uncertain until the nature of the transferred RP4 DNA is determined. Like the transfer of IncF and IncI<sub>1</sub> plasmids, RP4 transfer is a unidirectional process that is initiated at a specific site on the plasmid (26). It is known that a specific single strand of IncF and  $Incl<sub>1</sub>$  plasmids is transmitted during conjugation (26), and, if this also applies to RP4 transfer, then an obvious role of the plasmid primase is the generation of primers for complementary strand synthesis in the recipient cell. However, we calculate from the data of Lanka and Barth (17) that RP4 specifies synthesis of about 5,500 molecules of the 118-kDa *pri* polypeptide cell<sup>-1</sup>. This abundance suggests that the protein may perform another function apart from primer generation itself.

Using a different experimental approach, we showed previously that the DNA primase specified by the sog gene of ColIb acts, analogously in conjugation (8). Furthermore, there is functional and physical evidence that sog primase is transferred during conjugation to the recipient cell (9; A. Merryweather, manuscript in preparation). Although RP4 and ColIb primases are different polypeptides, they are similar in their abilities to act in vitro on the single-stranded DNA of phages fd, G4, and  $\phi$ X174, to suppress the *dnaG3* mutation in  $\overline{E}$ . coli, and to synthesize oligoribonucleotides of similar composition and size distribution (17, 18, 24). Despite these functional similarities, the two enzymes act specifically during bacterial conjugation since neither primase supplied in the donor cell functioned effectively in the nonhomologous conjugation system, even under conditions in which putative accessory proteins necessary for activity of the nonhomologous primase were likely to be present. However, sog primase in the recipient cell acted weakly on transferred RP4 DNA, provided that the enzyme was specified by a ColIb plasmid and not by a cloned fragment. Taken together, these results imply that sog primase can function on the transferred strand of RP4 in a process requiring other ColIb-encoded factors but that the various ColIb components cannot be transmitted from the donor cell via the conjugation apparatus of RP4.

There is no absolute requirement for RP4 plasmid primase in bacterial conjugation. In the absence of the enzyme, we assume that primer-generating enzymes native to the recipient cell can substitute, as indicated by the recipientdependent variation of the relative yield of pRP1 transconjugants. However, the genotype of the donor can influence this substitution, since one K-12 strain  $(W3110T^-)$  contributed to a relatively normal yield of pRP1 transconjugants in matings with K-12 recipients, whereas strains of the BW86 pedigree lacked this ability. The nature of these complex interactions is unknown, but the important point is that they are normally bypassed by plasmid primase. It may be significant that the chromosome of W3110T<sup>-</sup> carries an inversion between  $rrnD$  and  $rrnE$  (14) which may lead to altered levels of bacterial primase or accessory proteins.

Why is RP4 primase required for efficient bacterial conjugation yet essentially redundant for vegetative replication of the plasmid? These differences can be reconciled if it is assumed that, like plasmid  $F(26)$ , a specific strand of RP4 is transferred during conjugation with the <sup>5</sup>' terminus leading. As RP4 replication is unidirectional, only one strand will support formation of multiple primers for discontinuous DNA synthesis. Presumably, this strand contains sites that are recognized by bacterial primer-generating enzymes, possibly acting in conjunction with accessory proteins encoded by regions of RP4 that are nonessential for conjugation. The second strand is likely to be deficient in sites recognized by vegetative priming systems because it supports leadingstrand synthesis in DNA replication. Since the direction of RP4 transfer is opposite to that of DNA replication (1, 12, 21), it follows that this second strand is the one, as we have proposed, that is transmitted during conjugation. Hence, plasmid primase provides a conjugation-specific priming mechanism that ensures efficient conversion of the transferred strand into duplex DNA in the recipient cell, thus providing the substrate for vegetative DNA replication.

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