

Construction of Tn5 *lac*, a transposon that fuses *lacZ* expression to exogenous promoters, and its introduction into *Myxococcus xanthus*

(gene expression/bacteriophage λ /bacteriophage P1/multicellular development/5-bromo-4-chloro-3-indolyl β -D-galactoside toxicity)

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ABSTRACT A promoterless *trp-lac* fusion fragment was inserted near one end of the bacterial transposon Tn5 in the correct orientation to fuse *lacZ* gene expression to promoters outside Tn5. The resulting transposon, Tn5 *lac*, retains the kanamycin-resistance gene of Tn5 and transposes in *Escherichia coli* at 6% the frequency of Tn5 to many different sites in a bacteriophage λ target. Expression of β -galactosidase, the product of the *lacZ* gene, from Tn5 *lac* insertions in phage λ depends both on insertion into a transcription unit in the correct orientation and on the regulation of the promoter of the transcription unit, verifying that by transposition Tn5 *lac* can fuse *lacZ* expression to outside promoters. An insertion of Tn5 *lac* in bacteriophage P1 was isolated and used to introduce Tn5 *lac* into *Myxococcus xanthus*, a bacterium that undergoes multicellular development. Stable kanamycin-resistant transductants are obtained that contain no P1 DNA sequences but have Tn5 *lac* inserted at different sites in the *Myxococcus* chromosome. Individual transductants express different levels of β -galactosidase. A chromogenic substrate of β -galactosidase, 5-bromo-4-chloro-3-indolyl β -D-galactoside, is toxic in *Myxococcus* when cleaved in large amounts. In principle, Tn5 *lac* could be used to assay transcription in any bacterium in which Tn5 can transpose and β -galactosidase can be measured.

Myxococcus xanthus is a Gram-negative soil bacterium that undergoes a primitive multicellular development with true cellular differentiation (1). During that developmental process, which depends on interactions between cells (2), many new gene products are synthesized (3). To identify genes that change their level of expression during development and to study the regulation of those genes, we wish to make operon fusions by transposition. A general approach, which uses transposition to fuse the expression of an easily assayed gene to promoters whose gene products are unknown or are not readily assayed, has been developed to identify and study interesting promoters (4).

Transposon Tn5 can be introduced into *Myxococcus* from *Escherichia coli* by the specialized transducing phage P1::Tn5 (5). It was previously found that bacteriophage P1 injects its DNA into *Myxococcus*, but the P1 is not stably maintained (6). Tn5, which encodes kanamycin resistance, can transpose from P1 to apparently random sites in the *Myxococcus* chromosome rendering cells stably kanamycin-resistant (Km^r) (5). Studies on the functional organization of Tn5 had revealed that the left inverted repeat of Tn5 (IS50L) is not needed to encode proteins required for transposition, but that some sequences near the left end of IS50L are required for transposition of Tn5 (7). Therefore, it seemed feasible to insert a promoterless gene near the left end of Tn5 to construct a transposon for making operon fusions. We chose to insert a promoterless *trp-lac* fusion fragment (8) into Tn5 because it had been shown that the *lacZ* gene product, β -

galactosidase, can be synthesized in *Myxococcus* and detected by cleavage of the standard chromogenic substrates (R. Gill and L. Avery, personal communication).

MATERIALS AND METHODS

Construction of Plasmids. Cloning operations and plasmid DNA purification were done as described (9). DNA polymerase I and T4 DNA ligase were gifts from S. Scherer. BAL-31 exonuclease (Bethesda Research Laboratories), calf alkaline phosphatase (Boehringer Mannheim), and phosphorylated linkers (Collaborative Research, Waltham, MA) were used according to the manufacturers' instructions. Plasmids were transformed (10) into *E. coli* MC1061 [*araD139*, Δ (*ara*, *leu*)7697, Δ *lacX74*, *galU*, *galK*, *hsdR*⁻, *hsdM*⁺, *strA*] (11).

Transposition Assay. Plasmids were transformed into BNN45 (*hsdR*⁻, *hsdM*⁺, *supE44*, *supF*, *thi*, *met*) (10), which contains the amber suppressor *supF* to permit propagation of λ gt4 (Δ *lac5*, *ci857*, *nin5*, *S_{am}100*) (12) on LB plates (10) supplemented with 100 mM MgSO₄. The resulting phage stocks were titered for plaque formation on BNN45 at 37°C and for ability to form Km^r lysogens at 30°C when MC1061 was infected and plated on LB containing 30 μ g of kanamycin sulfate per ml (Sigma) (10).

Analysis of Km^r λ Lysogens from the Transposition Assay. The amber mutation S100 in λ gt4 facilitated analysis of Km^r lysogens obtained from the transposition assay by preventing lysis of cells by the *S* gene product. Lysogens were thermally induced to make phages and were lysed with chloroform (10). The lysates were titered for plaque formation on BNN45 growing on LB plates supplemented with 10 mM MgSO₄ at 37°C. Phage DNA was isolated from the lysogens by diluting (1:10) LB cultures grown at 30°C overnight into LB supplemented with 10 mM MgSO₄ and 30 μ g of kanamycin per ml, growing for 3 hr at 37°C, and using a rapid plasmid isolation procedure (10). Transposon insertions in λ gt4 were mapped by digesting phage DNA with appropriate restriction enzymes and by separating DNA fragments by horizontal gel electrophoresis on 0.5% agarose (MCB Chemical, Norwood, OH) gels (10).

To analyze β -galactosidase production by lysogens containing λ gt4::Tn5 *lac*, LB cultures grown at 30°C overnight were diluted 1:25 into LB supplemented with 0.4% glucose/10 mM MgSO₄/and 30 μ g of kanamycin per ml and grown for 2.5 hr at 30°C. Cells in the exponential growth phase were diluted into the same medium without kanamycin, to a density of 4×10^7 cells per ml, and β -galactosidase activity was quantitated in each of two cultures, one grown for 3 hr at 30°C and one shifted to 42°C for 0.5 hr followed by 2.5 hr at 37°C, using the *o*-nitrophenyl- β -D-galactoside (Calbiochem) cleavage assay (13).

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Abbreviations: Km^r , kanamycin-resistant; IS50L, the left inverted repeat of Tn5; X-Gal, 5-bromo-4-chloro-3-indolyl β -D-galactoside; kbp, kilobase pair(s); bp, base pair(s).

Table 1. Identification of Tn5 *lac*

Plasmid	Sequence remaining at left end, bp*	Relevant structure [†]	Lac phenotype [‡]	Transposition	
				Frequency [§]	% pRZ104
pRZ104	1534		-	5 × 10 ⁻⁶	100
pLRK32	187		++	3 × 10 ⁻⁷	6
pLRKΔ131	150 ± 15		++	3 × 10 ⁻⁷	6
pLRKΔ211	30 ± 10		+	3 × 10 ⁻⁷	6
pLRKΔ514	0		+	<1 × 10 ⁻⁹	<0.02
pLRK1	0		+	<1 × 10 ⁻⁹	<0.02
pLRK21	1534		+++	3 × 10 ⁻⁷	6

*The amount of IS50L to the left of the *trp-lac* fusion fragment was determined for deletion plasmids from the size of the smaller *EcoRI/BamHI* fragment as measured by gel electrophoresis with fragments of known sizes.

[†]The structure of the plasmid between the *EcoRI* site in ColE1 and the right end of the right inverted repeat of Tn5 is shown. Symbols are the same as in Fig. 1 and construction of plasmids is described in the legend of Fig. 1.

[‡]+, ++, and +++, increasing levels of β -galactosidase expression in *E. coli* MC1061 on MacConkey lactose indicator plates (13).

[§]The transposition frequency is the number of Km^r lysogens obtained at 30°C per plaque-forming unit.

between its ends. Decreased transposition with increased distance between the ends of transposable elements has been observed for transposons bordered by IS1 (22) or IS10 (23).

No transposition was observed for deletion 514 or pLRK1, as expected, because the left end of Tn5 was deleted from these plasmids. A careful deletion analysis of the left end of Tn5 has shown that transposition is abolished when less than 16–18 bp of this end remain (24). Treating the λ phage stocks with deoxyribonuclease I to digest all DNA not protected by phage heads did not change the transposition frequency, while incubation of phage stocks with λ antiserum abolished both the ability to obtain Km^r transductants and the ability to form plaques. These results and restriction mapping of phage DNA from Km^r lysogens obtained from the transposition assay, which is described below, indicate that the frequency of Km^r transductants is a valid measure of the frequency of transposition from plasmid to λ .

Tn5 *lac* Transposes to Many Sites in λ . Phage DNA was isolated from 11 Km^r lysogens obtained in the transposition assay of Tn5 *lac*, and the position and orientation of each insertion was mapped by restriction analysis. Also, the position of 10 Tn5 insertions in λ gt4, obtained similarly, were mapped as controls and the results are shown in Fig. 2. The sites of Tn5 *lac* insertions in λ gt4 appear to be distributed as widely as those of Tn5. When the Km^r lysogens were thermally induced, all but one produced infective phage, al-

though all insertions in the rightward transcripts reduced plaque size. The one Km^r lysogen that failed to produce plaque-forming phage appeared to have Tn5 inserted in the *J* gene of λ gt4. The manner in which the Km^r lysogens were obtained selected against insertions in regions of λ gt4 that are essential for either lytic growth or lysogeny, which accounts for the low frequency of insertions that map in essential genes. All λ gt4::Tn5 *lac* formed tiny plaques on BNN45 growing on LB plates not supplemented with 10 mM MgSO₄, presumably because λ gt4::Tn5 *lac* phage heads are unstable in the absence of added Mg²⁺. For this reason, plates were supplemented with 100 mM MgSO₄ for growth of λ gt4 on plasmid-containing strains in the transposition tests reported in Table 1.

β -Galactosidase Expression from Tn5 *lac* in λ Correlates with the Phage Transcription Map. Lysogens with different insertions of Tn5 *lac* in λ gt4 allowed testing of the ability of Tn5 *lac* to fuse β -galactosidase expression to exogenous promoters, because the regulation of transcription units in λ is well known both for the lysogenic state and after induction of lytic development (25). For example, Tn5 *lac* insertion 4 is in the *rex* gene in the correct orientation to fuse β -galactosidase expression to the P_M or P_E promoters (Fig. 2). Data in Table 2 show that β -galactosidase, like repressor and *rex*, is expressed in the strain 4 lysogen and that its expression falls when phage production is induced by transiently shifting the temperature of the (c1857) lysogen to 42°C. Insertion 3 is also

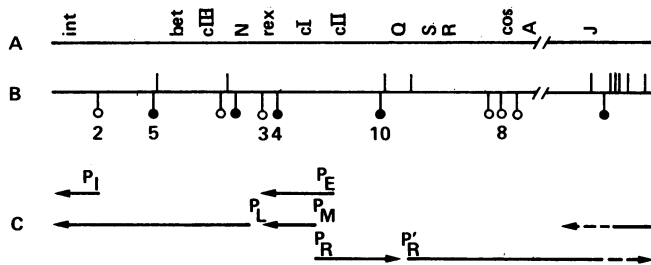


FIG. 2. Position of Tn5 and Tn5 *lac* insertions in λ gt4 (B) in relation to its known gene (A) and transcription (C) maps. (A) Partial gene map of λ gt4 prophage excluding most head and tail genes (—/—). (B) Tn5 insertions (—) were mapped by digestion of phage DNA with *Bam*HI, *Sma* I, and *Bam*HI followed by *Eco*RI. Tn5 *lac* insertions were mapped by digestion with *Bam*HI and *Eco*RI, allowing the orientation of Tn5 *lac* to be determined as that in which the *trp-lac* segment could be expressed by rightward (—) or leftward (—) transcription. β -galactosidase activity was determined for the numbered insertions (see Table 2). (C) Positions of known transcription units in λ gt4. —, Extent and direction of transcription; ---, readthrough (25).

Table 2. β -Galactosidase expression from Tn5 *lac* insertions in λ

Strain*	β -Galactosidase activity, units		Ratio [§]
	Lysogens [†]	After induction of λ [‡]	
wt	1100	1,100	1.0
4	1200	590	0.5
3	150	430	2.9
5	190	7,800	41
8	290	40,000	140
2	340	970	2.9
10	150	1,100	7.3

*The wild-type (wt) strain was MC1020 [*araD*139, Δ (*ara*, *leu*)7697, *lac*⁺, *galU*, *galK*, *strA*] (11); it was induced for β -galactosidase expression by including 1 mM isopropyl-thiogalactoside (Sigma) in the medium. Other strains were MC1061 (λ gt4::Tn5 *lac*) with Tn5 *lac* inserted in the position and orientation indicated in Fig. 2.

[†]The strain was grown at 30°C.

[‡]The strain was grown at 30°C then shifted to 42°C for 0.5 hr followed by 37°C for 2.5 hr.

[§]The ratio of β -galactosidase activity in the culture after induction of λ to that in the lysogen.

in the *rex* gene, but in the wrong orientation to fuse β -galactosidase expression to P_M or P_E . It shows 1/8th as much expression in the lysogen as insertion 4. Insertions 5 and 8, which are in the correct orientation to fuse β -galactosidase expression to P_L and P_R , respectively, show lower expression in their lysogens than insertion 4, with 41- and 140-fold increases in expression when the prophage is induced and phage growth ensues. Insertions 2 and 10, which are in the leftward and rightward transcription units (but in the wrong orientation to fuse β -galactosidase expression to P_L and P_R , respectively), display low levels of expression in their lysogens. The increase in β -galactosidase activity observed for insertions 3, 2, and 10 after induction of the prophage may be due to readthrough of antiterminated phage transcripts and the increased number of phage genomes. In summary, the level of expression of β -galactosidase from *Tn5 lac* insertions in λ gt4 depends on the position and orientation of the *Tn5 lac* insertion and correlates with the known regulation of transcription units in λ .

Isolation of $P1::Tn5 lac$. To introduce *Tn5 lac* into *Myxococcus*, an insertion of *Tn5 lac* in phage P1 was first obtained. Transposition of *Tn5 lac* from pLRK Δ 211 to P1 was detected by transduction into SF800 (P1). A P1 lysogenic recipient is necessary because insertion of *Tn5 lac* into P1 is expected to eliminate terminal redundancy and so prevent circularization of $P1::Tn5 lac$ when it enters *E. coli* (26); thus, homologous recombination between linear $P1::Tn5 lac$ and a (circular) resident P1 prophage can rescue the transposition product. The *polA* allele in SF800 was used to prevent maintenance of pLRK Δ 211 that might be introduced by P1 generalized transduction. Another undesired event, P1-pLRK Δ 211 cointegrate formation mediated by IS1 or Tn9 in P1, could be distinguished from *Tn5 lac* transpositions by screening transductants for colicin immunity encoded in the ColE1 portion of pLRK Δ 211. Cointegrations gave rise to Km^r transductants at about the same frequency as transposition of *Tn5 lac* from pLRK Δ 211 to P1.

Analysis of phage DNA from three colicin-sensitive transductants confirmed the insertion of *Tn5 lac* at different positions in P1. Structural analysis of only the $P1::Tn5 lac$ used in further studies is shown in Fig. 3. Since *Tn5 lac* has a single *EcoRI* site 4 kbp from one end and 8 kbp from the

other end, insertion of *Tn5 lac* into P1 should result in the disappearance of one normal P1 *EcoRI* restriction fragment and the appearance of two new fragments, one >4 kbp and one >8 kbp. The sum of the two new fragments should be 12 kbp longer than the missing fragment. The ethidium bromide-stained gel in Fig. 3A shows the expected pattern. In $P1::Tn5 lac$, a 6.5-kbp *EcoRI* restriction fragment is absent and two new fragments are present at 14 and 4.5 kbp. Southern blot hybridization using pLRK Δ 211 as the labeled probe (Fig. 3B) shows that only the new fragments contain *Tn5 lac* homology as expected.

Introduction of *Tn5 lac* into *Myxococcus*. When $P1::Tn5 lac$ infected *M. xanthus* strain DK101, Km^r transductants were obtained at a frequency of 72 ± 3 per 200 μ l of 25-fold-concentrated phage stock. Since insertion of *Tn5 lac* increases the genome length of P1 by $\approx 12\%$, the DNA of phage particles will lack terminal redundancy and consequently will be unable to recircularize upon infection of *E. coli*. Such particles are not expected to form plaques. Under our conditions, thermal induction of P1 *clr*100 lysogens generally yields $\approx 5 \times 10^9$ plaque-forming phage per ml. Assuming the $P1::Tn5 lac$ lysate contains this concentration of phage particles, the transduction frequency of $P1::Tn5 lac$ into *Myxococcus* is estimated to be 3×10^{-9} Km^r transductants per phage particle.

To verify that the Km^r transductants result from transposition of *Tn5 lac* to the *Myxococcus* chromosome, the experiment shown in Fig. 4 was carried out. DNA from the DK101 parent strain forms no hybrids with a *Tn5 lac* probe, while the *EcoRI* digests of nine independent Km^r transductants each have two hybridizable fragments, as expected, since there is one *EcoRI* site in *Tn5 lac*. No two transductants display the same set of fragments, showing that *Tn5 lac* has inserted at a different position in the *Myxococcus* chromosome in each transductant. Furthermore, comparing the sum of fragment lengths for individual transductants indicates that *Tn5 lac* had transposed to at least seven different regions of *Myxococcus* DNA. No hybridization was observed when P1 DNA was used as the labeled probe (data not shown).

Individual transductants express different levels of β -galactosidase as measured by colony color on CTT X-Gal, as if *Tn5 lac* had fused β -galactosidase expression to different *Myxococcus* promoters. The stability of *Tn5 lac* and its β -galactosidase expression were tested. Four transductants that were expressing β -galactosidase and two that were not were grown in the absence of kanamycin for 36 generations, plated on CTT, and 100 colonies for each were placed on CTT plates with *Km* and with X-Gal. All 100 colonies grew on the CTT *Km* plate and were the same color as originally

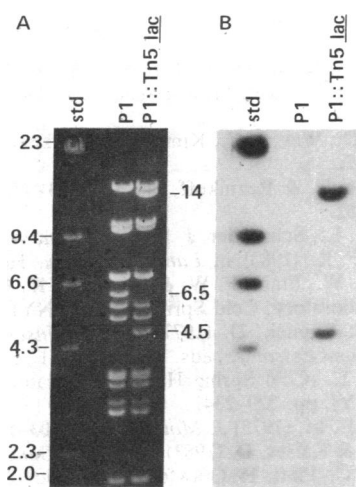


FIG. 3. Restriction (A) and Southern blot hybridization (B) analysis of $P1::Tn5 lac$. (A) DNA was digested with restriction enzyme, electrophoresed on a 0.5% agarose gel containing ethidium bromide, and photographed with UV illumination. Size standards (std) were *Hind*III-digested λ , and both P1 and $P1::Tn5 lac$ were digested with *EcoRI*. Numbers represent kbp. (B) Autoradiogram of a Southern blot (27) of the same gel as in A probed with a mixture of λ and pLRK Δ 211 DNA ^{32}P -labeled by nick-translation (10).

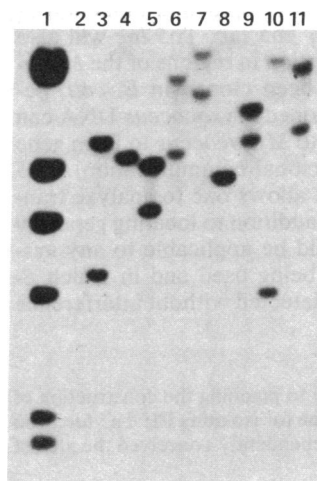


FIG. 4. Location of *Tn5 lac* in *Myxococcus* DNA. Chromosomal DNA was isolated (28), digested with *EcoRI*, electrophoresed on a 0.5% agarose gel, and Southern blot hybridization was carried out with a mixture of λ and pLRK Δ 211 DNA as the ^{32}P -labeled probe. Lanes: 1, *Hind*III-digested λ ; 2, DK101; 3-11, nine independent Km^r strains generated by infection of DK101 with $P1::Tn5 lac$.

determined on CTT X-Gal for each strain tested.

It was noted that *Myxococcus* colonies expressing a high level of β -galactosidase from Tn5 *lac* grew more slowly on CTT X-Gal plates than those expressing little or no β -galactosidase, and this effect was verified in liquid cultures supplemented with X-Gal (data not shown). Inclusion of X-Gal (50 μ g/ml) in CTT plates prevents growth of *Myxococcus* strains expressing β -galactosidase from Tn5 *lac* but does not prevent growth of nonexpressing strains when cells are plated for single colonies (D. Hodgson, personal communication).

DISCUSSION

A transposable promoter probe, Tn5 *lac*, was constructed by inserting a promoterless *trp-lac* fusion fragment near the left end of Tn5. When Tn5 *lac* inserts in a transcription unit in the correct orientation for the *trp-lac* fusion segment to be transcribed, a polycistronic messenger RNA is made. Stop codons in all three reading frames in the first 80 bp of the *trp-lac* fusion segment (29) prevent translation in the wrong reading frame of *trpA* and *lacZYA*, which retain their normal translation start signals. Thus, Tn5 *lac* is designed to make only transcriptional fusions to promoters outside Tn5. While this work was in progress, a modified Tn5 whose kanamycin resistance depends on outside promoters was described and was used to make operon fusions in *Caulobacter* (30).

We obtained an insertion of Tn5 *lac* in P1 that maps in the *EcoRI* fragment immediately downstream of the site at which P1 begins packaging its DNA into phage heads (31). This is the optimal position to ensure that almost every phage particle will contain Tn5 *lac*. This P1::Tn5 *lac* transduces *Myxococcus* to kanamycin resistance at $\approx 6\%$ the frequency observed for P1::Tn5 (5), consistent with the transposition frequency of Tn5 *lac* relative to Tn5 in *E. coli*. Southern hybridization of 9 Km^r transductants revealed that all had single insertions of Tn5 *lac* at different positions in the *Myxococcus* chromosome, and this pattern has been extended by tests of 20 other Km^r transductants (unpublished data). Because of the wide host range of P1 (32) and Tn5 (33), P1::Tn5 *lac* should be useful in a wide variety of bacteria.

By isolating large numbers of Tn5 *lac* insertions in the *Myxococcus* chromosome and screening them for β -galactosidase expression under growth and developmental conditions, we hope to identify genes that alter their expression during development. One way to study the regulation of developmentally induced genes is to devise a selection for mutations altering that regulation. We find that X-Gal is toxic to *Myxococcus* when it is cleaved by a strain expressing β -galactosidase, and it may be possible to select *cis*- and *trans*-acting mutations that eliminate developmental induction of β -galactosidase expression from Tn5 *lac*. Tn5 *lac* will also be useful for analysis of transcription in regions of the *Myxococcus* chromosome that have been cloned in *E. coli*, because insertions of Tn5 *lac* in cloned *Myxococcus* DNA can be obtained and reintroduced into *Myxococcus* to form gene replacements (K. Stephens, personal communication). Tn5 *lac* mutagenesis of cloned DNA allows one to analyze transcription in the cloned region in addition to locating genes by insertional mutation. This should be applicable to any system in which Tn5 is currently being used and in which β -galactosidase can be made and detected without interference from endogenous sources.

Ron Gill contributed substantially to planning the construction of Tn5 *lac* and helped devise the scheme for isolating P1::Tn5 *lac*. Ron Gill, Larry Shimkets, and D.K. independently conceived the idea of

constructing Tn5 *lac*. We thank M. Casadaban, R. Davis, S. Falkow, and J. Rosner for providing *E. coli* strains and bacteriophages. This investigation was supported by Grant AG02908 from the National Institute on Aging. L.K. was supported by a National Science Foundation Fellowship. This research was conducted in partial fulfillment of the requirements for the Ph.D. degree (to L.K.) from the Department of Biochemistry, Stanford University.

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