OVERPRODUCTION OF dna GENE PRODUCTS BY ESCHERICHIA COLI STRAINS CARRYING HYBRID Cole1 PLASMIDS

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SUMMARY
A bank of 2,000 E. coli strains carrying ColEl plasmids into which small random segments of the E. coli chromosome are inserted [Clarke, L., and Carbon, J. (1975) Proc. Nat. Acad. Sci. USA, 72, 4361-4365] was screened for those hybrids transferring thermoresistance to dna temperature-sensitive (ts) mutants. One or more was found to transfer dnaB, dnaC(D), dnaE, and dnaZ; none was found to transfer dnaG. These donor strains each overproduce by 3- to 10-fold the product of the gene whose defect is corrected, except in the case of dnaE. The strain donating dnaE does not overproduce DNA polymerase III.

The conversion of $\phi\chi 174$ single-stranded DNA to a double-stranded form by crude extracts of uninfected $E.\ coli$ (the " $\phi\chi$ system") (1, 2) provides an assay of the dnaB, dnaC(D), dnaE, dnaG, and dnaZ gene products by an $in\ vitro$ analog of the $in\ vivo$ complementation test (3-7). Using this assay, the dnaB, dnaC(D), dnaG, and dnaZ gene products have been purified (3-8). The dnaB gene product has ribonucleoside triphosphatase activity which is stimulated by single-stranded DNA (5). The dnaC(D) gene product interacts physically and functionally with the dnaB gene product (9). These two proteins function in the formation of an initiation complex in the $\phi\chi$ system (9, 10). The dnaE gene product is DNA polymerase III (11). The dnaZ gene product acts in the elongation of primed DNA in conjunction with other elongation factors and DNA polymerase II or III (7). In the $\phi\chi$ system, dnaZ and dnaG function in the reaction after formation of the initiation complex (7, 10).

Each of these proteins is present only in very small amounts in extracts of wild-type strains. The <u>dnaB</u> gene product required 40,000-fold purification (5) and DNA polymerase III required 4,500-fold purification (12) to reach homogeneity; <u>dnaC(D)</u>, <u>dnaG</u>, and <u>dnaZ</u> gene products have been purified 4,000-fold,

10,000-fold, and over 10,000-fold, respectively (6, 3, 7). Thus, it would be desirable to obtain strains which overproduce one or more of these proteins.

Molecular cloning facilitates the isolation of large amounts of the DNA, the RNA transcripts, and the protein gene products of the amplified chromosome segments (13-16). By linking random fragments of sonicated whole *E. coli* DNA to Colel DNA in vitro and transforming an F[†] strain to colicin El-resistance, Clarke and Carbon (16) have prepared a bank of *E. coli* strains, most of which carry hybrid Colel plasmids. The hybrid plasmids in the 2,000 strains in this bank probably cover most of the *E. coli* chromosome.

We have screened this bank for strains transferring the <u>dna</u> genes B, C(D), E, G, or Z, and report here the specific overproduction of <u>dnaB</u>, <u>dnaC</u>, or <u>dnaZ</u> proteins in different strains of this type.

MATERIALS AND METHODS

Bacterial strains are listed in Table 1. Strains from the Clarke and Carbon bank are indicated by MV12 (the recA F^+ derivative of C600 transformed by Clarke and Carbon with the ColEl hybrids) followed by an identification number.

Screening of E. coli strains containing hybrid ColEl plasmids for those carrying dna genes was accomplished as follows: Patches of each of the 2,000 F+ MV12/ColE1 streptomycin-sensitive strains were replica-plated to lawns (21) of each of the F- dna ts streptomycin-resistant strains on LB agar (1% tryptone, 0.5% yeast extract, 1% NaCl, 1.5% agar). After 30 minutes at 37° to allow transfer of potential hybrid plasmids in the F-mediated matings, plates were sprayed with 5% streptomycin sulfate to kill the donors and incubated for 24 hours at 42° to allow selective growth of recipients which had acquired a hybrid ColEl plasmid correcting the temperature-sensitive defect. Strains found to transfer dna+ were purified by single colony isolation and retested for transfer and other markers. In most cases, patches were first treated with colicin El to select cells which retained the ColEl hybrid plasmid from those which had lost it. Crude colicin El was the filter-sterilized supernatant from an overnight culture of the colicinogenic strain JF390 (obtained from John Foulds) grown in LB medum. In each case cells which had received the dna+ phenotype were repurified by single colony isolation and retested for streptomycin resistance, temperature resistance, and other markers.

For preparation of extracts, cells were grown in H-broth (8) with shaking at 37° to $A_{650} = 1.2$. They were collected by centrifugation and resuspended in 10% sucrose, 0.05 M Tris-Cl (pH 7.5) in a volume (ml) equal to the weight of the cells (gm), and frozen in dry ice-ethanol. Cells were lysed by incubating thawed cells for 30 minutes at 0° in centrifuge tubes with 0.2 mg/ml lysozyme, 0.01 M dithiothreitol, 0.005 M EDTA, and 0.15 M NaCl. Lysates were centrifuged for 45 minutes at 40,000 rpm in a Spinco 40 rotor. The clear supernatants were adjusted to 4% streptomycin sulfate with a 20% solution; the precipitate was collected by centrifugation. The supernatants were adjusted to 45% or 50% of saturation in ammonium sulfate by the addition of saturated neutralized ammonium sulfate, and the precipitates were collected. The precipitates for experiments 3, 5, 6, and 7 were resuspended in 0.02 M potassium phosphate (pH 6.5), containing 10% glycerol, 1 mM EDTA, and 1 mM dithiothreitol; those for experi-

TABLE 1. Strains of E. coli K12

Strain	Relevant										
Designation	Genotype Reference										
475	F thr leu strA dnaB391 ts	(17); J. Tomizawa									
PC221	F his thy arg malA xyl strA dnaC2 ts	(18)									
E486	F thr leu thy lac tonA met strA dnaE486 ts	(17)									
E293	F thr leu thy lac tonA strA dnaE293 ts	(17)									
BT308	F thr leu thy lac tonA strA dnaG308 ts	(19)									
NY731	F thy leu metE rif strA dnaG3 ts	(18)									
AX727	F lac strA dnaZ ts	(20)									
MV12/28	F ⁺ /AtrpE thr leu recA/ColEl-dnaB ⁺	This work and (16)									
MV12/29	F ⁺ /AtrpE thr leu recA/ColE1-dnaC ⁺	This work and (16)									
MV12/20	F ⁺ /AtrpE thr leu recA/Co1El-dnaC ⁺	This work and (16)									
MV12/26	F ⁺ /AtrpE thr leu recA/ColEl-dnaE ⁺	This work and (16)									
MV12/27	F ⁺ /AtrpE thr leu recA/Co1E1-dnaZ ⁺	This work and (16)									
MV12/16	F ⁺ / <u>AtrpE thr leu recA</u> /ColE1-dnaZ ⁺	This work and (16)									
MV12/15	F ⁺ /Δ <u>trpE</u> thr <u>leu recA</u> /ColEl α	This work and (16)									

 $[^]lpha$ This strain carries none of the dna genes on its ColEl plasmid.

ment 4 were resuspended in 0.02 M Tris-HCl (pH 7.5) containing 10% glycerol, 1 mM EDTA, and 1 mM dithiothreitol. Precipitates for use in experiments 1 and 2 were resuspended in 10% glycerol, 1 mM EDTA, 1 mM dithiothreitol, 0.05 M Tris-Cl (pH 7.5), and 40% saturated ammonium sulfate; precipitates were collected by centrifugation and resuspended in 10% glycerol, 1 mM EDTA, 1 mM dithiothreitol, and 0.02 M Tris-Cl (pH 7.5). All fractions within a given experiment were prepared at the same time in identical fashion.

Assays of dna gene products and DNA polymerase were performed as previously described (7, 8, 22).

RESULTS

The bank of 2,000 E. coli strains carrying random segments [average length 8 x 10^6 (16)] of the E. coli chromosome on hybrid ColEl plasmids was screened for transfer of temperature resistance to \underline{dnaB} ts, \underline{dnaC} ts, \underline{dnaE} ts, \underline{dnaE} ts, \underline{dnaG} ts, or \underline{dnaZ} ts recipients. One strain transferred \underline{dnaB}^+ , two transferred \underline{dnaC}^+ , one transferred \underline{dnaE}^+ , none transferred \underline{dnaG}^+ , and two transferred \underline{dnaZ}^+ (see Table 2). Single colonies which transferred temperature resistance were puri-

TABLE 2. Transfer of $\underline{dna}^{\dagger}$ markers from strains carrying ColE1 hybrids to \underline{dna} ts mutants a

											Recipient						
				Strain:					475	PC221	E486	BT308 C	AX727				
Donor Strain			dna ts:					В	С	Е	G	Z					
MV12/28		•						_			+	_	_	_	-		
MV12/29											-	+	-	-	-		
MV12/20		•									-	+	_	-	-		
MV12/26											-	-	+ ^b	-	_		
MV12/27		•									_	-	-	-	+		
MV12/16											-	-	_	_	+		

 $^{^{\}alpha}$ Ability to donate \underline{dna}^{+} markers was judged by appearance of thermoresistant recipients after matings with donors (see "Materials and Methods").

fied from each donor patch, in some cases by selecting for colicin El-resistance. Some patches contained a mixture of colicin-sensitive nontransferring and colicin-resistant transferring cells. Each patch transferred temperature resistance to only one of the five \underline{dna} ts mutants tested (Table 2). This was expected since the \underline{dna} genes are widely scattered on the $E.\ coli$ map (17) and each ColEl hybrid carries only a small part of the host genome. However, specific donation of thermoresistance to a \underline{dna} ts mutant is not proof that the \underline{dna}^+ allele has been donated. Further evidence for this can be obtained by examining donating strains for overproduction of the specific \underline{dna} gene product.

Partially purified fractions (prepared as described in "Materials and Methods") of strains donating thermoresistance to specific <u>dna</u> ts mutants were assayed for <u>dnaB</u>, <u>dnaC(D)</u>, and <u>dnaZ</u> gene products by *in vitro* complementation assays dependent on $\phi\chi174$ DNA and for DNA polymerase III with DNase-treated salmon sperm DNA as primer-template (Table 3). Fractions of strains correcting <u>dnaB</u> ts, <u>dnaC</u> ts, or <u>dnaZ</u> ts showed elevated levels (3- to 12-fold) only of the

 $^{^{}D}$ MV12/26 also transferred thermoresistance to E293, another dnaE ts mutant.

 $^{^{\}scriptscriptstyle \mathcal{C}}$ The same results were obtained with NY731, a different dnaG ts mutant.

TABLE 3. Elevated levels of <u>dna</u> gene products in extracts of strains carrying ColE1 hybrid plasmids

			Activity	Assayed (U/mg) b	
		· · · · · · · · · · · · · · · · · · ·		DNA Polymerase	
Strain;	ColEl Plasmid a	<u>dnaB</u>	dnaC	III	<u>dnaZ</u>
Experiment	1: 0-40% AS ^c				
MV12/15;	ColEl	8.3	-	-	44
MV12/16;	ColEl-dnaZ+	4.4	-	-	84
MV12/27;	ColE1-dnaZ+	4.9	-	-	130
Experiment	2: 0-40% AS				
MV12/26;	ColEl-dnaE+	_	-	200	19
MV12/27;	ColE1-dnaZ+	-	-	370	220
Experiment	3: 0-45% AS				
MV12/28;	ColEl-dnaB+	4.4	0.51	-	3.3
MV12/27;	ColEl-dnaZ+	0.9	0.36	-	11
Experiment	4: 0-45% AS				
MV12/29;	ColEl-dnaC+	-	1.4	-	1.1
MV12/27;	ColEl-dnaZ+	-	0.16	-	6.7
Experiment	5: 0-45% AS				
MV12/28;	ColEl-dnaB+	9.0	-	70	-
MV12/26;	ColEl-dnaE+	1.8	-	88	-
Experiment	6: 0-45% AS				
MV12/26;	ColEl-dnaE+	_	0.20	110	1.9
MV12/29;	ColE1-dnaC+	-	2.20	100	2.1
Experiment	7: 0-50% AS				
MV12/28;	ColEl-dnaB+	6.9	0.22	_	1.5
MV12/29;	ColEl-dnaC+	1.8	2.6	-	2.3
MV12/27;	ColE1-dnaZ+	0.9	0.32	-	4.9

 $^{^{}lpha}$ The <u>dna</u> gene indicates that whose ts phenotype is corrected by the particular ColEl hybrid plasmid (see Table 2).

product of the <u>dna</u> gene whose defect was corrected. Comparisons of specific activities between different experiments in Table 3 are not possible because of the different methods of fractionation used in different experiments.

 $[^]b$ One unit is the amount which stimulates the incorporation of 1 nmole of dTMP in 20 minutes at 30° under the conditions of the assays.

 $^{^{\}it C}$ Ammonium sulfate (AS) fractions were prepared as described in "Materials and Methods."

Calculations of total activity recovered per gram of cells indicated similarly elevated levels (3- to 10-fold) of dnaB, dnaC(D), and dnaZ gene products in extracts of the strains carrying the appropriate ColEl hybrid plasmids. Complementation assays of mixtures of fractions prepared from the various strains gave additive results, indicating that neither activators nor inhibitors could account for the observed differences.

Protein fractions of the strain donating \underline{dnaE}^+ did not have significantly elevated DNA polymerase III levels[†] when compared with fractions of strains donating \underline{dnaZ}^+ , \underline{dnaB}^+ , or \underline{dnaC}^+ . Thermoresistant DNA polymerase III activity was detected in extracts of E. coli E486 \underline{dnaE} ts carrying the ColEl plasmid which rendered this strain temperature-resistant for growth; no DNA polymerase III activity was detected in extracts of E486 \underline{dnaE} ts.

DISCUSSION

The strains described here will be useful in the purification of dnaC(D), and dnaZ gene products because each is overproduced 3- to 10-fold. Since overproduction occurs in exponentially growing cells and does not require any special maneuvers, it will be possible to use this approach in large-scale fermentations.

There are several possible explanations why elevated levels of DNA polymerase III were not detected in the strain transferring temperature-resistance to dnaE ts mutants. (a) The dnaE gene product may be only one of the subunits of DNA polymerase III [see (12)], and its overproduction would not necessarily yield more polymerase unless the other subunit were also overproduced. (b) The particular hybrid plasmid used may, for some reason, be present at a low copy number per genome. (c) Assay of DNA polymerase III in the relatively crude fractions examined is an unreliable indicator of actual polymerase

[†] The preliminary fractionation removed most of the DNA polymerase I, since over 90% of the DNA polymerase activity in these fractions was sensitive to N-ethylmaleimide. The polymerase activity was inhibited about 90% by the addition of 0.1 M KCl, indicating that DNA polymerase II was not a substantial portion of the total polymerase activity in these fractions (23).

content. (d) Overproduction of DNA polymerase III may be prevented by some regulatory mechanism. In this regard a specialized transducing λ phage carrying the dnaE gene also failed to overproduce DNA polymerase III on induction (24).

Our inability to find a hybrid plasmid carrying the $\underline{\operatorname{dnaG}}^+$ gene might be due to chance failure of this part of the $E.\ coli$ genome to be represented in the collection or inherent instability of plasmids carrying the dnaG region.

These strains, each carrying one of the <u>dna</u> genes, should also be useful in attempts to identify new <u>dna</u> genes; they could be used like small F' factors to rapidly map new <u>dna</u> ts mutants.

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