# Isolation and Characterization of a Temperature-Sensitive dnaK Mutant of Escherichia coli B

HIRAKU ITIKAWA\* AND JUN-ICHI RYU

Department of Biology, Tokyo Metropolitan University, Fukazawa, Setagaya-ku, Tokyo 158, Japan

**Received for publication 5 March 1979** 

A temperature-sensitive dnaK mutant (strain MT112) was isolated from Escherichia coli B strain H/r30RT by thymineless death selection at 43°C. By genetic mapping, the mutation [dnaK7(Ts)] was located near the thr gene (approximately 0.2 min on the map). E. coli K-12 transductants of the mutation to temperature sensitivity were assayed for their susceptibility to transducing phage  $\lambda$  carrying the dnaK and/or the dnaJ gene. All of the transductants were able to propagate phage  $\lambda$  carrying the dnaK gene. When macromolecular synthesis of the mutant was assayed at 43°C, it was observed that both deoxyribonucleic acid and ribonucleic acid syntheses were severely inhibited. Thus, it was suggested that the conditionally defective dnaK mutation affects both cellular deoxyribonucleic acid and ribonucleic acid syntheses at the nonpermissive temperature in addition to inability to propagate phage  $\lambda$  at permissive temperature.

It is known that phage lambda replication requires some host DNA synthesis function since the phage is unable to grow in certain Escherichia coli K-12 mutants containing temperature-sensitive (Ts) defects in DNA synthesis (4, 5, 8, 9; J. R. Walker, J. M. Henson, and C. S. Lee, Abstr. Annu. Meet. Genet. Soc. Am., 1976, 83:s80-81). In groP mutants, lambdoid phages were unable to multiply because the gene P product of the phages does not function (7). These groP bacteria were usually themselves conditionally defective in DNA synthesis and were identified as dnaB mutants (3). Saito and Uchida (12) isolated various grp mutants (grpA, -C, -D, and -E), in which the initiation of  $\lambda$  DNA replication is specifically affected. Furthermore, it was noted that grpC (or groPC) mutations were in the dnaK gene (12), which is located between thr and leu on the E. coli K-12 map (6, 12, 14). Recently, it was found that this class of mutation separates into two closely linked cistrons, dnaJ and dnaK, and the gene order is thr-dnaK-dnaJ-leu (15). Organization and expression of the dnaK and dnaJ genes were also characterized extensively in E. coli K-12 (13).

To study the mechanism of regulation of macromolecular synthesis, we used a temperaturesensitive mutant which was isolated in our laboratory from *E. coli* B stain H/r30RT from among the survivors of thymineless death at  $43^{\circ}$ C after mutagenic treatment (J. Ryu and H. Itikawa, Abstr. Annu. Meet. Genet. Soc. Jpn., 1975, 51:490). Genetic mapping data show that the mutation is located between the *thr* and *leu* genes.

Since *E. coli* B bacteria are unable to adsorb phage lambda, the temperature-sensitive mutation was transduced to *E. coli* K-12 strains for testing phage lambda propagation. The transductants were unable to propagate phage  $\lambda vh$ but were able to propagate transducing phage  $\lambda$ carrying the bacterial *dnaK* gene. Thus, the temperature-sensitive mutation of *E. coli* B is a mutation in the *dnaK* gene. It was noted that the *dnaK* mutant was defective in DNA synthesis and that RNA synthesis was also severely inhibited at 43°C.

### MATERIALS AND METHODS

**Bacteria and bacteriophages.** The bacterial and phage strains used in the present study are listed in Table 1.

Media. The following media were used. M9 minimal medium contained the following, in 1 liter of distilled water: Na<sub>2</sub>HPO<sub>4</sub>, 6g; KH<sub>2</sub>PO<sub>4</sub>, 3 g; NaCl, 0.5 g; NH<sub>4</sub>Cl, 1 g; MgSO<sub>4</sub> · 7H<sub>2</sub>O, 0.25 g; CaCl<sub>2</sub>, 11 mg; glucose, 2 g. M9CA contained 0.8 g of Casamino Acids (Difco; vitamin-free) per liter of M9 medium. Minimal medium (Davis) contained, per liter: K<sub>2</sub>HPO<sub>4</sub>, 7 g;  $KH_2PO_4$ , 2 g;  $MgSO_4 \cdot 7H_2O_1$ , 0.1 g;  $(NH_4)_2SO_4$ , 1 g; sodium citrate, 0.5 g; glucose, 2 g. L-broth medium contained 1% Polypepton (Daigo Eiyo Kagaku), 0.5% yeast extract (Difco), 0.5% NaCl, 0.1% glucose, and 2.5 mM CaCl<sub>2</sub>, (pH 7.0). λ-Broth contained 1% Polypepton and 0.25% NaCl (pH 7.0). Nutrient broth contained 1% meat extract (Kyokuto Seiyaku), 1% Polypepton, and 0.2% NaCl (pH 7.0). Brain heart infusion was obtained from Eiken Chemical Co. Solid medium contained 1.5% (minimal medium or nutrient broth) or 1.2% (L-

E. coli BH/r30RTthy argFS. KondoH/r30RTthy argF dnaK7H/r30RT, MNNG <sup>b</sup> MT112thy argF ilb dnaK7H/r30RT, MNNG <sup>b</sup> MT114thy argF ilb dnaK7ACB1 × MT114 $\rightarrow$ Leu <sup>-</sup> MT116thy ilb thrPloir(W3111) × MT116 $\rightarrow$ Thr <sup>-</sup> MT118ilb thr serBRB29 × MT117 $\rightarrow$ SerB <sup>-</sup> B1Hfr, wild typeS. KondoAC2519thr leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB29Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB85thr leu thi lacY rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB8511leu thi lacY rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB85112leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB11571thr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEHR4, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)W3111thrW3110Phage P1 wirFor E. coli K-12 WindletM. Abe	Strain	Genotype <sup>a</sup>	Origin/source
H/30RTthy argFS. KondoMT112thy argF dnaK7H/r30RT, MNNG <sup>b</sup> MT112thy argF dnaK7H/r30RT, MNNGMT114thy argF ilv dnaK7MT112, MNNGMT116thy ilv leu dnaK7ACB1 × MT114 → Leu <sup>-</sup> MT117thy ilv thrPlvir(W3111) × MT116 → Thr <sup>-</sup> MT118ilv thr serBRB29 × MT117 → SerB <sup>-</sup> B1Hfr, wild typeS. KondoAC2519thr leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr serB thi cet sup $\lambda^-$ FRB29Hfr serB thi cet sup $\lambda^-$ FRB85thr leu thi lacY rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lacY rpsL tonA supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851RB85112leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsxAB1157, MNNG <sup>b</sup> rpsL thy $\lambda^-$ rpsL tonA tryP1 (MT114) × AB1157T → TsAB11571leu pro his arg thi lac gal xyl mtl tsx rpsLP1 (MT114) × AB1157T → TsK1702grpC17 (dnaK170) leu pro his argE araAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lacHR4, H. Uchida (13)K7561groPAB756 [dnaK(Ts)] thr lac tonA thyAHR12, H. Uchida (12)W3111thrW3110P1For E. coli BP10irFor E. coli K-12M. AbeNuhVirulentM. Abe	E. coli B		
MT112thy argF dnaK7H/r30RT, MNNGbMT114thy argF ilv dnaK7MT112, MNNGMT116thy ilv leu dnaK7ACB1 × MT114 → Leu <sup>-</sup> MT117thy ilv leu dnaK7ACB1 × MT114 → Leu <sup>-</sup> MT118ilv thr serBRb29 × MT117 → SerB <sup>-</sup> B1Hfr, wild typeS. KondoAC2519th r leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr thr leu met arg pro trp gal lac rpsLB1 × AC2519 → HfrE. coli K-12RB29Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB85thr leu thi lac Y rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851R1leu thi lac Y rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 → Ts (dnaK7)RB851R2leu thi lac Y rpsL tonA supE dnaK7 $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsxAB1157, MNNGbK1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEHR4, H. Uchida (13)K2710grpC22 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)W3111thrW3110PhageF1For E. coli B NrbS. KondoP1 wir<	H/r30RT	thy argF	S. Kondo
MT114thy $argF$ ilv $dnaK7$ MT112, MNNGMT116thy ilv leu $dnaK7$ ACB1 × MT114 → Leu <sup>-</sup> MT117thy ilv thrPlvir(W3111) × MT116 → Thr <sup>-</sup> MT118ilv thr serBRB29 × MT117 → SerB <sup>-</sup> B1Hfr, wild typeS. KondoAC2519thr leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr thr leu met arg pro trp gal lac rpsLB1 × AC2519 → HfrE. coli K-12RB29Hfr serB thi cet $sup \lambda^-$ R. S. BuxtonRB851leu thi lacY rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lacY rpsL tonA supE $dnaK7\lambda^-$ P1 (MT114) × RB85 → Ts ( $dnaK7$ )RB851R1leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB11577thr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsLP1 (MT114) × AB1157T → Ts ( $dnaK7$ )K1702grpC17 ( $dnaK170$ ) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710groPAB756 [ $dnaK(Ts)$ ] thr lac tonA thyA thi supE44HR4, H. Uchida (12)W3111thrW3110PhageP1 P1virFor E. coli B Nr De K. coli K.12S. KondoP1MT114 × Koli L.12M. Abe	MT112	thy argF dnaK7	H/r30RT, MNNG <sup>b</sup>
MT116thy ilv leu dnaK7ACB1 × MT114 $\rightarrow$ Leu <sup>-</sup> MT117thy ilv thrPloir(W3111) × MT116 $\rightarrow$ ThrMT118ilv thr serBRB29 × MT117 $\rightarrow$ SerB <sup>-</sup> B1Hfr, wild typeS. KondoAC2519thr leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr thr leu met arg pro trp gal lac rpsLB1 × AC2519 $\rightarrow$ HfrE. coli K-12RB29Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB85thr leu thi lacY rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lacY rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB851R1leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>6</sup> K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710groPAB756 [dnaK(Ts)] thr lac tonA thyA thi supE44HR4, H. Uchida (12)W3111thrW3110Phage P1For E. coli B NuS. KondoP1For E. coli K-12 MuM. Abe	MT114	thy argF ilv dnaK7	MT112, MNNG
MT117thy ilv thrPluir(W3111) × MT116 $\rightarrow$ Thr <sup>-</sup> MT118ilv thr serBRB29 × MT117 $\rightarrow$ SerB <sup>-</sup> B1Hfr, wild typeS. KondoAC2519thr leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr thr leu met arg pro trp gal lac rpsLS. KondoRB29Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB85thr leu thi lacY rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lacY rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB851R1leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mt xyl gal lac thi supEHR4, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (12)W3111thrW3110Phage P1 WirFor E. coli B VinlentS. Kondo M. Abe	<b>MT</b> 116	thy ilv leu dnaK7	$ACB1 \times MT114 \rightarrow Leu^{-}$
MT118ilv thr serBRB29 × MT117 → SerB <sup>-</sup> B1Hfr, wild typeS. KondoAC2519thr leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr thr leu met arg pro trp gal lac rpsLB1 × AC2519 → HfrE. coli K-12RB29Hfr serB thi cet sup λ <sup>-</sup> R. S. BuxtonRB85thr leu thi lac Y rpsL supE λ <sup>-</sup> F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lac Y rpsL tonA supE dnaK7 λ <sup>-</sup> P1 (MT114) × RB85 → Ts (dnaK7)RB851R1leu thi lac Y rpsL tonA supE λ <sup>-</sup> Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx thy dnaK7 λ <sup>-</sup> AB1157, MNNG <sup>b</sup> K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)W3111thrW3110Phage P1For E. coli B VinlentS. Kondo M. Abe	<b>MT</b> 117	thy ilu thr	$Plvir(W3111) \times MT116 \rightarrow Thr^{-}$
B1Hfr, wild typeS. KondoAC2519thr leu met arg his pro trp gal lac rpsLS. KondoACB1Hfr thr leu met arg pro trp gal lac rpsLS. KondoB1 × AC2519 $\rightarrow$ HfrHfr thr leu met arg pro trp gal lac rpsLB1 × AC2519 $\rightarrow$ HfrE. coli K-12RB29Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB85thr leu thi lacY rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lacY rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB851R1leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)W3111thrW3110Phage P1For E. coli B VinlentS. Kondo M. Abe	<b>MT</b> 118	ilv thr serB	$RB29 \times MT117 \rightarrow SerB^{-}$
AC2519 ACB1thr leu met arg his pro trp gal lac rpsL ACB1S. Kondo B1 × AC2519 → HfrE. coli K-12 RB29 RB85 RB85 RB851 RB851 RB8511 RB851R1 RB851R2Hfr serB thi cet sup $\lambda^-$ RB8511 leu thi lacY rpsL supE $\lambda^-$ RB851R1 RB851R2 AB1157TR. S. Buxton F <sup>-</sup> C600, R. S. Buxton P1 (MT114) × RB85 → Ts (dnaK7) Spontaneous revertants from RB851 AB1157TAB1157T K1702thr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ R10 arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T → Ts (dnaK7)K1702 K2710 K2710 K7561 W3111grpC2 (dnaJ2) gal lac thi supE44 thi supE44HR4, H. Uchida (13) HR12, H. Uchida (12)Phage P1 P10For E. coli B For E. coli K-12S. Kondo M. Abe	<b>B</b> 1	Hfr, wild type	S. Kondo
ACB1Hfr thr leu met arg pro trp gal lac rpsLB1 × AC2519 $\rightarrow$ HfrE. coli K-12RB29Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB85thr leu thi lac Y rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lac Y rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB851R1leu thi lac Y rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)W3111thrW3110Phage P1For E. coli B VirulentS. Kondo M. Abe	AC2519	thr leu met arg his pro trp gal lac rpsL	S. Kondo
E. coli K-12 RB29Hfr serB thi cet sup $\lambda^-$ RB85 RB85 RB85 RB851 RB851 RE ut hi lacY rpsL tonA supE $\lambda^-$ RB8511 RB851R1 RB851R2R. S. Buxton F <sup>-</sup> C600, R. S. Buxton P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7) Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB11571Spontaneous revertants from RB851AB11571thr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB11571P1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13) HR12, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (12) thi supE44W3111thrW3110Phage P10For E. coli B VirulentS. Kondo M. Abe	ACB1	Hfr thr leu met arg pro trp gal lac rpsL	$B1 \times AC2519 \rightarrow Hfr$
RB29Hfr serB thi cet sup $\lambda^-$ R. S. BuxtonRB85thr leu thi lacY rpsL supE $\lambda^-$ F <sup>-</sup> C600, R. S. BuxtonRB851leu thi lacY rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB851R1leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)Phage P1For E. coli B For E. coli K-12S. KondoP1For E. coli K-12M. Abe M. Abe	E. coli K-12		
RB85thr leu thi lacY rpsL supE $\lambda^-$ RB851F^- C600, R. S. Buxton P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB851R1 RB851R2leu thi lacY rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEP1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)W3111thrW3110Phage P1For E. coli B For E. coli K-12S. Kondo M. Abe	RB29	Hfr serB thi cet sup $\lambda^-$	R. S. Buxton
RB851leu thi lac Y rpsL tonA supE dnaK7 $\lambda^-$ P1 (MT114) × RB85 $\rightarrow$ Ts (dnaK7)RB851R1leu thi lac Y rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEP1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13) HR12, H. Uchida (12)Phage P1For E. coli B For E. coli K-12S. Kondo M. AbeP1For E. coli K-12M. Abe	RB85	thr leu thi lacY rpsL supE $\lambda^-$	F <sup>-</sup> C600, R. S. Buxton
RB851R1 RB851R2leu thi lac Y rpsL tonA supE $\lambda^-$ Spontaneous revertants from RB851AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T → Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)K7561groPAB756 [dnaK(Ts)] thr lac tonA thyA thi supE44HR12, H. Uchida (12)W3111thrW3110Phage P10For E. coli B For E. coli K-12S. Kondo M. Abe	RB851	leu thi lacY rpsL tonA supE dnaK7 $\lambda^-$	P1 (MT114) $\times$ RB85 $\rightarrow$ Ts (dnaK7)
RB851R2Spontaneous reventants from RB851AB1157Tthe leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNG <sup>b</sup> AB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T → Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)K7561groPAB756 [dnaK(Ts)] thr lac tonA thyA thi supE44HR12, H. Uchida (12)W3111thrW3110Phage P10For E. coli B For E. coli K-12S. Kondo M. AbeVirulentM. Abe	RB851R1)	low thi looV mol ton A sum $F$ )-	Spontaneous revertants from DD951
AB1157Tthr leu pro his arg thi lac gal xyl mtl tsx rpsL thy $\lambda^-$ AB1157, MNNGbAB11571leu pro his arg thi lac gal xyl mtl tsx rpsL thy dnaK7 $\lambda^-$ P1 (MT114) × AB1157T → Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac thi supE44HR4, H. Uchida (13)K7561groPAB756 [dnaK(Ts)] thr lac tonA thyA thi supE44HR12, H. Uchida (12)Phage P1For E. coli B For E. coli K-12S. Kondo M. AbeVirulentM. Abe	RB851R2	teu thi taci TpsL tonA supE A	Spontaneous revertants from RD851
AB11571leu pro his arg thi lac gal xyl mtl tsx rpsLP1 (MT114) × AB1157T $\rightarrow$ Ts (dnaK7)K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lac groPAB756 [dnaK(Ts)] thr lac tonA thyA thi supE44HR4, H. Uchida (13)W3111thrW3110Phage P1P1For E. coli B For E. coli K-12S. Kondo M. AbeVirulentM. Abe	AB1157T	thr leu pro his arg thi lac gal xyl mtl tsx $\operatorname{rms} I$ thus $J^{-}$	AB1157, MNNG <sup>₺</sup>
AD11011       Int all pictulate gat Xyt mit tisk types       Int (mit Hit) A AD11011 - Fis         thy dnaK7 $\lambda^-$ (dnaK7)         K1702       grpC17 (dnaK170) leu pro his argE ara       AB1133, H. Uchida (13)         mtl xyl gal lac thi supE       HR4, H. Uchida (13)         K2710       grpC2 (dnaJ2) gal lac       HR4, H. Uchida (13)         K7561       groPAB756 [dnaK(Ts)] thr lac tonA thyA       HR12, H. Uchida (12)         thi supE44       W3111       thr         Phage       P1       For E. coli B       S. Kondo         P1vir       For E. coli K-12       M. Abe $\lambda vh$ Virulent       M. Abe	AB11571	ley nro his ara thi lac gal ryl mtl ter rnsl.	P1 (MT114) $\times$ AB1157T $\rightarrow$ Te
K1702grpC17 (dnaK170) leu pro his argE ara mtl xyl gal lac thi supEAB1133, H. Uchida (13)K2710grpC2 (dnaJ2) gal lacHR4, H. Uchida (13)K7561groPAB756 [dnaK(Ts)] thr lac tonA thyA thi supE44HR12, H. Uchida (12)W3111thrW3110Phage P1P1For E. coli BS. KondoP1virFor E. coli K-12M. Abe $\lambda vh$ VirulentM. Abe	ADITOTI	thy dnaK7 $\lambda^{-}$	(dnaK7)
Miloz       mil xyl gal lac thi supE       Milozofi (dividual)         K2710       grpC2 (dnaJ2) gal lac       HR4, H. Uchida (13)         K7561       groPAB756 [dnaK(Ts)] thr lac tonA thyA       HR12, H. Uchida (12)         thi supE44       W3111       thr         W3111       thr       W3110         Phage       For E. coli B       S. Kondo         P1       For E. coli K-12       M. Abe $\lambda vh$ Virulent       M. Abe	K1702	grnC17 (dnaK170) ley pro his argE ara	AB1133, H. Uchida (13)
K2710grpC2 (dnaJ2) gal lacHR4, H. Uchida (13)K7561groPAB756 [dnaK(Ts)] thr lac tonA thyAHR12, H. Uchida (12)thi supE44W3111thrW3111thrW3110PhageP1For E. coli BS. KondoP1virFor E. coli K-12M. Abe $\lambda vh$ VirulentM. Abe	111/02	mtl xvl gal lac thi sunE	1121100, 11. Comuu (10)
K7561groPAB756 [dnaK(Ts)] thr lac tonA thyAHR12, H. Uchida (12) thi supE44W3111thrW3110PhageP1For E. coli BS. KondoP1.virFor E. coli K-12M. Abe $\lambda vh$ VirulentM. Abe	K2710	grpC2 (dnaJ2) gal lac	HR4, H. Uchida (13)
thi supE44W3111thrW3110Phage $P1$ For E. coli BS. KondoP1 $vir$ For E. coli K-12M. Abe $\lambda vh$ VirulentM. Abe	K7561	groPAB756 [dnaK(Ts)] thr lac tonA thvA	HR12, H. Uchida (12)
W3111thrW3110PhageFor E. coli BS. KondoP1For E. coli K-12M. AbeλυhVirulentM. Abe		thi supE44	, (,
Phage     S. Kondo       P1     For E. coli B     S. Kondo       P1vir     For E. coli K-12     M. Abe       \u03c0vh     Virulent     M. Abe	<b>W3111</b>	thr	W3110
P1For E. coli BS. KondoP1 $vir$ For E. coli K-12M. Abe $\lambda vh$ VirulentM. Abe	Phage		
P1virFor E. coli K-12M. AbeλvhVirulentM. Abe	P1	For <i>E. coli</i> B	S. Kondo
λ <i>vh</i> Virulent M. Abe	P1 <i>vir</i>	For <i>E. coli</i> K-12	M. Abe
	λυ <b>h</b>	Virulent	M. Abe
$\lambda c$ 1857 <i>dnaJK</i> Transducing phage for <i>dnaJ</i> and <i>dnaK</i> H. Uchida (13)	λcI857dnaJK	Transducing phage for <i>dnaJ</i> and <i>dnaK</i>	H. Uchida (13)
$\lambda c I857 dna J \Delta 145$ Transducing phage for dna J H. Uchida (13)	λcI857dnaJ∆145	Transducing phage for dnaJ	H. Uchida (13)
$\lambda c I50 dnaK$ Transducing phage for $dnaK$ H. Uchida (13)	λcI50dnaK	Transducing phage for dnaK	H. Uchida (13)

 TABLE 1. Bacterial and phage strains

<sup>a</sup> Genotypes according to reference 1.

<sup>b</sup> MNNG, N-methyl-N'-nitro-N-nitrosoguanidine.

broth or  $\lambda$ -broth) agar. Supplements were added when needed, as follows: 20  $\mu$ g of thymine, 40  $\mu$ g of amino acids, or 2  $\mu$ g of thiamine per ml.

**Growth conditions.** Cells were incubated overnight in brain heart infusion broth at 37°C. Cells were washed with saline (0.85% NaCl) by centrifugation and then suspended in the fresh liquid medium and incubated with shaking at 37°C. Cell growth was monitored by measuring the turbidity of the cultures at 660 nm with a spectrophotometer (Hitachi FPO-3). Temperature-sensitive mutant cells were incubated at 30°C unless otherwise noted.

**Radioisotopes.** [6-<sup>3</sup>H]thymine, [6-<sup>3</sup>H]thymidine, [5-<sup>3</sup>H]uracil, and [U-<sup>14</sup>C]leucine were purchased from Daiichi Pure Chemicals Co. [2-<sup>14</sup>C]uridine was obtained from Amersham Products Inc.

Determination of DNA, RNA, and protein syntheses. For continuous labeling, cells grown in M9CA were divided into three portions. [<sup>3</sup>H]thymine (1.0  $\mu$ Ci/ml, 26.0 Ci/mmol), [<sup>3</sup>H]uracil (0.5  $\mu$ Ci/ml, 14.9 Ci/mmol) with 20  $\mu$ g of uracil per ml, or [<sup>14</sup>C] leucine (0.1  $\mu$ Ci/ml, 262 mCi/mmol) with 40  $\mu$ g of leucine per ml, respectively, was added to each of them. After five generation times of labeling, 0.4-ml samples were added to 0.4 ml of 10% cold trichloroacetic acid and chilled. After 1 h or more of chilling, they were washed with cold 5% trichloroacetic acid and then with cold 5% acetic acid on glass fiber filters (Whatman GF/C) and dried. Radioactivity in the acidinsoluble fraction was counted by using a Beckman liquid scintillation counter (LS-III) in glass vials containing 10 ml of 0.4% 2,5-diphenyloxazole in toluene.

For pulse-labeling, 0.5-ml portions were removed from the culture and incubated further with the addition of 0.5 ml of M9CA medium containing [<sup>3</sup>H]thymidine (1  $\mu$ Ci/ml, 10 Ci/mmol), [<sup>14</sup>C]uridine (50  $\mu$ Ci/ml, 62 mCi/mmol), or [<sup>14</sup>C]leucine (0.2  $\mu$ Ci/ml, 262 mCi/mmol) plus 4  $\mu$ g of leucine per ml. After 2 ([<sup>3</sup>H]thymidine or [<sup>14</sup>C]uridine) or 5 ([<sup>14</sup>C]leucine) min of labeling, acid-insoluble fractions were collected and assayed as described above.

Isolation of temperature-sensitive mutants. Parental cells (H/r30RT), grown exponentially in M9CA medium, were collected by centrifugation and suspended in 1 ml of acetate buffer (pH 5.0). N-methyl-N'-nitro-N-nitrosoguanidine was added (final concen-

#### Vol. 138, 1979

tration, 1 mg/ml), and cells were allowed to stand for 40 min at 37°C. They were washed twice with acetate buffer and once with M9CA medium, resuspended in M9CA medium, and incubated overnight at 30°C. The culture was diluted in the same medium, and, at midlog phase, cells were shifted from 30 to 43°C. After 1 h, they were washed on a membrane filter (Millipore HAWP, 0.45-um pore size) and incubated further for 2.5 h at 43°C in M9CA medium without thymine. Thus, growing cells under these conditions were subjected to thymineless death and viable cells were decreased to 0.1% of the intial cell number. Mutants defective in DNA synthesis at 43°C were resistant to thymineless death and were enriched by this treatment. One portion of the culture was then plated on nutrient broth plates and incubated at 30°C. Nine mutant strains unable to grow at 43°C but growing at 30°C on nutrient broth plates were obtained at a frequency of 1% among 10<sup>4</sup> cell survivors. One mutant [dnaK7(Ts)] was mainly studied in the present work.

# RESULTS

Genetic mapping of dnaK(Ts). Since preliminary mating experiments between Hfr CS101 and MT114 indicated that the mutation was located near leu on the E. coli K-12 map (data not shown), generalized transducing phage P1 was used with strain W3111 (thr) as the donor and MT116 (thy ilv leu dnaK7(Ts)) was used as the recipient. Resulting Leu<sup>+</sup> transductants were selected on a minimal medium agar plate supplemented with thymine, threonine, isoleucine, and valine. Among 578 Leu<sup>+</sup> clones, temperature resistance and Thr<sup>-</sup> were found at a frequency of 0.52% (3 of 578) and temperature resistance and Thr<sup>+</sup> were found at a frequency of 0.69% (4 of 578). The remaining 571 Leu<sup>+</sup> transductants were temperature sensitive and Thr<sup>+</sup>, indicating that dnaK7(Ts) is located near the thr gene. To determine the location of the mutation more precisely, further P1 transduction tests were carried out as follows.

(i) Transduction 1 was MT116 (temperaturesensitive Thr<sup>+</sup> donor)  $\times$  MT117 (temperatureresistant Thr<sup>-</sup> recipient) on minimal medium agar plate supplemented with thymine, isoleucine, valine, and leucine. Among 117 Thr<sup>+</sup> transductants, 108 (92.3%) were temperature sensitive, suggesting a very close linkage between the *thr* and *dnaK7*(Ts) genes.

(ii) Transduction 2 was MT114 (temperaturesensitive donor)  $\times$  MT118 (temperature-resistant recipient) on minimal medium agar supplemented with threonine, isoleucine, and valine. Among 85 SerB<sup>+</sup> transductants, 74.1% were Thr<sup>+</sup> and 61.2% were temperature sensitive (Table 2).

These results suggest that the gene order might be *serB-thr-dnaK7*(Ts). To confirm this order, Thr<sup>+</sup> transductants were selected from the same cross, and  $SerB^+$  and temperature-

 
 TABLE 2. P1 transduction between MT114 (donor) and MT118 (recipient)<sup>a</sup>

Cross	Genotype of transductant			N	
	serB	thr	dnaK7(Ts)	No. scored	70
A	1	1	1	52	61.2
	1	1	0	11	12.9
	1	0	1	0	0
	1	0	0	22	25.9
В	1	1	1	140	81.9
_	1	1	0	10	5.8
	0	1	1	16	9.4
	0	1	0	5	2.9

<sup>a</sup> P1 transduction was done by the method of Lennox (10). In cross A, SerB<sup>+</sup> colonies on the selective agar plate were subjected to single-colony isolation and assayed for their genotypes including threonine requirement and temperature sensitivity. In the table, all donor markers [ $serB^+thr^+$  dnaK7(Ts)] are designated as 1, and recipient markers (serB thr  $dnaK^+$ ) are designated as 0. In cross B, Thr<sup>+</sup> colonies on the selective agar plate were assayed for their requirement for serine and temperature sensitivity after single-colony isolation. Procedures and designation of the markers are the same as in cross A.

sensitive clones were scored as shown in Table 2. Again the results suggested that the gene order was serB-thr-dnaK7(Ts). The map position of the dnaK locus, therefore, may be estimated as follows. By using Wu's formula (1), the map distance of thr-serB corresponds approximately to 0.1 min (from 87.1% of the cotransduction frequency) and that of thr-dnaK corresponds to 0.06 min (91.7% cotransduction). Since the map distance of thr-serB is listed as 0.4 min on the E. coli K-12 map (1), a reasonable map distance for thr-dnaK might be proportionally calculated as around 0.24 min.

Identification of a temperature-sensitivity mutation in the dnaK gene. It is known that the dnaJ and dnaK genes are located near the thr gene and that conditionally defective mutations of these genes affect cellular DNA synthesis at nonpermissive temperatures, in addition to affecting the inability to propagate phage  $\lambda$  at permissive temperature (12, 13). Since the temperature sensitivity mutation in strain MT112 was cotransducible with the thr gene, the mutation was transferred to E. coli K-12 strains AB1157T and RB85 by P1 transduction. The transductants (AB11571 and RB851) were unable to propagate phage  $\lambda vh$ , in contrast to the parental cells, which grow  $\lambda vh$ . Next, an experiment was done to see whether the mutation resided in the dnaJ or the dnaK gene, using strains AB11571 and RB851 and transducing phage  $\lambda$  carrying the *dnaJ* and/or the *dnaK* chromosomal gene. Both of the strains are able to propagate phage  $\lambda$  carrying the *dnaK* gene but not the *dnaJ* gene (Table 3). This finding indicates that the mutation must be in *dnaK*. We designated the mutation *dnaK7*(Ts). Furthermore, it is suggested that the original mutant strain, MT112, carries the single mutation (*dnaK7*) related to its temperature sensitivity, because temperature-resistant revertants were obtained spontaneously in brain heart infusion broth culture at a frequency of about 10<sup>-9</sup>. This possibility is supported by another finding that the temperature sensitivity mutation and inability to propagate phage  $\lambda$  were not separable by P1 transduction as seen in the transductants, AB11571 and RB851 (see Table 3).

Effect of temperature shift on growth of the mutant. The doubling time of the parental strain (H/r30RT) in M9 medium at 30°C was 45 min, and that of the mutant (MT112) carrying dnaK7(Ts) was 60 min. To determine the effect of high temperature on the growth of the mutant, the cells were incubated at 30°C in M9CA medium and shifted to 43°C (nonpermissive temperature; Fig. 1). After a 30-min lag, the increase in optical density was severely inhibited. The residual increase after the temperature shift was about 80%. The viable cell number increased about 30% after 1 h at 43°C, and no decrease in colony-forming units was observed until 4 h at 43°C. Phase-contrast microscopy revealed that a small portion of the cells became very elongated (e.g., snake form).

Effect of the temperature shift on macromolecular synthesis. The effect of the temperature shift on DNA, RNA, and protein syntheses in mutant strain MT112 was examined by continuous or pulse-labeling of radioisotopes. Figure 2 shows the pattern of macromolecular synthesis after temperature shift by pulse-labeling. RNA synthesis decreased rapidly 5 to 10

min after the shift; DNA synthesis was first stimulated and then decreased rapidly. Protein synthesis decreased at a slower rate, and, surprisingly, it continued at more than 30% of the initial value even at 120 min. Thus, RNA synthesis seemed to be initially influenced most drastically by the temperature shift. We also examined the E. coli K-12 dnaK(Ts) mutant K7561 (carrying groPAB756 mutation) by the same procedure. Cellular RNA synthesis as well as DNA synthesis was inhibited after the temperature shift (Fig. 3). Two temperature-resistant revertants, RB851R1 and RB851R2, isolated spontaneously from strain RB851 seemed to regain the ability to synthesize DNA, RNA, and protein at 43°C, since these revertants showed similar patterns of growth at this temperature when compared with strain RB85. Thus, it is suggested that the dnaK7(Ts) mutation simultaneously affects cellular nucleic acid synthesis and propagation of phage lambda (see Table 3). It is possible that a second mutation coexisting with the temperature-sensitivite mutation [dnaK7(Ts)] might affect RNA polymerase. Therefore, the activity of RNA polymerase extracted from strains MT112 and H/r30RT by the method of Burgess (2), with minor modifications, was tested.

Enzyme activity was examined by incorporation of  $[5-{}^{3}H]$ UTP (1.0 Ci/mmol; Amersham Products Inc.) into cold acid-insoluble material (11). The results showed no alteration of activity of the enzyme from MT112 at 43°C compared with that from the parental strain, H/r30RT (data not shown).

# DISCUSSION

In the present study, a dnaK7(Ts) mutant was isolated from an *E. coli* B strain by thymineless death screening after mutagenic treat-

TABLE 3. Efficiency of plating of phage  $\lambda$  carrying the dnaK and/or the dnaJ gene on E. coli K-12 strains<sup>a</sup>

	Efficiency of plating of phage:			
Bacterial strain	λcI857dnaJdnaK	λcI857dnaJ∆145	λcI50dnaK	
AB1157T (temperature resistant)	1	1	1	
RB85 (temperature resistant)	1	1	1	
RB851R1 (temperature resistant)	0.77	1	1	
RB851R2 (temperature resistant)	0.90	1	1	
AB11571 [dnaK7(Ts)]	0.76	$<2.0 \times 10^{-3}$	0.48	
RB851 [dnaK7(Ts)]	0.64	$<2.6 \times 10^{-3}$	0.91	
K1702 (dnaK170)	0.80	$<2.0 \times 10^{-3}$	1	
K2710 (dnaJ2)	1	1	$<2.6 \times 10^{-3}$	

<sup>a</sup> Efficiency of plating denotes the number of plaques produced by a phage strain on a given bacterial host at 37°C relative to the number produced on RB85 cells. For the test, cells were grown in 3 ml of brain heart infusion broth at 37°C. After that, cells were collected by centrifugation and suspended in 1 ml of  $\lambda$  broth. Then 0.25 ml of the cell suspension (4 × 10<sup>8</sup> cells) and 0.1 ml of phage lysate were added to 2.5 ml of molten  $\lambda$  soft agar (0.5% agar). The mixture was overlaid onto a  $\lambda$  agar plate containing 10 µg of thymine per ml. After overnight incubation at 37°C, the number of plaques was counted.



FIG. 1. Cellular growth of strain MT112 (dnaK7). Growth in M9CA was monitored by measuring the optical density (OD). (A) Initially cells were grown at  $30^{\circ}C$  ( $\Box$ ); then one portion of the culture ( $\blacksquare$ ) was transferred to  $43^{\circ}C$  as indicated by the arrow. (B) Syntheses of DNA, RNA, and protein in cells of MT112. For continuous labeling, cells were grown in M9CA medium containing radioisotopes at 30°C for 5 h. One portion was shifted to 43°C at midlog phase, as indicated by an arrow. The radioisotope incorporation at the time of temperature shift ([<sup>3</sup>H]thymine, [<sup>3</sup>H]uracil, and [<sup>14</sup>C]leucine: 1,400, 3,800, and 2,100 cpm, respectively) was designated as 100%, and relative residual incorporation is shown. Symbols: (O)  $[^{3}H]$ thymine; ( $\bigcirc$ )  $[^{3}H]$ uracil; ( $\triangle$ )  $[^{14}C]$ leucine; (---) relative incorporation of radioisotopes at 30°C.

ment. The mutant is conditionally defective in DNA, RNA, and protein syntheses. The temperature sensitivity phenotype is considered to be due to a single mutation, judging from the following findings. (i) Spontaneous revertants were obtained from brain heart infusion broth cultures. (ii) Temperature sensitivity of the macromolecular synthesis and inability to propagate phage lambda were not separable by P1 transduction.

Another temperature-sensitive dnaK mutant (groPAB756) from E. coli K-12 was originally selected for inability to propagate phage lambda (7). It was confirmed in the present study that these mutant cells are conditionally defective in DNA and RNA syntheses as seen in our mutant (MT112). When E. coli K-12 dnaK7 transductants were tested, they were unable to propagate phage  $\lambda$ . Therefore, two different selective procedures are available for the isolation of dnaK mutants.

Since it is known in general that other dna(Ts) mutants are specifically defective in DNA synthesis at nonpermissive temperature, the block of RNA synthesis expressed by the dnaK(Ts) mutants seems to be very characteristic. The activity of RNA polymerase extracted from strain MT112 was normal at 43°C, excluding the loss of enzyme activity as responsible for the block of RNA synthesis in the dnaK mu-



FIG. 2. Rate of DNA, RNA, and protein syntheses of MT112 by pulse-labeling. Cells were grown in M9CA medium, and at midlog phase they were shifted up to  $43^{\circ}$ C (zero time). Aliquots were withdrawn and further incubated at  $43^{\circ}$ C in the medium containing radioisotopes. The incorporation at zero time ( $[^{3}H]$ thymidine, [ $^{14}$ C]uridine, [ $^{14}$ C]leucine: 112,200, 33,800, and 560 cpm, respectively) was designated as 100%, and the relative incorporation is shown. Symbols: (O) [ $^{3}$ H]thymidine; (**①**) [ $^{14}$ C]uridine; ( $\triangle$ ) [ $^{14}$ C]leucine.



FIG. 3. Rate of DNA and RNA syntheses of strain K7561 by pulse-labeling. Assays were carried out at 43.5°C as described in the legend to Fig. 2. The incorporation at zero time ( $[^{3}H]$ thymidine, 56,800;  $[^{14}C]$ uridine, 20,500 cpm) was designated as 100%, and the relative incorporation is shown. Symbols: ( $\bigcirc$ )  $[^{3}H]$ thymidine; ( $\bigcirc$ )  $[^{14}C]$ uridine.

tants. Saito and Uchida (13) estimated the size of the dnaK cistron as  $2.1 \pm 0.4$  kilobases, corresponding to a protein molecule as large as 77,000 daltons.

What is the function of the dnaK gene product? Elucidation of the function might lead us to an understanding of the mechanism of cellular nucleic acid synthesis and of the relationship between host function and multiplication of the lambdoid phages. Though various dna mutants have been isolated so far, the gene products still remain unknown, except in a few cases. DNA replication is currently thought of as a complex series of events in which many proteins operate together functionally and perhaps structurally. Thus, the dnaK gene product may operate as a part of such a complex for DNA synthesis of the cells and the lambdoid phages, in addition to cellular RNA synthesis.

#### ACKNOWLEDGMENTS

We are grateful to Yoichi Maruyama, Department of Biology, Tokyo Metropolitan University, for reading the manuscript and making useful comments. We also thank Sohei Kondo (Osaka University) for providing the strains of *E. coli* B used in this work and Hisao Uchida (The University of Tokyo) for providing the *dnaJ* and *dnaK* mutants of *E. coli* K-12 and phage lambda carrying the *dnaK* and/or the *dnaJ* gene.

This work was supported in part by a Grant-in-Aid for Scientific Research (238032) from the Ministry of Education of the Japanese Government.

#### LITERATURE CITED

- Bachmann, B. J., K. B. Low, and A. L. Taylor. 1976. Recalibrated linkage map of *Escherichia coli* K-12. Bacteriol. Rev. 40:116-167.
- Burgess, R. R. 1969. A new method for the large scale purification of *Escherichia coli* deoxyribonucleic acid

dependent ribonucleic acid polymerase. J. Biol. Chem. 244:6160-6167.

- D'Ari, R., A. Jaffé-Brachet, D. Touati-Schwaltz, and M. B. Yarmolinsky. 1975. A *dnaB* analog specified by bacteriophage P1. J. Mol. Biol. 94:341-366.
- Fangman, W. L., and A. Novick. 1968. Characterization of two bacterial mutants with temperature-sensitive synthesis of DNA. Genetics. 60:1-17.
- Filip, C. C., J. S. Allen, R. A. Gustafson, R. G. Allen, and J. R. Walker. 1974. Bacterial cell division regulation: characterization of the *dnaH* locus of *Escherichia coli*. J. Bacteriol. 119:443-449.
- Georgopoulos, C. P. 1977. A new bacterial gene (groPC) which affects λ DNA replication. Mol. Gen. Genet. 151: 35-39.
- Georgopoulos, C. P., and I. Herskowitz. 1971. Escherichia coli mutants blocked in lambda DNA synthesis, p. 553-564. In A. D. Hershey (ed.), The bacteriophage lambda. Cold Spring Harbor Laboratory, Cold Spring Harbor, N.Y.
- Hirota, Y., A. Ryter, and F. Jacob. 1968. Thermosensitive mutants of *E. coli* affected in the processes of DNA synthesis and cellular division. Cold Spring Harbor Symp. Quant. Biol. 33:677-693.
- Kohiyama, M. 1968. DNA synthesis in temperature sensitive mutants of *Escherichia coli*. Cold Spring Harbor Symp. Quant. Biol. 33:317-324.
- Lennox, E. S. 1955. Transduction of linked genetic characters of the host by bacteriophage P1. Virology 1:190-206.
- Ryu, J. 1978. Pleiotropic effect of a rifampin-resistant mutation in *Bacillus subtilis*. J. Bacteriol. 135:408-414.
- Saito, H., and H. Uchida. 1977. Initiation of the DNA replication of bacteriophage lambda in *Escherichia coli* K-12. J. Mol. Biol. 113:1-25.
- Saito, H., and H. Uchida. 1978. Organization and expression of the *dnaJ* and *dnaK* genes of *Escherichia coli* K-12. Mol. Gen. Genet. 164:1-8.
- Sunshine, M., M. Feiss, J. Stuart, and J. Yochem. 1977. A new host gene (groPC) necessary for lambda DNA replication. Mol. Gen. Genet. 151:27-34.
- Yochem, J., H. Uchida, M. Sunshine, H. Saito, C. P. Georgopoulos, and M. Feiss. 1978. Genetic analysis of two genes, *dnaJ* and *dnaK*, necessary for *Escherichia coli* and bacteriophage lambda DNA replication. Mol. Gen. Genet. 164:9-14.