804 Notizen

Host Factors Involved in the Growth of Microvirid Phage α 3

Akira Taketo

Department of Biochemistry, School of Medicine, Kanazawa University, Kanazawa

Z. Naturforsch. **33 c**, 804-805 (1978); received July 24, 1978

Microvirid Phage, Replication Factors, dna Mutants, Single-stranded DNA

Host factors involved in the growth of microvirid phage $\alpha 3$ were determined using various replication mutants of Escherichia coli. The viral multiplication was dependent on functional products of dnaE, dnaF(nrdA), dnaG, and dnaZ genes. Host functions directed by dnaA, so-called dnaH, dnaI, and dnaP genes were dispensable for the viral growth. In contrast with $\varphi X174$ and G4, $\alpha 3$ would grow sufficiently in dnaB and dnaC(D) mutants. The viral growth was not significantly affected by host polAts, seg, and groPC mutations.

Introduction

Recently, considerable diversities have been detected among microvirid (isometric) phages, concerning host factor dependence, immunological relationship, and host range. Thus $\varphi X174$ members (e. g. \varphi A, S13, G6) infect E. coli C and require host dnaB and dnaC(D) functions for their growth [1-3]. These two host functions are essential for immunologically unrelated phage G4 [4]. Moreover, G13 and G14, which can infect E. coli B as well, rely on dnaB and dnaC(D)activities [5]. On the other hand, neither dnaB nor dnaC(D) gene product is required for multiplication of K12-specific group (including φK and St-1) [6] which is immunologically remote from φX174. Assignment of host factors required for these phages is important for elucidation of mechanisms of viral replication and evolution. This report describes the host factor reliance of a3 [7] which is infective to E. coli C and B and immunologically somewhat related to St-1 but not to $\varphi X174.$

Materials and Methods

E. coli C-N27 polA4113 [8] was obtained from Dr. T. Okazaki. The sources of other E. coli strains

Requests for reprints should be sent to Dr. Akira Taketo, Department of Biochemistry II, School of Medicine, Kanazawa University, Kanazawa, Ishikawa 920, Japan. used were previously described [1-6, 9-12]. Unless otherwise specified, bacteria were grown in a nutrient broth, at 30 °C, with shaking. Phage a3, originally provided by Dr. D. E. Bradley, was propagated on $E.\ coli$ C and partially purified by differential centrifugation. Single-stranded viral DNA (SS) and double-stranded replicative-form DNA (RF) of a3 were prepared as previously described [13]. Infection experiments with a3 phage were performed as described for φ K[6]. Bacterial strains resistant to a3 were transfected with a3 SS or RF, after Ca²⁺-treatment [1]. The free phage titer was determined using $E.\ coli$ C as the indicator.

Results and Discussion

Effects of various host mutations on the growth of a3 are summarized in Table I. Like other microvirid phages, a3 could grow sufficiently in dnaA cells. Moreover, the viral growth proceeded normally in dnaB and dnaC(D) mutants, at 43 °C. On the other hand, multiplication of φ X174, φ A,

Table I. Growth of α3 in replication mutants of E. coli.

Strain	Phage yield		
	43 °	33 °	43 °/33 °
C dna ⁺	(2.4×10 ⁶)a	2.3×10 ⁶	1.0
C2307 $dnaA$	(4.2×10^7) a	3.2×10^{7}	1.3
LD312 $dnaB$	1.9×10^{7}	6.6×10^{6}	2.9
LD332 $dnaC(D)$	$(1.7 \times 10^6)^a$	1.6×10^{6}	1.1
LD301 dnaE	1.0×10^{4}	2.5×10^{5}	4.0×10^{-2}
$JG42 \ dnaF$	(1.1×10^5) b	(4.7×10^6) b	2.3×10^{-2}
C2309 $dnaG$	8.0×10^{3}	7.0×10^{5}	1.1×10^{-2}
HF4704S dnaH	(7.3×10^5) a	7.2×10^{5}	1.0
WM301-208 dnaI	$(1.9\times10^7)^{a}$	1.3×10^{7}	1.5
KM107 dnaP	$(7.7 \times 10^6)^{c}$	9.1×10^{6}	8.5×10^{-1}
AX727 dnaZ	(1.6×10^3) b	$(1.4 \times 10^6)^{\rm b}$	1.1×10^{-3}
C727 $dnaZ$	1.5×10^{4}	2.2×10^{6}	6.8×10^{-3}
C-N27 polA 4113	6.9×10^{6}	2.4×10^{7}	2.9×10^{-1}
BT4113 polAts	$(3.3\times10^5)^{\rm b}$	(5.2×10^5) b	4.4×10^{-1}
KS268 ligts7	(1.8×10^4) b	$(3.0 \times 10^6)^{b}$	6.0×10^{-3}
BW2001 xth-11	$(2.0 \times 10^5)^{\rm b}$	$(2.6 \times 10^6)^{\rm b}$	7.7×10^{-2}
PB213 seg ⁺	(3.4×10^3) b	(6.6×10^3) b	5.2×10^{-1}
PB1022 seg-2	(1.1×10^3) b	$(3.8 \times 10^3)^{\rm b}$	2.9×10^{-1}
PB1022 seg-2	$(2.3 \times 10^5) \mathrm{d}$	$(3.7 \times 10^4) d$	6.2
C600 gro+	(3.4×10^4) b *	$(4.1 \times 10^4)^{b}$	8.3×10^{-1}
MF634 groPC259	(1.3×10^4) b *	$(2.6 \times 10^4)^{\rm b}$	5.0×10^{-1}
C600 groPC756	(9.9×10^3) b *	$(7.1 \times 10^4)^{b}$	1.4×10^{-1}
C600 groPC756	$(1.9 \times 10^5)^{d}$ *	(5.3×10^5) d	3.6×10^{-1}

a Cells were grown for 60 min at 43 °C before infection.

b Phage yield was determined by transfection of SS DNA to the Ca²⁺-treated bacteria.

 $^{^{\}rm c}$ Cells were grown for 100 min at 43 $^{\circ}{\rm C}$ prior to infection. $^{\rm d}$ Phage yield was determined by transfection of RF DNA to

the Ca²⁺-treated cells.

* Incubation temperature was 43.5 °C.

Notizen 805

S13, and G6 was clearly restricted in these hosts, at the high temperature. In dnaE, dnaF(nrdA), and dnaG mutants, growth of a3 was thermosensitive, indicating involvement of host DNA polymerase III, ribonucleotide reductase, and primase in the viral replication. Yield of a3 was not reduced at 43 °C in HF4704S "dnaH" which was recently shown to carry double mutations in dnaA gene and in some step for utilization of exogenous thymine [14]. (Among enzymes participated in thymine utilization, thymidine phosphorylase and purine nucleoside phosphorylase were not particularly defective in HF4704S strain.) At 43 °C, replication of a3 was not significantly affected in strains WM301-208 dnaI and KM107 dnaP. Furthermore, growth of $\varphi X174$ was not thermosensitive in the dnaP strain (unpublished observation). In contrast, multiplication of a3 was distinctly thermosensitive in AX727 dnaZ mutant transfected with a3 SS, as well as in C727 dnaZ cells infected with intact a3 phage. Functional product of dnaZ gene is essential for all microvirid phages thus far tested. It must be noted here that host functions directed by dnaE, dnaF(nrdA), dnaG, and dnaZgenes are essential for λ phage as well.

Although multiplication of a3 was only slightly affected by $polA^{ts}$ mutation, the phage yield was markedly reduced at 43 °C in strain KS268 ligts7, indicating participation of host DNA ligase in the viral growth. In strain BW2001 xth-11, growth of a3 was considerably reduced at 43 °C. However, whether exonuclease III, product of the xth gene, is directly involved in a3 replication process (e. g. removal of primer RNA) or not is presently un-

known. Replication of $\alpha 3$ was not significantly affected at 43 °C in PB1022 seg-2 cells, whereas growth of λ phage was, as reported by Jamieson and Bergquist [15], abortive in this mutant at 42 °C -43 °C (data not shown). In contrast with λ , $\alpha 3$ could grow normally in groPC mutants, at 37 °C. Yield of $\alpha 3$ in strains MF634 groPC259 and C600 groPC756 was not particularly reduced even at 43.5 °C, as compared with the yield at 33 °C. Unlike λ phage, $\alpha 3$ requires host functions specified by rep gene (data not shown).

Host range of $\alpha 3$ is similar to that of G13 and G14: these phages can infect $E.\ coli$ BB and BB5 but not to BB2, BB1 BB20, BB7, BB9, BB4 and BB12 (unpublished observation). Host factor requirement of $\alpha 3$ nevertheless differs from that of G13 and G14, and closely resembles that of K12-specific phages St-1 and φK [6]. Moreover, φKh -1 (a host range mutant of φK) [2] can infect $E.\ coli$ K12, C, BB, BB5, BB1, and BB7, whereas host factor reliance of this phage is essentially similar to St-1, φK , and $\alpha 3$. These results are consistent with the fact that $\alpha 3$ is immunologically related to St-1 group [7] but quite different G13 and G14.

Dispensability of dnaB and dnaC(D) functions for a3 predicts that this phage, like G4 [16], may have a unique origin of synthesis of complementary (minus) DNA strand. Furthermore, in synthesis of viral (plus) DNA strand, a3 system is by far simpler than G4 which, like φ X174, requires dnaB and dnaC(D) genes products for this reaction [4]. Determination of nucleotide sequence of the origin of plus strand synthesis is essential for characterization of this unique replication system.

- [1] A. Taketo, Molec. Gen. Genet. 122, 15 (1973).
- [2] A. Taketo, J. Gen. Appl. Microbiol. 23, 29 (1977).
- [3] A. Taketo, J. Gen. Appl. Microbiol. 23, 85 (1977).[4] K. Kodaira and A. Taketo, J. Biochem. 83, 971 (1978).
- [4] K. Kodaira and A. Taketo, J. Biochem. 63, 971 (1978).
 [5] A. Taketo and K. Kodaira, Biochim. Biophys. Acta, in press.
- [6] A. Taketo, Molec. Gen. Genet. 148, 25 (1976).
- [7] D. E. Bradley, Canad. J. Microbiol. 16, 965 (1970).
- [8] T. Ogawa, S. Hirose, T. Okazaki, and R. Okazaki, J. Molec. Biol. 112, 121 (1977).
- [9] A. Taketo, Molec. Gen. Genet 139, 285 (1975).

- [10] K. Kodaira and A. Taketo, Biochim. Biophys. Acta 476, 149 (1977).
- [11] A. Taketo and K. Kodaria, Molec. Gen. Genet. 162, 151 (1978).
- [12] A. Taketo, Z. Naturforsch. 32 c, 429 (1977).
- [13] A. Taketo, J. Gen. Appl. Microbiol. 24, 51 (1978).
- [14] P. L. Derstine and L. B. Dumas, J. Bacteriol. 128, 801 (1976).
- [15] A. F. Jamieson and P. L. Bergquist, Molec. Gen. Genet. 150, 171 (1977).
- [16] J. C. Fiddes, B. G. Barell, and G. N. Godson, Proc. Nat. Acad. Sci. USA 75, 1081 (1978).