CHROMOSOME STRUCTURE IN PHAGE T4, II. TERMINAL REDUNDANCY AND HETEROZYGOSIS*

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In a previous communication¹ it was suggested that the chromosome of each particle of phage T4 is terminally redundant, and that after several rounds of replication (following infection), progeny chromosomes arise which have circularly permuted genetic sequences. Among the members of a population of phage particles, the ends of the chromosomes would be randomly distributed over the genome, and a phage particle carrying different alleles at one of its redundant loci would be heterozygous.

Nomura and Benzer² have reported that heterozygotes for rII deletion mutants (from mixed infections with r^+ phage) occur with about a third of the frequency of those for rII point mutants. It thus seemed likely that heterozygosity in phage T4 is of at least two kinds. One kind may be the result of heterozygosity within the molecule, as was first suggested by Levinthal,³ whereas another kind may be due to the terminal redundancy described above. Deletion mutants may be unable to form heterozygotes of the internal kind since these would involve violations of duplex complementarity (regardless of their precise molecular structure), whereas point mutants may form heterozygotes of both kinds.

The chromosome of a phage particle will be terminally redundant and heterozygous for marker a provided that (1) the chromosome terminates near marker a and includes marker a, and (2) the last recombinational event, occurring somewhere within the chromosome, has been such that the other end of the chromosome contains marker a^{+} . Terminal-redundancy heterozygotes are formed and lost by the process of recombination; their frequency (as a function of time after infection) will depend on the rate of recombination but not on the rate of replication. In a pool of vegetative chromosomes, the frequency of terminally redundant heterozygotes would be expected to change in time as does the frequency of recombination for very distantly linked markers; it would be expected to increase until an equilibrium value was reached and to remain constant after that. Because of the high frequency of recombination in phage T4, one would expect that the equilibrium value would be reached very early during the latent period.

Internal heterozygotes are most probably formed by an event that results in the recombination of markers on either side of the heterozygous region;³ thus, they may well represent the primary products of recombination. They would be expected to disappear upon the semiconservative replication of the chromosome bearing them. The chromosome of a phage particle will be internally heterozygous for marker a provided that (1) a recombinational event leading to internal heterozygosis will have occurred at the site of marker a, and (2) the chromosome is incorporated into a mature particle before it undergoes replication. Internal heterozygotes are formed by recombination and are lost by replication: their frequency (as a function of time after infection) will depend both on the rate of recombination and on the rate of replication.

The frequency of heterozygotes, as a function of time after infection, has been observed to remain constant.4 In a standard cross, the formation and disappearance of heterozygotes is at equilibrium throughout the period during which phage are withdrawn from the vegetative pool. If the rate of replication could be substantially decreased during this period, the frequency of internal heterozygotes would be expected to rise.

The thymidine analogue fluorodeoxyuridine (FUDR) inhibits DNA synthesis in phage-infected bacteria' but does not interfere with recombination.6 It would be expected that in phage-infected, FUDR-treated bacteria, the frequency of internal heterozygotes would increase as a function of time, whereas the frequency of terminal-redundancy heterozygotes would remain constant. A test of this prediction forms the substance of this communication.

Materials and Methods.—The phage strains used were T4B standard type and mutants r607, $r287$, $r205$, $r1364$, $rW8-33$, $rH23$, $r168$ (all obtained from Dr. S. Benzer); T4D standard type and mutants $tsN30$ and $rEDb42$ (obtained from Dr. R. S. Edgar); and T4 h^{2+} (obtained from Dr. G. Stent). Stent's strain was derived by him by crossing T4 and T2, crossing a progeny strain with the T2 host range to T4, and repeating the process a number of times. The h^{2+} locus of Stent's strain was incorporated by us into the appropriate T4B and T4D strains by exposing T4 h^{2+} phage to a dose of ultraviolet light sufficient to produce about 20 phage lethal hits, crossing the UV-treated T4 h^{2+} with the appropriate phage strain and selecting a (rare) progeny strain with the h^{2+} host range. The standard-type T4 strain is designated h^{4+} .

The bacterial strains used were *Escherichia coli* B, $B/2$, $B/4$, $S/6$, and K12 112-12(λ h), hereafter called K.

Media: Broth: H20, ¹ liter; bacto-tryptone, 10 gm; NaCl, 5 gm. Tryptone bottom agar: broth with 1.1% bacto-agar. Tryptone top agar: broth with 0.7% bacto-agar. EHA bottom agar: tryptone bottom agar, ¹ liter; sodium citrate -2H20, 1.1 gm; glucose, 1.3 gm. EHA top agar: tryptone top agar, 1 liter; sodium citrate $2H_2O$, 1.1 gm; glucose, 3.0 gm. M-9: H_2O , ¹ liter; Na2HPO4, 7 gm; KH2PO4, 3 gm; NH4Cl, ¹ gm; supplemented after autoclaving with glucose, 4 gm ; MgSO₄, $10^{-3} M$.

Fluorodeoxyuridine was a gift from Dr. R. Duschinsky of Hoffman-LaRoche, Inc.

Phage stocks were prepared by stabbing a 4- to 5-hr plaque with a needle and rinsing the needle into 20 ml of B cells grown to ^a concentration of about ¹⁰⁸ cells per ml in aerated M-9 at 37°. The stocks were lysed by the addition of chloroform after about 5 hr of aeration. Stocks of is mutants were prepared at 30°. DNase was added to the lysates, and the lysates were centrifuged at low speed to remove bacterial debris. Stocks of h^{2+} phage were incubated at 65° for $2^{1/2}$ min immediately after lysis and centrifugation.7

Assay of phage: For platings on K or on B, the bacteria were grown to a concentration of 2 \times $10⁸$ cells per ml in aerated broth at 37 $⁹$, and 0.5-ml amounts were used per 2.5 ml of tryptone top</sup> agar to overlay plates containing 36 ml of tryptone bottom agar.

For platings on $B/2 + B/4$, appropriate amounts of phage were added to B grown to a concentration of 2×10^8 cells per ml in aerated broth at 37°. After 5 min of incubation at 37°, one part of the phage-B mixture was added to two parts of an equal mixture of B/2 and B/4, each grown to a concentration of 2×10^8 per ml in aerated broth at 37°. One part of the final mixture was added to five parts of melted tryptone top agar at 45° , and 3.0-ml samples were immediately distributed to as many plates containing 36 ml of bottom tryptone agar as were necessary.

For platings on S/6, the bacteria were grown to a concentration of 2×10^8 cells per ml in aerated broth at 37°, chilled, and concentrated tenfold by centrifugation. Two drops of the concentrated bacteria were added to ² ml of EHA top agar, 0.05 ml of the appropriate dilution of phage was added, and the mixtures were poured on plates containing ³⁶ ml EHA bottom agar. In some cases phage were first adsorbed to concentrated B and were then diluted and plated on S/6.

Results.—The frequency of heterozygotes as a function of time: The frequency of

heterozygotes for point and deletion rII mutants was measured at various times after infection in the presence of FUDR.

Since little DNA synthesis takes place in the presence of FUDR, the removal, through maturation, of the few DNA molecules in the vegetative pool needs to be prevented. This can be accomplished by the addition of chloramphenicol (CAM) to the infected bacteria at about 9 min after infection. The chloramphenicol is removed at various times after infection, and the infected bacteria remain in the presence of FUDR for (usually) another ⁶⁰ min and are then lysed by the addition of chloroform. The infected bacteria are thus always in the presence of FUDR.

The frequency of heterozygotes as a function of time is illustrated in Figure 1. As expected, the frequency for point rII mutants increases, that for deletion rIl mutants stays constant.

FIG. 1.-The frequency of heterozygotes after various periods of incubation in FUDR. 8 Bacteria of strain B were grown to a concentration of ¹⁰⁸ per ml in aerated M-9 supple-mented with 0.5% bacto-casamino-acids, at ⁷ 37°. To these cultures 20 μ g of tryptophane
per ml, 4×10^{-6} M FUDR, and 2×10^{-4}
M uracil were added just before infection
with an average of five phage particles of each
type per bacterium. Nine min after infec added to the cultures, and at various times $\begin{array}{c} \n\ddots \\
\end{array}$ $\begin{array}{ccc} \n\ddots \\
\end{array}$ + chilled and washed twice by centrifugation, $\frac{1}{2}$ chilled and washed twice by centritigation, $\frac{1}{2}$ results and the first and the second results of $\frac{1}{2}$ r $\frac{1}{2}$ r being with M-9 containing bacto-casaminoacids, uracil, and FUDR, but no chlor-
amphenicol. The *time of removal of chlor*amphenicol refers to the time, in minutes, after infection at which the cultures were chilled,
before centrifugation. The cultures were before centrifugation. The cultures were \bigcup_{ρ^0} o \bigcup_{ρ^0} o then further incubated with aeration at 37° , and were subsequently lysed by the addition of chloroform, the time of lysis indicating the 10 30 60 120 180
time in minutes after infection (omitting the 10 35 90 120 180 240 time in minutes after infection (omitting the 35 90 ¹²⁰ ¹⁸⁰ 240 time required for centrifugation). The lysates were plated on $S/6$, and the frequency of π_{IME} or LYSIS mottled plaques was determined after overnight incubation.

The frequency of recombinants as a function of $\overline{}$ time: The frequency of recombinants is expected to increase in the presence of FUDR and chloramphenicol in the same way as did the $\frac{3}{5}$ are frequency of heterozy gotes. That this is so is pected to increase in the presence of FUDR and chloramphenicol in the same way as did the $\frac{2}{5}$ frequency of heterozygotes. That this is so is $\frac{1}{8}$ $\frac{1}{1}$ shown by the results presented in Figure 2.

The frequency of heterozygotes in the presence $\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{1+\frac{1}{$ and absence of $FUDR$: The frequency of heterozygotes formed by a number of different FIG. 2.—The frequency of recomdeletion rII mutants and by point rII mutants
was compared In each case FIIDR was added bation in FUDR. The procedure used was compared. In each case FUDR was added bation in FUDR. The procedure used
maar the time of infection, chloramphenicol was exactly as described for Fig. 1,
except that platings were on B and near the time of infection, chloramphenicol was added 9 min after infection and removed at \overline{X} . Plaques appearing on plates containing K represent r^+ ; those ap-
about 120 min after infection, and the infected pearing on B represent all types.

bacteria were lysed about 60 min after the removal of chloramphenicol. Control cultures were prepared in the absence of FUDR. The frequency of heterozygotes increased in the presence of FUDR for each of the point mutants examined, but remained the same as the control frequency for the deletion mutants (Table 1).

	FREQUENCY OF r/r ⁺ HETEROZYGOTES IN THE PRESENCE AND ABSENCE OF FUDR				
Nature of	Infecting	Mottled Plaques in the Absence of FUDR Number		Mottled Plaques in the Presence of FUDR Number	
r mutant	phages	Per cent	counted	Per cent	counted
Point	$r607 \times r^+$	1.0	45	7.1	142
	$r287 \times r$ ⁺	0.7	42	5.9	29
	$r205 \times r$ ⁺	0.9	45	3.0	91
Deletion	$r1364 \times r$ ⁺	0.5	24	0.6	44
	$rW8-33 \times r^+$	0.4	34	0.4	31
	$rH23 \times r^+$	0.5	42	0.4	34

TABLE ¹

Bacteria of strain B were grown to a concentration of 10^s per ml, in aerated M-9 supplemented with 0.5% bacto-casamino-acids, at 37°; 20 µg of tryptophane per ml were added to the bacteria just before the phage particle

The behavior of h^{2+}/h^{4+} heterozygotes: Genetic markers (other than rII deletions) that would not form internal heterozygotes seemed very desirable for a number of experiments described below and elsewhere. The host-range difference between phages T2 and T4 is under control of allelic markers; no recombinants possessing a combination of host ranges have been observed.8 That the host-range difference might be due to "compiex" genetic changes was suggested by the observations that a change from T2-like to T4-like host range (or vice versa) could not be induced by treatment with mutagens,⁹ and that the frequency of h^{2+}/h^{4+} heterozygotes was low⁸ and of the order observed by Nomura and Benzer² for deletion mutants.

It seemed important to determine whether the h^{2+}/h^{4+} heterozygotes were similar to those formed by deletion mutants. The frequency of heterozygotes among the progeny phage produced in bacteria mixedly infected with h^{2+} and h^{4+} phage and incubated with and without FUDR was therefore measured. As indicated in Table 2, the frequency was similar in the presence and absence of FUDR, indicating that h^{2+}/h^{4+} heterozygotes behave in the manner expected for terminal-redundancy heterozygotes. ¹⁰

Double heterozygosis: Plaques that contain two types of phage particles could be formed either by a heterozygous phage particle or else by a clump of several parti-

TABLE 2

The procedure was exactly as described for Table 1, except that the lysates were plated on $B/2 + B/4$ fewer than 75 plaques. The frequency of clear plaques was determined after overnight incubation. Each plate contained fe

cles. In order to determine whether clumps of particles are responsible for the formation of plaques containing both h^{2+} and h^{4+} particles (h^{2+}/h^{4+}) plaques), a cross of $h^{2+r} \times h^{4+r}$ r168 was performed. The marker r168 is a small deletion that is about 20 units distant from h^{2+} . Among the progeny of this cross, about 30 per cent of h^{2+}/h^{4+} plaques were found to contain both r and r⁺ phage (Table 3)

TABLE ³

ANALYSIS OF h^{2+}/h^{4+} Progeny Plaques after Fractionation through SEDIMENTATION IN A DEUTERIUM GRADIENT

Total number of plaques examined	Number of h^{2+}/h^{4+} plaques found	Frequency of h^{2+}/h^{4+} plaques	Fraction of h^{2+}/h^{4+} plaques that con- tained both r and r^+ phage
4.6×10^{4}	90	2.0×10^{-3}	$0.07\,$
2.3×10^3	3		
2.9×10^3	10	3.4×10^{-3}	0.10
2.1×10^3	15	6.7×10^{-3}	0.32
2.5×10^3	50	2.0×10^{-2}	0.72
2.0×10^4	53	2.7×10^{-3}	0.30

Fractions from the deuterium-gradient sedimentation of the progeny of a cross of $h^{2+}r^+ \times h^{4+}r168$ described
in Fig. 3 were plated on B/2 + B/4; each plate contained fewer than 40 plaques. Clear plaques were picked an

and could thus have been formed by clumps. As will be shown, the actual fraction of spurious heterozygotes is somewhat less than 30 per cent, too low to invalidate the experiment with FUDR presented above. For other purposes, however, it was necessary to isolate the true heterozygotes.

Since it was expected that clumps would sediment more rapidly than single phage particles, the progeny of the cross were centrifuged in a deuterium oxide $gradient^{11}$ and fractions were collected from the bottom of the centrifuge tube (Fig. 3). The frequency of h^{2+}/h^{4+} plaques and the proportion of these that contained both r and $r⁺$ phage were measured for the peak drop and for the more rapidly moving fractions. As shown in Table 3, the frequency of h^{2+}/h^{4+} plaques was con-

F FIG. 3.—The fractionation of r_{program} phage of a cross through sedimentation in a to⁶ deuterium gradient. Bacteria of strain B were grown to a concentration of 108 per ml in aerated broth at 370, chilled, and concentrated by centrifugation. The bacteria were infected at a concentration of 10^9 per ml with an $\qquad \qquad \big|$ average of six phage particles each of $h^{2+}r^+$
and $h^{4+}r168$. After 2 min incubation at 37°, $\qquad 0 \leq$ they were diluted in broth at 37° and were
incubated at that temperature: 30 min after incubated at that temperature; 30 min after
infection, the bacteria were lysed by the $\frac{1}{1}$ addition of chloroform.

A continuous gradient of D20 was pro-duced in a centrifuge tube by mixing broth prepared with D₂O and broth prepared with H_2O . An aliquot of broth containing the $\frac{10}{100}$ x progeny of the cross was delivered to the $\frac{100}{100}$ x x surface of the gradient and the tube was

centrifuged for 15 min at 20,000 rpm in the
SW39 head of a Spinco centrifuge. After 4 3 2 Peak drop x centrifugation, a hole was punched in the $\frac{1}{10}$ $\frac{1}{10}$ bottom of the tube, and individual drops were

collected in tubes containing 0.5 ml of broth. $\frac{10^{3} \left(\frac{1}{20^{3} \cdot 10^{3} \cdot 10^{3$ The contents of tubes were pooled as indicated DROP NUMBER in the figure to yield fractions 1-4.

siderably greater, and a greater proportion of them contained both r and r^+ phage in the more rapidly moving fractions than in the peak drop. In the most rapidly moving fraction examined, 72 per cent of the $h^{2+/h^{4+}}$ plaques contained both r and r^+ particles, whereas in the peak drop, only 7 per cent of the h^{2+}/h^{4+} plaques contained both r and r ⁺ particles.

The fast-moving fraction was heated for a length of time sufficient to inactivate 90 per cent of the phage particles. This treatment drastically reduced the frequency of apparent heterozygotes, an indication that at least a large proportion of them were clumps. Thus the rapidly sedimenting particles forming mixed plaques are mainly clumps, justifying the assumption that those sedimenting normally are authentic heterozygotes. If so, the data of Table 3 show that about 7 per cent of the authentic h^{2+}/h^{4+} heterozygotes are also heterozygous for the r marker, and that only about 25 per cent of the particles forming mixed plaques are clumps.

Recombination for neighboring markers among heterozygotes: Terminally redundant heterozygotes ought to be recombinant with respect to neighboring markers. In a cross of a b $c \times a^+b^+c^+$, for instance, terminal-redundancy heterozygotes that are b/b ⁺ would be expected to have either of two possible arrangements of markers:

$$
b\ c\qquad \quad a^+b^+\quad\quad {\rm or}\quad\quad b^+c^+\qquad\quad a\ b,
$$

both arrangements being recombinant with respect to the markers a and c .

In order to determine whether particles that are terminal-redundancy heterozygotes for a particular marker are recombinant for neighboring markers, a cross of $rEDb42 h^{4+}$ tsN30 $\times r+h^{2+}$ ts+ was performed. The r and ts markers are on opposite sides of the host-range marker and yield about 20 and 17 per cent recombinants, respectively, in crosses with the host-range marker. The progeny from the cross were centrifuged in a deuterium gradient and the contents of the peak drop were plated at 43[°] on $B/2 + B/4$. Only ts⁺ phage (or phage heterozygous for $ts/(s⁺)$ produce plaques under these conditions. Among the plaques appearing at 43° , h^{2+}/h^{4+} heterozygotes were scored with respect to the r marker. A majority of the h^{2+}/h^{4+} heterozygotes that were $ts+$ were r and thus recombinant, in contrast to the phage in the total population (Table 4). This cross was repeated with various other combinations of these same markers in order to show that the result did not depend on the particular alleles used (Table 4). We may thus conclude that h^{2+}/h^{4+} heterozygotes are recombinant with respect to neighboring markers

TABLE 4

ASSORTMENT OF OUTSIDE MARKERS AMONG h^{2+}/h^{4+} Heterozygotes		
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Bacteria of strain B were grown to a concentration of 10⁸ per m lin aerated broth at 37°, chilled, and concentrated by centrifugation. The bacteria were infected at a concentration of 109 per ml with an average of six p

as would be expected on the basis of the terminal-redundancy model. A similar result had been obtained previously by Doermann and Boehner,¹² who examined heterozygotes formed by an r-deletion mutant. In our crosses an appreciable fraction of the h^{2+}/h^{4+} heterozygotes were simultaneously heterozygous for r/r^+ even though clumps had presumably been eliminated by the centrifugal fractionation.

Discussion.-The experiments reported here demonstrate the existence of two types of heterozygotes and thus confirm observations of others.2 Heterozygotes of one type do not accumulate under conditions of limited DNA synthesis, and the behavior of this type of heterozygote confirms an important prediction of the terminal-redundancy model for the chromosome of phage T4.

An alternate model for this type of heterozygote may be considered: phage particles could consist of a whole chromosome and an additional small piece of a chromosome. This model is excluded by the observation presented here that h^{2+}/h^{4+} heterozygotes are recombinant for neighboring markers, and by the results of others12 that deletion-mutant heterozygotes are recombinant for neighboring markers.

Summary.-Point-mutant heterozygotes increase in frequency during phage growth under conditions of limited DNA synthesis but deletion-mutant heterozygotes do not. Heterozygotes for the h^{2+}/h^{4+} marker behave as deletion-mutant heterozygotes do and are recombinant with respect to neighboring markers. These observations support the terminal-redundancy model for the chromosome of phage T4.

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¹⁰ The frequency of h^{2+}/h^{4+} heterozygotes in this experiment is high owing to the presence of the long deletion rH23 in both parental stocks of the cross. The influence of deletions on the frequency of terminal-redundancy heterozygotes will be described in a subsequent communication.

¹¹ We are grateful to Professor Sidney Brenner, who suggested deuterium oxide as ^a convenient medium for this type of centrifugation.

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