# REPLICATION OF VIRAL RNA, XI. SYNTHESIS OF VIRAL "MINUS" STRANDS IN VITRO\*

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The replication of viruses containing single-stranded  $\rm RNA<sup>1</sup>$  as genetic material is accompanied by the formation, within the host cell, of substantial quantities of "minus" strands which are isolated predominantly in a double-stranded, RNaseresistant form (for references, see Billeter et  $al.^{2}$ ). Several lines of evidence indicate that a complex containing "minus" and "plus" strands is an intermediate in viral RNA synthesis. $3-8$ 

An enzyme preparation, RNA replicase, has been obtained from RNA phageinfected E. coli, by Haruna and Spiegelman.<sup>9, 10</sup> A remarkable specificity of the enzyme for the RNA of the inducing virus was demonstrated.<sup>10</sup> This preparation synthesized viral RNA ("plus" strands) in vitro when primed with viral "plus" strands.<sup>11</sup> Clearly, it was of considerable interest to determine whether or not "minus" strands were formed during the course of this synthesis. We have analyzed the RNA synthesized in vitro by  $Q_{\beta}$  replicase and have found that in the early phase of synthesis only "minus" strands could be detected, whereas at later times "plus" strands were synthesized. This sequence of events was similar to that found in vivo, in phage-infected bacteria.<sup>12</sup>

Materials and Methods. $-Q_\beta$  RNA: Phage  $Q_\beta$  was obtained from Professor I. Watanabe, Tokyo. It was grown on E. coli Q 13, obtained from Dr. S. Spiegelman, and purified by a procedure similar to that described for phage MS2.<sup>13</sup>  $Q_{\beta}$  differs in its behavior from MS2 inasmuch as it forms a precipitate band during the CsCl density gradient centrifugation; it redissolves on dilution with water. The RNA was extracted by the procedure of Gesteland and Boedtker<sup>14</sup> and stored as a precipitate in 66% ethanol at  $-70^{\circ}\text{C}$ . Q<sub>B</sub> RNA can be distinguished from MS2 RNA by virtue of its capacity to anneal specifically with denatured double-stranded  $Q_\beta$  RNA but not at all with double-stranded MS2 RNA.<sup>15</sup> For use as a template,  $Q_\beta$  RNA was fractionated on a sucrose gradient, and the peak fractions, corresponding to an  $S_{20,w}$  of about 28S, were pooled and utilized immediately. As shown in Figure 1, such peak fractions contain no detectable amounts of degraded RNA.

Double-stranded  $Q_\beta RNA$ : This was prepared as described for MS2 RNA,<sup>16</sup> with similar yields.  $Q_{\beta}$  replicase: The enzyme preparation used was a gift from Drs. Haruna and Spiegelman. When assayed under the conditions described by Haruna and Spiegelman,<sup>10</sup> 20  $\mu$ l of the solution  $(3.1 \text{ mg protein/ml})$  incorporated 0.45 m $\mu$ mole of UTP into acid-insoluble material, in 20 min at 35°. The same values were obtained both in Dr. Spiegelman's and in our laboratory. The specific template requirement of the  $Q_{\beta}$  enzyme<sup>10</sup> was confirmed.

Preparation of the product of  $Q_\beta$  replicase: Incubations were carried out under the conditions described by Haruna and Spiegelman.<sup>10</sup> The product was extracted twice with 2 vol of phenol (equilibrated with  $0.05$  M Tris HCl buffer, pH 7.6), and the phenol was removed by three ether extractions. The samples were freed of radioactive nucleotides by dialysis for 20 hr against several changes of  $1 \times SSC$  and finally,  $0.1 \times SSC$ . Determination of RNase-resistant radioactive RNA was carried out as described previously.<sup>16, 17</sup>

Double isotope specific dilution assay: This assay was used to determine radioactive viral "plus" and "minus" strands in the presence of radioactive nonviral RNA. The radioactive viral RNA is first converted into <sup>a</sup> double-stranded, RNase-resistant form by annealing with an excess of heat-denatured double-stranded viral RNA.'7 The product is then subjected to the specific dilution assay<sup>3, 17</sup> by heating and reannealing in the presence of an excess of unlabeled "plus"



FIG. 1.-Velocity centrifugation of purified  $Q_{\beta}RNA$ .  $Q_{\beta}RNA$  was purified by sucrose density centrifugation as described in *Methods*. The densitometer tracings were from photo-<br>graphs taken after (a) 8 min, and (b) 16 min of graphs taken after  $(a)$  8 min, and  $(b)$  16 min of centrifugation at 50,740 rpm;  $S_{20,w} = 27.5S$ .

strands. The radioactive "plus" strands are thereby displaced from the duplex and become RNase-sensitive. Radioactivity due to "minus" strands remains RNase-resistant.

In practice, the assay was carried out by distributing, into a series of tubes, aliquots of a mixture containing the C<sup>14</sup>-labeled sample to be analyzed, a small quantity of  $P^{32}$ -labeled viral "plus" strands (as an internal standard), and unlabeled double-stranded viral RNA in excess of the "plus" strands present in the sample mixture. Increasing amounts of unlabeled viral "plus" strands (from 0 to about a 20-fold excess over the double-stranded RNA) were then added to the tubes. The mixtures were heated, reannealed, and the RNase-resistant  $P^{32}$ - and  $C^{14}$ -radioactivities were determined. When a mixture of P<sup>32</sup>-labeled Q<sub>B</sub> RNA and the C<sup>14</sup>-labeled product of Q<sub>B</sub> replicase (40-min incubation) was annealed with double-stranded RNA, but without addition of unlabeled  $Q_0$  RNA, 85% of the C<sup>14</sup>-radioactivity and 80% of the P<sup>32</sup>-radioactivity were converted to an RNase-resistant form. When the reaction was carried out in the additional presence of excess nonlabeled  $Q_\beta$  RNA (Fig. 2), 53% of the C<sup>14</sup>-radioactivity and only 3% of the P<sup>32</sup>-radioactivity was resistant to RNase. A plot of the C<sup>14</sup>-values against the corresponding P<sup>32</sup>-values gives a straight line, with the slope indicating the fraction of C<sup>14</sup>-radioactivity in "plus" strands and the intercept giving the fraction of C<sup>14</sup>-radioactivity in "minus" strands (see *inset*, Fig. 2). The method underestimates "minus" strands by about 5-10%. A detailed description of the assay will be given elsewhere.<sup>18</sup>

*Results.—Synthesis of "minus" strands:*  $Q_{\beta}$  replicase was incubated with radioactive nucleoside triphosphates as described.<sup>10</sup> The integrity of the  $Q_{\beta}$  RNA used as a template was ascertained by ultracentrifugal analysis immediately prior to the experiment (cf. Fig. 1) . Samples of the reaction mixture were withdrawn at different times during incubation, purified, and analyzed by the double isotope specific dilution assay. As shown in Table <sup>1</sup> (expt. 1) and Figure 3, no radioactive "plus"



 $\mathbb{Q}_{\beta}$  replicase by the double isotope specific dilution assay. The product was prepared as RNA (0.1  $\mu$ g), 5  $\mu$ g of double-stranded  $Q_{\beta}$ <br>RNA, and unlabeled  $Q_{\beta}$  RNA as indicated.<br>After heating and annealing, the RNase-resist-<br>ant radioactivities were determined, and ant radioactivities were determined, blank values (RNase-resistant radioactivity are expressed as the fraction of input radioacto determine the fraction of C<sup>14</sup>-radioactivity<br>
in "plus" and "minus" strands (slope and in-<br>
tercept, respectively, of the resulting straight line).



## TABLE <sup>1</sup> ANALYSIS OF THE PRODUCT OF  $Q_\beta$  REPLICASE\*

\* The incubations were carried out as described by Haruna and Spiegelman,<sup>10</sup> using unlabeled ATP and C<sup>1</sup>-labeled UTP, CIP, and GTP, each of specific radioactivity 25,000 cpm/m<sub>*m*mole. In expt. 1,250  $\mu$ g of energine o</sub>

strands could be detected within the first 7 min of incubation even though the presence of 5-10 per cent of the radioactivity in "plus" strands would have been clearly detectable. Of the acid-insoluble radioactivity, 75-80 per cent was found in "minus" strands. Failure to account for all of the radioactive RNA as virus-spe-<br>cific material is probably due to incomplete annealing. Radioactive "plus" strands cific material is probably due to incomplete annealing. were first detected after 15 min of incubation and their amount exceeded that of "minus" strands at <sup>100</sup> min. An amount of RNA equivalent to that of the added template was synthesized by the twelfth minute. A similar experiment was carried out, using as template  $Q<sub>g</sub>$  RNA provided by Drs. Haruna and Spiegelman. This time; only radioactive "minus" strands were assayed for, by heat-denaturing and annealing the product with an excess of unlabeled  $Q_{\beta}$  RNA, and then determining the RNase-resistant radioactivity. The results (Table 1, expt. 2) were comparable to those of the first experiment.

Sucrose gradient centrifugation of the labeled product of  $Q_{\beta}$  replicase, isolated after 40 min of incubation (Fig. 4), showed that a large fraction of the radioactive RNA sedimented around 27S, as does intact viral RNA, in agreement with the report by Haruna and Spiegelman.<sup>19</sup> RNase-resistant RNA sedimented with an  $S_{20,w}$  around 14S, a value similar to that found for virus-specific, double-stranded RNA formed *in vivo*.<sup>4</sup> Annealing of the gradient fractions with "plus" strands Annealing of the gradient fractions with "plus" strands revealed "minus" strands not only in the region corresponding to double-stranded RNA, but also in the 27S region.

Discussion.--Earlier work<sup>3-8</sup> has led to the concept that double-stranded RNA (or a complex of "plus" and "minus" strands that gives rise to double-stranded RNA during the isolation procedure') is an intermediate in viral RNA synthesis, with "minus" strands serving as template for the formation of progeny "plus" strands. The hypothesis demanded that "minus" strand synthesis precede the formation of "plus" strands. This has now been shown to be the case, both in  $vivo^{12}$ and in vitro.





TIME (minutes)<br>
FIG. 4.—Sucrose gradient analysis of the product of Q<sub>B</sub> rep-<br>
licase. An aliquot of the Q<sub>B</sub> replicase product, described in the<br>
legend to Fig. 2, to which 20 A<sub>260</sub> units of MS2 RNA were ad-<br>
FIG. 3.—Sy and "minus" strands by  $Q_{\beta}$ -rep- linear sucrose density gradient (3-20% sucrose in SSC) and licase. The values are cal-<br>centrifuged for 16.5 hr at  $4^{\circ}$  and 21,000 rpm in the Spinco SW<br>culated from the data of Table 25.1 rotor. The gradient was syphoned off from the bottom<br>1, expt. 1. O. . O, Total in-<br>corpora into acid-insoluble material; acid-insoluble radioactivity  $(O---O)$ , (b) acid-insoluble ra-<br>x-x, radioactivity in dioactivity after RNase digestion (x---x), and (c) radioactive "minus" strands;  $\bullet$ -- $\bullet$ , "minus" strands( $\bullet$ ... $\bullet$ ); these were determined by heating radioactivity in "plus" strands. an aliquot of each fraction with 167  $\mu$ g/ml of  $Q_\beta$  RNA in 2.5 × SSC, for 3 min at 120°C and determining the acid-insoluble radioactivity after RNase di-gestion. The blank (RNase-resistant radioactivity of the enzymatic product after heat denaturation; less than  $1\%$  of the total radioactivity) was subtracted from all values.

Formation of "minus" strands in vivo was detectable several minutes prior to that of "plus" strands. Furthermore, the rate of "minus" strand synthesis approached a constant value at about 15 min, while the rate of "plus" strand formation increased up to 22 min after infection, suggesting separate control mechanisms for the two processes.12

The present experiments with  $Q_{\beta}$  replicase showed that in the first minutes of incubation predominantly, if not exclusively, "minus" strands were formed, and that subsequently "plus" strands were synthesized. The second phase of synthesis has been detected by Spiegelman et  $al$ .<sup>11</sup> with use of infectivity assays. It is remarkable that the major part of the "minus" strands formed early in the in vitro reaction occur in an RNase-sensitive form (Table 1) and therefore appear not to be part of a hydrogen-bonded double helix. The sedimentation profile of the "minus" strands (Fig. 4) suggests that free single-stranded "minus" strands may occur.

Replication of viral RNA thus appears to involve two steps: (a) synthesis of "minus" strands, with "plus" strands serving as template, and (b) synthesis of "plus" strands, with "minus" strands as template. A complex of "plus" and "minus" strands, detected as double-stranded RNA, has been shown to occur as intermediate in viral RNA synthesis in vivo.<sup>4, 7, 12</sup> Step (a) has been studied in vitro by August and his collaborators,<sup>20, 21</sup> who purified a virus-induced polymerase from  $E.$  coli infected with su-11,<sup>22</sup> a mutant of the RNA phage  $f_2$  which causes the formation of excessive amounts of "minus" strands (in a double-stranded form) in the infected host.<sup>23</sup> Step (b) was demonstrated with use of an enzyme-template complex RNA synthetase<sup>24</sup> that synthesized almost exclusively" plus" strands in vitro.<sup>17</sup> As shown in this paper, Haruna and Spiegelman's RNA replicase is capable of carrying out both synthetic steps. By their elegant analysis of RNA phage mutants, Lodish and Zinder<sup>8, 23</sup> have shown that the two steps of RNA replication can be genetically dissociated. It remains to be seen whether the two enzymatic activities can be separated.

Summary.—When incubated with nucleoside triphosphates and  $Q_6$  RNA as template,  $Q_{\beta}$  replicase synthesizes predominantly, if not exclusively, "minus" strands in the early phase of incubation and, later on, mainly "plus" strands. Part but not all of the "minus" strands occur in a double-stranded form.

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<sup>1</sup> Abbreviations: RNA, ribonucleic acid; ATP, GTP, UTP, and CTP, the <sup>5</sup>'-triphosphates of adenosine, guanosine, uridine, and cytidine, respectively; RNase A, pancreatic ribonuclease A; Tris, tris(hydroxymethyl)aminomethane; SSC, 0.15 M NaCl, 0.015 M sodium citrate, pH 7; cpm, counts per minute;  $A_{260}$ , absorbance at 260 m $\mu$ , 1.0 cm light path; "plus" strands are defined as viral RNA strands of the parental type as opposed to "minus" strands which have the complementary base sequence.

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