Sedimentation Coefficients of the Virions of Soil-Borne Wheat Mosaic Virus

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ABSTRACT


The sedimentation coefficient of virion I of soil-borne wheat mosaic virus was estimated to be 219S, the same, within error, as the sedimentation coefficient of the dimer of virion II, estimated to be 218S. The monomer of virion II sedimated at 177S, and was 138 nm long. Another strain of the virus had a virion II (designated IIb) that was 92 nm long and sedimated at 159S. The two virus strains coexist in some fields.

MATERIALS AND METHODS

Soil-borne wheat mosaic virus was obtained in April from infected wheat from several fields near Lincoln, Nebraska. One strain, SV17, was maintained on Michigan Amber wheat plants (Triticum aestivum L.) in growth rooms by mechanical inoculation by C. Powell and used by him (16). Virions of other strains were purified from infected wheat collected from the field in April and stored at −20 C until purification. Virions were purified by the following modifications of our earlier method (17). One percent Triton X-100 was used instead of 0.1% Igepon T-73, and high-speed pellets were suspended in pH 7.5, 0.05 M sodium orthoborate. In a final purification step, the virions in 0.05 M sodium orthoborate, pH 7.5, were layered over a gradient prepared from 4 ml each of 50, 100, 150, and 200 mg of sucrose/ml in 0.05 M, pH 7.5, sodium orthoborate. The gradients were centrifuged for 3 hr at 25,000 rpm (81,500 average g) at 5 C in the Beckman SW27 rotor. The pellet was suspended in pH 7.5, 0.05 M sodium orthoborate and stored at −20 C after addition of 5% ethylene glycol.

The sedimentation coefficients of individual virions were estimated essentially as described for barley stripe mosaic virus (2). The sucrose for the gradient columns for SBWMV was dissolved in 0.1% Igepon T-73 to inhibit virion aggregation (9). Linear-log gradients were centrifuged in either the Beckman SW41 or SW27 rotors at 5 C (3). Analysis of the gradients, collection of samples, gel electrophoretic analysis of the RNA, and interpretation of the results was as described earlier (2). Peacock and Dingman's (15) 0.09 M tris borate buffer was used for some of the RNA analyses.
Molecular weights of RNAs were estimated by gel electrophoresis of formaldehyde-treated RNA as described by Lane (13) except that gels contained 2.2% acrylamide and 0.5% agarose. After electrophoresis gels were soaked 1.0 hr in water, stained overnight in "stains-de-scribed by Lane (13) except that gels contained 2.2% all" (7), and destained in water in a dim light.

Tobacco mosaic virus RNA at 2.05 × 10^6, and E. coli RNA at 0.56 and 1.1 × 10^6 were molecular weight standards in gel electrophoresis of RNA (6, 12). Virions of TMV and southern bean mosaic virus with s_{20, w} values of 190S and 115S, respectively, were sedimentation standards in sucrose density-gradient centrifugation of SBWMV (3). Purified southern bean mosaic virus was used, but the TMV was a fresh, clarified extract of infected leaves at dilutions of 1:20 to 1:50 (g of leaves/ml).

Virions were prepared by two methods for length measurements by electron microscopy. In the first, a freshly cut leaf was immersed for a few seconds in a drop of water on a specimen grid. The drop was dried for 20-30 min, a drop of negative stain was applied, and the sample was examined in a RCA 3G electron microscope at about × 15,000. In the second procedure, a 0.5- to 1.0-cm portion of a leaf was chopped with a razor blade in a large drop of water or salt solution on a Parafilm membrane to give a slightly yellowish-green drop. A specimen grid with carbon-backed parlodion film was floated film side down on the drop for 10 min. Excess liquid was removed, and the grid was immediately stained and examined as before. A solution of 1.5% sodium phosphotungstate and 0.5% ammonium vanadomolybdate was usually used for the negative stain. Purified virus for width measurements was prepared according to Horne and Pasquali-Ronchetti (11). Widths were measured from edge to edge across three or more rods lying side-by-side. The magnification of the microscope was calibrated with a replica of a diffraction grating.

**RESULTS**

Virions from several sources of SBWMV were examined by sedimentation and the two showing least aggregation were selected for further investigation. The sedimentation of a typical two component strain of SBWMV is shown in Fig. 1(B, D). The RNA analysis by gel electrophoresis of the samples collected from the gradient column of Fig. 1-D are shown in Fig. 2. It is apparent that the two peaks on the absorbance scanning pattern of SBWMV are the monomer and dimer of virion II. Figure 1-C, shows a mixture of SBWMV and TMV cosedimented in the same column. The sedimentation coefficients of the monomer and dimer of virion II were calculated from depths measured from a gradient column such as Fig. 1-C, with TMV serving as an internal standard. The slope of the line of the log-log plot (3) was taken from the depths of southern bean mosaic virus and TMV measured in a companion gradient (Fig. 1-A). When values for the monomer and dimer of virion II had been accumulated for several gradients such as shown in Fig. 1-C, an average value for the sedimentation coefficient of these two particles was calculated (Table 1). The monomer and dimer peaks of virion II were then used as internal standards for the calculation of the sedimentation coefficient of virion I from patterns such as that of Fig. 1-D, with the position of virion I determined by RNA analysis (Fig. 2). Within the error of the method, virion I sedimented at the same rate (219S) as the dimer of...
virion II (218S) (Table 1).

Strain SV17, which had been maintained by mechanical inoculation for 3 yr, had a third RNA, smaller than RNA II, which was found in a virion sedimenting slower than virion II. This RNA is designated RNA Iib and the virion, virion Iib. The sedimentation pattern of the virions of strain SV17 is shown in Fig. 1-E. Because virion Iib could be detected on absorbance patterns, its sedimentation coefficient was estimated from depths of virions measured on absorbance patterns with the monomer and dimer of virion II serving as internal standards (Table 1).

Three RNAs have been found not only in strain SV17, but also in virions purified from infected plants from certain fields. The RNA pattern of virions purified from wheat from one such field in 1972 is shown in Fig. 3-A. RNA from virions purified from wheat collected from the same field in 1974 did not have a discrete third RNA peak (Fig. 3-B). The results of gel electrophoretic analysis of the formalinized SBWMV-RNA from this field, alone and in combination with TMV RNA and E. coli RNA as internal markers, are shown in Fig. 3-B and -C. Patterns of formaldehyde-treated RNA are shown for the comparison with TMV because untreated TMV RNA often showed multiple zones. Formaldehyde treatment did not change the number of components of SBWMV RNA nor their relative concentrations. Inspection of the stained gels showed a doublet zone of TMV-RNA and SBWMV-RNA I. The doublet was barely resolved in the scanning pattern (Fig. 3-C). From the relative amounts of the two components, the slower migrating component of the doublet was identified as SBWMV-RNA I. The apparent molecular weights of SBWMV-RNA's I and II calculated from gel electrophoretic mobility after formaldehyde treatment are given in Table 1. No difference could be detected in the molecular weights of SBWMV-RNA's I and II from different fields.

Rod-length measurements on strain SV17 confirmed the presence of virion Iib and, in fact, showed that some plants contained virion II and some contained virion Iib (Fig. 4). Measurements made on leaf dips prepared by procedure I (see Materials and Methods) gave a wide distribution of lengths. The results of these measurements did not correlate with virion lengths expected from RNA

![Graph](https://via.placeholder.com/150)

**Fig. 2.** Gel electrophoretic analysis of RNA from consecutive samples collected from the virion zone of the sucrose gradient column D of Fig. 1. Results of analysis of all samples except the first are shown. No RNA was detected in the first sample. The gel was 2.5% acrylamide with 0.5% agarose and the buffer was 0.09 M tris, 0.09 M borate, 2.7 mM EDTA. Gels were stained in Stains-all and scanned at 570 nm.

<p>| TABLE 1. Properties of components of soil-borne wheat mosaic and tobacco mosaic viruses |
|---------------------------------|-----------------|-----------------|----------------|-----------------|</p>
<table>
<thead>
<tr>
<th>Virions</th>
<th>Estimated S\textsubscript{w.20}</th>
<th>Component\textsuperscript{a}</th>
<th>TMV</th>
</tr>
</thead>
<tbody>
<tr>
<td>I</td>
<td>II</td>
<td>Iib</td>
<td>II-dimer</td>
</tr>
<tr>
<td>Estimation of determinations</td>
<td>6</td>
<td>5</td>
<td>4</td>
</tr>
<tr>
<td>Rod length</td>
<td>281 nm\textsuperscript{a}</td>
<td>138 nm</td>
<td>92 nm</td>
</tr>
<tr>
<td>No. of plants</td>
<td>134</td>
<td>714</td>
<td>1,546</td>
</tr>
<tr>
<td>No. of rods</td>
<td>15</td>
<td>9</td>
<td>15</td>
</tr>
<tr>
<td>No. of rods measured</td>
<td>20 nm</td>
<td>20 nm</td>
<td>20 nm</td>
</tr>
<tr>
<td>No. of rods measured</td>
<td>2.10 × 10\textsuperscript{4}</td>
<td>1.4 × 10\textsuperscript{6}</td>
<td>ND\textsuperscript{b}</td>
</tr>
</tbody>
</table>

\textsuperscript{a} The largest component is termed number I. Most strains have I and II, some I and IIb, and some all three.

\textsuperscript{b} Standard error of the mean. This indicates the reproducibility of the measurements, not the absolute accuracy. Based on 190S for TMV and 115S for southern bean mosaic virus.

\textsuperscript{c} Number of rods measured in each particular size class.

\textsuperscript{d} Number of plants from which virions were obtained for length measurements.

\textsuperscript{e} Based on electrophoretic mobility in gels after formaldehyde treatment.

\textsuperscript{f} The 281-nm rods were in the 15 plants containing virion Iib.

\textsuperscript{g} The abbreviation ND = not determined.

analysis. When the leaf-dip procedure was modified to procedure II, measurements showed a much narrower distribution of rod lengths (Fig. 4) and the results correlated with the RNA analysis. It should be noted that plants with virions IIb (92 nm long) contained as many rods that were 281 nm long as did plants with virion II of 138 nm. These 281-nm rods could not have been dimers of the 92-nm rods and are presumed to have been rods of virion I. In procedure II, the leaves were usually chopped in 0.5 M borate, pH 9.0, but 0.5 M borate pH 8.0, distilled

Fig. 3. Gel electrophoresis of RNA from purified SBWMV. Pattern A is of RNA of virus from infected wheat of 1972, not treated with formaldehyde and run in 2.4% acrylamide, 0.5% agarose, with 0.03 M NaH₂PO₄, 0.03 M tris, 1.5 mM EDTA buffer. Arrows indicate (from the left) the position of TMV-RNA, brome mosaic virus (BMV)-RNA II, BMV-RNA III, and BMV-RNA IV in a companion gel. Pattern B is of formaldehyde-treated SBWMV-RNA from virus purified from the same field in 1974. The gel was 2.2% acrylamide and 0.5% agarose and the buffer was 0.02 M NaH₂PO₄, 0.01 M NaH₂PO₄, and 1 mM EDTA. Pattern C is the same as pattern B, except that TMV-RNA and Escherichia coli rRNA were added as internal standards.

Fig. 4. Histogram of length distribution of rods from two plants infected with SBWMV strain SVI7 that had been maintained by mechanical inoculation. Grids were prepared by two procedures (see text for details) from each plant. One plant had 92- and 281-nm rods, the other, 138- and 281-nm rods.
The average lengths of virions I, II, and Ilb of SBWMV are given in Table 1; TMV, measured by chopping leaves as per procedure II, was 281 nm long. Because virion I of SBWMV is the same length as TMV, the more rapid sedimentation of SBWMV can be explained only if it has a larger diameter than TMV. The diameters of the virions of these two viruses were measured by electron microscopy and found to be 20 nm for SBWMV (Table 1) and 16.5 for TMV, which agree well with previously published values (4, 6, 18).

Strain SV17 was lost in a fire in August, 1975 that destroyed the growth rooms in which this strain was maintained. Attempts to recover a strain of the virus with virion Ilb have been unsuccessful. Therefore, it has been impossible to measure certain properties of this virion, such as the apparent molecular weight of RNA Ilb.

DISCUSSION

The fact that virion I of SBWMV sediments at almost exactly the same rate as the dimer of virion II suggests that virion I should be twice as long as virion II. These results and the RNA data support the conclusion that some of the rods of approximately 300 nm seen in the electron microscope are those of virion I.

The molecular weights of SBWMV-RNA's I and II determined by gel electrophoresis after formaldehyde treatment are larger than those determined by centrifugation of formaldehyde-treated RNA (9). The difference for RNA I (2.10 vs. 1.84 × 10^6 daltons) may not be outside the combined errors of the two methods, but the discrepancy for RNA II (1.4 × 10^6 daltons by gel electrophoresis vs. 0.95 × 10^6 daltons by centrifugation) is larger than the expected error. It is probable that formaldehyde did not destroy all the secondary structure, particularly that of RNA II, and that the correct value of its molecular weight is somewhere between 0.95 and 1.4 × 10^6 daltons. The relative sedimentation rates of the virions suggests that RNA II should have half the molecular weight of RNA I.

Present results confirm the report of Tsuchizaki et al. (19) of strains of SBWMV with short rods of different lengths. Such strains apparently coexist in some fields in Nebraska, with short rods (virion Ilb) being a minor and variable component. The results of Tsuchizaki et al. (19) and my results suggest that virion Ilb may have a selective advantage over virion II in virus maintained by mechanical transfer.

Present measurements give lengths of SBWMV virions shorter than those reported previously for the Nebraska virus (1) and the sedimentation coefficients are slightly higher than those reported earlier (9). The differences probably result from differences in procedures, standards, and in sedimentation coefficients assigned to the standards. Such variation is common. Even for TMV, careful electron microscopic measurements do not agree (e.g., see 10, 14). The 138- and 281-nm rods of SBWMV of the present report are probably the same as the 160- and 300-nm rods reported earlier (1), though they were obtained from measurements on virus from a different field and prepared by a different procedure.

The SBWMV resembles TMV in many respects and is slightly related serologically (16); it differs from TMV in having a larger diameter which results in faster sedimentation of rods of equal length. The weight that should be given these various differences and similarities when considering the taxonomy of these viruses is unclear.

LITERATURE CITED
