

Multiple Populations of Double-Stranded RNA in Two Virus-Harboured Strains of *Trichomonas vaginalis*

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ABSTRACT. The existence of six dsRNA segments of *Trichomonas vaginalis* virus was confirmed and the molar mass and relative abundance of these segments were determined by agarose gel electrophoresis with reovirus dsRNA serving as a standard. The M^r 's were 3.5, 3.4, 3.2, 2.5, 1.4 and 0.34 Mg/mol for the two strains studied, the relative abundances, however, were 1.0, 1.4, 3.0, 0.3, 2.7, 4.2 and 1.0, 0.6, 1.7, 0.5, 3.4 1.0 for these strains, respectively. Cell homogenate fractionation showed that all dsRNA segments were associated with viral particles. The data appeared to support the hypothesis of a relationship between viruses of the protozoan *T. vaginalis* and of the yeast *Saccharomyces cerevisiae*.

Trichomonas vaginalis, a sexually transmitted protozoan parasite of the human urogenital tract, has been found to harbour a dsRNA virus with an icosahedral capsid 33 nm in diameter (Wang and Wang 1986; Flegr *et al.* 1987a). The virus is rather common, its dsRNA being detectable in 45—95 % of *T. vaginalis* strains (Flegr *et al.* 1987a; Wang and Wang 1985). The presence of the virus appears to correlate with the mode of expression of a dominant *T. vaginalis* antigen (Wang *et al.* 1987). Trichomonads of virus-harboured strains express this antigen on their surface, trichomonads of virus-free strains contain it in their cytoplasm only. Absence of this antigen on the cell surface has been claimed to be a marker of certain virulence attributes, namely cytoadherence and cytotoxicity (Alderete *et al.* 1986; Alderete and Garza 1985). No differences, however, were found between dsRNA negative and positive strains if the virulence was judged on the basis of mouse assays and clinical and histopathological manifestations on female patients (Flegr *et al.* 1987a).

In the size and shape of the capsid, in the molar mass of its most abundant protein, as well as in the dsRNA nature of its genome, the *T. vaginalis* virus resembles the killer viruses of *Saccharomyces cerevisiae* (Tipper and Bostian 1984). To estimate the extent of this similarity, reliable data on the dsRNA molecules of the *T. vaginalis* virus are required. Wang and Wang (1985, 1986) reported the presence of a single species of dsRNA in *T. vaginalis*. In contrast, our previous results demonstrated the presence of multiple populations of dsRNA (Flegr *et al.* 1987b).

In this paper we confirm that several dsRNA species can be extracted from viral particles isolated from *T. vaginalis* and report the *M*'s and relative abundance of the six dsRNA species.

MATERIAL AND METHODS

Organisms. Axenic *Trichomonas vaginalis* strains A1 (Meingassner and Thurner 1979) and TV 10-02 MR5 (Čerkasovová *et al.* 1987) were used as sources of dsRNA. Trichomonads were stored as cryostabilates (5% dimethyl sulfoxide, liquid nitrogen) and cultivated without antibiotics in tryptose—yeast extract—maltose (TYM) medium (Diamond 1957) supplemented with 10% horse serum.

Isolation of viruses. Trichomonads (cell concentration 40/nL in 0.8% NaCl) were disintegrated in a Potter-Elvehjem homogenizer, diluted 1:20 with 0.8% NaCl and centrifuged (20 min., 15 000 *g*). The pellet was resuspended in the original volume of 0.8% NaCl and centrifuged as before. The supernatants were pooled and viral particles (VP) were obtained from this pool by ammonium sulfate fractionation (solid ammonium sulfate, saturation 20, 30, 40, 50 and 90%, each step lasting 2 h at 0 °C). The highest quantity of VP precipitated between 30 and 40% saturation (Plate 1).

Isolation of nucleic acids. Nucleic acids were extracted with a chloroform—3-methyl-1-butanol mixture (24:1) in the presence of guanidine hydrochloride (4 mol/L) and precipitated with a 0.75 volume of 2-propanol (Flegl 1987).

Electrophoresis. Nucleic acids were electrophoresed in 1% agarose gel in Tris-borate buffer and detected with ethidium bromide (Maniatis *et al.* 1983). Alternatively, nucleic acids were stained before electrophoresis with 4',6-diamidino-2-phenylindole (DAPI) or with a mixture of DAPI and ethidium bromide (3 and 15 ppm). Densitograms were read from negatives with a Magiscan 1 analyzer (Joyce-Loebl). The *M*'s of dsRNAs, supposed to be proportional to the inverse values of their electrophoretic mobility (Edmandson and Gray 1984), were estimated by electrophoresis with a dsRNA standard (reovirus type III, strain Dearing). The relative abundance of the segments were calculated from their *M*'s and from the intensity of fluorescence of the electrophoretic zones.

RESULTS

The extracted nucleic acids contained DNA (zone a), ssRNA (zone b) and six other distinct populations of nucleic acids (zones I—VI) (Plate 2, 3). Zones I—VI were susceptible to pancreatic ribonuclease (RNase A; EC 3.1.27.5) (Plate 4) but resistant to ribonuclease T₁ (RNase T₁; EC 3.1.27.3) (Plate 5) and deoxyribonuclease I (DNase I; EC 3.1.21.1) (Plate 6). When the nucleic acids were stained with DAPI, zones I—VI as well as the DNA zone emitted bluish fluorescence. When a mixture of DAPI and ethidium bromide was used, zones I—VI and the DNA zone emitted a red and a bluish fluorescence, respectively. All these results indicated that zones I—VI consisted of dsRNA.

The molar mass and relative abundances of dsRNA from strain A1 were 3.5, 3.4, 3.2, 2.5, 1.4, 0.34 Mg/mol and 1.0, 1.4, 3.0, 0.3, 2.7, 4.2, respectively.

The M 's of the dsRNA segments from strain TV 10-02 MR5 were the same, but their relative abundance were 1.0, 0.6, 1.7, 0.5, 3.4 and 1.0.

Extracts of nucleic acids from purified VP differed from those obtained from whole protozoans by the absence of DNA and by a lesser amount of ssRNA. The presence of all six dsRNA species, however, was clearly apparent. A maximum of dsRNA was detected in preparations of VP precipitated with 40 % saturation of ammonium sulfate; the relative abundances of particular dsRNA species, however, varied between the different fractions of VP obtained by ammonium sulfate fractionation (Plate 7).

DISCUSSION

In the present study, the existence of six dsRNA species in the viral particle fraction from *T. vaginalis* was confirmed. The M 's and the relative abundances of the particular species were estimated and an interstrain variability of the latter was demonstrated.

Wang and Wang (1985, 1986) report the presence of a single dsRNA species only. We suppose that differences in experimental methods rather than interstrain variability are responsible for this discrepancy, because: 1. No variability in the number of dsRNA species has been reported, in spite of the fact that large collections of *T. vaginalis* strains harbouring dsRNA from different parts of the world have been studied (Flegr *et al.* 1987a; Wang and Wang 1985; Wang *et al.* 1987). 2. Wang and Wang used guanidine isocyanate instead of guanidine hydrochloride for nucleic acid isolation. Both methods were compared and it was found that the former gives only scant yields of dsRNA so the less abundant species of dsRNA can easily escape detection (Flegr 1987). Furthermore, the electromobilities of the three largest dsRNA segments were very similar and long agarose gels or a polyacrylamide gel had to be used for their separation (compare Plate 2 with Plate 3). The different abundance of particular dsRNA segments and variation in the relative amounts of particular dsRNAs in different VP fractions or in different trichomonad strain suggested that at least some segments were packed separately, in distinct VP.

The existence of several dsRNA species of different M 's and abundance, the range of their M 's, as well as the existence of interstrain variability in the abundance, resembled the situation with killer viruses of *S. cerevisiae* (Tipper and Bostian 1984). However, direct evidence confirming the hypothesis of a relationship between the trichomonad and the yeast viruses suggested by Wang and Wang (1986) is still missing.

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The Plates will be found at the end of the issue.

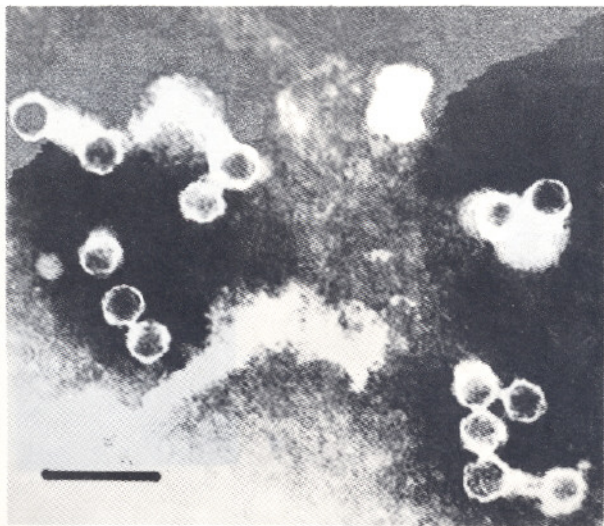


PLATE 1. Viral particles isolated by ammonium sulfate precipitation. Negative staining with phosphotungstic acid (Frank *et al.* 1978); bar represents 100 nm.

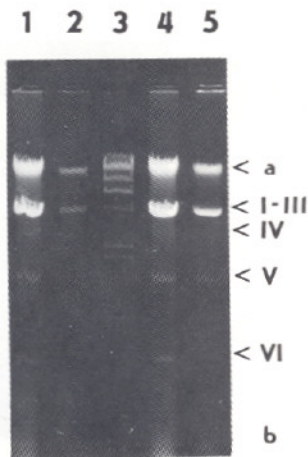


PLATE 2. Nucleic acids from two strains of *T. vaginalis*. a — DNA, b — ssRNA, I-VI dsRNA. Lanes 1,2 — strain TV 10-02 MR5, 4,5 — strain A1, 3 — DNA of phage λ digested with type III site-specific deoxyribonuclease *Hind*III (EC 3.1.21.5; formerly EC 3.1.23.21).

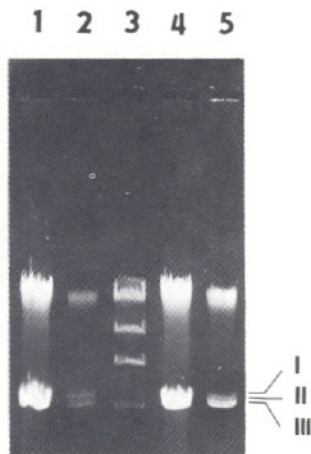


PLATE 3. The same electrophoretic gel as on Plate 2, after prolonged electrophoresis.

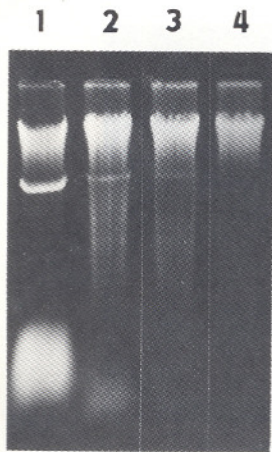


PLATE 4. Nucleic acids from *T. vaginalis* strain A1 digested with different concentrations of RNase A before electrophoresis. From left to right, 0, 0.13, 2 and 32 mg of enzyme per mL of medium incubation buffer (Maniatis *et al.* 1983).

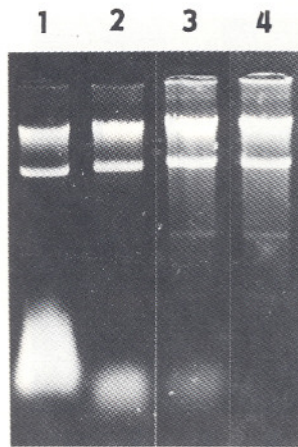


PLATE 5. Nucleic acids from *T. vaginalis* strain A1 digested before electrophoresis with different concentrations of RNase T1. From left to right 0, 2, 32 and 512 μ g of enzyme per mL of medium incubation buffer (Maniatis *et al.* 1983).

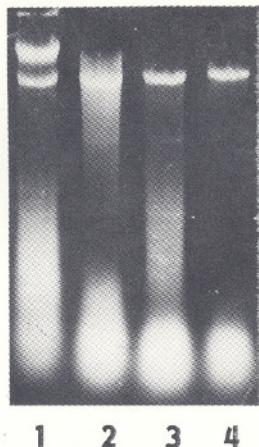


PLATE 6. Nucleic acids from *T. vaginalis* strain A1 digested before electrophoresis with different concentrations of DNase I. From left to right 3.2, 12.8, 51 and 205 μg of enzyme per mL of medium incubation buffer (Maniatis *et al.* 1983).

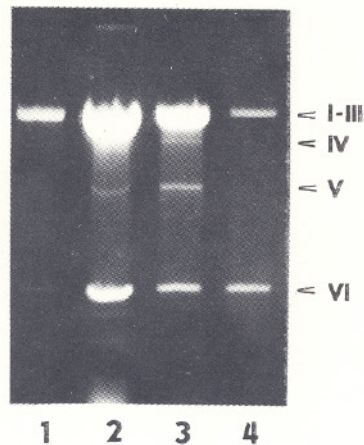


PLATE 7. Nucleic acids isolated from viral particles precipitated with 30, 40, 50 and 60% of ammonium sulfate (lanes 1, 2, 3 and 4, respectively).