ISOLATION AND CHARACTERIZATION OF SOME VIRUSES ASSOCIATED WITH WHEAT PATHOGEN GAEUMANNOMYCES GRAMINIS VAR. TRITICI

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Abstract

Isometric virus particles measuring 35-40 nm in diameter with double stranded RNA genomes of molecular weight $(1.48 - 6.1) \times 10^6$ dalton was detected in the field isolates of the wheat take all fungus Gaeumannomyces graminis var. tritici. Out of three isolates of Ggt four different viruses based on particle size, size of RNA and molecular weight of coat protein belonging to three groups of mycoviruses were detected.

Introduction

Lapierre et al., (1970) discovered the first isometric virus particles in two weakly pathogenic isolates of Gaeumannomyces graminis var. tritici (Sacc) Arx. oliv. (Ggt) the cause of take all disease of wheat. Previous work has established four groups of Ggt viruses based on physical and serological properties and on their double stranded RNA (dsRNA) and capsid polypeptide mol. wt. (Buck et al., 1981; Buck, 1984). The size distribution of these viruses within four group were: group I and II virus particles 35 nm, group III particles 40 nm and group IV particles 29 nm in diameter. These viruses are unusual in that they are transmitted only intracellularly, during growth in asexual and sometimes in sexual spores and via heterokaryosis. Pryor et al., (1988) have isolated a dsRNA mycovirus from maize rust fungus Puccinia sorghi. Presence of isometric virus particles in germinated uredospores of the fungus has presented the evidence of intracellular horizontal transmission of virus. Association of these virus particles with patho genecity of the fungi has not been established, although it is becoming increasingly clear that dsRNA can affect a fungus phenotype in other ways. For example, protein toxin secreted by Killer strains of Saccharomycescere visiae and of Ustilago maydis are encoded by specific dsRNA molecule (Bostian et al., 1980, Koltin & Kendel, 1978). Similarly Rogers et al., (1986) reported a clear association between specific segments of dsRNA and a cytoplasmically transmitted disease in the aggressive strain. Yeast Killer viruses, has been reported to be transmitted only by cytoplasmic mixing during division, mating or other induced forms of cell fusion. El-Sherbeni & Bostian (1987) have demonstrated the extracellular transmission of these virus particles. The present report describes the isolation and characterization of some viruses associated with Gaeumannomyces graminis var. tritici, the cause of take all disease of wheat.

Materials and Methods

Fungal Isolates: Isolates of G. graminis from wheat crops in Highfield, Rothamsted Experimental station, Harpenden, U.K. were kindly provided by Mr. D. B. Slope.

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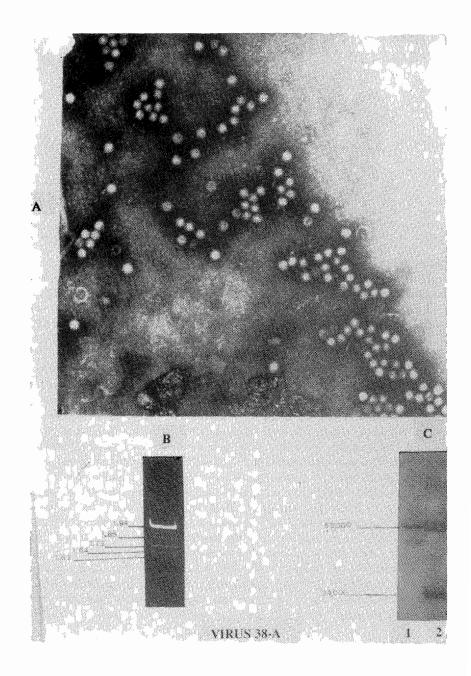


Fig.1. A. Electron Micrograph of Ggt virus 38-a (x 105.000 Magnification),
B. dsRNA banding pattern of virus 38-a in 46 Polyacrylamide gels (Sizes of dsRNA are in Kbp).
C. SDS Polyacrylamide gel Electrophoresis.
Lane 1 = Capsid polypeptide; mol. wt. 55.000.

Lane 2 = Human globulin: mol. wt. of heavy chain 55,000; mol. wt. of light chain 24,000.

Isolations and purification of virus: G. graminis isolates were grown in 60 litre fermenters for 3 days at 24°C in the glucose corn steep liquor medium (Jamil et al., 1984; Banks et al., 1971). Preparation and purification of virus by polyethylene glycol precipitation, ultracentrifugation and sucrose density gradient centrifugation were carried out as described by Buck et al., (1981). The two viruses in isolates 87-1 were separated by CsCl density gradient centrifugation as described by Jamil & Buck (1984).

Electron microscopy: Purified virus samples were negatively stained with 1% potassium phosphotungstate pH 7. The grids were examined in a Siemens Elmiskop Ia electron microscope and photographs were taken on Ilford EM plates. A magnification of x 32743 was used, followed by x 3.5 in printing to give an over all magnification of x 105000.

Isolation and analysis of dsRNA: Virus RNA was prepared by phenol extraction and shown to be double stranded by S1 nuclease treatment. Analysis and molecular weight determinations of dsRNA component were carried out by polyacrylamide gel electrophoresis (PAGE) as described by Buck & Ratti (1977) using internal dsRNA standards covering the mol. wt. range, 1.11 x 10 to 6.3x10⁶ from the following viruses: Aspergillus foetidus viruses S and F (Buck & Ratti, 1977); Helminthosporium maydis virus (Bozarth, 1977); Ggt 3 bla virus (Buck et al., 1981).

Analysis of capsid polypeptide: Viral proteins was denatured by heating viral suspensions in 1 % SDS and 0.1% 2-mercaptoethanol for 3 min, at 100 C, then analyzed by electrophoresis at 60-100V for 16-18 h in 4% - 10% SDS PAGE as described by Laemmli (1970). Denatured polypeptide samples were co-electrophoresed with denatured protein markers. The following standards were used; phosp horylase A subunit mol. wt. 94000; Bovine Serum albumin mol. wt. 68,000; Lactate dehydrogenase mol. wt. 36,000; glutamate dehydrogenase mol. wt. 55,490 human gamma globulin heavy chain mol. wt. 55,000 light chain 24,000; oval albumin mol. wt. 47,000 Pyrovate Kinase 60,000.

Results and Discussion

Buck *et al.*, (1981) defined the properties of *Ggt* groups I and II viruses as follows: group I particle diameter 35 nm, capsid polypeptide mol. wt. 54,000 to 60,000, two to four dsRNA segments 1.53 to 1.94 K6p; group II particle diameter 35 nm, capsid polypeptide mol. wt. 68,000 to 73,000, two to four dsRNA segments in the group II range. Confirmation of the grouping was obtained for isolated virus particles from four isolates (87-1, 38, 74 and 16).

Viruses 38-A and 87-1-L can be accommodated in group I, since their particle diameter, capsid polypeptide mol. wt. and size of dsRNA components are in good agreement with the other members of this group (Table 1). This was confirmed for 38-A virus which was found to be weakly serologically related to a previously described group I virus (F6-C) although no sequence homology could be detected between virus 87-1-L and several other group I viruses. However the study was not comprehensive and distant serological relationships cannot be excluded, since the titres of some of the sera used were only moderate(1:64 to 1:16). 38A and 87-1-L viruses are distinct viruses since no sequence homology was detected between their dsRNAs.

The number of dsRNA segments (Six) associated with 38-A virus (Fig.1) is greater than that of previously described group I viruses two to four (Buck *et al.*, 1981). Although the virus has a single capsid polypeptide mol. wt. species, it could still be a mixture of virus strains viruses 3bla-B1 and 3bla - B2, both present in isolates 3bla, have the same capsid polypeptide mol. wt. but have distinct dsRNA components (Mcfadden *et al.*, 1983;

Virus	Particle dia (nm)	Size of dsRNA components (kbp)	Mol. wt. of capsid polypeptide (x10 ⁻³)
38 - A	35	1.94, 1.85, 1.82, 1.73, 1.64, 1.62	55
74 - A	35	2.33, 2.24	73
87 - 1 - L	35	1.85, 1.76, 1.48	60
87 - 1 - H	41	6.1	84, 78

Table 1. Properties of viruses from Ggt isolates 34, 78, 87-1.

Stanway & Buck; 1984). Originally it was believed that they were a single virus (Buck et al., 1981). Furthermore in agarose gel electrophoresis intact particles of 38-A virus gave two closely spaced bands which stained with both coomassie Blue and ethidium bromide, suggesting the presence of two viruses (Fig. 2).

Virus 74-A has characteristic features of group II viruses. The capsid polypeptide mol. wt. of 73,000 and the size of the two genomic dsRNAs (2.33 Kbp and 2.24 Kbp). Fig.3 is the same as those of a previously described group II virus, 3 bla-B1 (Mcfadden et al., 1983; Stanway & Buck, 1984). Furthermore, gel transfer hybridization under conditions of both low and high stringency showed that the dsRNAs of the two viruses were closely related (unpublished). Both isolates come from Rothamsted: 3bla from Little Knottfield in 1970 and 74 from Highfield in 1981. Interestingly isolate 74 was infected with only one virus whereas isolate 3bla was infected with four viruses, encompassing group I, II, and III. However virus B1 was the only one to be transmitted into ascospores (Mcfadden et al., 1983).

Sequence homology was also detected between 74-A dsRNA and a dsRNA of 2.17 from virus OgA-A, another group II virus (unpublished). However in this virus the dsRNA was slightly smaller than either of those of 74-A. Isolate OgA came from a different location. Small differences of size in dsRNA segments are of doubtful significance since they can arise for example, during ascospore formation (Mcfadden et al., 1983). Hence of the five group II viruses described by Buck et al., (1981) and Mcfadden et al., 1983), virus 74-A is related to three closely to 3 bla -B1 and distantly to OgA-A and F6-B. No relationship was detected to group II virus T1-A, from an Australian isolate of Ggt, T1-A virus was previously found to be only distantly related to the group II viruses from European Ggt isolates.

Ggt virus groups I and II have been placed in the Partitiviridae family of dsRNA mycoviruses (Brown, 1985). Partitiviridae have genomes of two monocistronic dsRNA segments, one encoding the capsid polypeptide and the other probably encoding a dsRNA - dependent RNA polymerase. Some viruses in groups I and II have additional dsRNAs and these could be satellites, similar to a satellite dsRNA associated with a previously described Ggt virus (Romanos et al., 1981). Evidence to support came from

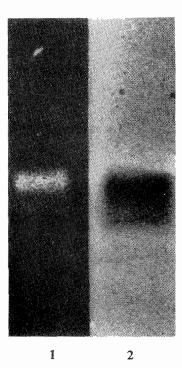


Fig.2. Agrose gel Electrophoresis of intact virus particles Lane 1 = Gel stained with ethidium bromide.

Lane 2 = Gel stained with Coomassie blue

individual dsRNAs. Virus 87-I-L Fig. 4, has three dsRNA and probes from each RNA did not hybridize with the other two (Jamil et al., 1984). Hence the third RNA is probably a satellite, rather than a defective RNA which would be expected to be related to one of the genomic RNAS (Huang & Baltimore, 1977). Similar results were obtained from the dsRNAs of virus 38-A, although complete separation of all the smaller dsRNA segments was not possible. Some or all of the dsRNAs with sizes below the group I i.e., 1.48 to 1.35 Kbp range was found in some of the field isolates.

Buck *et al.*, (1981) defined the properties of *Ggt* group III viruses as follows: Particles 40 nm diameter, two dsRNA components 4.7 to 6.3 KbP and three capsid polypeptides, mol. wt. 87,000, 83,000 and 78,000. Virus 87-I-H has similar properties (Fig. 5) to the group III viruses and was found to be closely related serologically to virus F6-A (antiserum to virus 3 bla A was not available) (unpublished). However, viruses 87-I-H had only two capsid polypeptide species with mol. wt. 84,000 (P1) and 78,000 (PII). It has been noticed that molar ratio of these polypeptides is 1:1 in fresh virus preparation purified by CsCl gradient centrifugation, but on storage, PI starts degrading into PII, hence the ratio of PI and PII in old preparation becomes 1:2. Tryptic and peptic fingerprints and amino acid analysis of PI and PII indicated that PI is derived from PII, either in *vitro* or *in vivo* (Jamil & Buck, 1984). The more rapid conversion of PI into PII on aging of 87-I-H virus preparation, purified by CsCl gradients compared to sucrose

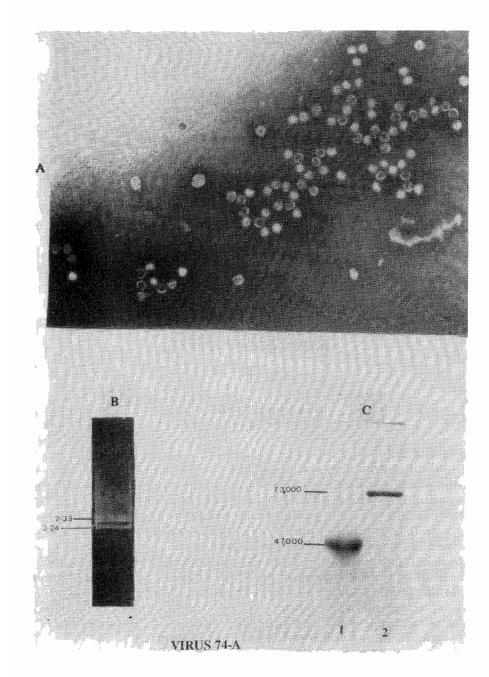


Fig.3. A. Electronmicrograph of Ggt virus 74-A (x 105,000 Magnification). B. dsRNA Banding Pattern of virus 74-A in $4^{\circ}6$ Polyacrylamide gels. (Sizes of dsRNA are in Kbp). Lane 1=EGG albumin Mol. Wt. 47,000 SDS Polyacrylamide gel Electrophoresis. Lane 2=Capsid Polyacrylamide of virus 74-A.

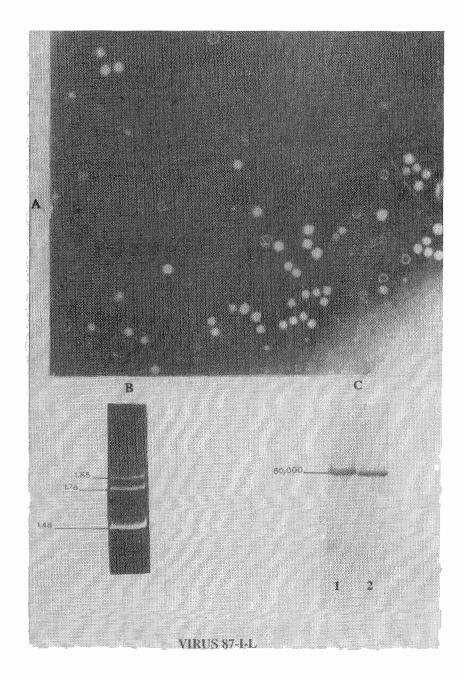


Fig.4. A. Electronmicrograph of Ggt virus 87-1 L (x 105,000 Magnification). B. dsRNA banding pattern of virus 87-1-L in 4°6 Polyacrylamide gel Electrophoresis. Lane 1 = Pyrovate Kinase mol. wt. 60,000.

Lane 2 = Capsid polypeptide of virus 87-1-L: mol wt 60,000.

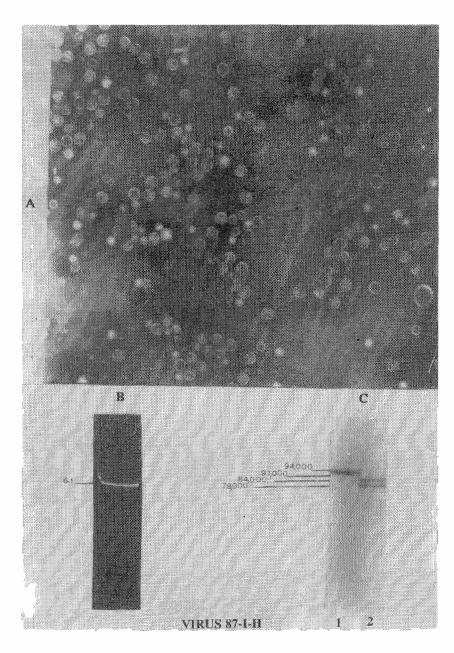


Fig.5. A. Electronmicrograph of Ggt Virus 87-1-H (x 105.000 Magnification).

B. dsRNA banding pattern of virus 87-1-H in 4% Polyacrylamide gels. (Size of dsRNA in Kbp)

C. SDS - Polyacrylamide gel Electrophoresis.

Lane 1 = Phosphorylase A. mol. wt. 94,000.

Lane 2 = Capsid polypeptides of virus 87-1-H Preparation purified by CsCl density gradient.

gradient purified virus, supports the possibility that PI cleaved into PII after virion disruption *in vitro*. Since the virus was less stable in CsCl Proteolytic cleavage of capsid polypeptide has been reported for *Aspergillus foetidus* virus S in which the major capsid polypeptide O-1 was converted to 0-2 by a protease both during virus preparation and on subsequent storage (Buck & Ratti, 1975). It is likely, therefore, that virus 87-I-H (and probably also viruses 3bla-A and F6-A) has only one major capsid polypeptide. However, it has been noticed that the molar ratio of PI and PII in 87-I-H virus, which was purified by sucrose gradient only, remained constant. Hence the possibility can not be eliminated that some conversion of PI into PII takes place *in vivo*, because of post translational modification, analog one to reovirus M2 protein which is a cleavage product of U2 *in vivo* (Silverstein *et al.*, 1976).

Another apparent difference between virus 87-I-H and the other two group III viruses, F6-A and 3bla-A, was that 87-I-H apparently had only one dsRNA species (Fig. 5). In this context it is noteworthy that dsRNA of 87-I-H has shown moderate and weak sequence homology with larger and smaller dsRNAs of 3bla-A virus respectively (unpublished). There are two possible interpretations of these results:

- 1) The two 3 bla-A dsRNAs could be related to each other, the larger being the virus genomic dsRNA and the smaller a defective yeast strains (Bruenn & Kane, 1978).
- 2) 87-I-H dsRNA could consist of two components which comigrate in gel electrophoresis and each of which is related to one of the 3bla dsRNAs. Some dsRNAs in yeast strains have been shown to consist of at least two different components with different nucleotide sequences, but which migrate as a single band in gel electrophoresis (Sommer & Wickner, 1982; Field *et al.*, 1982). Molecular cloning of 87-I-H dsRNA would resolve which of these two possibilities is valid.

Members of the *Totiviridae* family of dsRNA mycoviruses (Brown, 1985) have the following properties:

Virus particles, 40-43 nm; single molecule of dsRNA 4.8 to 6.1 KbP, capsid polypeptide mol. wt. 75 to 88 x 10⁻³, buoyant density in CsCl, 1.40-1.42 gm. If virus 87-I-H is confirmed to have only one dsRNA component and one major capsid polypeptide species as seems likely from the results presented here, it could be assigned to the *Totiviridae* family.

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