

Serological relationship of rice tungro spherical virus and bacilliform virus components associated with rice tungro disease

M. D. Mishra, F. R. Niazi and R. K. Jain

Division of Mycology and Plant Pathology, Indian Agricultural Research Institute, New Delhi 110 012, India

Immunosorbent electron microscopy revealed a degree of serological relationship between rice tungro spherical virus and bacilliform virus indicating the existence of some common epitopes on the capsid protein of the two particles.

ETIOLOGICAL studies on rice tungro disease led to the discovery of association of two morphologically and serologically distinct rice tungro spherical virus (RTSV) and bacilliform virus (RTBV)^{1,2}. However, a recent publication³ pointed out certain similarities in the biological and intrinsic structural properties of these two viruses. It was, therefore, considered worthwhile to reexamine serological relationship of these particles.

RTSV particles with and without RTBV particles respectively were purified following Mishra *et al.*⁴ and the modified Saito's procedure⁵ standardized in this laboratory (Niazi *et al.* unpublished). For preparing the antisera separately of RTSV with and without RTBV, rabbits were immunized separately with purified preparations by three intravenous injections followed by two intramuscular injections at weekly intervals. For intramuscular injections, antigen was mixed with equal volume of Freund's incomplete adjuvant. Rabbits were bled one week after the last injection for obtaining the antisera.

To study the serological relationship, combined immunosorbent electron microscopy (ISEM) and decoration technique⁶ was followed and various combinations of coating, trapping and decoration were tested (Table 1).

The present studies indicate that S+B antibodies trapped both S+B antigens in the ratio of 5:3 from clarified diseased sap, while S antibodies not only trapped S antigens but also B antigens from clarified diseased sap and purified preparations of S+B antigens in the ratio of 7:1 (Figure 1a). In further experimenta-

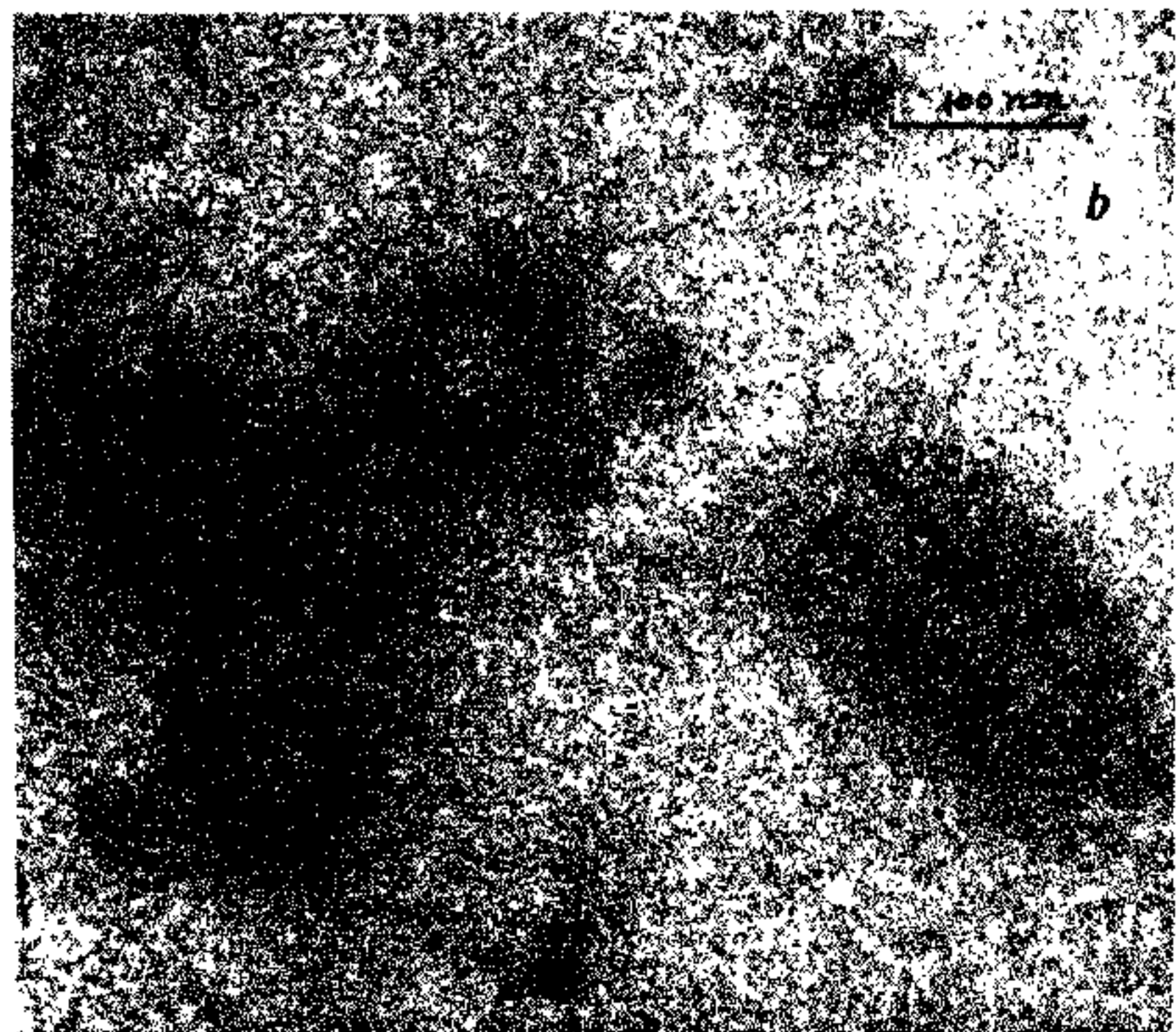
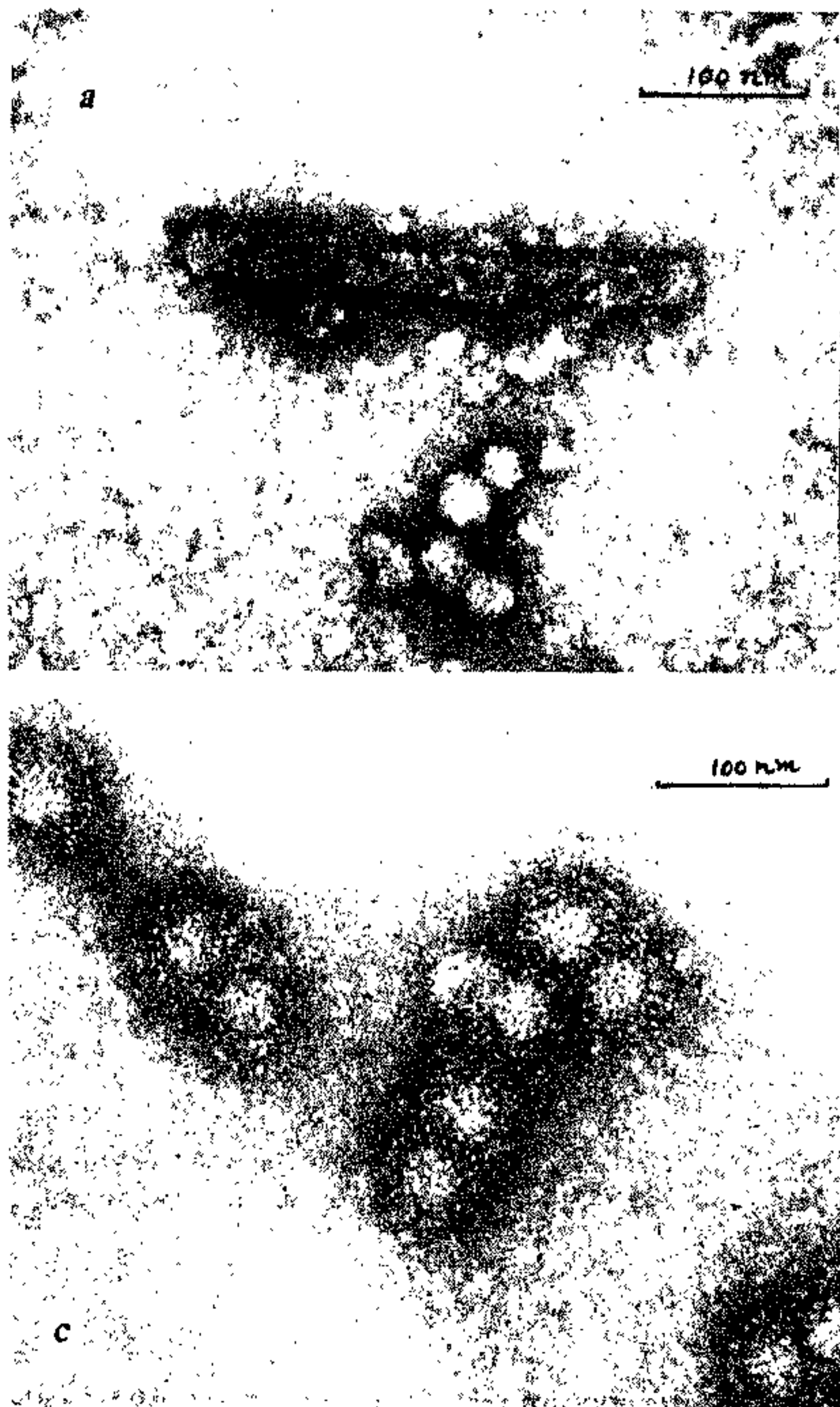


Figure 1. Transmission electron micrographs of RTV components subjected to combined ISEM and decoration tests. *a*, S+B antigens trapped by S antibodies. *b*, S+B antigens' decoration by S antibodies. *c*, S antigens trapped and decorated by S antibodies.

Table 1. Antigen-antibody combinations used for combined ISEM and decoration tests.

Combinations	Coating (1/320)	Trapping	Decoration (1/4)	Electron microscopic observations
1.	S antibodies	S antigens (purified)	S antibodies	Trapping and decoration of S antigens (Figure 1c)
2.	S antibodies	S+B antigens (purified and clarified diseased sap) (Figure 1a)	S+B antibodies	Trapping and decoration of S+B antigens
3.	S+B antibodies	S+B antigens (clarified diseased sap)	S antibodies	Trapping and decoration of S+B antigens (Figure 1b)

Antisera titre: S=1:1024, S+B=1:1024 (based on tube precipitin test).

tions of decorating the trapped antigens, S antibodies also decorated B antigens besides S antigens (Figure 1b). A point of interest here is the trapping and decoration of B antigens by S antibodies (Table 1), indicating some degree of positive serological relationship between the two morphologically distinct RTV components.

Earlier observations of Omura *et al.*¹ that the components of RTV are antigenically distinguishable were based on ouchtelony and clumping experiments. However, the present findings indicate antigenic relationship between the two particles at particulate level, probably due to the existence of some common

epitopes in the capsid protein of these two particles.

1. Omura, T., Saito, Y., Usugi, T. and Hibino, H., *Ann. Phytopath. Soc. Jpn.*, 1983, **49**, 73.
2. Cabauatan, P. Q. and Hibino, H., *Plant Dis.*, 1988, **72**, 526.
3. Jain, R. K. and Mishra, M. D., *Curr. Sci.*, 1989, **58**, 457.
4. Mishra, M. D., Niazi, F. R., Basu, A. N. and Nam Prakash, *Indian Phytopathol.*, 1981, **34**, 461.
5. Saito, Y., *Trop. Agric. Res. Ser., Jpn.*, 1977, **10**, 129.
6. Hill, S. A., *Methods in Plant Virology*, Vol. 1. Blackwell Scientific Publications, London, 1984, p. 167.

5 October 1989; revised 28 December 1989

Purification of β -trypsin by SP-Sephadex column chromatography

Y. S. Rajput and M. N. Gupta

Chemistry Department, Indian Institute of Technology, New Delhi 110 016, India

A method for purification of β -trypsin from commercial trypsin is described. The separation was based on the differential retardation of different forms of trypsin by SP-Sephadex column. β -trypsin was identified by SDS-gel electrophoresis.

TRYPsin undergoes autolysis in solution and results in a number of degraded products. Hence, attempts were made to stabilize this enzyme by simple chemical modification of amino groups^{1,2}. The extent of success varied with the type of trypsin used. β -Trypsin could be stabilized when treated with acetimidate whereas α -trypsin failed to stabilize³. Commercially available preparations of trypsin differ in the content of aggregated inert material, α -trypsin and β -trypsin⁴. The most commonly used method for separating unautolysed form of trypsin, β -trypsin, from other forms of trypsin is that of Schroeder and Shaw⁴ using sulphony ethyl-Sephadex C-50 (SE-Sephadex) column. Since Pharmacia Fine Chemicals no longer manufactures SE-Sephadex, efforts were made to purify β -trypsin on sulphony propyl-Sephadex (SP-Sephadex) column. The results are reported in this paper. SP-Sephadex C-50 (40–120 μ m) from Pharmacia Fine Chemicals, Sweden was used.

The modified method of Schroeder and Shaw⁴ for purification of β -trypsin is described below. SP-Sephadex column (1.8 \times 80 cm) was equilibrated with tris-HCl buffer (0.01 M, pH 7.0) containing 20 mM Ca^{2+} at 5°C. Trypsin (Sisco Research Laboratory) was dissolved in the above tris-HCl buffer and 10 ml of trypsin solution (10 mg/ml) was loaded on the Sephadex column and eluted using the same buffer. Fractions of 6.2 ml each at a flow rate of 15 ml/h were collected into tubes containing 0.25 ml sodium formate (1.25 M, pH 2.9). Sodium formate was used to immediately lower the pH of the eluted solution thus preventing autolysis of trypsin. Fractions (Nos 105–122, Figure 1) were pooled

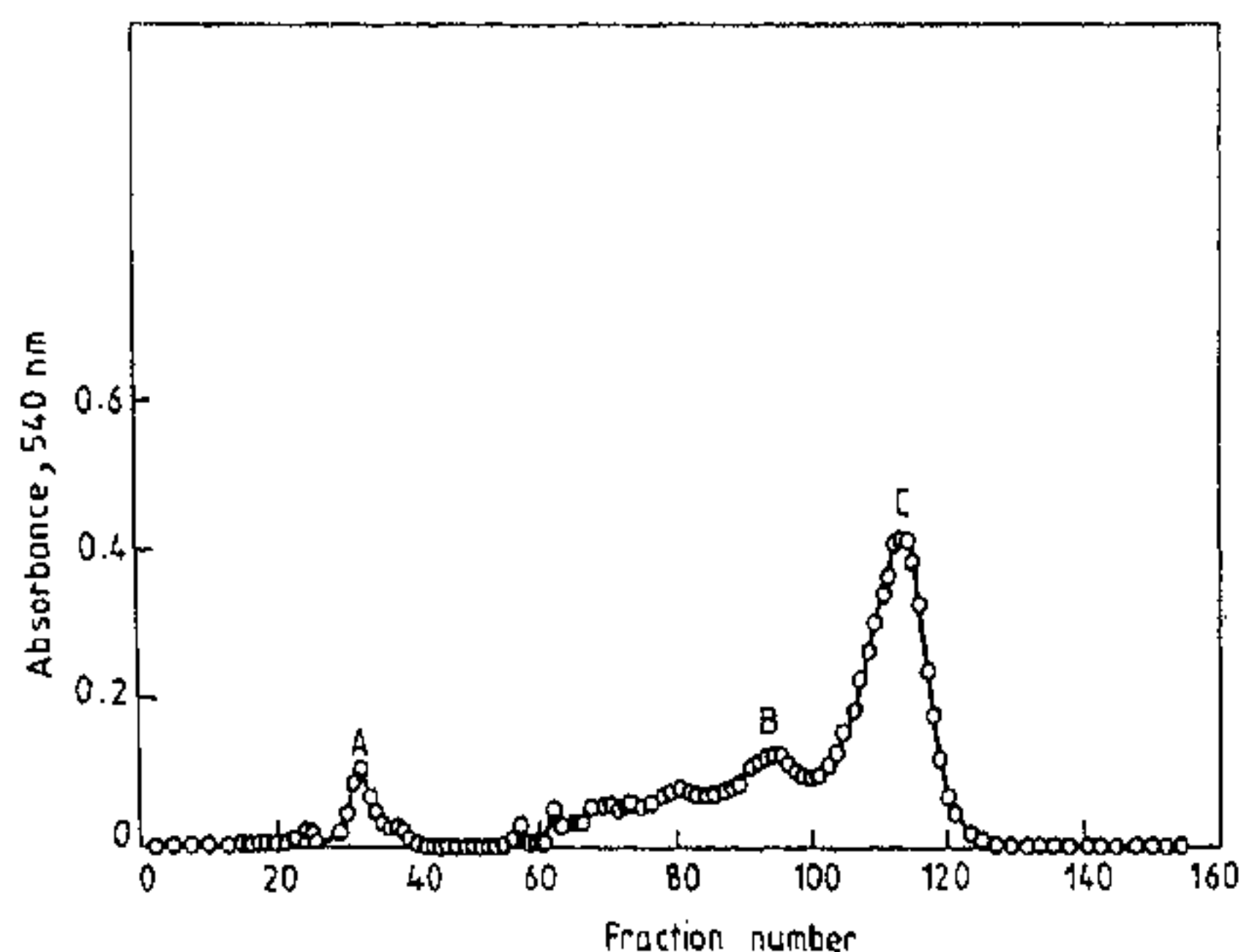


Figure 1. Fractionation of commercial trypsin on SP-Sephadex column.