

- of 40, 30, 20, and 10% sucrose in 10 mM Tris-HCl, pH 7.3; let it diffuse at 4°C overnight. Centrifuge at 82,000g in a Beckman SW40 rotor for 2 h.
- b. Remove equal volume fractions and analyze them by UV absorption and/or agarose electrophoresis (*see Note 2*).
 - c. Pool together fractions containing the virus, dilute them twofold with 10 mM Tris-HCl, pH 7.3, and centrifuge at 146,000g in a Beckman 50.2 Ti rotor for 2 h.
 - d. Resuspend the pellet in 10 mM Tris-HCl, pH 7.3. Remove unresuspended material as in **Subheading 3.1., step 7**.
6. There is an alternative method to the one described here that includes a polyethylene glycol (PEG 6000) precipitation step (**9**). If followed, supernatant obtained in **Subheading 3.1., step 4** is brought to 8% PEG 6000 and 200 mM NaCl, and stirred for 1 h at 0–4°C. Virions are then pelleted by centrifugation at 7700g for 20 min. Pellets are resuspended in 10 mM Tris-HCl, pH 7.3, and solution is finally ultracentrifuged at 146,000g for 2 h to pellet the virions again. We must stress, however, that in side-by-side experiments the sucrose cushion protocol described here rendered higher amounts (as much as twice) of better quality virus preparations, in a significantly shorter period of time, when compared to the PEG precipitation protocol.
 7. As reported previously (**1**), caution must be taken with carmoviruses that have been reported to be isoelectric at acidic pHs, since virus precipitation may occur during extraction. In these cases, tissue must be extracted in neutral phosphate buffers and clarification of the homogenate achieved with organic solvents (*n*-butanol and/or chloroform) (**1**). We do not predict any problems with the use of centrifugation through sucrose cushions in these examples.

References

1. Morris, T. J. and Carrington, J. C. (1988) Carnation mottle virus and viruses with similar properties, in *The Plant Viruses, vol. 3. Polyhedral Virions with Monopartite RNA Genomes* (Koenig, R., ed.), Plenum, New York, pp. 73–112.
2. Guilley, H., Carrington, J. C., Balázs, E., Jonard, G., Richards, K., and Morris, T. J. (1985) Nucleotide sequence and genome organization of carnation mottle virus RNA. *Nucleic Acids Res.* **13**, 6663–6677.
3. Carrington, J. C. and Morris, T. J. (1985) Characterization of the cell-free translation products of carnation mottle genomic and subgenomic RNAs. *Virology* **144**, 1–10.
4. Harbison, S. A., Davies, J. W., and Wilson, T. M. A. (1985) Expression of high molecular weight polypeptides by carnation mottle virus RNA. *J. Gen. Virol.* **66**, 2597–2604.
5. Carrington, J. C. and Morris, T. J. (1986) High resolution mapping of carnation mottle virus-associated RNAs. *Virology* **150**, 196–206.
6. Carrington, J. C., Heaton, L. A., Zuidema, D., Hillman, B. I., and Morris, T. J. (1989) The genome structure of turnip crinkle virus. *Virology* **170**, 219–226.
7. Hacker, D. L., Petty, I. T. D., Wei., N., and Morris, T. J. (1992) Turnip crinkle virus genes required for RNA replication and virus movement. *Virology* **186**, 1–8.