

Local Expression and Exocytosis of Viral Glycoproteins in Multinucleated Muscle Cells

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Abstract. We have analyzed the distribution of enveloped viral infections in multinucleated L6 muscle cells. A temperature-sensitive vesicular stomatitis virus (mutant VSV ts045) was utilized at the nonpermissive temperature (39°C). As expected, the glycoprotein (G protein) of this mutant was restricted to the ER when the multinucleated cells were maintained at 39°C. We demonstrate that this G protein remained localized when the infection was performed at low dose. By 4 h after infection the G protein patches spanned an average of 220 μm . The localization was independent of nuclear positions, showing that the ER was a peripheric structure. Thus, the infection did not recognize nuclear domains characteristic of nuclearly encoded proteins. After release of the 39°C block, transport through a perinuclear compartment into a re-

stricted surface domain lying above the internal G protein patch occurred. Accordingly, the transport pathway was locally restricted. After a 16-h infection the G protein spanned 420 μm , while the matrix protein occupied 700–800 μm of the myotube length. Double infection of multinucleated L6 muscle cells with Semliki Forest virus and VSV at high multiplicities showed that the glycoprotein of each virus occupied intracellular domains which were devoid of the other respective glycoprotein. Taken together, these findings indicate that the viral glycoproteins did not range far from their site of synthesis within the ER or other intracellular membrane compartments in these large cells. This result also suggests that relocation of viral RNA synthesis occurred slowly.

MULTINUCLEATED muscle cells form by fusion of myoblasts, resulting in a reorganization of the intracellular organelles and membranes of the parent cells. Experiments using artificial systems where a single nucleus encoded the protein of interest, have shown that in multinucleated muscle cells, membrane proteins are restricted to the vicinity of the nucleus of origin. A picture has emerged indicating that each nucleus directs the synthesis of a subset of myotube structures called a nuclear domain (Hall and Ralston, 1989). Thus in hybrid interspecies myotubes, a Golgi protein was targeted mainly into the Golgi region surrounding the nucleus encoding that protein (Pavlat et al., 1989). Similarly, the intracellular localization of a muscle adhesion protein 5.IH11 (Pavlat et al., 1989) or a transfected lymphocyte membrane protein CD8 (Ralston and Hall, 1989a) that was encoded by a single nucleus were found to be in proximity of the source nucleus. Proteins that naturally exist in myotubes like the acetylcholine receptor (Merlie and Sanes, 1985; Fontaine and Changeux, 1989) or acetylcholine esterase (Rotundo and Gomez, 1990) are encoded by a limited subset of nuclei.

Such a mode of expression results in mosaicism of protein localization. One explanation for the existence of nuclear domains is that certain gene products are effectively trapped into their target organelles remaining in the vicinity of the nucleus of origin. Another possibility is that the intracellular

organelles of fusing myoblasts do not intermingle but remain associated with their parent nuclei in the multinucleated cells. Both models suggest that there is little exchange of membrane material between distant parts of the myotube.

We analyze here how non-nuclearly encoded mRNA products distribute in multinucleated cells and whether they recognize nuclear domains. Enveloped RNA viruses were utilized for this purpose since the translation of the viral mRNA and the subsequent protein translocation and transport processes are totally dependent on cellular machinery. We found that a viral infection remained locally restricted inside the multinucleated cells whose dimensions are many orders of magnitude larger than those in mononucleated cells. The localization of the viral glycoprotein was independent of the nuclear positions. Our results suggest that in multinucleated muscle cells, membrane-associated viral mRNA products do not range further than $\sim 200 \mu\text{m}$. This behavior shows that the membrane structures in large cells are relatively immobile.

Materials and Methods

Cells and Viruses

Rat L6 myoblastic cells (Yaffe, 1968) were grown in DME (Gibco Laboratories, Grand Island, NY) containing 10% FCS (Gibco Laboratories). Sub-

confluent monolayers of L6 myoblasts were induced to fuse by growing the cells in DME containing 1% horse serum and 0.4 U/ml insulin. 4 d after the induction of fusion, the cells appeared as long, branched structures containing hundreds of nuclei which were gathered into groups. Wild-type vesicular stomatitis virus (VSV)¹ (Indiana serotype) was grown in BHK cells (Matlin et al., 1983). The mutant ts045 was a clone ts045-6 (Griffiths et al., 1985). Semliki Forest virus (SFV) was propagated in BHK cells as described (Kääriäinen et al., 1969). Virus titers were determined on 2-d-old L6 myoblast monolayers. In brief, infected cells were identified by immunofluorescence staining for the viral glycoprotein, while the number of cells was determined by staining the nuclei with Hoechst dye 33258 (Sigma Chemical Co., St. Louis, MO). Multiplicities so obtained were regarded equivalent to plaque forming unit (pfu). The multinucleated cells were usually infected at 0.01 pfu/cell with VSV ts045 for 1 h at 32°C and then grown at 39°C for indicated periods. Infections with wild-type VSV and SFV were done at 37°C.

Metabolic Labeling and Analysis of the Labeled Virions

The multinucleated muscle cells were infected with VSV at 2 pfu and grown for 4 h at 37°C. A 10-min pulse with 50 μ Ci/ml of [³⁵S]methionine (Amersham Corp., Bucks, England) was then given, followed by a 60-min chase. The chase medium was centrifuged (5,000 g, 10 min) to remove cellular debris, and virions were then pelleted (30,000 g, 60 min). The pellet was analyzed by SDS-PAGE (Laemmli, 1970).

Electron Microscopy

For thin section EM, myotubes were cultured on microscope object glasses. Cells were fixed with 3% glutaraldehyde in 0.2 M sodium cacodylate buffer, pH 7.2, for 4 h. The specimens were stored in the same buffer supplemented with 7.5% sucrose. Postfixation was for 1 h in 4% OsO₄. Specimens were dehydrated in ethanol and embedded in Epon. Thin sections were stained with uranyl acetate and lead citrate and examined with a Philips 410 electron microscope (Philips Electronic Instruments Co., Mahwah, NJ).

Antibodies

Polyclonal antibodies against VSV glycoprotein (G protein) were prepared by immunizing a rabbit with G protein liposomes. The liposomes were prepared by the octylglucoside dialysis method (Eidelman et al., 1984). A viral lipid mixture was used for the reconstitution as described (Metsikkö et al., 1986). About 100 μ g protein were injected intradermally three times at 2-wk intervals. The antiserum was collected 3 wk after the last booster and absorbed with a monolayer of multinucleated L6 muscle cells that were fixed and permeabilized. The preabsorbed antiserum specifically recognized the viral G protein when a cell lysate from BHK cells, infected with VSV, was analyzed by Western blotting at 1:200 antiserum dilution.

mABs against the VSV matrix (M) protein were prepared with some modifications as described (Galfré and Milstein, 1981). VSV nucleocapsids containing the M protein (skeletons) (Newcomb et al., 1982) were prepared as described by Petri and Wagner (1979), and injected with Freund's adjuvant intraperitoneally into BALB/c mice (100 μ g protein). During fusions, the ratio of spleen cells to P3-NSI-Ag4-1 myeloma cells was 5:1. Screening of the hybridomas was performed by Western blotting. In brief, polypeptides in cell lysates from BHK cells infected with VSV were separated by SDS-PAGE and transferred onto nitrocellulose (Risau et al., 1981). The culture medium of clone 2F9 specifically recognized VSV M protein and it was used for immunofluorescence studies.

A mAB against the VSV G protein was prepared by Henrik Garoff and Brian Burke at the European Molecular Biology Laboratory (EMBL). Ascites fluid was used at 1:400 dilution for immunofluorescence studies. Polyclonal antibodies against SFV p62/E2 glycoprotein have been described (Metsikkö and Garoff, 1990). They were directed against a peptide of 32 amino acids, corresponding to the cytoplasmic tail of p62 glycoprotein. Polyclonal antibodies against rat RER were prepared by Louvard et al. (1982). A mannosidase II specific mAb (53FC3) (Burke et al., 1982; Baron and Garoff, 1990) was used as a Golgi marker.

1. *Abbreviations used in this paper:* pfu, plaque forming unit; SFV, Semliki Forest virus; VSV, vesicular stomatitis virus.

Immunofluorescence Studies

Cells were fixed with 3% paraformaldehyde in PBS and permeabilized with Triton X-100 as described (Louvard, 1980). Polyclonal anti-G protein serum was used at 1:200 dilution, followed by TRITC-conjugated anti-rabbit IgG (DAKOPATTS, Glostrup, Denmark; 1:400). Culture medium from clone 2F9 containing monoclonal anti-M protein antibody was diluted 1:5. For double immunofluorescence staining, the polyclonal and monoclonal antibodies were mixed at appropriate dilutions, applied to fixed cells, and incubated. TRITC-conjugated swine anti-rabbit IgG (DAKOPATTS; 1:200) and FITC-conjugated goat anti-mouse IgG (DAKOPATTS; 1:200) were mixed and used as secondary antibodies. Nuclei were visualized with Hoechst dye 33258. Simultaneous visualization of the internal and cell surface G proteins was performed in two stages. First, the fixed cells were incubated with polyclonal antibodies, followed by the TRITC-conjugated IgG. The cells were then permeabilized with Triton X-100 and internal G protein was visualized with a monoclonal anti-G protein antibody and FITC-conjugated anti-mouse IgG. Specimens were mounted with Mowiol (Hoechst, Frankfurt, Germany) in PBS containing 2.5% 1,4-diazobicyclo-[2.2.2]-octane (Sigma Chemical Co.) and viewed under the Leitz Aristoplan fluorescence microscope (E. Leitz Inc., Rockleigh, NJ), using appropriate filters.

Results

Infection of Multinucleated L6 Cells

The L6 myoblasts were induced to fuse by growing them in a medium containing horse serum and insulin. Under these conditions, branched myotubes were generated which contained several groups of nuclei separated by stretches of cytoplasm. To examine whether the L6 myotubes were successfully infected with VSV or the temperature-sensitive mutant of VSV ts045, we performed immunofluorescence staining for the VSV G glycoprotein. We found that the myotubes were infected throughout their length at viral doses corresponding to 0.5 pfu/cell in the parental mononucleated L6 myoblasts. Furthermore, by using [³⁵S]methionine pulse-chase labeling of a myotube culture where 90% of the nuclei were counted in myotubes, we observed that the label was transferred into the medium in the form of virions (Fig. 1 A). The viral titer of a VSV-infected myotube culture medium was 1.7×10^6 pfu/ml when determined after a 6-h infection period. Thus, viral particle formation occurred in the multinucleated muscle cells. Examination of the infected cells by EM indicated that budding occurred at the cell surface (Fig. 1 B). Rounding and detachment of the myotubes was not observed after a 24-h infection period with the wild-type VSV. While the VSV M protein probably disrupts the cytoskeleton and causes a cytopathic effect in mononucleated cells (Blondel et al., 1990), it appears that myotubes are more resistant to this effect.

Infection at Low Dose Remains Local

We used the mutant VSV ts045 at 39°C to analyze whether the infection generated by a single viral particle remained local in the large multinucleated cells. At this temperature the mutant G protein should remain restricted to the ER and no infective particles should be formed (Schnitzer et al., 1979; Zilberstein et al., 1980; Metsikkö and Simons, 1986). We determined that the conditioned medium from multinucleated muscle cells infected with ~ 2 pfu/cell of VSV ts045 was unable to infect a fresh culture of muscle cells, indicating that no infective progeny was generated to propagate the infection. At pfu < 0.01 the ts045 infection was local as evaluated by indirect immunofluorescence for the G protein in

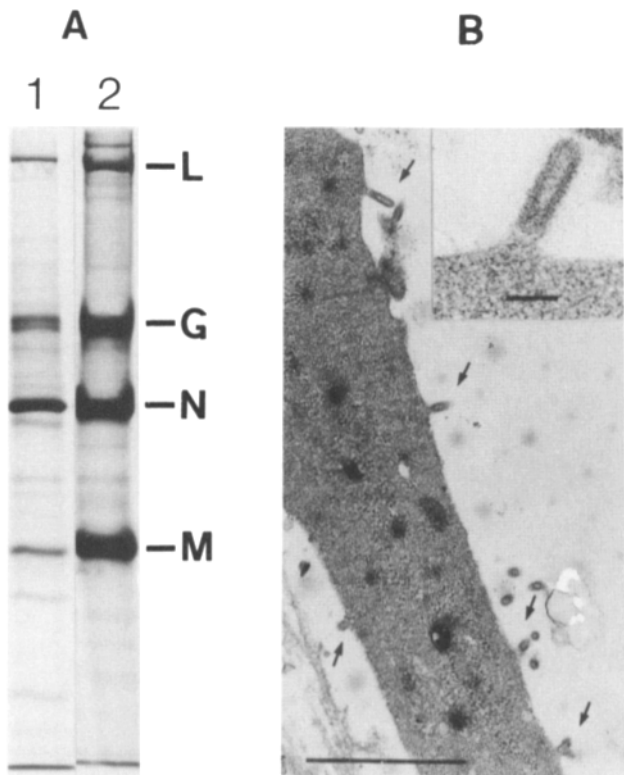


Figure 1. Viral particle formation in multinucleated L6 myotubes. The L6 multinucleated muscle cells were infected with VSV (2 pfu and grown for 4 h). In *A*, the cells were pulsed with [³⁵S]methionine and chased for 1 h. Electrophoretic analysis followed by autoradiography shows that viral proteins (L, G, N, M) were synthesized (lane 1). 5% of the postnuclear cell lysate was applied to the gel. Lane 2 shows analysis of the virions in the medium (see Materials and Methods). All viral proteins were identified, indicating viral particle formation. *B* shows an electron micrograph of budding viral particles (arrows) on a multinucleated muscle cell surface. The inset shows a budding virion in detail. Bars: (*B*) 1 μ m; (inset) 100 nm.

permeabilized cells. An example is shown in Fig. 2 *A*. The intracellular G protein localization showed an intense central region and the intensity disappeared with distance. The infected area generally varied between 5 and 25% of the total area of a multinucleated cell. A fine granular staining pattern typical of the myotube ER (Gu et al., 1989) was observed as shown in detail in Fig. 2 *C*. Double immunofluorescence staining for the ER (Fig. 2 *D*) strongly suggests that the mutant G protein in cells incubated at 39°C was located at the ER. In nonpermeabilized cells no viral products were visualized.

By utilizing double immunofluorescence staining for the VSV G and M proteins, we found that these proteins colocalized during early (<4 h) infection (not shown). After a 4-h infection period, the average extent of the intracellular staining was $220 \pm 64 \mu\text{m}$ (SD) when measured on five myotubes, all of which were longer than 1 mm. These infection patches generally did not colocalize with nuclei or nuclear groups, and areas devoid of nuclei were often found to locally express G protein. When an infection center colocalized

with a group of nuclei, fluorescent dots were often observed to surround vicinar nuclear groups.

To analyze whether the dispersion of the G protein was because of the exchange of ER membranes between nuclear groups, we stopped protein synthesis with cycloheximide (0.4 mM). Cycloheximide was added at 4 h after infection and the cells were fixed at 5 h after infection. Visualization of the G protein showed that the intense central staining disappeared (Fig. 3), suggesting that the G protein was part of a diffusible vesicle population. However, we could not demonstrate an increase in the size of the fluorescent G protein patches during the 1-h cycloheximide treatment, indicating that the G protein did not migrate far from the site of synthesis.

At 16 h after infection G protein localization spanned $420 \pm 76 \mu\text{m}$ (SD) when measured on 10 myotubes. In the same myotubes, the viral matrix protein showed smooth staining spanning 700–800 μm on the average (Fig. 4). Phase contrast microscopy revealed usually that the M protein staining did not cover the entire length of the myotubes. These measurements show that the G protein dispersed at a rate of $\sim 26 \mu\text{m}/\text{h}$ in multinucleated muscle cells while the corresponding rate for M protein was 40–50 $\mu\text{m}/\text{h}$. At this speed the viral infection should fill the cytoplasm of a 20 μm diameter mononuclear cell within 45 min.

Exocytic Transport Is Vertical in Multinucleated Muscle Cells

Incubation of mononucleated cells at 20°C blocks exocytic membrane traffic into a compartment called the *trans*-Golgi network (Matlin and Simons, 1983). We applied this block on myotubes infected with the VSV ts045. Accordingly, after a 3-h infection period at 39°C the locally infected myotubes were transferred to 20°C for 2 h. A prominent discontinuous perinuclear staining with diffuse dots in the cytoplasm was seen (Fig. 5 *A* and *C*). The infection remained local but in the majority of cases vicinar groups of nuclei shared the perinuclear G protein. A Golgi-specific marker partially colocalized with the G protein in the infected area after the 2-h incubation at 20°C (Fig. 5, *B* and *D*). Some G protein was probably left at the ER, explaining why the colocalization was not complete. It seems plausible that the perinuclear and dotted cytoplasmic staining represented the myotube *trans*-Golgi network. The presence of Nocodazole during the 20°C block did not scatter the observed circumnuclear ring structures (data not shown). Thus, depolymerization of microtubuli had little effect on the 20°C compartment. Furthermore, microtubuli were not essential for the transport from the ER into the 20°C compartment. This situation is thus comparable to that in the Cos cells where Nocodazole does not inhibit exocytic transport of the G protein (Featherstone et al., 1985).

Incubation for 30 min at 32°C after the 20°C block resulted in the appearance of the G protein on the cell surface. The surface staining was located over the sites where the internal G protein was seen, as shown by double immunofluorescence staining for the G protein before and after permeabilizing the cells (Fig. 6). Thus the G protein was transported from the presumed *trans*-Golgi network to the cell surface along the shortest route. After 2 h at 32°C, the surface G protein dispersed to a larger area on the myotubes, suggesting that at least a portion of the total G protein frac-

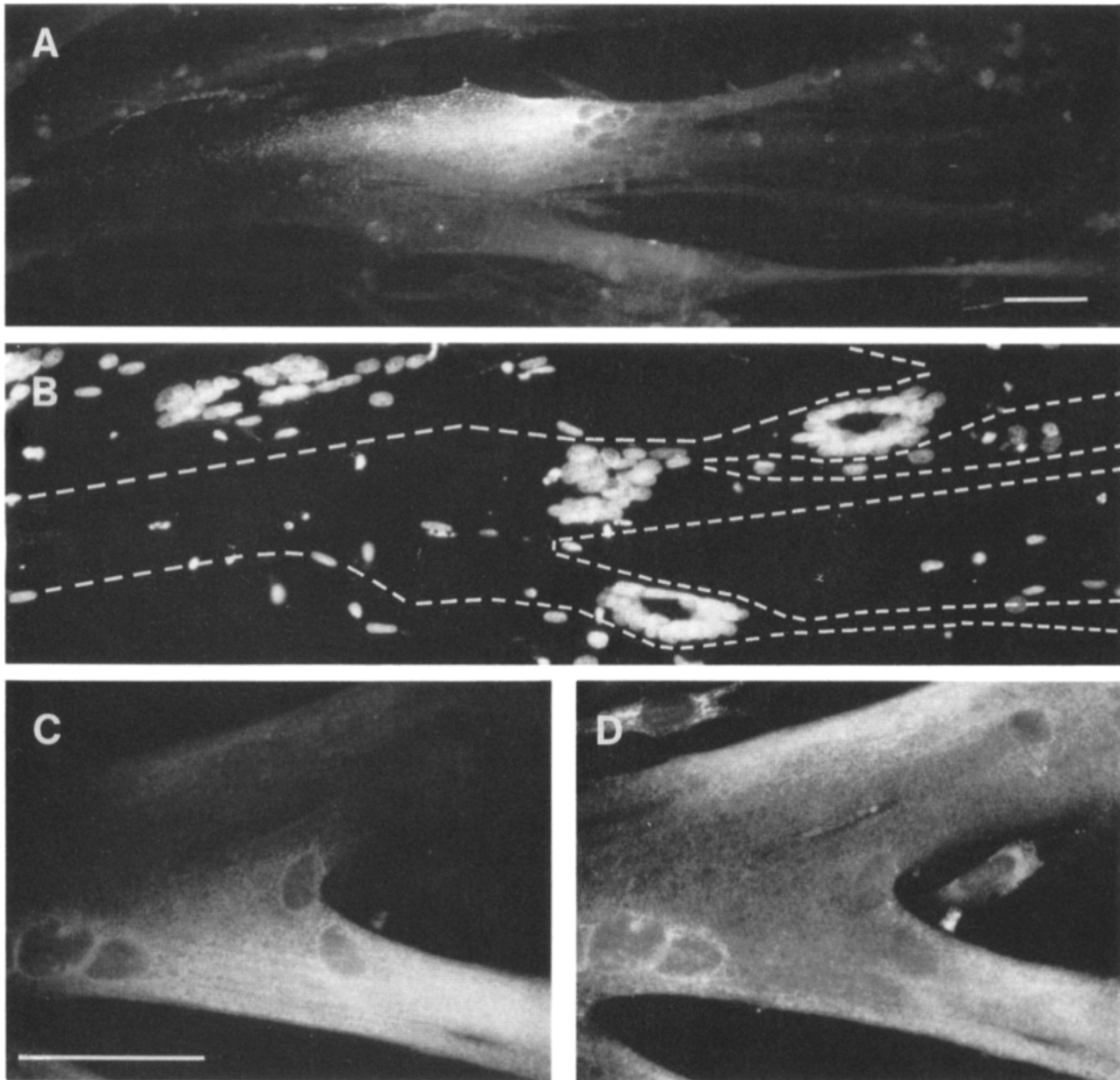


Figure 2. Distribution of VSV ts045 G protein in locally infected L6 multinucleated muscle cells at the nonpermissive temperature. In *A*, a local infection patch is shown. Polyclonal anti-G protein antibodies and TRITC-conjugated anti-rabbit IgG were used for the visualization. Nuclei in the same microscope field were visualized by Hoechst 33258 dye and are shown in *B*, together with the cell borders. The total length of the cell shown was 1.8 mm. *C* and *D* show double immunofluorescence staining with a mAb against the G protein and polyclonal anti-ER antibodies, followed by the appropriate FITC- and TRITC-coupled secondary antibodies. Note that the infection (*C*) remains local. Bars, 50 μ m.

tion was mobile. In mononucleated cells 75% of the surface G protein has been shown to be mobile (Scullion et al., 1987).

Double Infection in the Multinucleated Cells

Lytic viruses such as SFV or VSV cut off cellular protein synthesis and generally will not grow in the same cell (Met-sikkö and Garoff, 1989). To establish whether multinucleated cells could be simultaneously infected by two lytic viruses, we infected myotubes with VSV and SFV and followed the progress of this double infection. We found that

double infection in a multinucleated muscle cell was possible but that the two viral glycoproteins generally did not intermingle. As shown in Fig. 7, intracellular domains of VSV G protein were separated from those containing SFV p62 glycoprotein.

Extent of the mutual exclusion of VSV and SFV infections in multinucleated L6 muscle cells (as shown below). Infected areas for the VSV and SFV glycoproteins (see Fig. 7) were counted in five microscope fields. Percentages of the total number of the infected areas are given. The means and SD are shown:

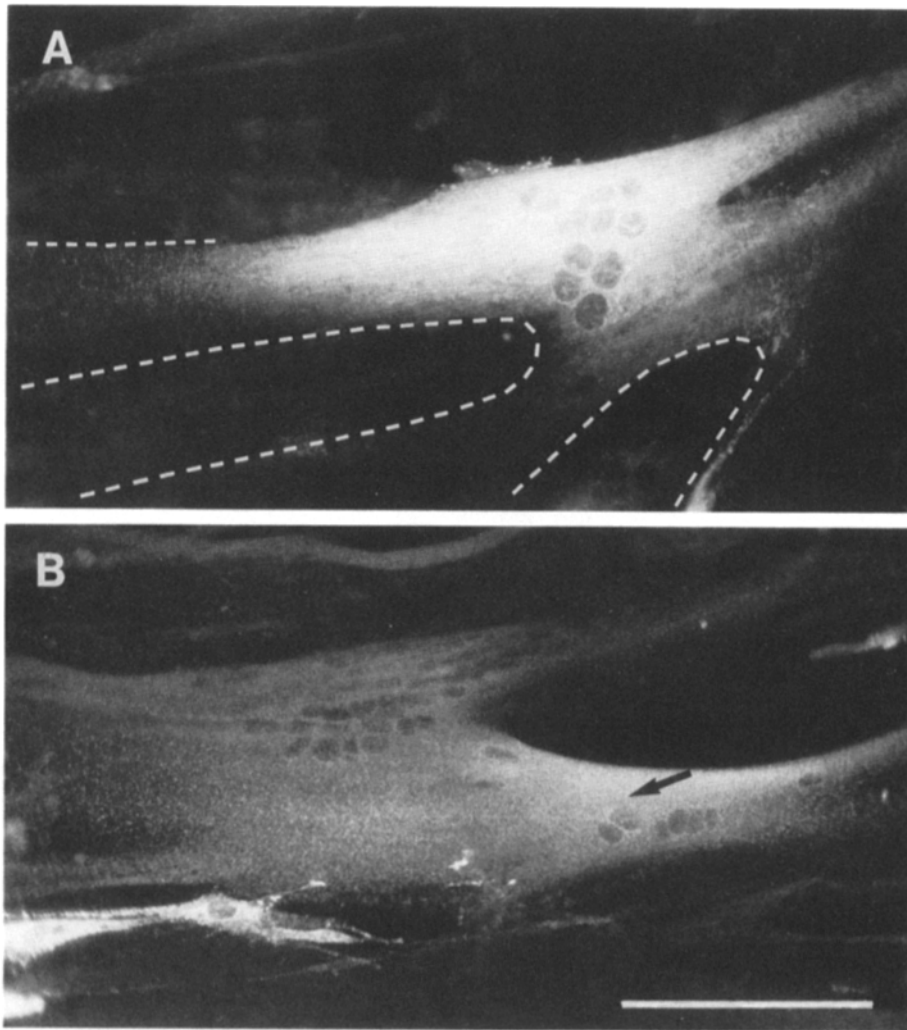


Figure 3. Dispersion of the viral glycoprotein in the ER. The VSV ts045 was grown for 3 h at 39°C, then fixed and processed for immunofluorescence. A localized distribution is seen (A). Cycloheximide was added after a 3-h growth period at 39°C to stop protein synthesis, and after 1 h treatment at 39°C the cells were fixed. The typical gradient-like distribution of the G protein disappears and a dot-like staining at 150 μm from the presumed infection center (arrow) is seen (B). The cells studied were >1 mm long. Bar, 100 μm .

only VSV G	only SFV p62/E2	VSV G and SFV p62/E2
41.6 \pm 4.8	41.3 \pm 15.0	17.1 \pm 10.6

The results indicate that exclusion occurred in 83% of the infected domains. The mean dimensions of these domains were 200–300 μm . Providing that the viral glycoproteins as such were not mutually exclusive, we conclude that the glycoproteins did not migrate far from their sites of synthesis. Immunofluorescence staining for the VSV G protein in non-permeabilized, doubly infected muscle cells showed uniform distribution on the cell surface (not shown), indicating that exocytosis occurred. Since the patches of intracellular G protein staining represented the viral glycoprotein in both the ER and the Golgi network, we conclude again, that the transport pathway from the ER to the Golgi network was locally restricted in the myotubes, and that lateral diffusion in the plasma membrane occurred.

Discussion

Our data indicate that the viral G protein was locally synthesized and inserted into the ER. This finding implies that the mRNA for the glycoprotein remained locally restricted, suggesting that the intracytoplasmic translocation of viral repli-

cation occurred slowly. Because the VSV ts045 G protein was blocked at the ER, we could follow the time course of its subsequent distribution within the ER. Since the VSV G protein remained localized for hours, we conclude that exchange of ER material within these large cells is slower than that of G protein degradation. Our results with cycloheximide suggest, however, that the protein in the ER could migrate short distances (Fig. 3). The average span of the G protein was 220 μm at 4 h after infection and 420 μm at 16 h after infection. For the CD8 membrane protein a 68 μm intracellular range was found (Ralston and Hall, 1989a). The reason for the discrepancy between these two results is probably that the viral infective RNA is more mobile than that of the nuclei. Moreover, CD8 was not blocked at the ER and therefore did not have time to disperse within the ER. We conclude that the longitudinal exchange of ER material occurred slowly. Thus, the ER in these large cells appears to be a relatively static compartment.

The infection patches were constantly found at areas without nuclei. This supports the idea that rough ER structures in myotubes do not belong to individual nuclei. Thus, the localization of the ER in multinucleated muscle cells is independent of nuclear positions. It appears that after myoblast fusion, their respective ER membranes completely intermin-

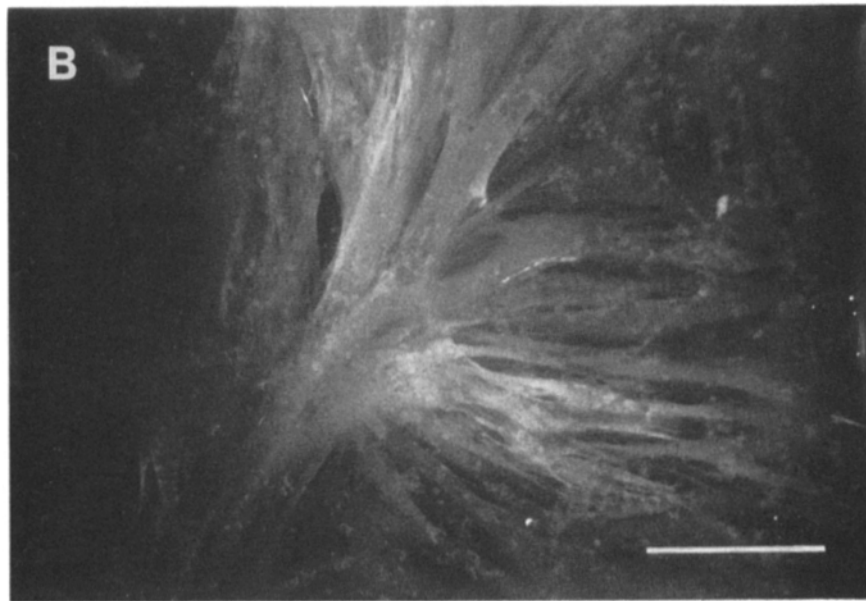
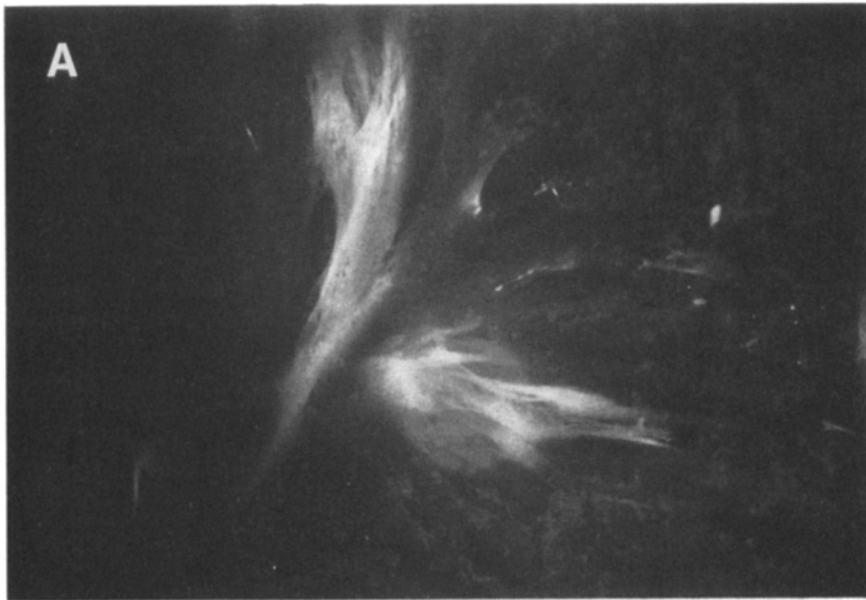
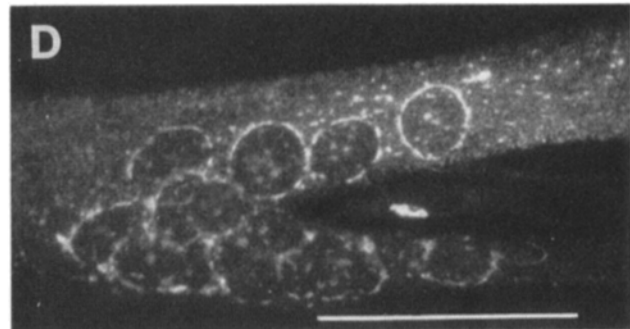
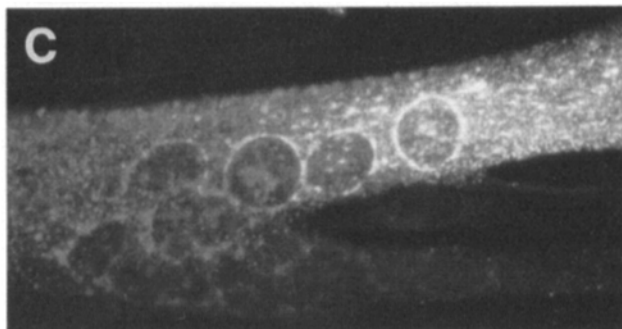
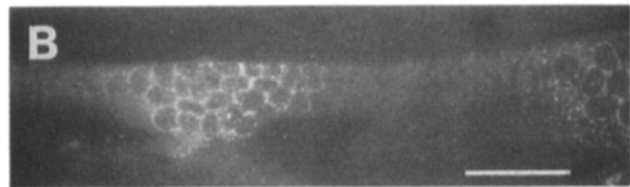
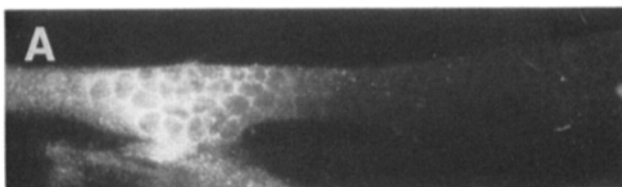


Figure 4. Double immunofluorescence staining of VSV G and M proteins at 16 h after infection in locally infected multinucleated cells. The cells were infected with VSV ts045 at 0.003 pfu for 1 h, then incubated at 39°C for 15 h, and fixed. *A* shows G protein distribution. Two local infections are seen. *B* shows the M protein distribution. Bar, 300 μ m.



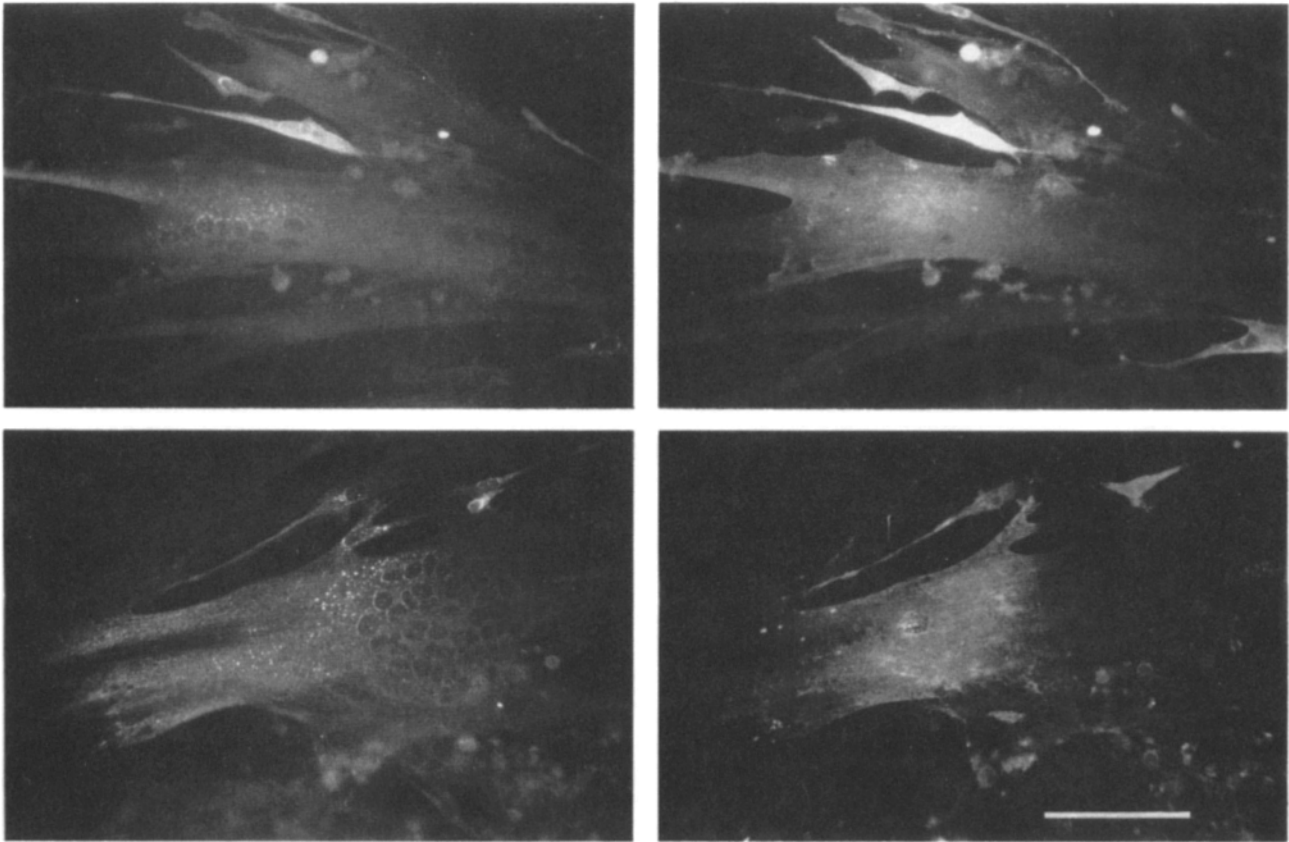


Figure 6. Examples of the surface appearance of the G protein in locally infected multinucleated muscle cells. Muscle cells were infected at 0.01 pfu and then grown for 3 h at 39°C, followed by a 2-h growth period at 20°C. To facilitate transport to the cell surface, the temperature was shifted to 32°C for 30 min. The left panels show internal staining while the corresponding surface stainings are shown on the right. Bar, 100 μm .

gle. Furthermore, the ER in myotubes is associated with microtubuli which also exhibit nuclear-independent localization (Tassin et al., 1985a).

We demonstrated myotube membrane structures where the G protein accumulated at 20°C. The structures were mainly perinuclear and resembled the Golgi apparatus previously described in myotubes (Tassin et al., 1985b; Gu et al., 1989). A Golgi-specific marker showed similar localization in double immunofluorescence staining (Fig. 5). Occasionally, dot-like staining after the block at 20°C was found at areas devoid of nuclei. Hence, the 20°C compartment was not totally perinuclear. However, in contrast to the situation with the ER, most of the 20°C compartment showed a structural association with the nuclei, i.e., the nuclear periphery. The localization of the 20°C compartment in myotubes differed from that of the *trans*-Golgi network described in mononuclear cells (Matlin and Simons, 1983; Griffiths and Simons, 1986). Whether or not this compartment functionally corresponds to the Golgi or the *trans*-Golgi network remains to be seen. The perinuclear staining was seldom restricted to a single group of nuclei but was generally also found as

a faint fluorescence in nearby groups of nuclei. This finding again suggests that the nuclei and their associated Golgi regions in the multinucleated cells share a common ER. In summary, we show that after the release of a 39°C block and subsequent incubation at 20°C, the locally expressed G protein in the ER was transported into circumnuclear structures surrounding several nuclear groups. At 32°C, transport into the overlying plasma membrane was observed. This situation is comparable with that found for the T lymphocyte antigen CD8, which is expressed in a single nucleus in multinucleated cells. This antigen may be visualized both as a localized intracellular patch and as being diffusely distributed over the entire cell surface (Ralston and Hall, 1989a). In contrast, an isoform of N-CAM (5.1H11) generally remains as a patch on the cell surface near the nucleus encoding it (Pavlath et al., 1989).

The VSV ts045 G protein dispersed with time in the multinucleated cells albeit slowly (26 $\mu\text{m}/\text{h}$). This also suggests that the intracytoplasmic relocation of the viral RNA replication was slow. SFV polymerase which is not packaged into viral particles is apparently located on cytopathic vacuoles

Figure 5. Distribution of the VSV ts045 G protein in multinucleated L6 muscle cells during a 20°C transport block. The G protein was visualized using polyclonal anti-G protein antibodies and a TRITC-conjugated anti-rabbit IgG. Perinuclear G protein staining is shown in A and C. The corresponding double immunofluorescence stainings with a monoclonal anti-Golgi antibody and a FITC-conjugated anti-mouse IgG are shown in B and D. The infections (A and C) are local. Bar, 50 μm .

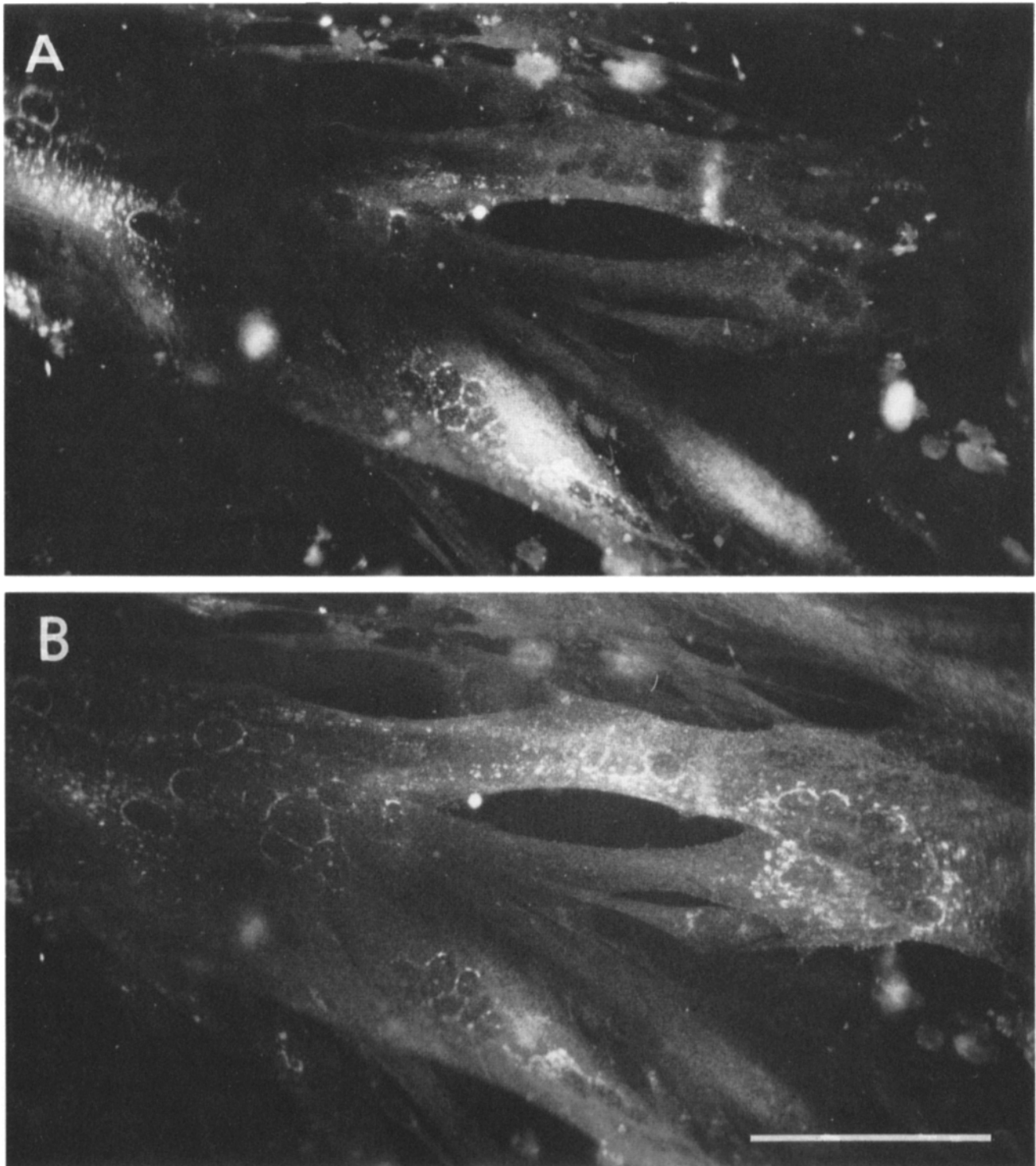


Figure 7. Double infection of multinucleated muscle cells with VSV and SFV. Cells were infected at 2 pfu of both VSV and SFV simultaneously. At 5 h after infection the cells were fixed and subjected to double immunofluorescence staining for the VSV G protein (*A*) or the SFV p62 glycoprotein (*B*). Bar, 100 μm .

in BHK or CHO cells (Froshauer et al., 1988). It seems that the replication of the SFV RNA is restricted to the sites of viral entry. However, Peränen and Kääriäinen (1991) found that one incoming virion induced the formation of numerous cytopathic vacuoles, suggesting that viral RNA can move from membrane to membrane. The situation with the VSV

polymerase remains to be elucidated. We think that the dispersion of the G protein was not exclusively due to the movement of ER membranes. Rather, viral replication moved short distances. The finding that the M protein dispersed faster than the G protein could be because of the cytoplasmic nature of the M protein. It is known that a fraction of the M

protein is soluble in infected cells (Knipe et al., 1977). Furthermore, cytoplasmic proteins have been shown to diffuse freely in myotubes (Minz and Baker, 1967; Ralston and Hall, 1989b).

We show here that SFV and VSV can be propagated in a single multinucleated cell but that both viruses occupy their own domains of glycoprotein expression (Fig. 7). It is likely that the mRNA synthesis of both viruses was also restricted to such domains. It is reasonable to think that neither the glycoprotein nor the replicative viral RNA ranged more than $\sim 200 \mu\text{m}$ in the doubly infected cells. We think that the first virus to be uncoated by endocytosis initiated local mRNA synthesis and occupied the protein synthesis machinery in a domain spanning $\sim 200 \mu\text{m}$.

In summary, in multinucleated L6 myoblast-derived muscle cells viral protein synthesis did not recognize nuclear domains at the ER level but still remained localized for a considerable period of time. This behavior reflects a static characteristic of the ER. Thus, there was little exchange of material between the ER membranes at different longitudinal parts of myotubes. At later steps of exocytosis, the membrane compartments were at least to some extent associated with individual nuclei. Thus there is strong supporting evidence for the functionality of nuclear domains. First, there is restricted longitudinal transport and second, organelles such as the Golgi network or the 20°C compartment are preferentially associated with individual nuclei. This situation provides the basis for positional information in multinucleated cells.

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