



Mosquito cells infected with Japanese encephalitis virus release slowly-sedimenting hemagglutinin particles in association with intracellular formation of smooth membrane...

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博士論文

Mosquito Cells Infected with Japanese Encephalitis Virus Release
Slowly-Sedimenting Hemagglutinin Particles in Association with
Intracellular Formation of Smooth Membrane Structures

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2 Slowly-Sedimenting Hemagglutinin Particles in Association with
3 Intracellular Formation of Smooth Membrane Structures

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14 Running head: SMS-SHA association in JEV-infected mosquito cells

15 Key words: Encephalitis Virus, Japanese: ultrastructure: Virus Particle: Morphogenesis

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1 **Abstract**

2

3 Arthropod-borne flaviviruses can grow in both arthropod and mammalian cells.

4 Virion morphogenesis, though well studied in mammalian cells, is still unclear in

5 arthropod cells. Here, we compared a mosquito cell line C6/36 and a mammalian cell

6 line Vero in extracellular virus particles and intracellular ultrastructures triggered by

7 infection with Japanese encephalitis virus (JEV). Sedimentation analyses of virion and

8 slowly-sedimenting hemagglutinin (SHA) particles released by infection with the

9 Nakayama strain revealed that C6/36 cells produced higher envelope (E) antigen levels

10 in the SHA than the virion fraction in contrast to Vero cells that showed the opposite

11 pattern. Specific infectivities per ng of E were similar in both cells, whereas specific

12 hemagglutinating activities in the SHA fraction were lower in C6/36 than Vero cells.

13 The precursor membrane protein was less efficiently cleaved to the membrane protein in

14 SHA particles released from C6/36 than Vero cells. Ultrastructural studies showed more

15 remarkable production of smooth membrane structures (SMSs) in C6/36 than in Vero

16 cells. The differences in sedimentation patterns of extracellular virus particles between

17 Nakayama-infected C6/36 and Vero cells were consistently observed in 5 other strains

18 (Beijing P1, Beijing P3, JaTH-160, KE-093 and JaGAR-O1), except for KE-093-infected

1 C6/36 cells which exhibited the Vero-type sedimentation profile under conditions of
2 open cultivation. By electron microscopy, the production of SMSs from
3 KE-093-infected C6/36 cells under open conditions was markedly less than that under
4 closed conditions where the cells exhibited the C6/36-type sedimentation profile. Thus,
5 intracellular SMS formations were associated with extracellular SHA production in
6 JEV-infected mosquito cells.

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8 Key words: Encephalitis Virus, Japanese: ultrastructure: Virus Particle: Morphogenesis

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1 Introduction

2

3 Japanese encephalitis virus (JEV) is a member of the genus *Flavivirus* family
4 *Flaviviridae*. *Flavivirus* virions, 40-60 nm in diameter, contain a nucleocapsid about 30
5 nm in diameter that is surrounded by an envelope. The nucleocapsid is composed of a
6 capsid (C) protein and a single strand of positive-sense RNA, while the envelope is
7 composed of a lipid bilayer derived from the host cell, an envelope (E) protein and a
8 membrane (M) protein. A small amount of the precursor of M (prM) is usually
9 contained in the envelope. In most cases, flavivirus-infected cells release not only
10 virions but also empty particles that do not contain a nucleocapsid. These particles
11 sediment more slowly than do virions in sucrose density gradient centrifugation and are
12 named slowly-sedimenting hemagglutinin (SHA) particles.

13 *Flavivirus* virion morphogenesis has been extensively studied (35). Structural
14 proteins C, prM and E, and seven nonstructural (NS) proteins are synthesized in order
15 from the start of a single long open reading frame encoded by the flavivirus genome.
16 Transmembrane domains consisting of hydrophobic amino acids at the carboxy-termini
17 of C and prM play a role as the signal that directs translocation of prM and E to the
18 lumen of the endoplasmic reticulum (ER). Carboxy-termini of C, prM and E are cleaved

1 on the luminal side of the ER membrane by the host enzyme signalase, and prM and E
2 are fixed on the ER membrane as an N-linked type I membrane glycoprotein with the
3 transmembrane domain as a membrane anchor. Cleavage of the junction of the
4 cytoplasmic and transmembrane domains of C is mediated by the viral protease
5 NS2B/NS3. Budding from the prM/E-accumulated ER membrane into the lumen to
6 envelope a nucleocapsid produces an immature virion, whereas particle formation
7 without nucleocapsids is considered to produce SHA particles (2, 6). These viral
8 particles are transported through a secretory pathway to the trans-Golgi network where
9 prM is cleaved to M by a protease furin (41). Cleavage of prM to M that converts
10 prM/E heterodimers to E homodimers results in the acquisition of infectivity and
11 hemagglutinating (HA) activity. When exposed to low pH, E homodimers are
12 irreversibly converted to a homotrimer form, resulting in the acquisition of the
13 membrane fusion competence (1). Structural analyses of the surface of flavivirus
14 particles by X ray crystallography (5, 39) and cryo-electron microscopy (11, 24, 34, 36,
15 44, 45) have shown E homodimers on the particles and low pH-induced conversion to E
16 homotrimers; however, it is suggested that a part of E remains to form a heterodimer
17 with prM.

18 In morphological studies, virions and recombinant subviral particles that are

1 similar to SHA particles have mainly been observed in the lumen of the ER and the
2 Golgi apparatus in both mammalian and mosquito cells (4, 15, 16, 27, 29, 37). On the
3 other hand, the budding intermediates into the lumen of the ER to form virion or SHA
4 particles have not been frequently observed, probably because the budding process is
5 rapid. The most prominent features observed in flavivirus-infected cells are induction of
6 membrane proliferation particularly in the perinuclear region. Characteristic membrane
7 structures and the dilated ER have been observed in flavivirus-infected mammalian and
8 mosquito cells (10, 14, 26, 30, 43).

9 JEV is a mosquito-transmitted agent and has the ability to propagate in both
10 mammalian and mosquito cells. However, the course of infection is different between
11 the two cell species, as can be seen in other mosquito-borne flaviviruses (35). Most
12 remarkably, JEV infection induces cytopathic effects and cytolysis in most of
13 mammalian cells, whereas the infection of mosquito cells is usually noncytopathic and
14 may establish persistent infections (19, 38). Since the mechanism of flavivirus virion
15 morphogenesis described above has mainly been established using mammalian cells, it
16 is important to study mosquito cells as another host of JEV infection. The mechanism of
17 virion maturation in mosquito cells may contribute to the development of a strategy for
18 the prevention of JEV transmission, and could include a possible new type of vaccine

1 targeting this invertebrate vector.

2 The limited electron microscopic studies conducted to date on virion
3 morphogenesis in JEV-infected mosquito cells have shown virus particles in the ER
4 lumen, Golgi apparatus and coated vesicles; as well, characteristic membrane structures
5 named smooth membrane structures (SMSs) were also observed (14, 16, 26). In this
6 study, as our initial approach to the mechanism of virion maturation in mosquito cells,
7 we performed basic biochemical and biophysical analyses of virus particles released
8 from JEV-infected cells, as well as morphological studies of infected cells utilizing
9 electron microscopy. Our results indicated that JEV-infected mosquito cells released
10 higher levels of E antigen in SHA than virion particles, and that the intracellular
11 formation of SMSs was associated with the release of SHA particles from infected
12 mosquito cells.

13

14 **Materials and Methods**

15

16 **Cells.** Vero (22) and C6/36 (18) cells were used as mammalian and mosquito
17 cell lines, respectively. Vero cells were cultivated in a growth medium composed of
18 Eagle's minimal essential medium (MEM) supplemented with 10% heat-inactivated

1 fetal bovine serum (FBS) at 37 C in a humidified atmosphere of 5% CO₂-95% air.
2 C6/36 cells were grown under the same conditions as in Vero cultures except for the
3 addition of 1 mM nonessential amino acids to the medium and that the cultivation
4 temperature was 28 C.

5 **Viruses.** Six strains of JEV were used in this study. The Nakayama and Beijing
6 P3 strains have been described (23). The Beijing P1 strain, which had been passaged
7 eleven times through suckling mouse brains (SMBs) and one time through Vero cells
8 was provided by Dr. Robert E. Shope of Yale Arbovirus Research Unit, Yale University
9 School of Medicine, CT, USA. The JaTH-160 strain, which had been passaged three
10 times through SMBs, and the KE-093 strain, which had been passaged four times
11 through SMBs, were provided by Dr. Ichiro Kurane of National Institute of Infectious
12 Diseases, Japan. The JaGAR-O1 strain, which had been passaged twelve times through
13 SMBs, was provided by Dr. Eiji Kimura of Hyogo Prefectural Institute of Health, Japan.
14 In our laboratory, these strains were passaged one or two times through C6/36 cells and,
15 the infected culture fluid was used in this study.

16 **Virus infection.** C6/36 and Vero monolayer cells were infected with each
17 strain of JEV at a multiplicity of infection (m.o.i.) of 5 plaque-forming units (PFU)/cell.
18 After 1-hr adsorption, cells were rinsed three times with phosphate-buffered saline

1 (PBS), and incubated at temperatures corresponding to each cell line in a maintenance
2 medium that was the growth medium replacing 10% FBS to 0.075% bovine serum
3 albumin. Culture fluids were harvested at 24, 48 and/or 72 hr postinfection (p.i.).

4 Infected cells were cultivated in culture flasks or dishes (including 6-well
5 microplates). In some experiments, the cap of the culture flask was tightly closed to
6 eliminate gas exchange with the 5% CO₂-95% air environment (closed cultivation),
7 different from the other conditions where the cap was loosened enough to allow gas
8 exchange (open cultivation). Cultivations using culture dishes were regarded as an open
9 condition.

10 **Precipitation with polyethylene glycol (PEG).** Culture fluids harvested from
11 infected cells, 40 ml in volume from 4 75-cm² culture flasks, were clarified and
12 incubated in 10% PEG and 1.9% NaCl for 2 hr at 4 C. Following centrifugation at
13 10,000 rpm for 30 min in the R22A rotor of a Himac CR22E (Hitachi Koki Co. Ltd.,
14 Ibaraki, Japan), the pellet was suspended in 1 ml TN buffer (10 mM Tris-HCl pH 7.5,
15 100 mM NaCl) and clarified. This suspension was further concentrated by one more
16 PEG precipitation in a similar way, to obtain 200 µl of the concentrate suspended in TN
17 buffer.

18 **Sucrose density gradient centrifugation.** Culture fluids or PEG-concentrated

1 culture fluids were applied to a 10 to 40% (wt/wt) continuous sucrose gradient prepared
2 in TN buffer. Following centrifugation for 90 min at 55,000 rpm at 4 C in the S55S rotor
3 of a Himac CS100GX microultracentrifuge (Hitachi Koki Co. Ltd., Ibaraki, Japan),
4 fractions were collected from the bottom.

5 **ELISA for measuring E antigen levels.** E antigen levels were measured by a
6 sandwich ELISA (21). Briefly, 96-well microplates sensitized with rabbit anti-JEV
7 polyclonal antibody were incubated with samples (sucrose gradient fractions) at 1:10
8 dilution, a monoclonal antibody J3-11B9 (33) recognizing E antigen, alkaline
9 phosphatase-conjugated anti-mouse IgG and then p-nitrophenyl phosphate. The E
10 protein amount was calculated by comparing absorbances at 415 nm with those
11 obtained from a recombinant E protein standard prepared from prM/E-expressing
12 mammalian cells.

13 **Plaque assay and HA test.** Infective titers were determined on a monolayer of
14 Vero cells as previously described (22). HA titers were determined by a
15 micro-modification of the method described by Clarke and Casals (9): titers obtained
16 with 25 µl of the sample were expressed in HA unit (HAU). Specific infectivity and HA
17 activities were obtained by calculating infective and HA titers per ng of the E protein,
18 respectively.

1 **Endoglycosidase treatment.** The virion and SHA fractions obtained by
2 sucrose density gradient centrifugation from PEG precipitated culture fluids were
3 centrifuged at 55,000 rpm for 90 min at 4 C in the S55S rotor (see above) and the pellets
4 were then dissolved in 20 µl of the “sample” buffer (0.1 M Tris-HCl (pH 6.8), 4%
5 sodium dodecyl sulfate (SDS), 4 mM ethylenediamine-N, N, N', N'-tetraacetic acid,
6 30% glycerol, 0.2 g/ml bromophenol blue). A half (10 µl) of this sample solution were
7 mixed with 2 µl of the “reaction” buffer (10% SDS, 10% 2-mercaptoethanol) and
8 incubated at 100 C for 5 min. After cooling, one unit of N-glycosidase F (PNGase F;
9 Roche diagnostics, Basel, Switzerland) was added to the sample and incubated at 37 C
10 for 24 hr. For inactivation of PNGase F, the samples were incubated at 100 C for 5 min.

11 **Western blot analysis.** The analysis was performed essentially as previously
12 described (21). Briefly, serial 2-fold diluted PNGase F-treated samples were separated
13 by 14% SDS-polyacrylamide gel electrophoresis at 150 V for 75 min and transferred to
14 polyvinylidene difluoride membrane at 200 mA for 120 min. The membrane was
15 incubated with a monoclonal antibody J2-2F1 (33) recognizing prM/M, alkaline
16 phosphatase-conjugated anti-mouse IgG and then nitroblue tetrazolium and
17 5-bromo-4-chloro-3-indolyl phosphate substrate. The levels of prM and M were
18 expressed as the reciprocal highest dilution representing a visible band (see Fig. 3). To

1 compare viral particles obtained from JEV-infected C6/36 and Vero cells, prM:M ratios
2 were calculated from the levels of prM and M.

3 **Electron microscopy.** C6/36 and Vero monolayer cells prepared in culture
4 dishes were infected at an m.o.i. of 5, incubated for 48 hr and fixed with 2%
5 paraformaldehyde-0.25% glutaraldehyde in PBS at room temperature for 4 hr.
6 Following rinsing with PBS, cells were further fixed with 2% osmium tetroxide in PBS
7 at room temperature for 1 hr. Cells were dehydrated with a graded ethanol series and
8 embedded with Epok 812 (Okenshoji Co. Ltd., Tokyo, Japan). Sections were obtained
9 by cutting Epok-embedding samples with a Leica UltracurR (Leica Microsystems
10 GmbH, Vienna, Austria) and stained with uranyl acetate and then with lead citrate.,
11 Sections were observed under a JEM-1220 electron microscope (JEOL Ltd., Tokyo,
12 Japan).

13

14 **Results**

15

16 **Comparison of virus particles released from C6/36 and Vero cells infected**
17 **with the JEV Nakayama strain.** To perform a basic characterization of the virus
18 particles released from C6/36 and Vero cells infected with JEV and to assess the effect

1 of the harvest time point, we first used a prototype Nakayama strain. The
2 Nakayama-infected C6/36 and Vero cells were cultivated in a tissue culture flask under
3 the closed condition and culture fluids obtained at 24, 48 and 72 hr p.i. were analyzed
4 on sucrose density gradients in terms of their E antigen levels, and infective and HA
5 titers (Fig. 1). At any time point, the sedimentation profiles obtained with infected
6 C6/36 cells indicated higher levels of E antigen in the SHA than the virion fraction,
7 whereas infected Vero cells released higher E antigen levels in the virion than the SHA
8 fraction. Based on this profile characteristic of C6/36 cells, subsequent analyses were
9 focused on SHA particles, as well as virions, for directing the events occurring in
10 infected mosquito cells.

11 The results obtained in Fig. 1 were further analyzed to obtain time courses of
12 the ratio of E antigen levels between the virion and SHA fractions (virion:SHA ratio;
13 Fig. 2a), specific infectivities and HA activities (Fig. 2b, c). The virion:SHA ratios
14 obtained from infected C6/36 cells were <1 and those obtained from infected Vero cells
15 were >1 with differences of >10 -fold during the experimental period of 24-72 hr (Fig.
16 2a). Specific infectivities of the virion fraction obtained with infected C6/36 cells were
17 similar to those obtained with infected Vero cells (Fig. 2b). Specific HA activities of the
18 virion and SHA fractions from infected C6/36 cells were lower than those of the

1 corresponding fractions from Vero cells (Fig. 2c). In addition, specific HA activities of
2 the virion fractions were higher than those of the SHA fractions in both cells.

3 Furin-mediated cleavage of prM to M is a critical step for the flavivirus virion
4 to subsequently acquire infectivity and related viral activities. We therefore compared
5 levels of prM and M in each of the virion and SHA fractions. For this purpose, a
6 semiquantitative method based on Western blotting analysis was used. A typical result
7 of this method is shown in Fig. 3, which represents prM and M bands of virion and SHA
8 fractions obtained from JEV-infected C6/36 and Vero cells at 48 hr p.i. Although all
9 24-hr samples showed undetectable levels and one 72-hr sample showed levels of
10 $\geq 2,048$ for both M and prM, the 48-hr samples allowed us to compare 4 data (Fig. 2d)
11 and showed that the prM:M ratios obtained with the SHA fraction were higher than
12 those obtained with the virion fraction in both C6/36 and Vero cells. In addition, the
13 prM:M ratio in the SHA fraction obtained from C6/36 cells was higher than that
14 obtained from Vero cells. Although only the 48-hr samples were used for comparison in
15 Fig. 2d, similar results were obtained in a repeated experiment (data not shown).

16 These results indicate that virus particles released from JEV-infected C6/36 and
17 Vero cells had different natures and that their surface proteins had different functions.

18 **Ultrastructural changes induced in Nakayama-infected C6/36 and Vero**

1 cells. To compare the ultrastructural changes induced by JEV infection between C6/36
2 and Vero cells, we used Nakayama-infected C6/36 and Vero cells at 48 hr p.i. for
3 electron microscopic examination. Although culture dishes were used for preparation of
4 cell samples and therefore cells were cultivated under an open condition, the culture
5 fluids obtained from both cell lines showed consistent sedimentation profiles (data not
6 shown) with those of the corresponding harvest time (48 hr) shown in Fig. 1.

7 At a low magnification, numerous characteristic vesicles of varying sizes were
8 distributed in the whole cytoplasmic area of infected C6/36 cells (Fig. 4a), whereas
9 infected Vero cells did not contain such large-sized vesicles (Fig. 4e). At a high
10 magnification, the dilated ER and SMSs were observed in both infected C6/36 and Vero
11 cells (Fig. 4b, f, respectively), neither of which were found in uninfected cells (data not
12 shown). However, C6/36 cells exhibited a higher degree of membrane proliferation than
13 Vero cells. The production of SMSs was limited mainly to the perinuclear regions in
14 infected Vero cells (Fig. 4g, h), whereas these abnormal membrane structures were
15 observed at similar frequencies both in the perinuclear and peripheral regions of
16 infected C6/36 cells (Fig. 4c, d). Even in the perinuclear regions, SMSs were more
17 remarkably observed in C6/36 than Vero cells (Fig. 4c versus 4g). Virions were
18 observed in the ER in both C6/36 and Vero cells (Fig. 4d, g) and, though in a small

1 number, in SMSs in C6/36 cells (Fig. 4b, c; arrow heads). No virions were found in
2 SMSs in Vero cells, as far as the samples examined in this study were concerned.

3 **Comparisons of virus particles released from C6/36 and Vero cells infected**

4 **with other strains of JEV.** To examine if the difference between extracellular viral

5 particles released from Nakayama-infected C6/36 and Vero cells shown in Fig. 1 would

6 be a common phenomenon in JEV-infected cells, 5 additional strains of JEV were

7 examined under the same characterization methods. The strains we used were Beijing

8 P1, Beijing P3, JaTH-160, KE-093 and JaGAR-O1: for convenience, infected cells were

9 cultivated in a 6-well culture plate. Since similar results were obtained at any time point

10 after infection (24, 48 and 72 hr) using the Nakayama strain (Fig. 2), culture fluids

11 harvested at 48 hr p.i. were used for sedimentation analyses of the 5 strains (Fig. 5).

12 Infected C6/36 cells released higher levels of E antigens in the SHA than the virion

13 fraction and the virion:SHA ratios were <1 in most strains (Table 1), consistent with the

14 results obtained from Nakayama-infected C6/36 cells. In all cases, infected Vero cells

15 released higher levels of E antigens in the virion than the SHA fraction and the

16 virion:SHA ratios were >1 (Table 1), again consistent with the results obtained from

17 Nakayama-infected Vero cells. However, C6/36 cells infected with the KE-093 strain

18 released higher levels of E antigens in the virion than in the SHA fraction with the

1 virion:SHA ratio being >1, similar to the pattern obtained with Vero cells infected with
2 Nakayama and the 5 other strains.

3 Specific infectivities of the virion fraction obtained from infected C6/36 cells
4 ranged from 5.0 to 24.3×10^4 in 6 strains including the Nakayama and those obtained
5 from infected Vero cells varied from 2.1 to 15.3×10^4 (Table 2), similar to previously
6 obtained specific infectivities from Nakayama-infected C6/36 and Vero cells (Fig. 2b).
7 Comparison of specific HA activities (Table 3) indicated that the activity of the virion
8 and SHA fractions obtained from infected C6/36 cells were lower than the
9 corresponding fractions of Vero cells with some exceptions including the virion fraction
10 obtained with KE-093-infected cells. In all strains and both cell species, specific HA
11 activities were higher in the virion than the SHA fractions. Further, comparison of the
12 prM:M ratios (Table 4) indicated that the ratios obtained with the SHA fraction were
13 higher in C6/36 than Vero cells in any strain and those obtained with the virion fraction
14 were higher in C6/36 than Vero cells in 4 of the 6 strains. In all strains for C6/36 cells
15 and in almost all strains for Vero cells, the prM:M ratios obtained with the SHA fraction
16 were higher than those obtained with the virion fraction: however, there was a low level
17 of prM and an extremely high level of M in the SHA fractions from KE-093-infected
18 C6/36 and Vero cells, respectively.

1 These results indicated that quantitative and qualitative differences in virus
2 particles released from C6/36 and Vero cells infected with the Nakayama strain were a
3 common feature of JEV infections, with the exception of the KE-093 strain.

4 **Comparison of virus particles released from KE-093-infected C6/36 cells**
5 **under the open and closed cultivating conditions.** To further examine the exceptional
6 nature of the virus particles released from KE-093-infected C6/36 cells, we attempted to
7 compare sedimentation profiles of culture fluids obtained from cells cultivated in
8 culture flasks under the open and closed conditions (Fig. 6). Under the closed
9 cultivating condition, KE-093-infected C6/36 cells released higher levels of E antigens
10 in the SHA than the virion fraction, showing the original C6/36-type sedimentation
11 profile shown by the other strains of JEV. On the other hand, the open cultivation of
12 KE-093-infected C6/36 cells released higher levels of E antigens in the virion than the
13 SHA fraction, showing the Vero-type sedimentation profile. The virion:SHA ratio
14 obtained with the open and closed cultivating conditions were 2.73 and 0.34,
15 respectively; comparable to the results obtained with Vero and C6/36 cells infected with
16 the other strains.

17 Specific infectivities of the virion fraction obtained under the open (48.73) and
18 closed (67.23) cultivating conditions were equivalent. Although specific HA activities

1 of the virion fraction obtained from the open (3.00) and closed (2.69) cultivating
2 conditions were higher than those of the SHA fraction (<0.1 under both the open and
3 closed cultivating conditions), the values obtained under both cultivating conditions
4 were similar. In addition, the prM:M ratios obtained with virion (0.06) and SHA (1.00)
5 under the closed cultivating condition did not differ from previously obtained values
6 (0.031 for virion, 1.00 for SHA, Table 4). These results indicated that although particle
7 formation was affected by the cultivating condition in KE-093-infected C6/36 cells as
8 shown by sedimentation profiles, the effect of the cultivating condition was not large
9 enough to detect the difference in the nature of virus particles.

10 **Ultrastructural changes of KE-093-infected C6/36 cells under the open and**
11 **closed cultivating conditions.** In addition to the above morphological approach using
12 different cell species (C6/36 and Vero cells), a study of comparisons in a single cell
13 species (C6/36) showing differences in extracellular virus particles under different
14 cultivating conditions is another model appropriate for studying associations of
15 extracellular virus particles with intracellular morphological changes. We therefore
16 compared the ultrastructural features of KE-093-infected C6/36 cells under the open and
17 closed cultivating conditions (Fig. 7). Although SMSs were observed throughout the
18 cytoplasmic area under both conditions, remarkably reduced SMS production was noted

1 under the open cultivating condition. The formation of the dilated ER was also reduced
2 in KE-093-infected C6/36 cells under the open cultivating condition. Virions were
3 observed in the ER and much less in SMSs under both open and closed cultivating
4 conditions.

5

6 **Discussion**

7

8 This study demonstrated that virus particles released from JEV-infected cells
9 have different natures, respectively, in two host cell species, C6/36 and Vero. C6/36
10 cells released higher levels of E antigens in the SHA than the virion fraction, whereas
11 Vero cells released higher levels of E antigens in the virion than the SHA fraction. The
12 difference between these mosquito and mammalian cells was also observed in
13 ultrastructural approaches, which indicated that SMSs were more markedly observed in
14 C6/36 cells than in Vero cells. Thus, the present comparative studies using C6/36 and
15 Vero cells showed preferred release of SHA particles from JEV-susceptible cells in
16 which increased intracellular SMS formation occurred. Further, the association of SMS
17 formations with SHA releases was strongly supported by comparisons between open
18 and closed cultivating conditions using a single cell species (C6/36) infected with the

1 KE-093 strain: the use of the single cell species could rule out any possible effect
2 derived from the difference of host cells.

3 Virus-induced membrane structures have generally been observed in
4 flavivirus-infected cells. Smooth membrane vesicles similar to SMSs were shown in
5 dengue virus-infected mammalian and mosquito cells (4, 29). In Kunjin virus-infected
6 mammalian cells, vesicle packets (VPs) containing SMS-like vesicles, convoluted
7 membranes localized adjacent to VPs and paracrystalline arrays, which were similar to
8 the structures observed when microtubules were disrupted , have been reported (28, 32,
9 43). In hepatitis C virus-infected cells, contiguous vesicles and membranous webs are
10 observed (10). Moreover, such membrane proliferations have been found in cells
11 contained in the midgut epithelium, midgut muscles, and salivary glands in West Nile
12 virus-infected mosquitoes *in vivo* (12). Thus, flavivirus infections can induce a variety
13 of intracellular membrane structures in mammalian and mosquito cells.

14 The roles of these induced membrane structures are still unclear. However, the
15 association of virus-induced membranes with viral RNA and nonstructural proteins has
16 been reported. dsRNA, NS1 and NS3 were localized in association with the induced
17 membrane structures of Kunjin virus-infected cells (43). In dengue virus-infected cells,
18 the induced membrane structures were related with the localization of genome RNA

1 (13), dsRNA and NS1 (30) and NS3 (8). In JEV-infected cells, NS3 was found localized
2 in association with virus-induced microtubules (7). These results suggest that the
3 induced membrane structures are related to particle formation and the subsequent
4 release of virion and SHA particles in flavivirus-infected cells.

5 In the present study, JE virions were observed in SMSs in infected C6/36 cells,
6 although the number was much smaller than that in the ER that is considered as a major
7 factory of viral particles. The presence of virions in SMSs suggests that this type of
8 membrane structure is another factory related to both virion and SHA formations. Since
9 it is unlikely that viral particles produced in SMSs are transported through a normal
10 intracellular secretory pathway, these particles are considered different in nature from
11 those released by an anterograde transport from the ER through the Golgi apparatus to
12 the periphery, particularly in terms of furin-mediated prM cleavage. Consistent with this
13 hypothesis, extremely higher prM:M ratios were shown in SHA particles released from
14 C6/36 than Vero cells. Further, specific HA activities in SHA samples obtained from
15 C6/36 cells were lower than those from Vero cells in most of the JEV strains. It is
16 probable that preferred release of SHA to virion particles in C6/36 cells with increased
17 SMS formation may be attributed to an inefficient envelopment of the nucleocapsid in
18 an abnormal factory (SMS), resulting in the marked differences in the prM:M ratio

1 between SHA particles released from C6/36 and Vero cells. The potential production of
2 viral particles in induced membrane structures and their subsequent secretion in C6/36
3 cells may be supported by reports that an anterograde protein transport inhibitor,
4 brefeldin A that inhibits transport from the ER, inhibited the release of West Nile virions
5 from Vero but not C6/36 cells (37). Further, the two sizes of subviral particles reported
6 in tick borne encephalitis virus (TBEV)-infected mammalian cells (3) may relate to the
7 existence of at least two factories and their own secretory pathways.

8 JEV prM had the highest cleavability to pr and M among 4 representative
9 flaviviruses, as demonstrated in a study using a full-length cDNA clone of dengue type
10 2 virus in which the 13 sequential amino acids at the pr-M junction was replaced by the
11 corresponding region of JEV, TBEV and yellow fever virus (20). This study also
12 reported that virions containing a JE sequence at the pr-M junction showed the most
13 delayed export outside infected cells (20): one reason for the export retardation would
14 be the premature expression of fusion-competent virions that may fuse intracellular
15 membranes. In the present study, induction of membrane structures was observed upon
16 infection with the Nakayama strain in C6/36 cells, but much less in Vero cells. Since
17 Vero cells are known to be resistant to JEV-induced polykaryocyte formation (17) in
18 contrast to C6/36 cells that are highly sensitive (42), it is probable that the fusion of

1 intracellular membranes mediated by fusion-competent virions induced membrane
2 structures like the SMSs in JEV-infected C6/36 cells.

3 The present system utilizing mosquito and mammalian cells demonstrates a
4 unique approach to the morphogenesis of flavivirus virions. Although we analyzed only
5 virus particles released from infected cells, analyses of intracellular virus particles
6 should provide information about production level of virion and SHA particles and their
7 export efficiency, which may cause differences in extracellular particles. In addition,
8 further studies using gene engineering technologies in the present comparative system
9 will contribute to elucidating functions of each of the viral proteins related to flavivirus
10 virion morphogenesis.

11

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13

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4

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6

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4 T.S., Strauss, J.H., Kuhn, R.J., and Rossmann, M.G. 2003. Structures of immature
5 flavivirus particles. EMBO J. 2: 2604-2613.

1 Table 1. Virion:SHA ratios obtained from infected C6/36 and Vero cells using six strains
2 of JEV.

JEV strain	Virion:SHA ratio ^{a)}	
	C6/36	Vero
Nakayama ^{b)}	0.18	1.98
Beijing P1	0.51	2.66
Beijing P3	0.13	8.00
JATH-160	0.25	7.01
KE-093	6.50	14.79
JaGAr-O1	0.60	3.76

3 ^{a)}Calculated based on E antigen levels contained in each of the virion and SHA fractions
4 obtained from culture fluids of JEV-infected cells by sucrose density gradient
5 centrifugation.

6 ^{b)}Data of the Nakayama strain are in Fig. 2.

- 1 Table 2. Specific infectivities of the virion fraction obtained from C6/36 and Vero cells
 2 infected with one each of the six strains of JEV.

JEV strain	Specific infectivity (x 10 ⁴) ^{a)}	
	C6/36	Vero
Nakayama ^{b)}	13.6	7.9
Beijing P1	7.2	8.8
Beijing P3	5.0	6.2
JaTH-160	10.7	15.3
KE-093	24.3	4.4
JaGAR-O1	16.3	2.1

- 3 ^{a)}Defined as infective titers per ng of E.

- 4 ^{b)}Data of the Nakayama strain are in Fig. 2.

- 1 Table 3. Specific HA activities of the virion and SHA fractions obtained with C6/36 and
 2 Vero cells infected with one each of the six strains of JEV.

JEV strain	Specific HA activity (x 10) ^{a)}			
	C6/36		Vero	
	Virion	SHA	Virion	SHA
Nakayama ^{b)}	1.09	<0.024	5.06	0.16
Beijing P1	0.82	<0.052	2.83	0.23
Beijing P3	0.50	0.033	1.59	0.40
JaTH-160	0.09	0.029	0.07	<0.01
KE-093	1.94	<0.039	0.62	0.29
JaGAr-O1	0.40	0.001	0.90	0.05

3 ^{a)}Defined as the HA titer per ng of E.

4 ^{b)}Data of the Nakayama strain are in Fig. 2.

1 Table 4. prM:M ratios in the virion and SHA fractions obtained with C6/36 and Vero
 2 cells infected with one each of the six strains of JEV.

Cell	JEV strain	Virion			SHA		
		prM level ^{a)}	M level ^{a)}	prM:M ratio	prM level ^{a)}	M level ^{a)}	prM:M ratio
C6/36	Nakayama ^{b)}	4	128	0.031	128	64	2.00
	Beijing P1	<1	16	<0.063	512	216	2.00
	Beijing P3	1	16	0.063	64	128	0.50
	JaTH-160	1	64	0.016	128	256	0.50
	KE-093	4	128	0.031	8	8	1.00
	JaGAR-O1	2	128	0.016	16	32	0.50
Vero	Nakayama ^{b)}	32	256	0.125	8	16	0.50
	Beijing P1	<1	128	<0.008	<1	16	<0.06
	Beijing P3	16	≥1024	<0.016	4	64	0.06
	JaTH-160	<8	512	<0.016	4	64	0.06
	KE-093	32	≥2048	<0.016	8	1024	0.06
	JaGAR-O1	8	128	0.063	8	32	0.25

3 ^{a)}Determined by semiquantitation on Western blots (see Materials and Methods).

1 ^{b)}Data of the Nakayama strain are in Fig. 2.

1 **Figure legend**

2

3 Fig. 1. Sedimentation profiles of viral particles released from Nakayama-infected C6/36
4 and Vero cells. Culture fluids harvested at 24, 48 and 72 hr p.i. were fractionated
5 following sucrose density gradient centrifugation. Each fraction was examined for the E
6 antigen level (closed circle with solid line), infective titer (dotted line) and HA titer
7 (broken line).

8

9 Fig. 2. Analysis of the virion and SHA fractions released from Nakayama-infected
10 C6/36 and Vero cells. (a) The ratio of the E protein amount contained in the virion to
11 SHA fractions obtained from infected C6/36 (open circle) and Vero (closed circle) cells.
12 (b) The specific infectivity of the virion fraction obtained from C6/36 (open circle) and
13 Vero (closed circle) cells. Since the E antigen contained in the virion fraction obtained
14 from C6/36 cells at 24 hr p.i. was undetectable, the specific infectivity was not
15 calculated. (c) The specific HA activity of the virion (triangle) and SHA (square)
16 fractions obtained from C6/36 (open symbol) and Vero (closed symbol) cells. The E
17 antigen contained in the virion fraction obtained from C6/36 cells at 24 hr p.i. was
18 undetectable, and so the specific HA activity was not calculated. (d) The ratio of the

1 prM to M antigen level of the virion (triangle) and SHA (square) fractions obtained
2 from C6/36 (open symbol) and Vero (closed symbol) cells.. The prM:M ratio was not
3 available in all samples at 24 hr p.i. in which prM and M levels were undetectable and
4 the SHA sample obtained from C6/36 cells at 72 hr p.i. in which both of the prM and M
5 levels were ≥ 2048 .

6

7 Fig. 3. Semiquantification of prM and M levels based on Western blotting analysis. Four
8 examples of the results are shown, which were obtained with virion and SHA fractions
9 from Nakayama-infected C6/36 and Vero cells at 48 hr p.i. The levels of prM and M
10 were determined by the highest dilution showing a visible band. Thus, the prM and M
11 levels were 4 and 128 in the virion sample from C6/36 cells (C6/36 virion), 128 and 64
12 in the SHA sample from C6/36 cells (C6/36 SHA), 32 and 256 in the virion sample
13 from Vero cells (Vero virion), and 8 and 16 in the SHA sample from Vero cells (Vero
14 SHA), respectively.

15

16 Fig. 4. Electron micrographs of Nakayama-infected C6/36 (a-d) and Vero (e-h) cells at
17 48 hr p.i. (a, e) Observations at a low magnification. Due to a larger size of Vero than
18 C6/36 cells, a portion of a Vero cell was shown. (b, f) a cytoplasmic area containing

1 SMSs and the dilated ER. (c, g) Observations of the perinuclear region. (d, h)
2 Observations of the peripheral region. Magnification, x 4,000; scale bar = 2 μ m (a, e):
3 magnification, x 50,000; scale bar = 100 nm (b-d, f-h). Arrows indicate SMSs, and
4 arrow heads indicate virions in the ER (d, g) and in SMSs (b, c). Abbreviation: N,
5 nucleus; PM, plasma membrane.

6

7 Fig. 5. Sedimentation profiles of viral particles released from infected C6/36 and Vero
8 cells using six strains of JEV. Culture fluids harvested at 48 hr p.i. from C6/36 and Vero
9 cells infected with the JEV Nakayama, Beijing P1, Beijing P3, JaTH-160, KE-093 and
10 JaGAR-O1 strains were fractionated following sucrose density gradient centrifugation.
11 Each fraction was examined for the E antigen level (closed circle with solid line),
12 infective titer (dotted line) and HA titer (broken line). Data of the Nakayama strain are
13 in Fig. 2.

14

15 Fig. 6. Sedimentation profiles of viral particles released from KE-093-infected C6/36
16 cells under both the open and closed cultivating conditions. Culture fluids harvested at
17 48 hr p.i. were fractionated following sucrose density gradient centrifugation. Each
18 fraction was examined for the E antigen level (closed circle with solid line), infective

1 titer (dotted line) and HA titer (broken line).

2

3 Fig. 7. Electron micrographs of KE-093-infected C6/36 cells of the perinuclear (a) and

4 peripheral (b) regions under the closed cultivating condition, and the perinuclear (c) and

5 peripheral (d) regions under the open cultivating conditions at 48 hr p.i. Arrows indicate

6 SMSs. Open and closed arrow heads indicate virions in the ER and SMSs, respectively.

7 Magnification, x 50,000: scale bar = 100 nm. Abbreviation: N, nucleus; PM, plasma

8 membrane.

1 **Abbreviations**

2

3 C, capsid; E, envelope; ER, endoplasmic reticulum; FBS, fetal bovine serum;
4 HA, hemagglutination; HAU, hemagglutinating unit; JEV, Japanese encephalitis virus; M,
5 membrane; MEM, minimal essential medium; m.o.i., multiplicity of infection; N,
6 nucleus; NS, nonstructural; PBS, phosphate-buffered saline; PFU, plaque-forming unit;
7 p.i., postinfection; PM, plasma membrane; prM, precursor of M; SHA,
8 slowly-sedimenting hemagglutinin; SMB, suckling mouse brain; SMS, smooth
9 membrane structure; TBEV, tick-borne encephalitis virus; VP, vesicle packet.

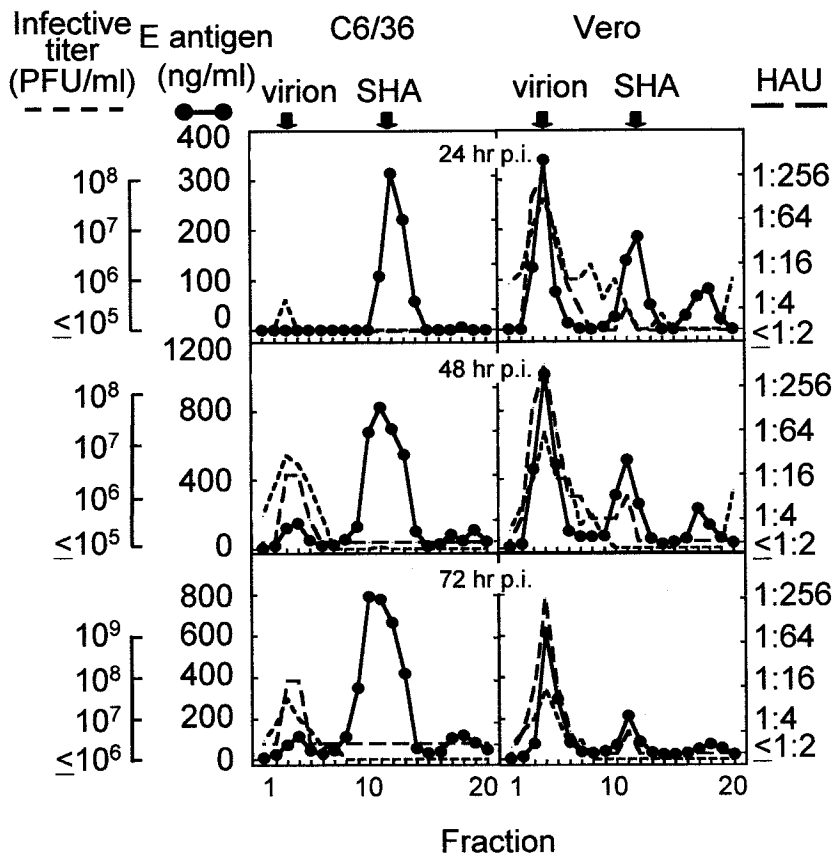


Fig. 1

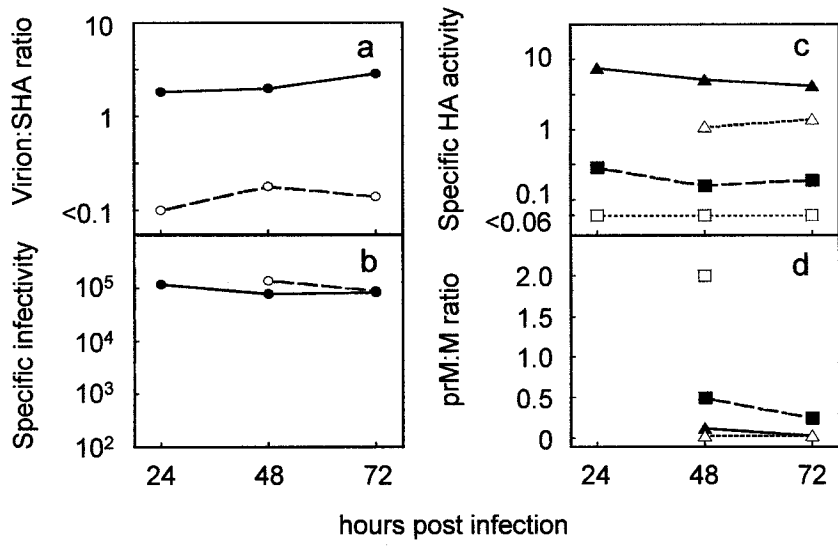


Fig. 2

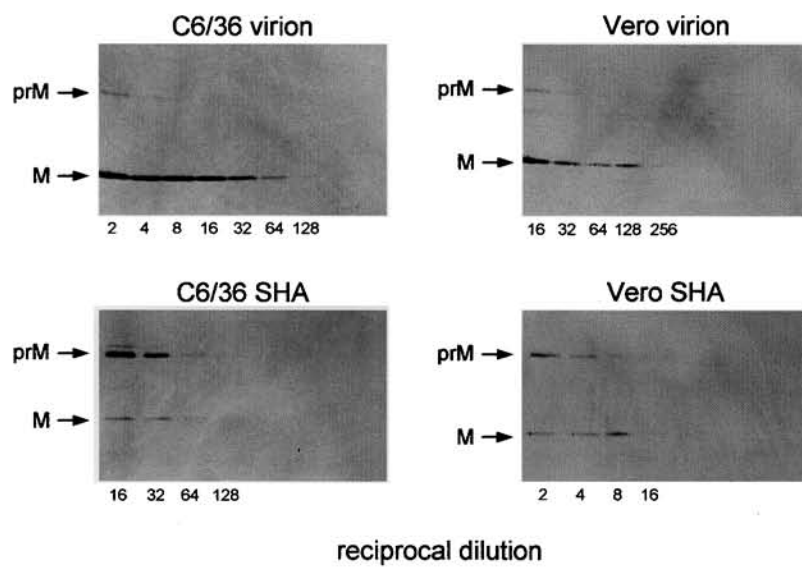


Fig. 3

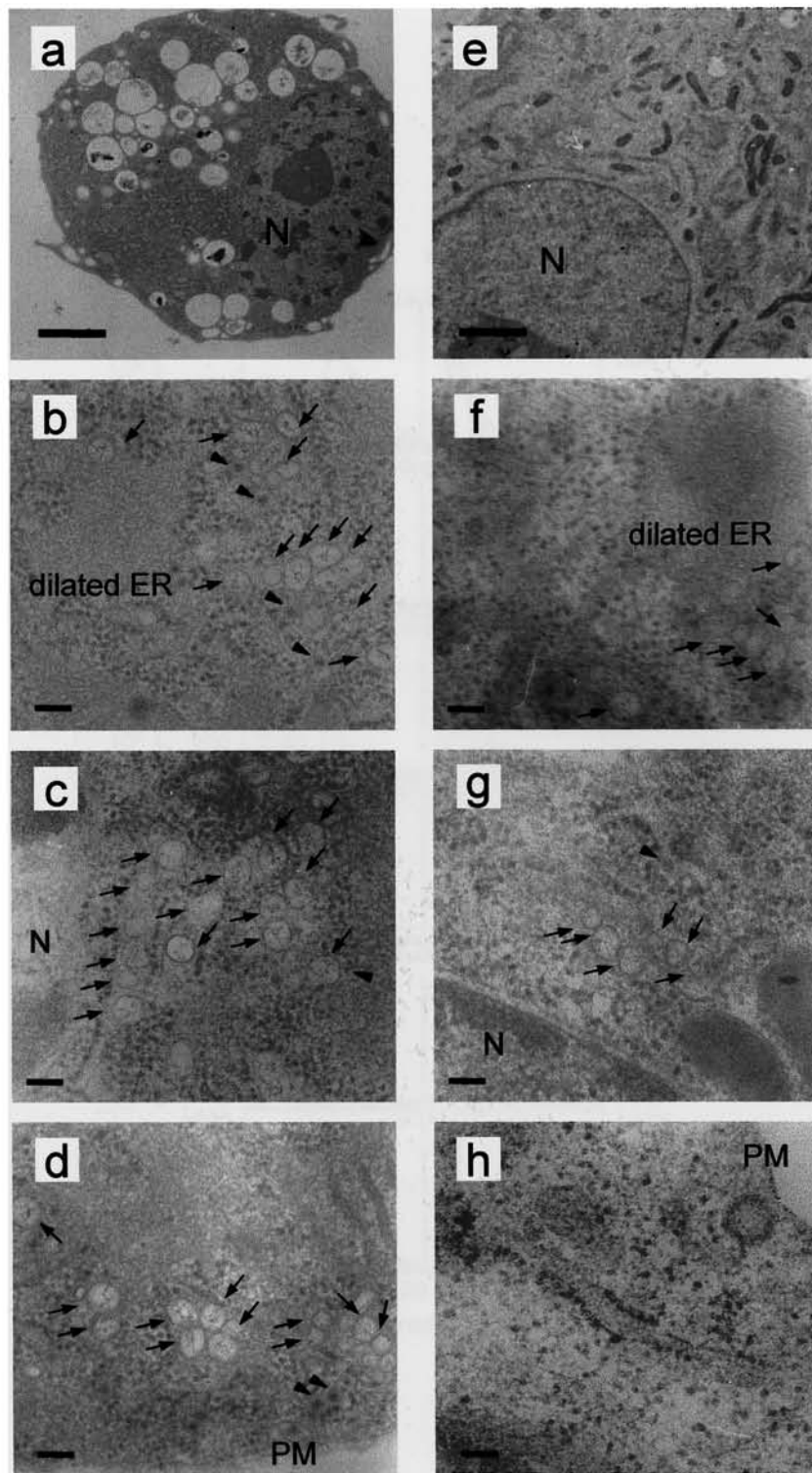


Fig. 4

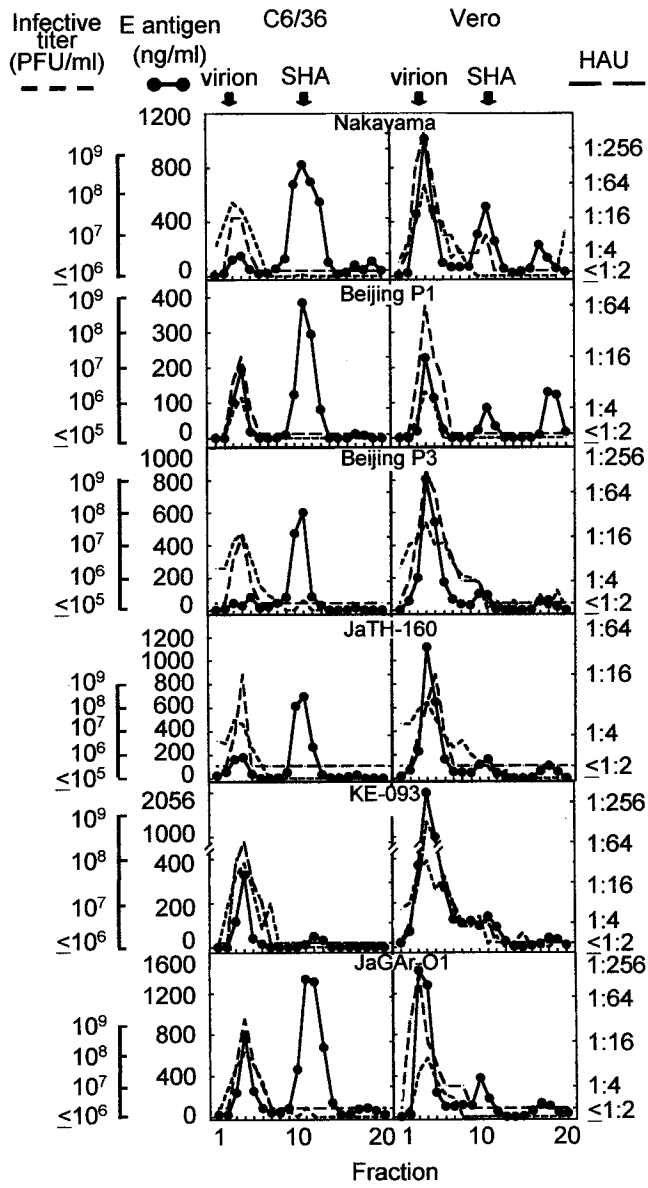


Fig. 5

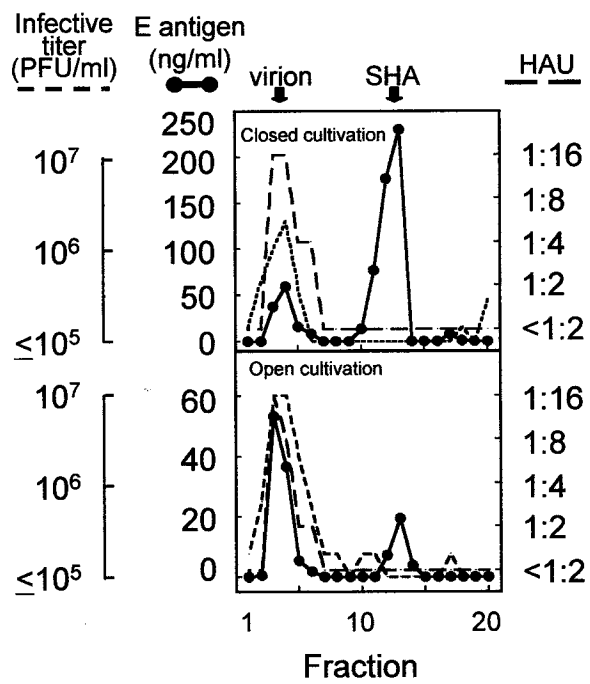


Fig. 6

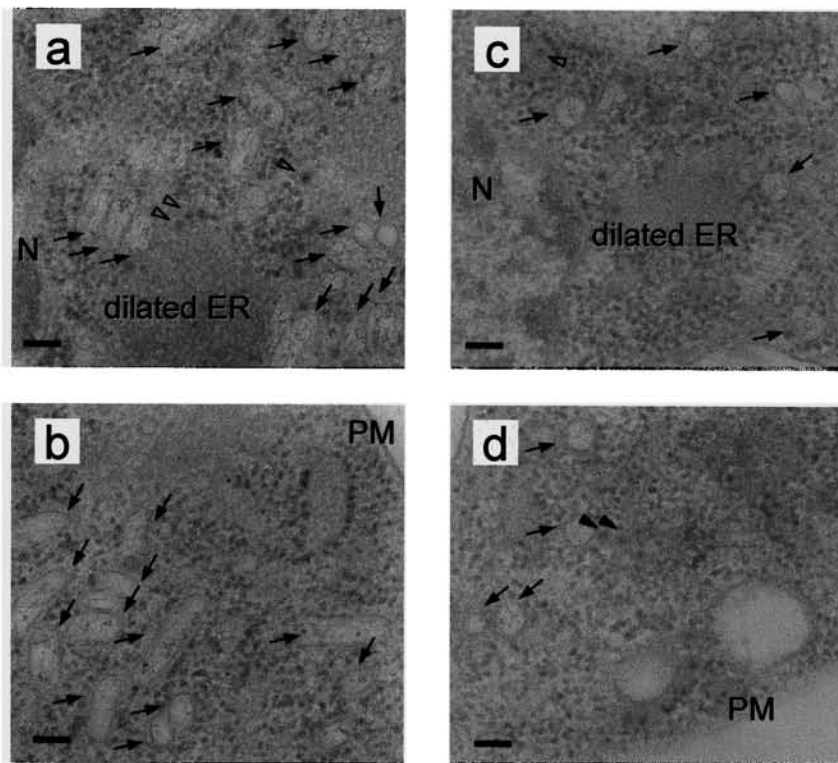


Fig. 7