Dynamics of foot-and-mouth disease virus replication in cells at different phases of the cell-division cycle

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Abstract Foot-and-mouth-disease virus (FMDV) replicates in epithelial cells. The restriction of FMDV RNA to the basal cell layer of epithelia suggests a possible link between FMDV replication in vivo and the cell status. This paper describes in vitro studies in which FMDV infection was investigated in cells that were held at various cell division phases using cell cycle inhibitors. The results suggest that when cells were arrested at the G1 or G1/S phase, high levels of viral RNA were detected by quantitative real-time reverse transcription PCR and viral protein synthesis was observed by specific labeling techniques. In contrast, when cells were arrested at the G2/M phase, reduced or no viral RNA synthesis was detected.

Keywords foot-and-mouth disease virus, cell cycle, replication

1 Introduction

Foot-and-mouth disease virus (FMDV), a member of the family Picornaviridae, genus Aphthovirus, is a small, non-enveloped RNA viruses containing a positive single strand RNA genome, which is the causative agent of foot-and-mouth disease (FMD) characterized by the appearance of vesicles on the feet and in and around the mouth.

Studies on the cellular site of FMDV replication have shown that FMDV RNAs were predominantly localized in the basal cells of both mouth and foot epithelium during disease with visible cytopathology [unpublished data]. During persistence, the viral RNA was localized in the basal cell layer of epithelia from the dorsal soft palate of animals not showing cytopathology [1]. Within a normal epithelium, the stratum basale cells are the cells involved in DNA synthesis and cell division. Some of the daughter cells of the stratum basale cells leave this cell cycle and differentiate into the spinosum, granulosum and cornear cells. Previous studies on the reproduction of picornaviruses indicated the relationship of the kinetics of viral RNA synthesis in coordinated cells to the stage of the cell-division cycle [2–5]. For example, a link between the cell-division cycle and virus replication is seen with coxsackievirus B infection, where low-level of the virus reproduction occurred in quiescent cells but rapid viral protein expression was seen at the wound margin of the monolayer cells, resulting in the production of infectious virus. It appears that viral RNA synthesis is strongly related to the stages of the cell-division cycle, particularly at the later stages [2].

The cell-division cycle consists of four different phases: G0 phase is a resting phase where the cell has left the cycle and has stopped dividing; G1 phase where cytoplasmic materials are accumulated; S and G2 phases in where DNA replication occurs; and the M phase, where cell duplication occurs. It is unclear if FMDV synthesis in vivo is related to the cell-division cycle, but the restriction of FMDV to the certain cells of epithelia [6] suggests a possible link between FMDV replication in vivo and the status of cells at infection. To address this question, an in vitro epithelial cell model was developed to investigate if there was an association between the rate of virus replication and the type/status of infected epithelial cells. The results demonstrated that when cells were arrested at the G1 or G1/S phase, high levels of viral RNA were detected by quantitative real-time reverse transcription PCR and extensive viral protein synthesis was observed by specific labeling techniques. When cells were arrested at the G2/M phase, however, reduced or no viral RNA synthesis was detected or viral protein synthesis observed.

2 Materials and methods

2.1 Cells and virus

Primary cultures of epithelial cells from tissues of ovine
dorsal soft palate, tongue and kidney as well as bovine thyroid (BTY) were prepared according to standard procedures. Briefly, tissues were collected post mortem and transported in ice-cold Dulbecco’s modified Eagle’s medium (DMEM). After washing with phosphate buffered saline (PBS), the epithelial cell layer was separated from other contaminants, cut into 3–4 mm pieces and washed in 5 mmol·L⁻¹ DTT in ice-cold PBS. Tissues were then incubated overnight in protease (0.4 mg·mL⁻¹ in PBS) at 4°C. Protease was inactivated by fetal calf serum. Cells were pelleted and washed several times in DMEM before being filtered through a 100 μm·L⁻¹ strainer followed by a 40 μm·L⁻¹ filter. Cells were pelleted and re-suspended in DMEM/F12, supplemented with 5% fetal calf serum (FCS), and seeded at a concentration of 10⁵ cells per milliliter. Cultures were incubated at 37°C, under 5% CO₂. Cell line ZZ-R127 (supplied by the Collection of Cell Lines in Veterinary Medicine of the Friedrich-Loeffler-Institute), was grown in DMEM and Ham’s F12 (DMEM/ F12) supplemented 1:1 with 5% FCS, at a concentration of 10⁵ cells per milliliter. Cells were incubated at 37°C, under 5% CO₂.

2.2 Virus isolation

The infectivity of collected samples was assayed by inoculation of BTY cells [7].

2.3 Quantitative real-time reverse transcription PCR (qRT-PCR)

An automated extraction procedure was used for nucleic acid extraction of cell supernatants [8]. Quantitative PCR (qRT-PCR) were performed as previously described on a StratageneMxPro 3005P QPCR System [9]. Results were analyzed by means of MxPro QPCR software and genome copy numbers per reaction were also calculated according to Quan et al. [10]. Statistical analyses were carried out by using a non-parametric test (independent samples t-test using Excel). P < 0.05 was considered statistically significant.

2.4 Cell wounding

Monolayers of primary cells from ovine dorsal soft palate, tongue and kidney as well as BTY and ZZ-R127 monolayers were seeded onto glass coverslips and grown to 90% confluence. Monolayers were then wounded with a fine needle 2 h before inoculation with FMDV type OUKG34/2001 at multiplicity of infection of approximately 0.1 M.O.I. After incubation for 1 h at 37°C and 5% CO₂, cells were washed three times with PBS and returned to an incubator. Cell supernatants were collected and glass coverslips were fixed at 0, 4, 8 and 24 h post inoculation (hpi). Cell supernatants were stored at −70°C for qPCR analyses and coverslips were fixed with 4% paraformalde-
ture, on a shaker and blocked for 30 min with 0.5% bovine serum albumin (BSA) in PBS. Cells were washed three times with PBS and then incubated with pre-diluted primary antibodies for 1 h at room temperature, on a shaker. Cells were washed three times with PBS and then incubated with secondary, isotype specific antibodies conjugated to Alexa dyes (Molecular Probes) for 1 h at room temperature, on a shaker. Cells were then washed with deionised water and mounted onto glass slides using Vectashield Mounting Medium (Vector Laboratories). Slides were observed with a Leica TCS/SP2 confocal microscope.

All primary antibodies were diluted in 0.5% BSA in PBS. IB11, mouse monoclonal, raised against 146S FMDV type O1 antigen, specific for conformational, non-neutralising epitopes of the FMDV capsid, diluted at 1:10, kindly provided by Nicholas Jullef, Immunology Department, Pirbright Institute. Ki67 (mouse anti-human IgG1, clone M1-B1, Dako) diluted at 1:100. Zo-1 (anti-mouse IgG1, Invitrogen) diluted 1:50. αβ6 (mouse anti human integrin, IgG2a, Chemicon) and β1 (mouse anti-human IgG1, Chemicon) both diluted at 1:100. Cytokeratin (pan anti-mouse IgG1, Sigma) diluted at 1:100. Secondary antibodies were diluted in 0.5% BSA (in PBS), at 1:200. Goat anti-mouse IgG2a or IgG2b 488 and goat anti-mouse IgG1 568 were obtained from Alexa Fluor (Molecular Probes).

3 Results

3.1 No correlation of the quiescent status of the cells (G0 phase) with FMDV RNA synthesis

The quiescent ZZ-R 127 cells were generated by starvation of FCS (Quiesc) 24 h before inoculation with FMDV. Some cells (Quiesc/continual G0 cells) remained at G0 phase by having no addition of FCS to their medium throughout the experiment, whereas others (Quiesc/cycling cells) were allowed to cycle by the addition of FCS after 24 h starvation and virus inoculation. The virus replication in the quiescent cells at 0, 6 and 24 hpi was assessed by qRT-PCR.

As shown in Fig. 1, FMDV replication occurred in the quiescent cells. At 6 and 24 hpi, there was no difference in the amount of viral genomes in cells between cells that were starved prior to inoculation (Quiesc/continual G0 cells; 5.4 and 9.1 log10 copies per milliliter at 6 and 24 hpi, respectively) and control cells that were not starved (No Quiesc/cycling cells; 6.4 and 9.2 log10 copies per milliliter at 6 and 24 hpi, respectively). Further study showed that release of the cells (starved cells before virus inoculation) from G0 phase by the addition of serum in the media after inoculation with FMDV had no impact on FMDV RNA synthesis, even though a higher level of viral RNA was detected in these Quiesc/cycling cells [6.7 log10 (copies per milliliter)] at 6 hpi compared to those at Quiesc/Continual G0 [5.4 log10 (copies per milliliter) (P = 0.11, P > 0.005)]. Similarly, the viral RNA level in Quiesc/continual G0 cells was higher than in cells that were starved after inoculation [No Quiesc/continual G0; 6.1 and 9.8 log10 (copies per milliliter) at 6 and 24 hpi, respectively]. Qualitative immunofluorescence supported the data obtained with qRT-PCR (data not shown). These results indicated that FMDV RNA synthesis occurs in quiescent cells (G0 phase).

3.2 Viral replication does not preferentially occur in proliferating cells

It has been suggested that proliferating cells near the wounded area preferentially support Coxsackievirus protein production [11]. Therefore wounding the cell monolayer to induce proliferating cells was carried out to investigate if FMDV replication preferentially occur in proliferating cells. The level of viral RNA synthesis was analyzed by qRT-PCR. As shown in Fig. 2, the difference in number of viral genome copies between injured and uninjured monolayers was negligible for the majority of samples analyzed (Fig. 2). Confocal microscopy also showed that there was no trend in relation to the labeling of viral protein when comparing injured with uninjured
monolayers (data not shown). The pattern of labeling observed was not restricted to and it was no more intense at or near the injury site as has been reported for other viruses [11,12]. These observations agree with findings of other FMDV workers [13], indicating proliferating cells do not preferentially support viral replication.

3.3 The stages of the cell-division cycle influence viral propagation

To investigate the impact of cell cycle status on viral RNA synthesis, cell cycle inhibitors were used to arrest cells at different stages and then infected with FMDV. As shown in Fig. 3, when the ZZ-R127 cells were blocked at G1 phase with inhibitors mimosine or rapamycin with or without 5% FCS in medium, the level of viral RNA synthesis at 12 hpi was not significantly different from that in the control cells. Similarly, when the cells were arrested at G1/S phase using hydroxyurea as inhibitor, there was no significant difference in the level of viral RNA synthesis at 12 hpi from those control cells. However, when the cells were blocked at G2/M phase with the inhibitor taxol with or without FCS (data not shown) in the medium, no viral RNA was detected, which was significantly different from...
those control cells. Similarly, the level of viral RNA synthesis at 12 hpi [4 log (copies per milliliter)] in the cells treated with another inhibitor Nocodazol with CFS in medium was considerably lower than those control cells [7.4 log10 (copies per milliliter); \( P = 0.018, P < 0.05 \)]. To test if taxol has a direct inhibitory effect on viral RNA synthesis, an experiment where cells were inoculated with FMDV and exposed to taxol for a period up to 72 h was conducted. The results showed virus genome was detected and viral protein was observed at 18 hpi and the difference between samples treated with and without the taxol [9.7 and 9.8 log10 (copies per milliliter), respectively] was negligible (Fig. 4). Further study showed that taxol was unable to inactive FMDV (data not shown). The result indicated that there is an association of FMDV RNA synthesis with the stages of the cell-division cycle.

Ki67 (a nuclear protein) is expressed in proliferating cells during late G1, S, G2 and M phases but not in resting cells in the G0 phase of the cell-division cycle. To perform an assessment of the influence of these cell cycle inhibitors on cell proliferating function, the ZZ-R127 cells treated with the inhibitors was stained with Ki67 antibody. Quantification of the proliferation rate can then be determined by the Ki67 index which is the number of

**Fig. 3** Virus replication during cell cycle arrest. \( \log_{10} \) (FMDV genome copy numbers per milliliter) detected from different cells arrested at \( G_1, G_{1/S} \) or \( G_{2/M} \) phase of the cell cycle after a 12 h exposure to FMDV. Cells were cultured in medium containing 5% (Ser+) of fetal calf serum. Inhibitors used were mimosine (Mim), rapamycin (Rap), hydroxyurea (HyU), aphidicolin (Aph), nocodazole (Noc), taxol (Tax) and blank control (Ctrl). Error bars represent standard deviation of average of QPCR results.

**Fig. 4** Comparison of virus kinetics between cultures of medium containing taxol and cultures without this inhibitor. Taxol (Tax) was added to monolayers 24 h before virus inoculation; control monolayers (Ctrl) were maintained inhibitor-free. Samples were collected at 18, 24, 48 and 72 h post inoculation. Taxol half-life was assessed by replacing medium at 48 and 72 h post-inoculation (48Xge and 72 Xge, respectively) and comparing with those samples where medium was not replaced.
Ki67 positive cells among the total number of resting cells (Fig. 5). Confocal microscopy studies (Fig. 6) showed that the highest number of Ki67 labeled nuclei was observed in FMDV inoculated monolayers which were arrested with nocodazole (plus FCS) or hydroxyurea (without FCS). However, there was no relationship between FMDV RNA synthesis (Fig. 3) and the number of proliferating cells (Figs. 5 and 6) which further indicates that viral replication do not preferentially occur in proliferating cells.

4 Discussion

Replication of FMDV in cells in vitro in quiescent cells, five different types of proliferating cells and cells arrested at different phases of the cell division cycle was investigated in this study. The results suggest that when cells are arrested at the G2/M phase, reduced or no viral RNA synthesis was detected, indicating a possible link between FMDV replication and the cell-division cycle status.

Our studies have demonstrated that, when cells were first synchronized at G2/M phase with Nocodazole and then inoculated with FMDV, the amount of viral genome detected decreased greatly or was not detected at all (taxol). Cells synchronized at either G1/S or G1 phase showed a high number of viral genome copies, similar to those observed in control samples, i.e., where cells were not exposed to inhibitors. This result corroborates previous findings which suggest that certain viral and cellular IRES elements are active during the G2/M phase of the cell cycle. The protein synthesis of positive-strand RNA viruses is under the control of a defined IRES. There might be an advantage for a virus to arrest cells in the G2/M stage of the cell-division cycle because at these stages IRES-dependent translation of some (but not all) cellular and certain viral mRNAs appears to be optimal in comparison to other stages of the cell-division cycle [14]. Furthermore, the eIF4E subunit of eIF4F is dephosphorylated at the G2/M stage, and this is correlated with diminished cap-dependent translation [9]. On the other hand, cells forced into G0, by either serum starvation or wounding of monolayers, did not show an increase in copy numbers of viral genome, as observed with other Picornaviruses [12] in which viral replication does not preferentially occur in proliferating cells.

Taxol gave rise to the most intriguing data of the two G2/M inhibitors chosen for the experiments. Taxol is a plant alkaloid commonly used to perturb microtubule dynamics [15]. As a part of the cell’s apparatus, microtubules are essential for cell division and replication, hence inhibition of these structures will lead to cell death. It has been demonstrated that taxol inhibits HeLa cell proliferation by inducing a sustained mitotic block at the metaphase/anaphase boundary [16]. However, our results suggest that taxol did not inhibit cell proliferation of both normal or FMDV inoculated ZZ-R127 cells. We have also demonstrated that taxol did not inactivate FMDV. However, there seemed to be a delay on the development of CPE of taxol diluted samples which remains to be explained. One hypothesis is that a taxol induced modification of UTRs, located at each end of the viral RNA, could impact on the control of gene expression and viral replication and, consequently, on inefficient viral replication. Nevertheless, further studies are required to understand the effects of taxol on viral-specific RNA
structures which work in close association with host and viral proteins.

5 Conclusions

Reduced or no viral RNA synthesis was detected in cells arrested at the G2/M phase, indicating there is a possible link between FMDV replication and the cell-division cycle status.

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References