Sites of feline coronavirus persistence in healthy cats

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Introduction

Feline infectious peritonitis (FIP) is a fatal disease of cats, caused by feline coronavirus (FCoV). FIP is currently the leading infectious cause of death in cats (Pedersen, 2009). Despite the generally high prevalence of FCoV infection among the cat population (up to 90% seropositive animals depending on environment and geographical area), FIP morbidity is low and rarely surpasses 5% of infected cats, mostly affecting young animals (Pedersen, 2009). This is probably due to the fact that FIP only develops with the occurrence of virulent FCoV mutants, possibly either generated within the individual infected host or acquired externally (Poland et al., 1996; Vennema et al., 1998; Brown et al., 2009; Pedersen et al., 2009; Chang et al., 2010). Two FCoV serotypes, I and II, can be distinguished; these show different geographical prevalence, but, so far, no evident differences in their pathogenic potential (Kummrow et al., 2005; Lin et al., 2009). FCoV is transmitted via the faecal–oral route and primarily infects enterocytes (Pedersen, 1995). In the literature, two FCoV biotypes have been distinguished: feline enteric coronaviruses (FECVs), which are endemic in cat populations and generally not associated with clinical disease, and FIP viruses (FIPVs), which might arise from endemic FECVs either as an in vivo mutation or as virulent strains and are responsible for the development of FIP (Vennema, 1999; Brown et al., 2009; Pedersen et al., 2009; Chang et al., 2010). Regardless of the development of FIP, FCoV spreads from its initial site of infection within the intestine via monocyte-associated viraemia (Gunn-Moore et al., 1998; Kipar et al., 1999, 2005; Meli et al., 2004) and can replicate within monocytes in healthy cats (Can-Sahna et al., 2007). The pathogenesis of FIP is not fully understood, but it has been shown that a monocyte-triggered vasculitis, in association with systemic monocyte and endothelial cell activation, represents the crucial event (Kipar et al., 2005), probably in combination with some antibody-mediated enhancement and complement activation (Dewerchin et al., 2006). Cats can be FCoV carriers and generally remain healthy, despite systemic infection (Herrewegh et al., 1995; Addie et al., 1996; Gunn-Moore et al., 1998; Kipar et al., 1999; Meli et al., 2004). Previous studies indicate that FCoV persists within the intestine and is shed persistently or intermittently with the faeces (Foley et al., 1997; Herrewegh et al., 1997; Harpold et al., 1999; Meli et al., 2004). Therefore, persistently infected, healthy carriers are believed to play a key role in the epidemiology of FIP (Foley et al., 1997; Meli et al., 2004).
The aim of the present study was to identify the sites of FCoV persistence in healthy carriers. As FCoV can spread systemically within monocytes, we surmised that the virus could persist in both intestinal and extra-intestinal sites. To investigate this hypothesis, specific-pathogen-free (SPF) cats were infected experimentally via the oral route with high doses of infectious, non-pathogenic FCoV serotype I field isolates. Cats were examined at different time points between 14 and 80 days post-infection (p.i.) for viraemia, viral shedding and viral loads, as well as viral antigen, in selected organs.

**RESULTS**

**Oral infection with non-virulent FCoV serotype I field strains remains clinically inapparent but leads to viraemia**

FCoV serotype I is known to be the most common serotype in field cases (Hohdatsu et al., 1992; Addie et al., 2003; Benetka et al., 2004; Kummrow et al., 2005; Lin et al., 2009). It differs from serotype II FCoV in that it hardly grows in tissue culture (Jacobse-Geels & Horzinek, 1983); however, oral administration of faeces or gut homogenates from shedding cats leads to intestinal infection and monocyte-associated viraemia (Meli et al., 2004).

For the present study, 30 SPF cats were infected orally with different doses of previously described non-virulent FCoV serotype I field strains (FECV biotype) prepared from faeces or gut homogenates of infected, clinically healthy serotype I field strains (FECV biotype) prepared from different doses of previously described non-virulent FCoV serotype I field isolates. Cats were examined at different time points between 14 and 80 days post-infection (p.i.) for viraemia, viral shedding and viral loads, as well as viral antigen, in selected organs.

<table>
<thead>
<tr>
<th>Cat group and no.</th>
<th>Virus strain and origin</th>
<th>Infectious dose (RNA copy numbers)</th>
<th>Time of euthanasia (day p.i.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.1, 1.2</td>
<td>FCoVZu1 (FE)*</td>
<td>8.0 x 10^7</td>
<td>14</td>
</tr>
<tr>
<td>2.1–2.3</td>
<td>FCoVZu1 (FE)</td>
<td>8.0 x 10^6</td>
<td>17</td>
</tr>
<tr>
<td>3.1, 3.2</td>
<td>FCoVZu2+Zu5 (FE*)</td>
<td>1.0 x 10^6</td>
<td>28</td>
</tr>
<tr>
<td>4.1, 4.2</td>
<td>FCoVZu3 (FE*)</td>
<td>3.0 x 10^5</td>
<td>28</td>
</tr>
<tr>
<td>5.1–5.15</td>
<td>FCoVZu1 (GH)</td>
<td>1.5 x 10^6</td>
<td>48</td>
</tr>
<tr>
<td>6.1, 6.2</td>
<td>FCoVZu1 (GH)</td>
<td>3.0 x 10^6</td>
<td>80</td>
</tr>
<tr>
<td>7.1, 7.2</td>
<td>FCoVZu1 (GH)</td>
<td>3.0 x 10^8</td>
<td>80</td>
</tr>
<tr>
<td>8.1, 8.2</td>
<td>FCoVZu1 (GH)</td>
<td>3.0 x 10^7</td>
<td>80</td>
</tr>
</tbody>
</table>

Viraemia was detected at 1 week p.i., and most cats (20/30; 67%) were positive at one to several, often not consecutive, time points, with a peak at day 7 p.i. (14/30; 47%) and decline thereafter (day 14 p.i.: 12/30 (40%), day 17 p.i.: 4/28 (14%), day 28 p.i.: 1/25 (4%)), day 35 p.i.: 1/21 (5%), day 42 p.i.: 0/21, day 48 p.i.: 3/21 (14%); days 58, 65 and 80 p.i.: 0/6). No cat proved to be viraemic over the entire test period, but three showed recurrent viraemia. Four cats were viraemic at the time of death (day 17 p.i. (n=1), day 48 p.i. (n=3), 13.3%). There were no differences in the course and level of viraemia depending on virus isolate or infectious doses. These results showed that the virus isolates generally led to viraemia within a week of infection, and that viraemia could recur at later time points.

Cats were necropsied and examined grossly and histologically to identify any potential pathological effects of the infection. There were no changes indicative of FIP (Kipar et al., 2005). Histological findings were restricted to moderate lymphatic hyperplasia in spleen and mesenteric lymph nodes, some degree of lymph node sinus histiocytosis, no or minimal thymic involution, and moderate to high bone marrow activity. This was consistent with previous findings in clinically healthy, FCoV-infected cats and confirmed that FCoV infection induces an intense systemic immune response (Kipar et al., 1999, 2001; Meli et al., 2004).

**Spread of FCoV to all organs after oral infection and persistence in the absence of detectable viraemia**

Having confirmed that oral infection did generally induce viraemia, we wanted to assess (i) whether all intestinal compartments become infected and where in the gut the virus persisted, and (ii) whether the virus established itself in other organs. For this purpose, we performed FCoV real-time RT-PCR on a wide range of tissues (see below) collected immediately after death, a time when only four cats (13.3%) were confirmed to be viraemic. FCoV RNA was detected in the organs of all animals. All organs were positive, regardless of detectable viraemia, in at least some cats (183 positive samples; 42.4%), with the following frequencies: colon 28 (15.3%), liver 21 (11.5%), mesenteric lymph nodes 19 (10.4%), ileum 14 (7.7%), thymus 13 (7.1%), jejunum 12 (6.6%), kidney 12 (6.6%), tonsil 11 (6.0%), lung 11 (6.0%), spleen 10 (5.5%), duodenum 9 (4.9%), bone marrow 8 (4.9%), brain 6 (3.8%), skin 5 (2.7%) and skeletal muscle 2 (1.1%) (Fig. 1). Overall, the colon tested positive significantly more frequently than any other tissue. The liver was positive significantly more often than all other organs, except the mesenteric lymph nodes. The frequencies of a test positive organ or tissue outside the alimentary tract and lymphatic tissues were as follows: liver 64 % (28/43), kidney 40 % (10/25), lung 36 % (9/25), brain 20 % (5/25), skin 22 % (4/18) and skeletal muscle 11 % (2/18). These findings confirmed that FCoV can infect the entire intestine, but indicated the colon as the
main site of viral persistence. They also showed that FCoV viraemia generally leads to widespread organ infection and virus persistence in organs and tissues when virus cannot be detected in the blood.

Viral shedding is consistent in the early phase of infection and is seen with infection of several intestinal compartments; the colon is the main site of viral persistence

Having found evidence of viral persistence within the intestine, we wanted to identify any possible association with viral shedding, and tested faecal samples from all cats by FCoV PCR over the time periods between infection and death (Meli et al., 2004). All animals started to shed on day 2 or 3 p.i. and shed relatively consistently up to day 17, followed by intermittent shedding (data not shown). Independent of the level of viral shedding, the cats remained clinically healthy, confirming the FCoV isolates as non-pathogenic FECV biotypes. Also, virus isolates and infectious dose did not have any effect on onset and duration of viral shedding (Meli et al. 2004).

At the time of death, all cats euthanized at days 14, 17 and 28 p.i. did shed virus. On days 14 and 17 p.i., the colon and ileum were always positive, and the duodenum and jejunum each in most (4/5) cats. On day 28 p.i., the duodenum, ileum and colon were always positive, and the jejunum in most (3/4) cats. On day 48 p.i., the cats did not shed virus. The duodenum, jejunum and ileum were only positive in one, two and three of the 15 cats, respectively, but the colon yielded a positive result in 14 cats. On day 80, 50 % (3/6) of cats did shed virus. Whilst the duodenum was always negative, the jejunum was positive in three, the ileum in two and the colon in five of the six cats. One cat tested negative in the entire intestine. Comparison of relative viral loads in each intestinal compartment over the entire study identified the significantly highest loads in the colon on day 14 (Fig. 2b), and in the jejunum and ileum on day 17 (not shown).

We looked at the association between viral shedding and the presence of viral RNA in the different intestinal compartments at the time of death (Table 2). From the 12 cats that exhibited FCoV RNA only in the colon, only the one with the highest relative viral load was shown to shed virus. When the colon plus at least two small intestinal compartments were positive, most cats (10/12) did shed virus. The two non-shedders (days 28 and 48 p.i., respectively) tested positive in all intestinal compartments, but with relatively lower overall intestinal viral loads than the five shedding animals that showed viral RNA in all intestinal compartments.

Overall, relative viral loads in the colon were significantly higher [mean $1.66 \times 10^5$-fold higher (range $11–1.3 \times 10^5$)]
than in any other organ, and were a mean of 122-fold higher (range 11–327) than in the other intestinal compartments. There was also a significant difference between positive and negative faecal shedders in terms of cumulative viral intestinal load, whereby cats that were found to shed virus had, on average, ten times higher cumulative viral loads than cats that did not shed. These results indicated that oral FCoV uptake led to initial infection of all intestinal compartments. The virus was cleared frequently from all compartments but the colon at later stages (Fig. 1), which confirmed the colon as the main site of virus persistence and the source of recurrent shedding. However, virus spread to and colonization of the small intestine appeared to be essential for viral shedding, even when the colon exhibited relatively high viral loads.

**Table 2.** Correlation between viral shedding (faecal swabs, detection of FCoV by real-time RT-PCR) and the detection of FCoV RNA (real-time RT-PCR) in different intestinal compartments of FCoV serotype I-infected, healthy cats on the day of euthanasia (14–80 days p.i.)

<table>
<thead>
<tr>
<th>Intestinal compartments positive for FCoV RNA (n)</th>
<th>Faecal swab</th>
<th>FCoV RNA-positive</th>
<th>FCoV RNA-negative</th>
</tr>
</thead>
<tbody>
<tr>
<td>None (2)</td>
<td>–</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>Colon (12)</td>
<td>1</td>
<td>11</td>
<td></td>
</tr>
<tr>
<td>Colon, jejunum (2)</td>
<td>–</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>Colon, ileum (2)</td>
<td>–</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>Colon, jejunum, ileum (3)</td>
<td>3</td>
<td>–</td>
<td></td>
</tr>
<tr>
<td>Colon, duodenum, ileum (2)</td>
<td>2</td>
<td>–</td>
<td></td>
</tr>
<tr>
<td>Colon, duodenum, jejunum, ileum (7)</td>
<td>5</td>
<td>2</td>
<td></td>
</tr>
</tbody>
</table>

**Fig. 2.** Mean relative viral loads [in relation to glyceraldehyde-3-phosphate dehydrogenase (GAPDH) in organs of healthy cats after experimental oral infection with FCoV type I strains for 14–80 days. Box and whisker plots demonstrating overall mean relative viral loads in the different organs tested at 14–80 days p.i. (a) and in selected organs at days 14, 17, 28, 48 and 80 after infection (b). The y-axis shows the viral load relative to GAPDH. LN, Lymph nodes.

Table 2. Correlation between viral shedding (faecal swabs, detection of FCoV by real-time RT-PCR) and the detection of FCoV RNA (real-time RT-PCR) in different intestinal compartments of FCoV serotype I-infected, healthy cats on the day of euthanasia (14–80 days p.i.)

At different time points p.i., the range of organs that harbour FCoV varies

Having identified FCoV in a wide range of organs in infected cats, we evaluated whether the extent and pattern
of organ infection varied over time (Fig. 1). Overall, colon, liver and mesenteric lymph nodes were positive with the highest frequency.

Cats examined on days 14, 17 and 28 p.i. all exhibited systemic infection, as FCoV RNA was demonstrated in several organs. On days 14 \((n=2)\) and 17 \((n=3)\), 6–11 of the 13 organs tested were positive (mean 70.8 %), with intestines and mesenteric lymph nodes positive in all cats. On day 28 \((n=4)\), 8–12 of the 13 organs were positive (mean 73.1 %).

On day 48 \((n=15)\), the number of positive organs had dropped significantly (mean 26.2 %; 1–10 of 15). In one cat, only the colon was positive at that time point without detectable viraemia; thus, this cat was most likely not systemically infected at this stage, although it had been viraemic on days 7 and 14 p.i. (data not shown).

On day 80 \((n=6)\), the number of organs harbouring virus increased again (mean 42.2 %; 3–10 of 15). The one cat that did not harbour virus in its intestine (see above) was nonetheless systemically infected, as its tonsils, lung and kidney were positive.

These results showed that oral FCoV infection with non-pathogenic strains led to viraemia and subsequent persistent systemic infection in the absence of detectable viraemia. There was no distinct organ infection pattern or a correlation between the number and distribution of positive organs and the presence of viraemia. The higher number of FCoV-positive organs at day 80 p.i. compared with day 40 p.i. indicated that organs can become reinfected with recurrent viraemia. In general, however, at later time points (days 48 and 80 p.i.), the colon, liver, mesenteric lymph nodes and tonsils were the organs positive with the highest frequency (Fig. 1).

**Organs exhibit the highest relative viral loads at earlier time points after infection but can remain persistently infected for a longer period**

The presence of FCoV in organs without detectable viraemia indicated that the virus can infect and persist in cells other than monocytes and enterocytes. As a basis to identify candidate host cells, we identified the organs with the highest relative viral loads by means of FCoV real-time RT-PCR. In general, mean relative viral loads varied considerably (Fig. 2a). However, there were significant differences in overall relative viral loads between the different time points after infection \((14, 17, 28, 48 \text{ and } 80 \text{ days p.i.}; P=0.002)\), and Tukey’s grouping for multiple comparisons identified significant differences in cats euthanized at days 14 and 17 compared with other time points, but not significantly different from each other. Thus, there was a trend for higher viral recoveries at 14 and 17 days p.i. compared with the later time points. For example, viral recoveries at day 14 were on average 41-fold higher \((\text{range } 1.1–152\text{-fold})\) than at other time points. Day 17 viral recoveries were 49-fold higher \((\text{range } 3.9–138\text{-fold})\) than day 28, 48 and 80 recoveries.

There were significant differences between viral recovery and the organ examined \((P<0.001)\). In particular, the colon and jejunum showed significantly higher viral recoveries than all other organs but not than each other. When the jejunum with a particularly high viral load (day 17 p.i.) was excluded, the colon showed the significantly highest loads. An organ group comparison on the lymphatic tissues revealed a significantly higher relative viral load in mesenteric lymph nodes compared with other lymphatic tissues \([e.g. \text{spleen } (P=0.021; 25\text{-fold higher}), \text{tonsil } (P=0.017; 74\text{-fold higher}) \text{ and thymus } (P=0.016; 146\text{-fold higher}])\]. These findings indicated viral spread from the intestine to the regional lymph nodes via the lymphatics, with only limited further spread. No other organ group comparison revealed significant differences. Relative viral loads within the liver, which was the organ with the second highest frequency of test positives, were not particularly high, and on average were lower than, for example, in the lung \((9.3\text{-fold lower})\) (Fig. 2a).

We compared relative viral loads in selected organs (liver, mesenteric lymph nodes and lung) and the colon over time. Relative viral loads in the colon and liver were significantly higher on day 14. For the mesenteric lymph nodes, day 14 displayed significantly higher viral loads than days 48 and 80, but not days 17 and 28 (Fig. 2b). The lung showed no significant differences in viral loads over time. These results revealed a general peak in organ viral loads in the first weeks after infection, ebbing off with time, similar to the intestine.

**Viral persistence is mediated by columnar epithelial cells in the colon and by tissue macrophages in other organs**

FCoV is known to infect intestinal epithelial cells (Pedersen, 1983; Herrewegh et al., 1997; Kipar et al., 1998). In an attempt to identify the cells in which FCoV persisted in the gut, we performed immunohistology for FCoV antigen \((Kipar et al., 1998, 2005)\) on all intestinal compartments (FCoV RNA-positive or -negative) of the 15 cats euthanized at day 48 p.i. without evidence of virus shedding. FCoV antigen was detected in epithelial cells of the colon in two animals. Staining was restricted to several columnar epithelial cells on the mucosal surface in one animal (Fig. 3a) and a single columnar epithelial cell on the mucosal surface of the second. In both cats, the colon had been the only intestinal compartment that tested positive for FCoV RNA, with comparatively high relative viral loads. These results confirmed the colonic columnar epithelial cells as the site of viral persistence in the intestine of non-shedding cats.

Circulating monocytes mediate FCoV viraemia and are responsible for the granulomatous vasculitis and focal granulomatous lesions in organs that are characteristic for FIP and harbour virus in macrophages (Kipar et al., 2005). Having demonstrated viral RNA in organs regardless of detectable viraemia, we hypothesized that the virus could
also infect resident tissue macrophages and we performed immunohistology for FCoV antigen on some tissues that had exhibited comparatively high relative virus loads or were consistently FCoV RNA positive: the mesenteric lymph nodes as they drain the intestines, and the liver and lung, which both contain specific tissue macrophages in constant contact with blood, i.e. hepatic Kupffer cells and pulmonary intravascular macrophages (Brain et al., 1999; Bilzer et al., 2006). FCoV antigen was detected in several sinus macrophages in the mesenteric lymph nodes with the highest relative viral load (day 28 p.i.), in association with intense sinus histiocytosis with several macrophage aggregates (Fig. 3b). In those with the second- and fourth-highest viral loads (days 14 and 17 p.i., respectively), sinus macrophages were found to express viral antigen. All three cats exhibited high viral loads within all intestinal compartments, and virus shedding, which suggests virus uptake by macrophages in the intestine and subsequent transport to the regional lymph nodes. Among the three examined lungs was one (day 28 p.i.) where viral antigen was observed in scattered pulmonary intravascular macrophages (Fig. 3c). FCoV antigen was not demonstrated in the livers, suggesting that the amount of virus per cell was too low to be detected by immunohistology, a relatively insensitive method compared with real-time RT-PCR.

**DISCUSSION**

The present study investigated SPF cats that had been infected orally with high doses of non-virulent FCoV type I field strains for evidence of systemic virus infection, virus persistence and viral shedding in relation to viraemia over a period of up to 80 days after infection.

All animals remained clinically healthy. Initially, viral RNA was found in all intestinal compartments, the blood and several organs of most animals, confirming viraemic spread and systemic infection.

In the intestine, FCoV was detected most consistently and at highest levels in the colon. Virus shedding, however, was generally observed when both colon and small intestine harboured virus, with significantly higher cumulative intestinal viral loads, and in the early phase after infection. It was occasionally also seen at later time points and in cats with recurrent shedding. These findings suggested that the virus persists in the colon, from which it can reinfect the small intestine at any time. The immunohistological demonstration of viral antigen within columnar epithelial cells in the colon of cats that did not shed virus but harboured FCoV RNA in the intestine confirmed the colonic enterocytes as the major site of FCoV persistence in the gut (Herrewegh et al., 1997). The presence of viral antigen in intestinal epithelial cells without evidence of cell loss and faecal shedding suggested that non-virulent FCoV persists by establishing a non-lytic infection in epithelial cells, which is different from FCoV that induces enteritis in cats with enterocyte degeneration (Kipar et al., 1998). With regard to persistence in epithelial cells, FCoV shows similarities to foot-and-mouth disease virus and coxsackievirus B3 (Harrath et al., 2004; Zhang & Alexandersen,
Carrasco virus, in acute infections (Thanawongnuwech respiratory syndrome virus and African horse sickness virus (Bingen infected with human and feline immunodeficiency virus (Jiang, 2004). The question arises as to how the colonic columnar epithelium remains infected, considering that viral antigen appears only to be present in the superficial epithelium, which is sloughed off in the course of normal epithelial turnover, and not within replicating cells. Cell-to-cell spread between epithelial cells may be the mode of transmission to ensure persistence. Furthermore, the mechanism of FCoV entry into monocytes/macrophages within the gut has so far not been identified, nor whether it occurs within specific intestinal compartments. The latter seems unlikely, as resident macrophages are generally numerous throughout the intestine (Platt & Mowat, 2008).

The mesenteric lymph nodes were shown to frequently harbour FCoV RNA and at relatively high levels. Considering that some also exhibited viral antigen within sinus macrophages, it appears likely that virus from enterocytes is taken up by macrophages in the intestinal mucosa and transported to the regional lymph nodes. From there, further spread via the lymphatics and ultimately the blood, as a potential additional mode of virus distribution, is possible.

We also identified FCoV RNA in a large range of organs and tissues, which must be a consequence of systemic viral spread via the blood, i.e. monocyte-associated viraemia, at some point. In viraemic cats, organs could harbour viral RNA in infected monocytes within the vasculature. In addition and in the absence of detectable viraemia, however, infection of parenchymal cells and/or resident macrophages has to be considered. Previous in vitro studies have shown that FCoV can infect specialized macrophages, such as peritoneal macrophages (Stoddart & Scott, 1989). Indeed, we were able to identify viral antigen in scattered pulmonary intravascular macrophages (PIMs) in a non-viraemic cat with a relatively high viral titre in the lung. PIMs, like hepatic Kupffer cells (HKCs) are bone marrow-derived macrophages that form part of the mononuclear phagocyte system. They are resident macrophages anchored to endothelial cells within pulmonary capillaries. Together with HKCs, the resident macrophages of the liver that adhere to endothelial cells in the sinusoids, PIMs have been shown in cats to be the main cells that rapidly phagocytose particles carried in the blood (Brain et al., 1999). PIMs have been identified as target cells for viruses, such as classical swine fever virus, porcine reproductive and respiratory syndrome virus and African horse sickness virus, in acute infections (Thanawongnuwech et al., 1997; Carrasco et al., 1999, 2001), whilst HKCs can become infected with human and feline immunodeficiency virus and African swine fever virus (Bingen et al., 2002; Ciborowski & Gendelman, 2006; Sánchez-Cordon et al., 2008). Although we were not able to demonstrate FCoV antigen within HKCs, HCKs may be responsible for the presence of FCoV RNA in the liver, particularly in non-viraemic cats. This is supported by a previous study that provided ultrastructural evidence of FCoV in HKCs in cats that had developed FIP after intra-peritoneal application of highly virulent FIPV (Pedersen, 1976). Consequently, PIMs and HKCs could represent sources of recurrent viraemia in FCoV-infected cats via release of virus into the blood and/or transmission to monocytes. In acute African swine fever, PIMs have been shown to become activated (Carrasco et al., 2002) and it needs to be determined whether, in FCoV infection, PIMs, due to their location in pulmonary capillaries, can contribute directly to the development of the granulomatous vasculitis that is often seen in the lung in cats with FIP (Kipar et al., 2005).

FCoV was demonstrated in all organs/tissues tested both in viraemic cats and in cats without detectable viraemia. The number of infected organs and overall viral loads were significantly higher in the early phase after infection (days 14 and 17). However, over time, most animals remained systemically infected and FCoV persisted in several organs, although in a more sporadic manner and with evidence of reinfection in the course of intermittent viraemia. The latter is a well-known feature of FCoV infection (Foley et al., 1997; Herrewegh et al., 1997; Harpold et al., 1999; Meli et al., 2004) and is also suggested by our real-time RT-PCR results on the blood. Alternatively, infected cats might exhibit highly variable blood viral loads over time, which drop below detection levels in several cases. In the present study, besides the mesenteric lymph nodes, the liver was shown to harbour the virus most frequently and at comparatively high levels. This finding is not surprising considering that the liver directly drains the blood from the intestines via the portal vein. Also, HKCs, should they indeed become infected with FCoV, are very numerous in the liver and represent 80–90% of tissue macrophages in the body (Bilzer et al., 2006), which renders them a potential source of high viral burdens in the liver. The presence of FCoV RNA in tissues without specialized vessel-associated resident macrophages, such as the skeletal muscle, brain or skin, in non-viraemic cats cannot readily be explained. However, circulating monocytes have been shown to leave the blood stream and differentiate to a large proportion into long-lived tissue macrophages (Randolph et al., 1999). It might therefore be possible that FCoV-infected monocytes migrate into any tissue where they then differentiate into (persistently infected) resident macrophages.

In conclusion, our data identify the colon as the major site of FCoV persistence and provide convincing evidence that the differentiated columnar epithelial cell is the target cell of persistence. However, our results also identify other organs/tissues as sites of persistent infection and potential sources for recurrent viraemia. Organs with specialized resident macrophages that either directly drain the gut, such as the mesenteric lymph nodes, or serve to filter the blood, such as liver (HKCs) and lung (PIMs), are the main candidates for this. These findings imply that clearance of the virus from the gut does not necessarily protect FCoV-infected cats from recurrent viraemia and, ultimately, the development of FIP at some stage.
**METHODS**

**Virus preparation.** For experimental infections, FCoV serotype I field strains isolated from faecal samples of naturally infected cats (FCoVZu1, GenBank accession no. DQ256137; FCoVZu2, DQ256138; FCoVZu3, DQ256139; FCoVZu5, DQ256140) or from the intestines (gut homogenates) of cats experimentally infected with FCoVZu1 (DQ256137) were used. All isolates have been shown to have similar effects with regard to infectivity, induction and rate of viral shedding, development of viraemia and pathological effects (Meli et al., 2004).

Briefly, faecal samples were prepared by dilution in RPMI 1640 containing L-glutamine and 10% fetal calf serum, incubation for 10 min at 4°C with occasional shaking, and centrifugation (2 x 10 min at 900 g). Gut homogenates were prepared from snap-frozen intestinal segments of kittens experimentally infected with the FCoVZu1 strain, using an Ultra Turrax homogenizer (Polytron PT; Kinematica AG) at 20,000 r.p.m. for 15 min at 5–6°C intervals. Faecal supernatants and intestinal homogenates were analysed to determine the FCoV load (RNA copy number; see below; Meli et al., 2004).

**Animals.** In total, 30 SPF kittens were provided by IFFA-Credo (Saint-Germain sur l’Arbresle, France) or Liberty Research Laboratories (Waverly, NY, USA) at either 6 or 16 weeks of age. The animals were first acclimatized by keeping them together for 4 days and later separated into groups (Table 1). After infection, all cats underwent daily clinical examinations. At the end of each experiment, all animals were euthanized and necropsied within 2 hr of death.

**Experimental infection.** Experimental studies were officially approved by the Swiss Veterinary Office (66/2000). Cats were kept in groups under optimal ethological conditions. Each animal was infected per-orally twice within 24 hr, under anaesthesia, by application of 2 ml of the total infectious dose using an oesophageal tube. Different virus isolates, infection doses and formulations were used (Table 1).

**Sample preparation for determination of viraemia, faecal shedding and viral loads.** Viral shedding was identified from purified viral RNA extracted from faecal samples collected during the examination period and immediately after death. Viraemia was determined based on a standard RNA template (Gut et al., 1999). To normalize the extracted RNA to the same cell number in all samples, expression levels of the housekeeping gene GAPDH were used to determine viral load value of

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\text{Viral load} = \frac{10^{2}}{N_{GAPDH}}
\]

Relative FCoV viral loads were determined by one-tube real-time RT-PCR, using an automated fluorimeter (TaqMan, ABI 7700; Applied Biosystems) to detect a 102 bp fragment of the conserved FCoV 7b gene (Gut et al., 1999).

Expression levels of the housekeeping gene GAPDH were used to normalize the extracted RNA to the same cell number in all samples, and viral loads were recalibrated to the GAPDH content in the respective tissues. For samples where an FCoV signal was not observed after 45 cycles, a plausible viral load was created. This value was ascertained from the lowest RNA signal for GAPDH found in this study. Thus, a test result in an organ was defined as positive with a viral load value of

\[
>0.000000000027776
\]

The infectious doses for the experimental infection were calculated as the RNA copy number present in the volume of the infectious inoculum given to the cats (Table 1). RNA copy number was determined based on a standard RNA template (Gut et al., 1999).

**Immunohistological demonstration of FCoV antigen in tissues.** The intestines (duodenum, jejunum, ileum, caecum, colon, and rectum) from cats euthanized on day 48 p.i. (5.1–5.15) and the seven livers [cats 1.1 and 1.2 (day 14 p.i.), 3.1 and 4.2 (day 28 p.i.), and 5.4, 5.14 and 5.15 (day 48 p.i.)], ten mesenteric lymph nodes [cats 1.1 and 1.2 (day 14 p.i.), 2.1–2.3 (day 17 p.i.), 3.1, 4.1 and 4.2 (day 28 p.i.), and 5.4 and 5.14 (day 48 p.i.)] and three lungs [cats 1.1 (day 14 p.i.), 2.3 (day 17 p.i.) and 4.1 (day 28 p.i.)] with the highest relative FCoV viral loads were examined for the presence of FCoV antigen by immunohistochemistry, using a mouse monoclonal antibody (clone FCV3-70; Custom Monoclonals) as described previously (Kipar et al., 1998, 1999, 2000, 2005).

**Statistical analysis.** All analyses were performed using SAS v9.1 computer software (SAS Institute). Statistical associations were explored between the viral loads in cats after infection and the characteristics of the experiment (e.g. dose of virus, viral load in organs and days from challenge to euthanasia). Initial assessment of the data was carried out using descriptive statistics. Comparisons between the two groups of cats were carried out using a Mann–Whitney test with unpaired ties. Comparisons between organs (i.e. more than two groups) were carried out using a Kruskal–Wallis ANOVA. For instances of four or more group comparisons, an additional statistical test (e.g. two-way ANOVA and multiple comparisons using Tukey’s test for unequal groups) was also employed and results were compared. Comparisons between the frequencies of positive organs were performed using Fisher’s exact test. The exact \( P \) values for all non-parametric statistics were calculated using Monte Carlo estimation.

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