Campylobacter Species and Neutrophilic Inflammatory Bowel Disease in Cats


Background: Inflammatory bowel disease (IBD) is a common cause of signs of gastrointestinal disease in cats. A subset of cats with IBD has neutrophilic inflammation of the intestinal mucosa.

Hypothesis: Neutrophilic enteritis in cats is associated with mucosal invasion by microorganisms, and specifically Campylobacter spp. infections might also induce neutrophilic infiltrates because of the loss of mucosal barrier integrity. If no etiological agent is identified, directed or appropriate treatment is unclear and empirical antibacterial treatment might be prescribed.

Using more traditional diagnostic techniques, the evidence for infectious agents causing neutrophilic intestinal inflammation is poor. The clinical relevance of identifying bacteria in fecal samples is unclear as they might be found in the feces of healthy cats as well as in those with diarrhea.6–9

The use of fluorescent in situ hybridization (FISH) has enabled detection and visualization of intact bacteria within tissues and their localization as part of understanding the ecology of a disease process.10 In small animal gastrointestinal research, FISH has been used to demonstrate gastric Helicobacter spp. infections and clearance of these bacteria with treatment,11 and that attaching and invasive E. coli (AIEC) are associated with granulomatous colitis in Boxers and can be eradicated with prolonged antibiotics.12,13 Janeczko14 used FISH and showed higher number of mucosa-associated

Abbreviations:

12-LOX 12-lipoxygenase
AIEC attaching and invasive E. coli
DNA deoxyribonucleic acid
FCEAI feline chronic enteropathy activity index
FISH fluorescence in situ hybridization
H & E hematoxylin and eosin
HPF high-powered field
IBD inflammatory bowel disease
IL-8 interleukin 8
NF-κB nuclear factor kappa beta
PCR polymerase chain reaction
rRNA 16S ribosomal ribonucleic acid
WSAVA World Small Animal Veterinary Association

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The work in this study was performed at the School of Veterinary Sciences, University of Bristol and supported by a grant from the Langford Trust. The abstract from this paper was presented at the European College of Veterinary Internal Medicine Congress in Lisbon, Portugal 2015.

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Submitted October 21, 2015; Revised February 19, 2016; Accepted June 11, 2016.

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DOI: 10.1111/jvim.14374
Enterobacteriaceae in cats with gastrointestinal signs compared with healthy cats, although it was not clear whether this association was the cause of the inflammation or a consequence of the mucosal disruption and altered environment selecting for bacterial dysbiosis.

The aims of this study were: (1) to determine the presence, and identity of intact bacteria within feline intestinal biopsy specimens using FISH and (2) to determine the location of neutrophils and their proximity to any bacteria to demonstrate an association and possible bacterial etiology.

**Materials & Methods**

**Study Population**

Cases from which there was histopathologic assessment of endoscopic intestinal biopsy specimens at the University of Bristol, School of Veterinary Sciences, were identified from the pathology archive and the biopsy specimens were reviewed. The inclusion criteria were that duodenal biopsy specimens had a diagnosis of neutrophilic or lymphoplasmacytic inflammation and were of adequate quality and size for FISH to be performed requiring at least 3 well-oriented villi with subvillus lamina propria extending to the muscularis mucosa in each section.

Samples from 15 cats (7 with neutrophilic IBD and 8 with lymphoplasmacytic IBD) met the inclusion criteria for the study and also had sufficient intestinal tissue archived as formalin-fixed and paraffin-wax-embedded material. The signalment of these cats was documented, but the clinical severity and response to treatment was not uniformly reported. Ethical approval for the study was granted by the University of Bristol Ethics Committee. Samples from all cats were retrospectively analyzed for intestinal bacterial pathogens by PCR as detailed below.

**Histopathologic Analysis**

In every case, hematoxylin and eosin-stained sections of the duodenal biopsy specimens were reviewed, without knowledge of the case history, by a Board-certified veterinary pathologist (MJD). The histopathologic changes were graded according to the scoring system developed by the World Small Animal Veterinary Association (WSAVA) Gastrointestinal Standardization Group. Five parameters of inflammation (ie, intraepithelial lymphocytes, lamina propria lymphocytes, plasma cells, eosinophils and neutrophils, and other inflammatory cell types within the lamina propria) were evaluated. Morphologic features of mucosal inflammation (villus stunting, epithelial injury, crypt dilatation/distortion, lacteal dilatation, and mucosal fibrosis) were also evaluated. Changes in morphological features were noted, but there was no appreciable difference between groups, with a range of none–moderate within each group. The final diagnosis was categorized as 1 of 8 different histopathologic diagnoses: no abnormalities detected (NAD), lymphoplasmacytic inflammatory, eosinophilic inflammatory, neutrophilic inflammatory, lymphangiectasia, lymphoma, mucosal atrophy/fibrosis (noninflammatory) and other. Only samples from cats with lymphoplasmacytic inflammatory histopathologic changes and neutrophilic inflammatory histopathologic changes were included in the study.

**Fluorescence In Situ Hybridization—Bacteria**

Fluorescence in situ hybridization (FISH) was performed on formalin-fixed and paraffin wax-embedded duodenal biopsy specimens using a method slightly modified from that previously reported in the literature, in that hybridization buffer containing 25% formamide was used as opposed to buffer containing 30% formamide in the previous study. 16S ribosomal RNA (rRNA)-directed fluorescence-labeled oligonucleotide probes were used (Table 1). Successive sections from each biopsy specimen were analyzed in steps, each employing a different probe mix. Steps one and two were general probes for Eubacteria and Firmicutes using three different probes for each bacterial group. A mixture of probes with slight base variations were used to maximize the spectrum of coverage of each probe mix. The third step used specific probes for Campylobacter species. These were used in combination with one probe labeling C. upsaliensis, another probe for C. jejuni, and a final probe labeling all thermophilic Campylobacter spp. Bacteria labeled solely by this last probe could either be C. lari or C. coli and were speciated by subsequent polymerase chain reaction (PCR). We did not examine the sections for E. coli and would not, using these methods, have been able to detect AIEC.

Processed slides were mounted with glass cover slips with hard-set Vec-tashield mounting medium containing DAPI. Once set, slides were examined with a Leica DMR-A Microscope equipped with a monochrome Hamamatsu 8-bit digital camera. Bacterial cells in the epithelium and lamina propria were counted at ×400 magnification by a single blinded observer across 10 fields of view, and an average count was obtained. Fields of view were chosen as being central to a biopsy and including epithelium and so that not more than 2 were viewed for a single biopsy.

**Polymerase Chain Reaction (PCR)**

DNA was extracted from pooled multiple sections of tissue using a DNeasy blood and tissue kit (Qiagen, Manchester, UK). This was screened by Langford Veterinary Services (Bristol, UK) for the presence of Salmonella, pathogenic E. coli, Campylobacter spp., and Clostridium spp. A 153-bp DNA fragment unique to Campylobacter was amplified by PCR and sequenced to confirm the Campylobacter species present.

| Table 1. FISH oligonucleotide probes. Details of sequences of oligonucleotide probes used for in situ localization of bacterial species within duodenal biopsy tissues. |
|----------------|----------------|-----------------|----------------|
| **Probe name** | **Bacterial target** | **Sequence (5’–3’)** | **5’-Modification** |
| EUB338 | Eubacteria | GCTGCCTCCCGTAGGAGT | FITC |
| EUB338-II | Eubacteria | GACGCCACGGCTAGGTT | FITC |
| EUB338-III | Eubacteria | CGCTGCCACGGCTTAGGTT | FITC |
| LGC354A | Firmicutes | TGGAAATTCCTACTGC | Texas Red |
| LGC354B | Firmicutes | CGGAAGATCTCCATGC | Texas Red |
| LGC354C | Firmicutes | CGGAAGATCTCCATGC | Texas Red |
| Cathem | Thermophilic Campylobacter | GCCCTAAGGCTCCTCCA | Texas Red |
| Cajej | Campylobacter jejuni | AGCTACAACACCTTATACCG | FITC |
| CaUp | Campylobacter upsaliensis | CTCAGAGATTTGTTGAT | AF350 |
**Fluorescent In Situ Hybridization—Colocalization of Bacteria and Neutrophils**

*Campylobacter jejuni* and thermophilic *Campylobacter* spp. probes were used in combination with a FISH probe designed to detect feline neutrophil elastase mRNA. The feline neutrophil elastase probe 5’CAGAGGCTGCTGAACGACATCGTGATTCTC CAGCTCAAT3’ was used concurrently with *Campylobacter* spp. probes (Thermophilic *Campylobacter* spp. 5’GCCCTAAGCGTCC TTCCA 3’ and *C. jejuni* 5’AGCTAACCACCTTATACCG 3’). Slides were viewed under fluorescence using a DMRB microscope (Leica) equipped with a Retiga EXi camera (QImaging) and Volocity imaging software (PerkinElmer) and searched for the expression of feline neutrophil elastase and *Campylobacter*. Images were taken when a field of view (×400 objective) contained a neutrophil elastase expressor and one or both of the *Campylobacter* expressors. A record was made of any neutrophil elastase expressors that did not have a *Campylobacter* expressor in the same field, as well as the overall presence of *C. jejuni* and thermophilic *Campylobacter*. Image J software (http://rsb.info.nih.gov/ij) was used after images had been captured to allow automated measurement of the distance between two points selected by the user—neutrophil and Campylobacter. The distance was measured in pixels on the digital images. The maximum field size used was 1850 pixels. The use of this software allowed multiple accurate measurements to be taken and compiled for statistical analysis.

**Statistics**

Analysis was performed using statistical software; GraphPad Prism version 5.03. The presence of bacteria in the cats with neutrophilic and lymphoplasmacytic IBD was compared and tested for statistical significance using Chi-squared testing.

**Results**

Fifteen cats met the inclusion criteria: 8 were domestic short hair (DSH), 1 was domestic long hair (DLH), and 5 were pure breeds (Maine Coon, Birman, Oriental, and 2 Siamese). There was no breed recorded for one of the biopsy specimens. The age range was 2–15 years (median 9 years). There were 11 male cats and 4 female cats. Bacterial DNA extraction results for all the cats showed than none were positive for *Salmonella* or pathogenic genotypes of *E. coli* (a definition which excludes AIEC). One animal in each group was positive for *Clostridium* spp. All were *Campylobacter* positive. Sequencing for *Campylobacter* to separate the thermophilic species showed that *C. lari* and *C. upsaliensis* could not be detected in tissues (data not shown). Only the species *C. coli* and *C. jejuni* were present in tissues. The PCR speciation results concurred with FISH results (Fig 1).

**Presence of Bacteria in the Cats with Neutrophilic IBD and the Cats with Lymphoplasmacytic IBD**

Bacteria were identified within the mucosa of cats with either neutrophilic or lymphoplasmacytic IBD.
inflammation. There was no significant difference in the number of Eubacteria seen in specimens from either of the two groups. There were few *C. upsaliensis* and further statistical analysis was not possible. *C. jejuni* and *C. coli* were present in greater number and so the distribution of these organisms between the neutrophilic group and lymphoplasmacytic group of cats was compared. There was no significant difference between the presence of *C. jejuni* within the mucosa of the cats with neutrophilic or lymphoplasmacytic inflammation. However, *C. coli* was more prevalent in the mucosa of cats with neutrophilic inflammation (6/7) compared with cats with lymphoplasmacytic inflammation (1/8) and this distribution was significant \( (P = .009) \) (Fig 1).

**Number of Bacteria in the Cats with Neutrophilic IBD and the Cats with Lymphoplasmacytic IBD**

The number of *C. jejuni* and *C. coli* within the mucosa of the cats with neutrophilic inflammation and cats with lymphoplasmacytic inflammation were then compared. For *C. jejuni* the numbers were similar in both groups and there was no significant difference in distribution. However, for *C. coli* the numbers were greater within the cats with neutrophilic inflammation (median number of bacteria per high-powered field of view [×400 magnification] was 0.7) compared to the cats with lymphoplasmacytic inflammation (median number of bacteria per high-powered field of view was 0) and this difference was significant \( (P = 0.002) \). (Fig 1).

**Coocalization of Bacteria with Neutrophils**

There was an insufficient number of neutrophils within the mucosa of cats with lymphoplasmacytic inflammation to investigate the association with microorganisms, as 62 of the 63 fields of view examined did not show *Campylobacter* and a neutrophil in the same field of view. Within the mucosa of cats with neutrophilic inflammation, only 6 of the 63 fields of view examined did not show *Campylobacter* and a neutrophil in the same field of view. Cells expressing neutrophil elastase had typical PMN morphology on light microscopy.

Within the mucosa of cats with neutrophilic inflammation the co localization of *C. jejuni* and *C. coli* was analyzed; *C. upsaliensis* numbers were again too few for statistical analysis. For *C. jejuni* there was no evidence of an association between location of the bacteria and that of the neutrophils and often no organism was seen within the same field of view as a neutrophil (>1850 pixels). For *C. coli*, there was an association between the location of the organism and that of the neutrophils \( (P < .001) \) (Fig 2).

**Discussion**

Inflammatory bowel disease is one cause of chronic signs of gastrointestinal disease in cats, others include adverse food reactions, antibiotic responsive enteropathy, and infection (virus, fungi, bacterial, protozoal, or parasitic). Cases with histopathologic changes consistent with neutrophilic inflammation may result from mucosal invasion by microorganisms, develop secondary to mucosal defects, or *Trichomonas foetus* infection or be idiopathic. The aims of this study were to use FISH to elucidate the presence of mucosal bacterial invasion in feline IBD, and to identify any association between *Campylobacter* invasion and neutrophilic inflammation (Picture 1).

Bacteria were identified within the mucosa of cats with both neutrophilic and lymphoplasmacytic inflammation, suggesting mucosal disruption. There was no clear pattern to their localization (mucosa/submucosa/lamina propria) within the tissue. The total numbers of bacteria did not differ between groups and for two

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**Fig 2.** Proximity of *Campylobacter coli* is closer to neutrophils as compared to *C. jejuni*. Data show pixel distance of neutrophils to *Campylobacter* as measured by ImageJ software. The maximal pixel distance per field is 1850 pixels; when a field of view did not contain both a *Campylobacter* and a neutrophil distance was recorded as 1850 pixels.

**Picture 1.** Fluorescence in situ hybridization (FISH) demonstrating red fluorescence for a thermophilic *Campylobacter* species (*C. coli*) and blue fluorescence for neutrophil elastase. Not all bacteria are highlighted in a single sectional view as organisms outside the plane of focus are not visible in a single image.
bacterial species (C. jejuni and C. upsaliensis) there was also no difference in distribution between the two groups. A significant difference was found between the two groups regarding the presence of C. coli. This in itself is simply an observation and merely demonstrates an increased presence of C. coli in cats with neutrophilic inflammation. However, FISH neutrophil elastase probes demonstrated close proximity to neutrophils and confirmed the possible association with C. coli. No association between neutrophil location and other species of bacteria was demonstrated. This association supports the hypothesis that C. coli is particularly associated with a specific neutrophilic form of inflammation in the duodenal mucosa of cats with enteric disease compared with other microorganisms.

Campylobacter species are known to be powerful chemoattractants for neutrophils. Previous in vitro studies with porcine and human epithelial cell lines have shown that C. jejuni and C. coli induce neutrophilic intestinal inflammation via activation of NF-κB and subsequent production of the neutrophil chemokine IL-8 and also induce neutrophilic inflammation in human and feline epithelium via the production of chemotactic n-formyl peptides in concert with metabolites of 12-LOX. In the present retrospective study there was evidence that C. coli, but not C. jejuni, appears to colocalize with neutrophils and therefore might be acting as the etiological agent in this particular subset of feline inflammatory bowel disease, although other cell types and mechanisms could also be involved. In vitro studies using feline neutrophils similarly to previous work on porcine neutrophils would be necessary to determine a mechanism for chemoattraction.

Limitations of the study mostly relate to its retrospective nature and the lack of clinical data. Once the inclusion criteria were met, the numbers recruited to each group were small because samples had to be of sufficient quality for histopathological review and of adequate volume for the subsequent molecular diagnostic tests to be performed.

We were unable to document any differences in clinical presentation between the two groups or to assess data relating to treatment and outcome as the cases were not always treated in our hospital; these were drawn from a pathology archive so the remit of this study is specifically to associate bacteria with pathology. The lack of clinical data means we are unable to determine if any of the cats received antibiotic treatment before endoscopic biopsy collection. This is a potential confounding effect but prior antibiotic treatment would be expected to reduce bacterial numbers and the likelihood of finding a significant difference between the groups, except in the unlikely event that the antibiotic treatment before endoscopic biopsy induced the neutrophilic IBD. This preliminary findings from this retrospective study highlight a need for prospective investigation to understand the relevance of the C. coli tissue localization. This would allow investigation into whether this subset of cats have a different clinical presentation and FCEAI score or different endoscopic lesions identified on endoscopy. Identification of C. coli as a potential etiological agent for this subset of cats would allow directed treatment and would potentially improve the outcome and also reduce any zoonotic risk. Further in vitro molecular studies would also help determine by what means C. coli induces the neutrophilic response; whether it produces compounds which stimulate neutrophils or induces feline intestinal cells to produce neutrophil chemoattractants. A prospective study would also allow an epidemiological study of risk factors and the interaction of these cats with their owners.

Acknowledgments

Conflict of Interest declaration: Authors declare no conflict of interest.

Off-label Antimicrobial Declaration: Authors declare no off-label use of antimicrobials.

References