

Effect of Incubation Temperature on the Detection of Thermophilic *Campylobacter* Species from Freshwater Beaches, Nearby Wastewater Effluents, and Bird Fecal Droppings

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This large-scale study compared incubation temperatures (37°C versus 42°C) to study the detection of thermophilic *Campylobacter* species, including *Campylobacter jejuni*, *C. coli*, and *C. lari*, in various surface water samples and bird fecal droppings around Hamilton Harbor, Lake Ontario. The putative culture isolates obtained from incubation temperatures of 37 and 42°C were confirmed by *Campylobacter* genus- and species-specific triplex PCR assays targeting the 16S rRNA gene and the 16S-23S rRNA gene internal transcribed spacer (ITS) region. A total of 759 water, wastewater, and bird fecal dropping samples were tested. Positive amplification reactions for the genus *Campylobacter* were found for 454 (60%) samples incubated at 37°C, compared to 258 (34%) samples incubated at 42°C. *C. jejuni* (16%) and *C. lari* (12%) were detected significantly more frequently at the 42°C incubation temperature than at 37°C (8% and 5%, respectively). In contrast, significantly higher rates of *C. coli* (14%) and other *Campylobacter* spp. (36%) were detected at the 37°C incubation temperature than at 42°C (8% and 7%, respectively). These results were consistent across surface water, wastewater, and bird fecal dropping samples. At times, *Campylobacter* spp. were recovered and detected at 37°C (3% for *C. jejuni*, 10% for *C. coli*, and 3% for *C. lari*) when the same samples incubated at 42°C were negative. A significantly higher rate of other *Campylobacter* spp. was detected only at 37°C (32%) than only at 42°C (3%). These results indicate that incubation temperature can significantly influence the culturability and detection of thermophilic and other fastidious *Campylobacter* spp. and that a comprehensive characterization of the *Campylobacter* spp. in surface water, wastewaters, or bird fecal droppings will require incubation at both 37 and 42°C.

Campylobacter species have been recognized as one of the leading causes of bacterial gastroenteritis in humans (1–3). The genus *Campylobacter* currently comprises 22 species and 8 subspecies (4), but from a public health perspective, *Campylobacter jejuni*, *C. coli*, and *C. lari* are the most frequently implicated species in human infections (5). These species were first isolated on a selective agar incubated microaerobically at 43°C (6, 7), and that led to the name thermophilic campylobacters. These three species can commonly occur in the gastrointestinal tracts of humans and other animals, including bovines, pigs, and birds. Sea gulls and several other wild birds can be an important source of *Campylobacter* spp., such as *C. jejuni* and *C. lari* (3, 8, 9). *Campylobacter* spp. from the feces of birds or domestic and wild animals, municipal sewage discharges, or agricultural runoff can contaminate water (3), and water is an established vehicle for the transmission of these species to human and domestic animals, leading to outbreaks of waterborne disease.

Several *Campylobacter* isolation procedures have been developed for water, including an ISO standard method (10) using concentration of cells by membrane filtration or centrifugation, as well as enrichment regimens using different selective growth media. These procedures are used widely to isolate these species from food and water (11, 12). There are numerous challenges in recovery of *Campylobacter* spp. from water, such as the frequently small numbers of cells, low growth rate, intrinsic fastidious growth requirements, and presence of a significant proportion of organisms that may be injured or have difficulty in adapting to *in vitro* conditions. Incubation temperature can play a vital role in the culturability and detection of thermophilic *Campylobacter* spp. (13–15).

Traditionally, procedures to isolate these organisms from food,

water, and feces have most commonly been conducted at a 42°C incubation temperature (16–18). Many investigations of *Campylobacter* spp. in water have used a 42°C incubation temperature (9, 19–25). However, other water studies have used a 37°C incubation temperature (26–28). Several studies have used the Cape Town method with an incubation temperature of 37°C for detection of *Campylobacter* from food and human feces, suggesting this method as better for detecting a wide range of *Campylobacter* spp. (14, 29–32). Although the Cape Town method has been considered a useful tool for the isolation of *Campylobacter* spp. from clinical samples (31, 32), other studies have reported it as insufficient for the recovery and detection of thermophilic *Campylobacter* spp. in aquatic environments, where species such as *C. jejuni* and *C. lari* could be present in small numbers (26, 27). Another approach in water studies has been to have a 2- to 4-h preenrichment step at 37°C before subsequent enrichment at 42°C (33–38). However, studies that used a preenrichment step of 37°C followed by enrichment at 42°C have often observed a low detection rate for *C. coli*, suggesting that a 42°C incubation temperature might not support the growth of stressed and injured cells of *C. coli* present in small numbers in water. Phillips (39) suggested that more than one incubation temperature might substantially improve the iso-

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lation of thermophilic *Campylobacter* spp. without diminishing the isolation of other fastidious *Campylobacter* spp.

To our knowledge, no large-scale study has previously compared *Campylobacter* detection rates at 37 and 42°C across diverse environmental matrices, such as surface water samples, municipal wastewater samples, and bird fecal droppings common to urban environments. It is important to assess the limitations of different incubation temperatures regarding the culturability and detection of thermophilic and other fastidious *Campylobacter* spp. from various environmental matrices. This study compared incubation temperatures of 37 and 42°C for detecting *Campylobacter* spp., including *C. jejuni*, *C. coli*, and *C. lari*, in water samples collected from two freshwater beaches and offshore harbor water, as well as municipal wastewater and bird fecal dropping samples from around Hamilton Harbor, Lake Ontario. We refer to campylobacters other than *C. jejuni*, *C. coli*, and *C. lari* as “other *Campylobacter* spp.”

MATERIALS AND METHODS

Collection of surface water and bird fecal dropping samples. Surface water samples were collected from two freshwater beaches (Bayfront Park and Pier4 Park) in Hamilton Harbor, Lake Ontario, from 2007 to 2009. Sampling at the beaches was carried out along a single transect at each beach, at three depth zones, including sand pore water and ankle- and chest-depth waters. Sand pore water was collected by digging a hole in the wet foreshore sand about 1 meter inland from the water's edge and collecting the water that seeped into the hole. Offshore surface water samples were collected by boat from the middle of the harbor, near a wastewater treatment plant offshore outfall. Municipal wastewater samples were obtained from the final effluents of four municipal sewage treatment plants (STPs) that discharge into the harbor area and a combined sewer overflow (CSO) storage tank located at Bayfront and Pier4 Park beaches that occasionally overflowed during storm events. Fresh bird fecal dropping samples from ring-billed gulls (*Larus delawarensis*) and Canada geese (*Branta canadensis*) were collected on the beaches within 2 m of the waterline. All water, wastewater, and bird fecal samples were collected on a biweekly basis between April and December over 3 years (2007 to 2009). Water and wastewater samples were collected in 2-liter sterile bottles, whereas fecal samples were collected in sterile tubes containing 2 ml phosphate-buffered saline (1× PBS) solution. The water, wastewater, and fecal samples were returned on ice to the laboratory and processed on the same day of their collection.

Isolation and culture conditions. Samples were processed following a protocol described by Khan et al. (12). Briefly, 1 liter of each water or wastewater sample was centrifuged at 14,000 × *g* for 20 min (Beckman, Indianapolis, IN), and the pellet was resuspended in 3 ml saline (0.85%) solution for concentration of *Campylobacter* cells. Bird fecal samples collected in PBS solution were homogenized by vortexing and analyzed to detect the presence/absence of thermophilic *Campylobacter* spp. To estimate the number of *Campylobacter* cells per liter, the resuspended pellet obtained from 1 liter of centrifuged water was analyzed by a minimum probable number (MPN) method using a 10-fold serial dilution approach for a semiquantitative analysis of *Campylobacter* occurrence. One-milliliter aliquots of each suspended pellet and fecal sample were inoculated into two sets of Bolton broth (Oxoid) tubes containing a selective antibiotic (cefoperazone, cycloheximide, trimethoprim, and vancomycin) supplement. The inoculum was serially diluted in the two sets of tubes, and one set each was incubated at 37 and 42°C under microaerophilic conditions (5% O₂, 85% N₂, and 10% CO₂) for 48 h in an MCO-18 M multigas incubator (Sanyo, Tokyo, Japan). The semiquantitative enumeration was carried out by assessing turbidity and subculture confirmation. The cultures from each tube were further streaked with a sterile loop onto modified Karmali agar (MKA) (Oxoid) containing a selective supplement including antibacterial and antifungal (amphotericin B, cefoperazone,

sodium pyruvate, and vancomycin) agents, and plates were incubated at 37 and 42°C under microaerophilic conditions for 24 to 48 h. The putative *Campylobacter* cultures were selected based on their growth characteristics and colony morphology, i.e., smooth, shiny, and convex with defined or flat edges, transparent or translucent, colorless to grayish or light cream (24). The plates that contained such colonies were further analyzed by DNA extraction and PCR assays.

DNA extraction and genus-specific PCR amplification. DNA extractions from scraping of multiple isolates recovered from an MKA growth medium plate were carried out using a boiling protocol as previously described (40). Briefly, putative *Campylobacter* colonies were resuspended in 75 μl 1× TE buffer (10 mM Tris-HCl, 1 mM EDTA, pH 8.0). The suspended solution was boiled for 10 min and centrifuged. The supernatant containing DNA was transferred to a 1.5-ml sterile microcentrifuge tube and kept at –20°C for further analysis. For the confirmation of putative cultures to genus-level identification, a DNA-based PCR amplification assay was performed using *Campylobacter* genus-specific oligonucleotide primers (41). The 25-μl reaction mixture contained 50 to 70 ng of template DNA, 0.15 unit of *Ex Taq* DNA polymerase (TaKaRa, Shiga, Japan), 1× *Ex Taq* buffer with MgCl₂, a 200 μM concentration of each deoxynucleoside triphosphate (dNTP), and 50 ng each of forward (5'-GGA TGA CAC TTT TCG GAG C-3') and reverse (5'-CAT TGT AGC ACG TGT GTC-3') primers. The amplification was performed by an initial template denaturation step at 94°C for 3 min, followed by 30 cycles of amplification by repeating denaturation at 94°C for 60 s, annealing at 55°C for 60 s, and extension at 72°C for 60 s, with a final 5-min incubation at 72°C, using a Mastercycler gradient PCR system (Eppendorf, Hamburg, Germany). The PCR amplicons were electrophoresed in a 1% agarose gel matrix (Fisher Scientific, NJ) with 1× (Tris-acetate-EDTA) TAE buffer, using a 100-bp DNA size marker (PGC Scientific, MD). The gels were stained with ethidium bromide (0.5 μg/ml), visualized on a UV transilluminator, and photographed using an Ingenius Syngene Bioimaging gel documentation system.

Species-specific PCR amplification. A species-specific PCR amplification was further performed to identify *Campylobacter* spp. by a triplex PCR assay using oligonucleotide primer pairs for three thermophilic *Campylobacter* spp. (*C. jejuni*, *C. coli*, and *C. lari*) as described by Khan and Edge (42). The triplex PCR amplification reaction was carried out in a Mastercycler Gradient PCR system (Eppendorf) with a 25-μl reaction mixture containing 50 to 70 ng of template DNA, 1.25 units of *Ex Taq* DNA polymerase (TaKaRa), 1× *Ex Taq* buffer with MgCl₂, a 200 μM concentration of each dNTP, and 80 ng each of the forward and reverse primer pairs for each target *Campylobacter* spp., including *C. jejuni* (forward, ACT AAA TGA TTT AGT CTC A; and reverse, CTT AGA TTT ATT TTT ATC TTT AAC T), *C. lari* (forward, AAA TAT ATA CTT GCT TTA GAT T; and reverse, CAA TAA AAC CTT ACT ATC TC), and *C. coli* (forward, GAA GTA TCA ATC TTA AAA AGA TAA; and reverse, CTT ACT TTA GGT TTT AAG ACC). The final volume (25 μl) was adjusted with filtered, sterile distilled water. The PCR was performed using an initial template denaturation step of 94°C for 3 min followed by 30 cycles of amplification (denaturation at 94°C for 30 s, annealing at 46°C for 45 s, and extension at 72°C for 30 s) and ending with a 5-min extension at 72°C. Due to the expected small amplicon fragment size, the PCR amplicons were electrophoresed in a 2% agarose gel matrix, stained, and scanned as described in the preceding section.

Statistical analysis. Statistical analyses were performed using Statistica 10.0 (StatSoft Inc.). McNemar chi-square contingency tests were applied to test for significant differences in the recovery and detection of thermophilic and other *Campylobacter* spp. between 37 and 42°C. Differences were considered significant if the *P* value was <0.05.

RESULTS

A total of 759 surface water, wastewater, and bird fecal dropping samples were collected around Hamilton Harbor between 2007 and 2009. These samples included 288 beach water samples, 89

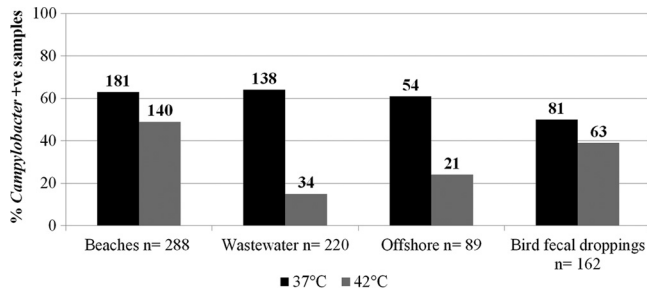


FIG 1 Percentages of beach water, wastewater, offshore water, and bird fecal dropping samples positive for campylobacters at two different incubation temperatures (n = total number of collected samples). The number of positive samples is presented above each bar.

offshore water samples, 220 wastewater samples, and 162 bird fecal dropping samples. All putative *Campylobacter* culture isolates observed at incubation temperatures of 37 and 42°C on MKA medium showed typical growth patterns and were consistently confirmed by the *Campylobacter* genus-specific 16S rRNA gene PCR assay, with an expected amplicon size of 816 bp. Species-specific detection of *Campylobacter* spp. in water and fecal samples was further confirmed using the triplex PCR assay, with expected amplicon sizes of 349, 279, and 72 bp for *C. jejuni*, *C. lari*, and *C. coli*, respectively.

Analysis of all 759 beach water, wastewater, offshore, and bird fecal samples indicated that *Campylobacter* was detected significantly more frequently at an incubation temperature of 37°C (60%) than at one of 42°C (34%) ($P < 0.05$). These findings were consistent across beach, wastewater, offshore water, and bird fecal dropping samples. *Campylobacter* recoveries from beach water (63%), wastewater (63%), offshore water (61%), and bird fecal (50%) samples were significantly higher ($P < 0.05$) at 37°C than at the 42°C incubation temperature (49%, 15%, 24%, and 39%, respectively) (Fig. 1).

A comparison of the two incubation temperatures revealed differences in the ability to recover and detect *Campylobacter* spp. across all 759 beach and offshore water, wastewater, and bird fecal samples (Table 1). Thermophilic *Campylobacter* spp., including *C. jejuni* (16%) and *C. lari* (12%), were detected significantly more often using the 42°C incubation temperature, whereas *C. coli* (14%) and other campylobacters (36%) were detected significantly more often using the 37°C incubation temperature ($P < 0.05$).

These differences between incubation temperatures in the ability to recover and detect *Campylobacter* spp. were consistent across all beach water, offshore water, wastewater, and bird fecal samples (Fig. 2A to D). Based on the number of samples collected from each sampling location and type, *C. jejuni* and *C. lari* were always more commonly detected at 42°C than at 37°C, and they were more common in beach water samples (24 and 18%, respectively) and bird fecal droppings (26 and 20%, respectively) than in offshore (7 and 11%, respectively) and wastewater (3 and 1%, respectively) samples (Fig. 2A and C). On the other hand, *C. coli* and other *Campylobacter* spp. were always more commonly detected at 37°C than at 42°C, and they were more common in beach water (20 and 30%, respectively), wastewater (17 and 45%, respectively), and offshore (11 and 35%, respectively) samples than in bird fecal droppings (2 and 32%, respectively) (Fig. 2B and D).

Interestingly, no *C. coli* was detected in bird fecal droppings at the 42°C incubation temperature.

Further analysis of the recovery and detection of multiple thermophilic *Campylobacter* spp., including *C. jejuni*, *C. coli*, and *C. lari*, in a single sample revealed an overall low frequency of co-occurrence of *Campylobacter* spp. in water and bird fecal samples. *C. jejuni* and *C. lari* were detected together more commonly at 42°C (6%) than at 37°C (2%) (Table 2). On the other hand, *C. coli* was rarely recovered together with other *Campylobacter* spp. ($\leq 1\%$) at both incubation temperatures (Table 2). A higher frequency of the samples yielded only *C. coli* (13%) as opposed to only *C. jejuni* (5%) or *C. lari* (3%) at 37°C.

From 759 water, wastewater, and bird fecal dropping samples, a significantly higher frequency of campylobacters (35%; $P < 0.05$) was detected only at 37°C than only at 42°C (9%), whereas 25% of samples were positive for *Campylobacter* spp. at both incubation temperatures (Table 3). Significantly higher frequencies ($P < 0.05$) of *C. jejuni* (11%) and *C. lari* (10%) were detected only at 42°C than only at 37°C (3% for each species). In contrast, higher frequencies of *C. coli* (10%) and other *Campylobacter* spp. (32%) were detected only at 37°C than only at 42°C (Table 3).

An analysis of the frequency of detection of *Campylobacter* at different sampling locations at 37°C and/or 42°C was performed, and the results showed that the highest frequencies of *C. jejuni* and *C. lari* were detected at 42°C for beach water (17% and 16%) and bird fecal dropping (16% and 15%) samples (Table 4). In contrast, *C. coli* and other *Campylobacter* spp. were detected most frequently at 37°C for wastewater (15% and 43%) (Table 4).

The concentration of *Campylobacter* cells (number of cells/liter) in water samples was estimated for both incubation temperatures by using an MPN method (Fig. 3). The highest *Campylobacter* cell concentrations ($\geq 10,000$ cells/liter) were commonly obtained from wastewater samples, where most of the samples with very high MPN values were *Campylobacter* negative at the 42°C incubation temperature but were found to contain *C. coli* or other *Campylobacter* spp. at 37°C. The lowest *Campylobacter* cell concentrations, ranging from 10 to 1,000 cells/liter, were commonly obtained from beach and offshore water samples that were positive for *C. jejuni* and *C. lari* at 42°C.

DISCUSSION

Method comparative studies often require intensive and long-term sampling in order to capture the variation and to perform comprehensive comparative analyses as well as assessments of different parameters used in the study. This allows researchers to improve understanding and to recommend optimal parameters

TABLE 1 Numbers of beach water, wastewater, offshore water, and bird fecal dropping samples ($n = 759$) positive for thermophilic *Campylobacter* spp., using two different incubation temperature conditions

Campylobacter(s)	No. (%) of samples	
	37°C	42°C
<i>Campylobacter</i> spp.	454 (60)	258 (34)
<i>C. jejuni</i>	61 (8)	124 (16)
<i>C. coli</i>	107 (14)	60 (8)
<i>C. lari</i>	40 (5)	94 (12)
Other <i>Campylobacter</i> spp.	270 (36)	52 (7)

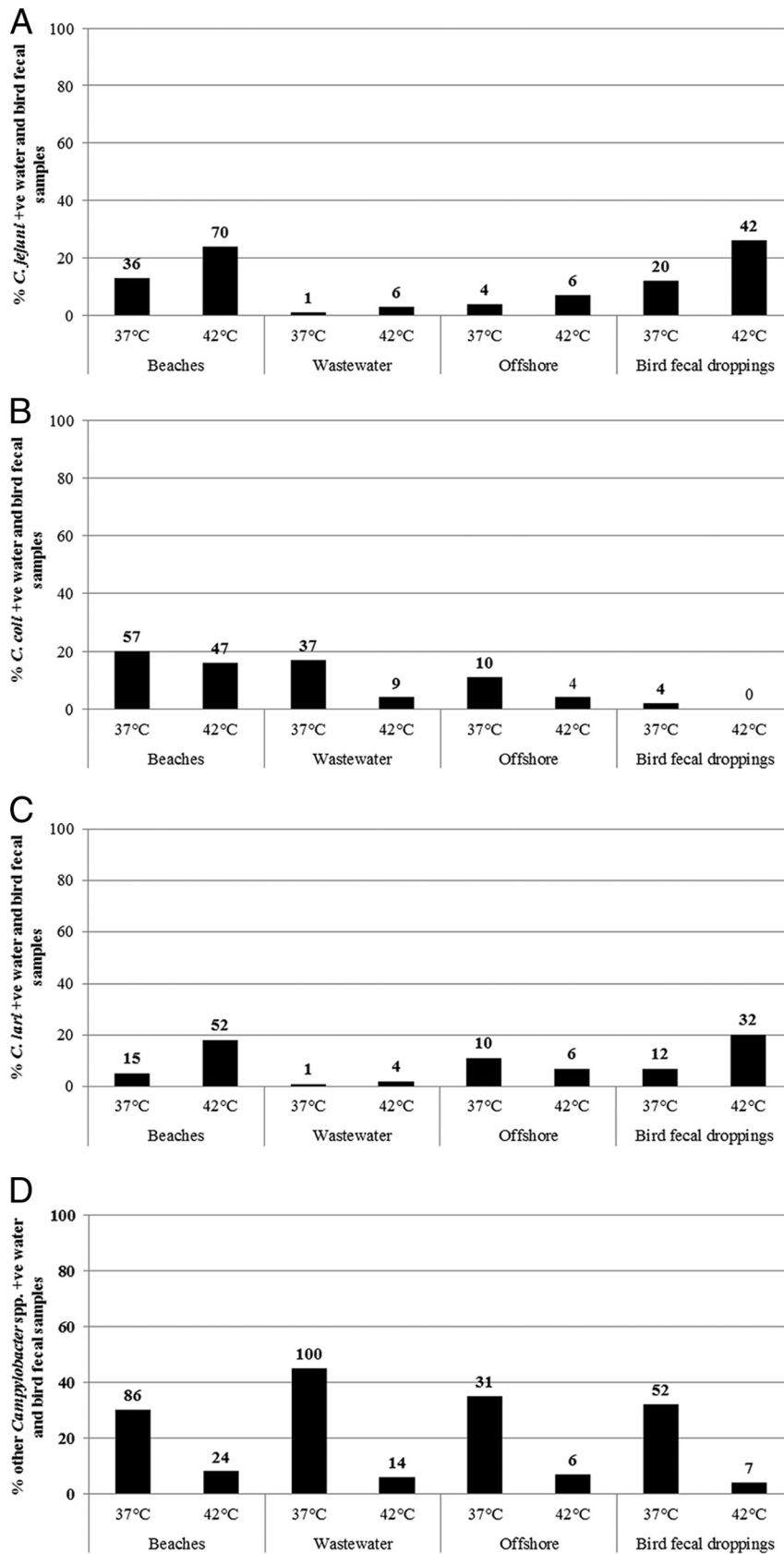


FIG 2 Percentages of recovery of *C. jejuni* (A), *C. coli* (B), *C. lari* (C), and other *Campylobacter* spp. (D) from various water and bird fecal dropping samples at two different incubation temperatures. The number of positive samples is presented above each bar.

TABLE 2 Recovery of multiple *Campylobacter* species from beach water, wastewater, offshore water, and bird fecal dropping samples ($n = 759$) at two different incubation temperatures

<i>Campylobacter</i> spp.	No. (%) of samples	
	37°C	42°C
<i>C. jejuni</i> only	40 (5)	63 (8)
<i>C. coli</i> only	98 (13)	46 (6)
<i>C. lari</i> only	24 (3)	37 (5)
<i>C. jejuni</i> and <i>C. coli</i>	7 (1)	5 (1)
<i>C. jejuni</i> and <i>C. lari</i>	14 (2)	48 (6)
<i>C. coli</i> and <i>C. lari</i>	2 (<1)	1 (<1)
<i>C. jejuni</i> , <i>C. coli</i> , and <i>C. lari</i>	0 (0)	8 (1)

that can be used for future applications. Several comparison studies have been conducted for enhanced recovery and isolation of thermophilic *Campylobacter* spp., such as *C. jejuni*, from food, feces, and water by using different cell concentration approaches (filtration versus centrifugation), enrichment and growth media, and incubation durations (24 versus 48 h). However, samples processed for the isolation of thermophilic *Campylobacter* spp. have typically been incubated at 42°C (12, 43, 44). Preenrichment at 37°C for 2 to 4 h before exposure to selective agents and incubation at 42°C has also been reported as a recovery method that resulted in an increased isolation of *Campylobacter* spp. from natural waters (33–35, 37). Many previous studies on the occurrence of *Campylobacter* spp. in various waters, ranging from drinking water to river, lake, pond, urban, and agricultural watersheds, used a 42°C incubation temperature (9, 18–21, 23, 24, 45). These studies detected *C. jejuni* more commonly than other thermophilic species, such as *C. coli*. These results could reflect actual differences in species occurrence, or they could reflect the fact that a 42°C incubation temperature may not provide a comprehensive recovery of all campylobacters, including *C. coli* and other *Campylobacter* spp. Since *C. coli* has also been significantly implicated in human infections, it is important to consider the growth requirements of *C. coli* and other *Campylobacter* spp. at 37°C. Therefore, this large-scale 3-year study was designed with a goal to compare two incubation temperatures (37 versus 42°C) to determine if incubation temperature significantly influenced the detection of thermophilic *Campylobacter* spp. (*C. jejuni*, *C. coli*, and *C. lari*) that are often present in small numbers, with low growth rates, and with cells that may be stressed as well as sublethally injured. The comparative analysis was conducted on environmental samples collected from 597 water and wastewater samples and 162 bird fecal droppings, using an initial enrichment step with Bolton broth and incubation for 48 h at 37 and 42°C.

TABLE 4 Frequency of recovery of thermophilic and other *Campylobacter* spp. from beach water, wastewater, and offshore water samples and bird fecal droppings at 37 and/or 42°C^a

<i>Campylobacter</i> (s)	No. (%) positive samples											
	Beach water ($n = 288$)			Wastewater ($n = 220$)			Offshore water ($n = 89$)			Bird fecal droppings ($n = 162$)		
	Both temps	Only 37°C	Only 42°C	Both temps	Only 37°C	Only 42°C	Both temps	Only 37°C	Only 42°C	Both temps	Only 37°C	Only 42°C
<i>Campylobacter</i> spp.	107 (37)	73 (25)	33 (11)	22 (10)	116 (52)	11 (5)	18 (20)	37 (41)	4 (4)	44 (27)	37 (23)	19 (12)
<i>C. jejuni</i>	22 (8)	14 (5)	48 (17)	1 (≤1)	0 (0)	5 (2)	1 (1)	3 (3)	5 (6)	16 (10)	4 (2)	26 (16)
<i>C. coli</i>	24 (8)	33 (11)	23 (8)	4 (2)	33 (15)	5 (2)	1 (1)	9 (10)	3 (3)	0 (0)	4 (2)	0 (0)
<i>C. lari</i>	5 (2)	10 (3)	47 (16)	0 (0)	1 (≤1)	4 (2)	4 (4)	6 (7)	2 (2)	7 (4)	5 (3)	25 (15)
Other <i>Campylobacter</i> spp.	13 (5)	73 (25)	11 (4)	6 (3)	94 (43)	8 (4)	1 (1)	30 (34)	5 (6)	5 (3)	47 (29)	2 (1)

^a n values are the total numbers of collected samples for the different sample types.**TABLE 3** Frequency of recovery of *Campylobacter* spp. from water and bird fecal dropping samples ($n = 759$) at 37 and/or 42°C

<i>Campylobacter</i> (s)	No. (%) positive samples at:		
	Both temps	Only 37°C	Only 42°C
<i>Campylobacter</i> spp.	191 (25)	263 (35)	67 (9)
<i>C. jejuni</i>	40 (5)	21 (3)	84 (11)
<i>C. coli</i>	29 (4)	78 (10)	31 (4)
<i>C. lari</i>	16 (2)	24 (3)	78 (10)
Other <i>Campylobacter</i> spp.	26 (3)	244 (32)	26 (3)

Campylobacter was detected significantly more frequently in water samples (including beach water, wastewater, and offshore water samples) and bird fecal droppings at 37°C than at the 42°C incubation temperature. In addition, the water samples collected from two different beaches showed a similar trend of significantly higher recovery of campylobacters at 37°C than at 42°C. Similar results were reported by Bolton et al. (46), where the majority of campylobacters showed better recovery and culturability at 37°C than at 42°C.

Interestingly, *C. jejuni* and *C. lari* were found to be detected more commonly at 42°C in our water and bird fecal dropping samples than at the 37°C incubation temperature. In contrast, *C. coli* and other *Campylobacter* spp. were significantly more commonly detected at 37°C than at 42°C. Studies of *C. jejuni* and *C. fetus* subsp. *jejuni* in human fecal samples, using incubation temperatures of 35 or 37 and 42°C and performed by Gee et al. (47) and Janssen and Helstad (48), found that 42°C resulted in recovery of more isolates of *C. jejuni* and *C. fetus* subsp. *jejuni*. Conversely, Bolton et al. (46) showed that *C. fetus* subsp. *fetus* grew better at 37°C than at 42°C. In our Hamilton Harbor study, the most notable difference between the two incubation temperatures was that more “other *Campylobacter* spp.” were recovered at 37°C than at 42°C. Other *Campylobacter* spp. were recovered from only 52 (7%) water and bird fecal dropping samples at 42°C, compared to 270 (36%) isolates that were recovered and detected at the 37°C incubation temperature. Although our environmental samples may have amplified the problem of recovery due to sublethal injury, cell stress, and small numbers of campylobacters, it is possible that these results are also applicable to clinical samples. For example, the occurrence of *Campylobacter* spp. such as *C. coli* may be underestimated in stool samples by using only a 42°C incubation temperature for recovering thermophilic *Campylobacter* spp.

It was observed that using a 37°C incubation temperature led to detection in water samples of large numbers of campylobacters (MPNs of $\geq 10,000$ to 1,000,000 cells/liter) in 104 samples which were negative for campylobacters at 42°C. Most of the water sam-

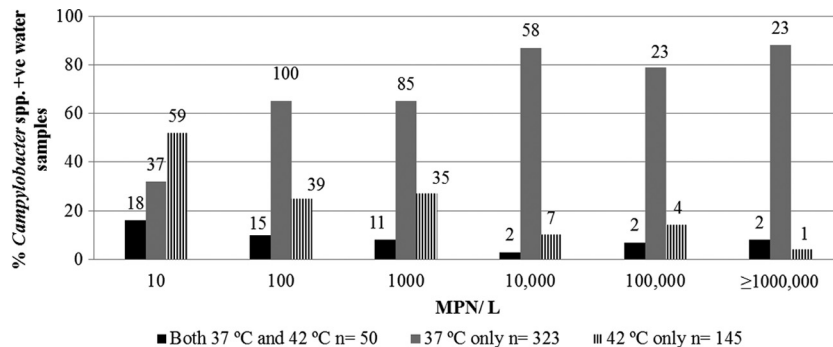


FIG 3 Effect of incubation temperature (37 and/or 42°C) on the percentage of *Campylobacter* spp.-positive beach water, wastewater, and offshore samples at various MPNs. The number of positive samples is presented above each bar.

ples with very high MPNs were found to contain *C. coli* or other *Campylobacter* spp. A 37°C incubation temperature may facilitate detection of cells of *C. coli* and other *Campylobacter* spp. in water samples by allowing growth of stressed and sublethally injured cells. It is also possible that some of the putative campylobacters detected at 37°C are as yet undescribed and potentially novel species within the *Campylobacter* genus or may also be closely related *Campylobacter*-like species from genera such as *Arcobacter* (24). However, Humphrey (33, 34) reported that preenrichment at 37°C may also increase isolation rates of *C. jejuni* from food and water, and we found that *C. jejuni* (3%) and *C. lari* (3%) cells could also be recovered at 37°C when the same samples were negative at 42°C. These results suggest that 37°C may facilitate the recovery and culturability of classical thermophilic *Campylobacter* spp., such as *C. coli*, *C. jejuni*, and *C. lari*, and also enhance the isolation of a wider range of other fastidious *Campylobacter* spp. that could not grow at 42°C.

Similarly, the growth of other fastidious *Campylobacter* spp. that may not be recovered at 37°C can be facilitated and recovered at a 42°C incubation temperature. The study suggests that 37°C could have advantages over 42°C in providing an optimum environment for enhancing culturability of *C. coli* and the diverse *Campylobacter* spp. present in water, wastewater, and bird fecal dropping samples. This appears to be particularly the case for investigating the occurrence of *Campylobacter* in municipal wastewaters, where an incubation temperature of 42°C may not detect many *C. coli* isolates. From a human health perspective, the objective of many previous studies was focused mainly on detecting clinically important species such as *C. jejuni*, and therefore, the 42°C incubation temperature has been used widely for the recovery and isolation of thermophilic *Campylobacter* spp. However, the presence of *C. coli* and multiple types of *Campylobacter* spp. in human infections has been recognized and is seen as a significant epidemiological problem, since diagnostic laboratories may only isolate and characterize a single colony. This has led to some studies suggesting that to obtain a wide range of *Campylobacter* spp., it may be necessary to enrich water, food, and fecal samples at incubation temperatures of both 37 and 42°C (15, 49).

In conclusion, this large-scale long-term study shows that incubation temperatures of both 37 and 42°C influence the ability to recover thermophilic and other *Campylobacter* spp. from water, wastewater, and bird fecal droppings, especially if unusual fastidious campylobacters are believed to be a significant problem and possible threat to human health. It is recommended that for a

comprehensive characterization of the *Campylobacter* spp. in surface waters, wastewaters, or bird fecal droppings, incubation at both 37 and 42°C will likely be needed. Application of molecular methods would also be recommended to provide a more comprehensive characterization of campylobacters, including those that may occur in a viable but nonculturable condition. Further research on subtyping of thermophilic *Campylobacter* sp. isolates recovered from 37 and 42°C is needed in order to compare the strains recovered from water samples at these two temperatures. This subtyping might help in assessing strain diversity and in identifying the source of contamination of campylobacters in freshwater and bird fecal droppings.

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