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1. Introduction

Following emergence as an enteric pathogen in 1970s (Skirrow, 1977), Campylobacter has been a major concern worldwide. In the UK, Campylobacter is the most common bacterial cause of gastrointestinal infection recorded in the last two decades (Adak et al., 2005; Food Standards Agency, 2013). The total number of cases of Campylobacter infection during 2000–2012 was 781,581, from 1,052,581 laboratory confirmed cases of foodborne disease (Food Standards Agency, 2013). Campylobacteriosis is the most frequently reported foodborne disease but these figures belie actual unreported caseloads that are estimated to be 9 million and 1.3 million cases per year within the EU and USA, respectively (Centers for Disease Control and Prevention, CDC, 2014; EFSA, 2015).

The primary source of the major pathogenic species, C. jejuni and C. coli, are contaminated chicken and cattle meat (Adak et al., 2005; Suzuki and Yamamoto, 2009; Wilson et al., 2008), whereas less frequently they arise from wildlife (Hughes et al., 2009; Sippy et al., 2012), water, sewage and the environment (Jones, 2001; Waage et al., 1999). These bacteria are prevalent in offal, and in particular chicken liver (Cornelius et al., 2005, Kenar et al., 2009; Noormohamed and Fakhr, 2013; Strachan et al., 2012; Vashin et al., 2009; Whyte et al., 2006). Dishes such as liver paté and liver parfait have been reported as potential transmission vehicles for outbreaks of foodborne disease (Centers for Disease Control and Prevention, CDC, 2013; Edwards et al., 2014; Hope et al., 2014; Inns et al., 2010; O’Leary et al., 2009; Wensley and Coole, 2013) and the number of cases is increasing (Little et al., 2010). Moreover, their presence could pose a risk to animal welfare as Campylobacter species have been associated with a disease affecting poultry liver termed vibriobiotic hepatitis (Crawshaw et al., 2015; Jennings et al., 2011; Stephens et al., 1998).

In some cases, the occurrence of Campylobacter in liver may be the result of contamination from the intestinal contents during processing (Barot et al., 1983). Nonetheless, isolation from the internal tissue of
liver samples indicated that *Campylobacter* can be present in these organisms (Cox et al., 2007). It has been recognised that bacteria can cross the intestinal barrier of animals and humans, a process known as bacterial translocation. In general, the lymphatic pathway is perceived as the more convincing primary route of the translocation as compared with the venous system (Balzan et al., 2007). In vitro studies have demonstrated that *Campylobacter* can translocate using either transcellular passage through the enterocytes or paracellular routes via the tight junctions (Backert et al., 2013). Specific translocation mechanisms have been elucidated for enteric pathogens such as *Salmonella*, which uses several routes to pass through the intestinal barrier to inhabit systemic organs (Watson and Holden, 2010). However, further studies are required to obtain evidence of the translocation mechanisms operating for *Campylobacter* in humans and animals (Backert et al., 2013). For example, the capacity of *C. jejuni* to colonise particular tissues is affected by the organism’s ability to utilise specific nutrients—asparagine utilisation has been reported to improve the ability of the pathogen to colonise liver (Hofreuter et al., 2008).

Thorough cooking is the key to eliminating the risk of *Campylobacter* enteritis from poultry dishes. However, recipes for meats such as liver paté indicate minimal treatment to preserve the sensory properties and retain a pink appearance inside. To safely cook such dishes, critical core temperatures of 68–70 °C must be reached and held for periods as long as 45 min (Hutchison et al., 2015), which can result in unacceptable sensory characteristics (Whyte et al., 2006). Pre-cooking treatments could be applied to lower the initial contamination level, for example by freezing and washing of the liver using organic acid (Harrison et al., 2013; Hutchison et al., 2015). However, the use of organic acid was found to cause a colour change or bleaching of the liver surface, and may not be effective for *Campylobacter* naturally present within the internal structures of the liver.

Bacteriophages have gained recognition as therapeutic agents to control pathogens in livestock and poultry (reviewed by Johnson et al., 2008), and represent a potential approach to control *Campylobacter* in livers. *Campylobacter* bacteriophages can be isolated from chicken meat and chicken excreta (Atterbury et al., 2003, 2005; El-Shibiny et al., 2005; Loc Carrillo et al., 2007) but to date attempts to isolate *Campylobacter* phages from chicken liver have not been reported. The application of a single dose or mixtures of *Campylobacter* phages have been reported to be effective in reducing the intestinal colonisation of chickens by *C. jejuni* and *C. coli* (El-Shibiny et al., 2009; Kiltner et al., 2013; Loc Carrillo et al., 2005). The efficacy of the treatment varies depending on the phage type and dose, the phage-sensitivity of the host, the time interval post administration (Loc Carrillo et al., 2005) and the route of administration, i.e. by oral gavage or via chicken feed (Carvalho et al., 2010). Phage resistant *Campylobacter* have been reported post-treatment at relatively low frequencies of 2–4% (El-Shibiny et al., 2009; Hammel et al., 2014; Loc Carrillo et al., 2005).

In this study, *Campylobacter* and their phages were isolated from retail chicken liver. *Campylobacter* isolates were tested for their ability to re-colonise extra-intestinal organs of chickens in order to identify *Campylobacter* isolates able to inhabit the liver of broiler chickens. Finally, virulent bacteriophages were applied to *Campylobacter* contaminated chicken liver homogenates to provide proof of principle that bacteriophages can reduce *Campylobacter* contamination within the liver matrix.

2. Material and methods

2.1. Bacterial strains and bacteriophage

*Campylobacter jejuni* PT14 (Brathwaite et al., 2013) was used as a reference strain and also for phage isolation and propagation. *Campylobacter jejuni* HPCS (Loc Carrillo et al., 2005) and *C. jejuni* 81–170 (Korlath et al., 1985) were used as controls in the chicken colonisation experiments and the phage treatments of contaminated chicken livers. All *Campylobacter* isolates were cultured on blood agar base no. 2 CM0271 (Oxoid, Basingstoke, United Kingdom) supplemented with 5% defibrinated horse blood (TCS, Buckingham, United Kingdom) under microaerobic conditions (5% O2, 5% H2, 10% CO2, 80% N2) at 42 °C for 18–24 h. *Campylobacter* phages CP30A (GenBank accession number JX569801) and CPX (GenBank accession number JN132397) were propagated on *C. jejuni* PT14 or a contemporary *Campylobacter* isolate using the soft agar overlay method (Atterbury et al., 2003). Phages from the UK typing scheme (φ1 to φ16) were propagated as described by Frost et al. (1999). In order to obtain high titre stocks of bacteriophage, 30 ml volumes of plate lysates were centrifuged at 40,000g for 2 h at 4 °C. The pellets obtained were re-suspended in 1 ml of SM buffer (50 mM Tris·HCl (pH 7.5), 100 mM NaCl, 8 mM MgSO4, 0.01% Gelatin) to give a phage suspension containing approximately 10 log10 PFU/ml.

2.2. Preparation of chicken liver

Chicken liver samples were purchased from local supermarkets in Nottingham and Loughborough in the UK. Samples were kept at 4 °C and analysed before their expiry date as stated on the packaging. Each package contained 5–9 livers which were divided into two halves. Half of the liver was transferred into a stomacher bag (Seward Ltd., Worthing, UK) and 10 ml of Maximum Recovery Diluent CM733 (MRD; Oxoid; Basingstoke, UK) was added. The liver was gently massaged to re-suspend *Campylobacter* on the liver surface. To recover *Campylobacter* from internal tissues, the other half of liver was sterilised by dipping the liver into boiling water for 20–30 s (Whyte et al., 2006) and then tissue was excised with hot scalpel before being stomached with the addition of MRD (1:1 dilution ratio).

2.3. Isolation of *Campylobacter* from chicken liver

A 4 ml aliquot of suspension from the liver surface sample or the stomached internal tissue was transferred into 4 ml of enrichment media. This consisted of 2× *Campylobacter* Enrichment Broth Lab135 (Lab M, Heywood, UK) made up with the addition of: 10% lyed horse blood (TCS), 0.25 g/l each of sodium pyruvate, sodium metabisulphite and ferrous sulphate (each from Sigma Aldrich, Poole, UK) and *Campylobacter* Enrichment Selectivat CM 739 (Mast, Bootle UK), in a bijou bottle. The total volume of 8 ml resulted in limited airspace in the bottle, hence maintaining microaerobic conditions during incubation at 37 °C for 48 h. Five 10 μl aliquots from each bijou were dispensed onto mCCDA CM739 agar (Oxoid) prepared with the addition of mCCDA CM739 agar (Oxoid) prepared with the addition of *Campylobacter* selective supplement code (SR155, Oxoid) and additional Agar No. 1 (Oxoid) added to give 2% and then incubated at 42 °C for 48 h under microaerobic conditions. *Campylobacter* were confirmed after subculture, using microscopic observation of Gram stained cells, together with catalase and oxidase tests.

2.4. Enumeration of *Campylobacter*

*Campylobacter* was enumerated using the Miles and Misra technique, with serial dilutions prepared in MRD and 10 μl aliquots spotted in quintuplicate on 2% mCCDA before incubating under microaerobic conditions at 42 °C for 48 h. Typical *Campylobacter* colonies were counted and the total number calculated as either log10 CFU/g for internal tissue samples or log10 CFU/cm² for surface liver samples.

2.5. Species identification and Fla-typing using PCR methodologies

*Campylobacter* DNA was isolated using the GenElute™ Bacterial Genomic DNA Kit according to manufacturer’s instructions for Gram negative bacteria (Sigma-Aldrich, UK). The PCR methodology was based on conditions previously described by Linton et al. (1997) for species identification and by Elvers et al. (2008) for FlaA SVR-typing. The oligonucleotides were purchased from Eurofins (Ebersberg, Germany) and consisted
of the primers HIP400F (5′-GAAGAGGTCTGGAGTGTTG-3′) and HPI134R (5′-ACCTGGTGACGTTTACATGG-3′) targeting the C. jejuni hippuricase gene, CC18F (5′-GTTTGCTTACACGAGGAG-3′) and CC51R (5′-ATAAAAAAGCCTAGTCGCGTG-3′) specific for the C. coli aspartokinase gene, and FLA4R (5′-GGATTTCTTCAAAACATATGGTCG-3′) and FLA625R (5′-CAAGGTTTCAATATAGGGAAG-3′) for the Campylobacter flaA gene. To determine the fla type of the Campylobacter isolates, the PCR products were purified using the Wizard® SV Gel and PCR Clean-Up System (Promega, Southampton, UK) and the DNA sequences were sequenced using the Eurofins MWG Value Read service.

2.6. Isolation and characterisation of bacteriophages from chicken liver

Bacterial lawns were prepared, plaque purified and the plaque forming units per ml (PFU/ml) were determined as previously described (Atterbury et al., 2003). The bacteriophages were diluted to the routine test dilution of approximately 6 log_{10} PFU/ml. Ten microliters of phage suspensions were dispensed onto the surface of the lawns of the test Campylobacter strain and then incubated under microaerobic conditions at 42 °C for 18–24 h. The lysis profiles of the isolates produced by each phage were scored according to the protocol described by Frost et al. (1999) for the UK phage typing scheme. Phage genomic DNAs were prepared as previously described (Loc Carrillo et al., 2007). PCR amplification of Campylobacter phage DNAs was performed using group III-specific primers CPB353B (5′-CGGGTTATACCGGATATAG-3′) and CPB354B (5′-ATATAGGGCTTCTGGAATG-3′), the amplification products of which can discriminate CP30A and CP8-like bacteriophages (Siringan et al., 2014).

2.7. Colonisation of chickens with Campylobacter

Procedures for the chicken colonisation experiments were carried out as previously described (Loc Carrillo et al., 2005). For each Campylobacter isolate, a suspension of 7 log_{10} CFU was administered by oral gavage to seven 16-day-old broiler chickens (male Ross 308) reared under strict biosecure conditions. The birds were killed after 7 days, and the caecal content, liver, spleen, heart, kidney and breast meat were examined for the presence of Campylobacter by direct plating on mCCDA and by enrichment as described in Section 2.3. These animal studies were conducted under the Animals Scientific Procedures Act (1986) and were approved by the University of Nottingham local ethical review committee.

1.1. Recovery and survival of C. jejuni in fresh and frozen chicken liver during storage

Campylobacter-free chicken livers were harvested from Campylobacter-negative broiler chickens reared under strict biosecure conditions. Fresh chicken livers were divided into sections weighing approximately 10 g. Each section was placed into a stomacher bag and weighed. The liver was then inoculated with 5 ml C. jejuni suspension containing 3 or 7 log_{10} CFU/ml, and replaced with sterile water for negative controls. The samples were stomached and stored at 4 °C for 72 h. Aliquots of 200 μl were taken for Campylobacter enumeration at 0, 8, 24, 32, 48, 56, and 72 h time intervals over the period. Frozen chicken liver was defrosted for 18 h at 4 °C prior to inoculation and storage at 4 °C or −20 °C. The subsequent steps followed the same protocol as fresh liver samples but at daily intervals over 7 days. Three independent replicate experiments were performed with fresh and frozen livers.

2.8. Phage treatment of Campylobacter contaminated chicken livers

Campylobacter-free chicken livers (10 g) were stomached before the addition of C. jejuni suspensions to inoculum densities of approximately 3 log_{10} CFU/g (low inoculum) or 5 log_{10} CFU/g (high inoculum). The liver stomachates containing C. jejuni were treated with either a phage suspension at 8 log_{10} PFU/ml or with an equivalent volume of SM buffer (mock treatment). Campylobacters were enumerated and the phage titres as indicated above, following incubation at 4 °C over 48 h. All experiments were performed in triplicate.

2.9. Statistical treatment of data

Statistical differences between paired control and treatment groups (using log_{10}-transformed Campylobacter counts) were assessed by using the Student’s t-test with significance p < 0.05. Differences between experimental groups were analysed by analysis of variance.

3. Results

3.1. Prevalence of Campylobacter in retail chicken liver

A total of 109 samples of retail chicken liver were analysed for the presence of Campylobacter recoverable from surface or internal tissues. Isolation was performed on 7 different batches within a 2 month period. There was a high prevalence of Campylobacter with 87.2% and 82.6% of samples positive from surface and inner tissues respectively (Table 1). Most samples contained low numbers of Campylobacter that were only recoverable by enrichment. Samples that could be enumerated contained Campylobacter in the range of 1.8–3.8 log_{10} CFU/cm² for surface samples and 3.0–3.8 log_{10} CFU/g for internal tissue samples. Three surface samples and 5 internal tissue samples contained Campylobacter ≥3 log_{10} CFU/g, which would be considered to pose a significant risk to consumers (Food Safety Agency UK 2014).

3.2. Frequency and characteristics of Campylobacter phages in retail chicken liver

Three Campylobacter-specific bacteriophages were isolated from 109 retail chicken livers (2.7%). One of the phage originated from a surface sample (CLP6), while the other two were from the internal tissues of the livers (CLP47 and CLP63). Phages CLP47 and CLP63 exhibited similar lytic abilities against the C. jejuni liver isolates (64%), whilst phage CLP6 was virulent against more of the C. jejuni isolates (88%). However, the host ranges of the three new liver isolates were more specific than phages CP30A and CPX previously isolated from chicken intestinal contents or chicken meat respectively. None of the phage isolated from chicken liver infected the C. coli isolates.

Campylobacter bacteriophages possess double-stranded DNA genomes that are classified into three groups according to their genome size and head diameter, i.e. group I with genome sizes of 320 kb and head diameters of 140.6 and 143.8 nm; group II, with genome sizes of 184 kb and head diameters of 99 nm; and group III with genome sizes of 138 kb and head sizes of 100 nm (Sails et al., 1998). PFGE analysis of bacteriophage genomic DNA revealed that the three phages isolated from chicken liver were approximately 140 kb in size, which is typical of group III bacteriophages and similar to the reference phages CP30A and CPX. PCR amplification of the phage DNAs with group III-specific primers confirmed the classification of the liver phages.

<table>
<thead>
<tr>
<th>Table 1</th>
<th>Prevalence and concentration of Campylobacter in retail chicken liver.</th>
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<tbody>
<tr>
<td>Sample</td>
<td>Samples containing Campylobacter</td>
</tr>
<tr>
<td>Liver surface</td>
<td>95/109 (87.2%)</td>
</tr>
<tr>
<td>Internal tissue</td>
<td>90/109 (82.6%)</td>
</tr>
</tbody>
</table>
3.3. Characterisation of Campylobacter isolates

A combination of Fla-typing and phage typing was used to discriminate the Campylobacter isolates, which enabled them to be placed into five groups that are summarised in Table 2. The C. coli isolates represent a single Fla-type that could not be distinguished with the phage used in this study. C. jejuni isolates could be placed in four groups where concordance was observed between the Fla-types and the phage sensitivity profiles. One group, represented by isolate CLB104, were recovered exclusively from the internal tissues of retail chicken liver with counts ≥3 log\textsubscript{10} CFU/g, and therefore represents a significant risk to human health. C. jejuni isolates CLB44, CLB68 and CLB104 that originated from the internal tissues of retail chicken were selected for further study.

1.2. Persistence of Campylobacter liver isolates in addition to the liver isolates (control strains) representing either low (3 log\textsubscript{10} CFU/g) or high (7 log\textsubscript{10} CFU/g) target inoculums. Error bars represent the standard deviations for n = 3.

3.4. Phage treatment of Campylobacter contaminated liver

As noted above the bacteriophages isolated from liver have a restricted host range amongst the Campylobacter liver isolates compared to those of chicken intestinal (CP30A) or chicken meat (CPX) origin or the typing phages φ3 and φ15 (Table 2). Phages φ3 and φ15 have tailed morphologies and are classified as group II based on their genome sizes of 180 and 190 kb (Sails et al., 1998). However, φ3 and φ15 were able to lyse 3 of the 5 groups of Campylobacter liver isolates in addition to the control C. jejuni strains HPC5 (original source chicken intestine) and 81–176 (original source human with campylobacteriosis), and were therefore selected for phage therapy (biosanitization) applications with chicken liver to enable comparisons of the effect between C. jejuni strains. Phages φ3 or φ15 were added at 8 log\textsubscript{10} PFU/g to liver stomachates containing either low (3 log\textsubscript{10} CFU/g) or high (5 log\textsubscript{10} CFU/g) target Campylobacter inoculums and stored at 4 °C over 48 h. Fig. 2 presents the viable counts of five C. jejuni strains following either mock or phage treatments of chicken liver suspensions. All the phage treated C. jejuni strains showed a significant reduction in the viable count compared to the control for low and high inoculums (p < 0.05). However, the reductions observed were modest. For example, the reductions in viable count recorded for the high risk livers represented by the high inoculum series of 5 log\textsubscript{10} CFU/g ranged between 0.7 log\textsubscript{10} CFU/g for the chicken liver isolate CLB68 treated with φ15 and 0.2 log\textsubscript{10} CFU/g for the chicken intestinal isolate HPC5 treated with either φ3 or φ15. The phage recovered from these experiments showed minimal variation in titre and showed no significant difference to the initial inoculum titre (p > 0.05).
3.5. Capability of Campylobacter isolates to colonise broiler chickens

Cell suspensions of five C. jejuni cultures in physiological phosphate buffered saline (approximately 7 log_{10} CFU/ml) were administered orally, to 6 or 7 broiler chickens and colonisation was established after 7 days by examining post-mortem Campylobacter counts from chicken caecal contents and from extra-intestinal organs, i.e. liver, heart, spleen, breast muscle and kidney. There was no observable pathology for any of the organs. All chickens contained high counts of Campylobacter in their caecal contents (>7 log_{10} CFU/g) based on enumeration on mCCDA plates (Table 3). However, Campylobacters could only be recovered from the extra-intestinal organs of chickens colonised by the liver isolates, and only by enrichment. C. jejuni isolate CLB104 was detected in the liver and kidney 3 of 7 chickens, while being recovered from all of the extra-intestinal organs of one bird. No Campylobacters were recovered from the excised breast meat of any chicken. No Campylobacters were recovered from the extra-intestinal organs of the control C. jejuni strains HPC5 or 81–176.

4. Discussion

Campylobacter was found in the majority of retail chicken liver samples at varying levels of contamination. Of concern are chicken meat samples containing >3.0 log CFU/g, which pose a disproportionately high risk to consumer health (Food Standards Agency, 2015). However, we recorded Campylobacter counts >3.0 log CFU/g for 2.8% of the surface and 4.6% of the internal tissue samples from retail chicken livers. A compilation of findings presented here with those available in the literature, are presented in Table 4, which demonstrates that Campylobacter contamination of livestock liver is prevalent with surveys recording that 66–100% of the samples tested were positive. In the majority of cases the livers showed a low level contamination, for example, Cornelius et al. (2005) and Whyte et al. (2006) found that 83–88% of internal tissues of livers harboured 10^8 MPN (most probable number) per g, while the remaining samples contained 10^2–10^3 MPN/g. A clear dose–response relationship between consumption of chicken liver paté and the risk of infection with Campylobacter has been demonstrated (Edwards et al., 2014). A low level of contamination does not eliminate the risk of Campylobacter infection since the infective dose can be as low as 500 cells (Robinson, 1981).

Whilst Campylobacters are frequently reported from liver, this is the first study to report the isolation of Campylobacter-specific bacteriophage. The isolation frequency was comparatively low at 2.7% but the phages recovered were generally able to infect the Campylobacters recovered from liver suggesting they are replicating in the source tissues. This would offer the prospect that phage therapy could be applied to control Campylobacters in vivo or on retail liver. The application of Campylobacter-specific bacteriophages has been demonstrated to successfully reduce contamination levels in chicken skin and meat (Atterbury et al., 2003, Bigwood et al., 2008; Goode et al., 2003). Similarly the application of phages ɸ3 and ɸ15 to chicken liver stomachates containing C. jejuni resulted in significant reductions in the viable counts of all five strains tested. However, the reductions observed post phage treatment in this study are unlikely to have a universal impact on the risk imparted by the consumption of chicken liver. As discussed above meat containing >3 log_{10} CFU/g represents a disproportionate risk, and viable count reductions in the range of 0.2 to 0.7 log_{10} CFU/g for the high level contamination series of 5 log_{10} CFU/g would not be sufficient to reduce the risk of infection. Whereas reductions of 0.7 log_{10} CFU/g in the viable count for lower levels of contamination, as demonstrated in the 3 log_{10} CFU/g series experiments in this study, could be of benefit. For the application to be of general use the levels of pathogen reduction need to be uniformly at the higher levels observed here, and the application would have to be on the liver before any processing for cooking and consumption. Bacteriophages in general have gained support for food sanitisation applications since bacteriophage capable of lysing the foodborne pathogens Listeria monocytogenes or Salmonella have been approved in the USA (US Food and Drug Administration and the Food Safety Inspection Service of the US Department of Agriculture) for use on retail food products.

The genome sizes of the chicken liver bacteriophages were estimated to be 140 kb using PFGE, which places them as group III Campylobacter bacteriophages (Loc Carrillo et al., 2007; Sails et al., 1998). Recently a new sub-family of the T4-like phage super family, the Eucamphyvirinae, has been proposed for Campylobacter bacteriophages based on their genomic DNA sequences/sizes and particle morphologies (Javed et al., 2014). Group III bacteriophages

<table>
<thead>
<tr>
<th>Campylobacter isolates</th>
<th>Caecal content (log_{10} CFU/g)</th>
<th>Recovery by enrichment</th>
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<tbody>
<tr>
<td></td>
<td>Liver</td>
<td>Heart</td>
</tr>
<tr>
<td>CLB44</td>
<td>7.7 ± 0.82</td>
<td>2/6 (33%)</td>
</tr>
<tr>
<td>CLB68</td>
<td>8.0 ± 0.37</td>
<td>1/7 (14%)</td>
</tr>
<tr>
<td>CLB104</td>
<td>7.5 ± 0.43</td>
<td>3/7 (43%)</td>
</tr>
<tr>
<td>HPC5</td>
<td>7.4 ± 0.70</td>
<td>0/7 (0%)</td>
</tr>
<tr>
<td>81–176</td>
<td>7.2 ± 0.47</td>
<td>0/7 (0%)</td>
</tr>
</tbody>
</table>
constitute the genus Cp8unilikeviruses with genome sizes in the range of 130–140 kb. The typing phages φ3 and φ15 used for phage therapy in this study are group II but also fall within the €ucampyvirinae as members of the genus CP220likeviruses. CP220likeviruses and Cp8unilikeviruses have been used successfully for active phage therapy in chickens against Campylobacters (El-Shibiny et al., 2009; Loc Carrillo et al., 2005; Hammerl et al., 2014; Scott et al., 2007), where there are sufficient densities of host bacteria to support phage replication (Cairns et al., 2009). Below the phage proliferation threshold requires that the bacteriophage encounter, adsorb and penetrate the target bacteria - a process that has an intrinsic requirement for high phage titres. It is likely that some phages are better suited to this purpose in terms of achieving high titres, maintaining stability and retaining activity. In this application the phage titres applied to chicken liver would have to remain high at retail and post disruption of the liver when internalised bac-

### References


