

# Evaluation of malacosporean life cycles through transmission studies

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**ABSTRACT:** Myxozoans, belonging to the recently described Class Malacosporea, parasitise freshwater bryozoans during at least part of their life cycle, but no complete malacosporean life cycle is known to date. One of the 2 described malacosporeans is *Tetracapsuloides bryosalmonae*, the causative agent of salmonid proliferative kidney disease. The other is *Buddenbrockia plumatellae*, so far only found in freshwater bryozoans. Our investigations evaluated malacosporean life cycles, focusing on transmission from fish to bryozoan and from bryozoan to bryozoan. We exposed bryozoans to possible infection from: stages of *T. bryosalmonae* in fish kidney and released in fish urine; spores of *T. bryosalmonae* that had developed in bryozoan hosts; and spores and sac stages of *B. plumatellae* that had developed in bryozoans. Infections were never observed by microscopic examination of post-exposure, cultured bryozoans and none were detected by PCR after culture. Our consistent negative results are compelling: trials incorporated a broad range of parasite stages and potential hosts, and failure of transmission across trials cannot be ascribed to low spore concentrations or immature infective stages. The absence of evidence for bryozoan to bryozoan transmissions for both malacosporeans strongly indicates that such transmission is precluded in malacosporean life cycles. Overall, our results imply that there may be another malacosporean host which remains unidentified, although transmission from fish to bryozoans requires further investigation. However, the highly clonal life history of freshwater bryozoans is likely to allow both long-term persistence and spread of infection within bryozoan populations, precluding the requirement for regular transmission from an alternate host.

**KEY WORDS:** Myxozoa · Malacosporea · Proliferative kidney disease · Freshwater bryozoans · Transmission · Potential hosts · Life cycles

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## INTRODUCTION

The Phylum Myxozoa is comprised of about 55 genera and over 1300 species (Canning & Okamura 2004) primarily parasitic in marine and freshwater fish. The phylum contains 2 classes: the Myxosporea (Bütschli 1881) and the Malacosporea (Canning et al. 2000, Kent et al. 2001). The former includes the majority of species, some of which cause economically important diseases, such as *Myxobolus cerebralis*, the causative agent of salmonid whirling disease (Bartholomew & Reno 2002). Only some 25 myxosporean life cycles are

known to date, and these involve annelid worms and fish as primary and secondary hosts (Okamura & Canning 2003). Direct transmission has been demonstrated between fish (Diamant 1997, Redondo et al. 2002, Yasuda et al. 2002), but whether this process precludes the facultative incorporation of invertebrate hosts and represents a reduction in life cycle complexity is unknown (Canning & Okamura 2004). Malacosporeans parasitise freshwater bryozoans (Bryozoa: Phylactolaemata) during at least part of their life cycle, but no complete malacosporean life cycle is known.

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Two species of malacosporean have been described so far. One is *Buddenbrockia plumatellae* (formerly *Tetracapsula bryozoides*; Canning et al. 2002), which occurs as both sac-like and vermiform stages in freshwater bryozoans (Monteiro et al. 2002, Okamura et al. 2002). The other member of the Class Malacosporea is *Tetracapsuloides bryosalmonae* (formerly *Tetracapsula bryosalmonae*; Canning et al. 2002). Only sac-like stages of *T. bryosalmonae* have been encountered in bryozoan hosts to date. *T. bryosalmonae* is the causative agent of salmonid proliferative kidney disease (PKD) (Anderson et al. 1999, Canning et al. 1999, Feist et al. 2001) and was, until recently, referred to as the PKX organism (Seagrave et al. 1980). The identification of bryozoans as hosts of *T. bryosalmonae* (Anderson et al. 1999) represented a breakthrough, as it finally determined the source of a disease which causes significant financial losses to fish farms and hatcheries of Europe and North America (Hedrick et al. 1993, Bromage 1999). PKD has also been implicated in the decline of wild salmonid populations in Switzerland (Wahli et al. 2002).

Freshwater bryozoans are sessile, colonial invertebrates that are common in both lotic and lentic habitats. They undergo prolific colony growth in summer and produce dormant, seed-like propagules (statoblasts) that overwinter and hatch into small colonies when favourable conditions return (Okamura & Hutton-Ellis 1995). The freshwater bryozoan *Fredericella sultana* can also overwinter as colonies (Raddum & Johnsen 1983, Gay et al. 2001). Malacosporeans develop into freely circulating sacs and active vermiform stages (Canning et al. 1996, 1999, 2002), which can be observed within the body cavity of their bryozoan hosts by inspection using a dissection microscope. Infective spores develop within sacs and worms. There is relatively little host specificity. Molecular data have identified *Tetracapsuloides bryosalmonae* infections in 5 bryozoan species (Anderson et al. 1999, Longshaw et al. 1999, Okamura & Wood 2002), 2 of which have been confirmed as hosts through transmission studies (Feist et al. 2001, Gay et al. 2001). The vermiform stage of *Buddenbrockia plumatellae* has been found in 3 species (Canning et al. 2002, Monteiro et al. 2002), but the sac-like stages have only been found parasitising 1 bryozoan host—*Cristatella mucedo* (Okamura 1996, Canning et al. 2002).

Knowledge of the ecology, development and life cycles of malacosporeans is limited. Transmission studies from bryozoans to fish have confirmed bryozoans as true hosts of *Tetracapsuloides bryosalmonae* (Feist et al. 2001, Gay et al. 2001), but whether stages that develop in fish are capable of infecting new hosts has so far not been demonstrated. It is, however, known that PKD is not transmitted from fish to fish (Ferguson & Ball

1979, D'Silva et al. 1984). In addition, the presence of *T. bryosalmonae* in bryozoan populations from sites devoid of salmonids (Okamura et al. 2001) suggests that these fish are, at best, facultative hosts. D. C. Morris et al. (2002a) and D. J. Morris et al. (2002) reported possible transmission of *T. bryosalmonae* and *Buddenbrockia plumatellae* from fish to bryozoans. However, these results may have reflected previous development of infection in bryozoans in the field, or PCR amplification of residual DNA that had adhered to bryozoans or of DNA from undetected, infected invertebrates.

Investigations into the pathogenesis of PKD have shown that *Tetracapsuloides bryosalmonae* has 2 developmental stages in salmonids (Hedrick et al. 1993). The extrasporogonic stage, found in the blood and kidney interstitium of infected fish, produces the chronic inflammatory response which is characteristic of the disease (Kent & Hedrick 1986). The sporogonic stage is found in the kidney tubule lumen 2 to 3 wk after the extrasporogonic stages are first observed. Sporogonic stages can persist in the kidney long after the recovery of the fish from clinical disease (Kent & Hedrick 1986, Morris et al. 2000). Development of sporogonic stages in at least some fish hosts proceeds to the production of malacosporean-like spores containing 2 capsulogenic cells and 1 sporoplasm surrounded by 2 unstrengthened (soft) valve cells (Kent & Hedrick 1986, Hedrick et al. 2004).

Certain observations have been cited as evidence that salmonids may be aberrant hosts for *Tetracapsuloides bryosalmonae*, including the severe inflammatory response of fish and incomplete spore development (Kent & Hedrick 1986). However, before description of the Class Malacosporea (Canning et al. 2000), researchers expected final spore development to terminate in stages with typical hardened valves consistent with those observed in other myxosporeans. The possibility that fish should be considered as potential hosts is supported by the release of malacosporean-like spores in fish urine, as similar and functional spores of *T. bryosalmonae* develop in bryozoan hosts (Hedrick et al. 2004).

In view of our incomplete understanding of malacosporean life cycles, we undertook a series of studies to investigate potential routes of transmission of *Tetracapsuloides bryosalmonae* and *Buddenbrockia plumatellae*. In particular we focused on 2 unresolved issues: whether transmission can occur from fish to bryozoan and from bryozoan to bryozoan.

## MATERIALS AND METHODS

We undertook the following types of transmission: (1) Fish to bryozoan: exposure of bryozoans to

*Tetracapsuloides bryosalmonae* stages in fish kidney; (2) fish to bryozoan: cohabitation of bryozoans with fish infected with *T. bryosalmonae*; (3) bryozoan to bryozoan: exposure of bryozoans to *T. bryosalmonae*-infected *Fredericella sultana*; (4) bryozoan to bryozoan: exposure of bryozoans to *Buddenbrockia plumatellae*-infected *Plumatella fungosa* and *Cristatella mucedo*.

Below we describe the material used for transmission studies and the protocols for each type of transmission. Details of the individual trials are summarised in Table 1.

**Bryozoans used as targets for infection.** Three species of freshwater bryozoan (*Fredericella sultana*, *Plumatella fungosa* and *Plumatella emarginata*) were cultured in aquarium tank systems composed of smaller aquaria, housing the bryozoans, connected by recirculation to separate, larger aquarium tanks containing goldfish, according to the methods of Wood (1971) in conditions of constant light and temperature (20°C). The tank systems were topped up with deionised water every 2 to 3 d when required and periodically seeded with water from a small, concrete-lined pond devoid of both bryozoans and fish with PKD. The incorporation of goldfish and periodic seeding with pond-water creates suitable environmental conditions and food, allowing bryozoans to be maintained indefinitely in culture. Bryozoans in such culture systems have been grown for many years (T. Wood pers. comm., S. Tops & B. Okamura pers. obs.) with no sign of malacosporean infections being transferred between bryozoans and goldfish. This cultured bryozoan material provided potential target hosts for infection in our transmission studies and was regarded as infection-free, as described below. The material was maintained in independent culture systems and was thus never in contact with infected bryozoan material collected from the field for use in transmission studies (described later).

***Fredericella sultana*:** Transmission trials were performed using *F. sultana* material from several different derivations. The original material was collected from the Kennet and Avon Canal (51°24'N, 01°08'W) in October 2000. Colonies were induced to attach to surfaces of water-filled petri dishes or to submerged plastic discs (5.4 cm diameter) via their sticky, newly formed tips (Wood 1973, S. Tops pers. obs.). Subcultured material was derived by removing branches from original material established in culture, and inducing the branches to attach to new surfaces. Statoblast-hatched material was obtained from statoblasts produced by subcultured *F. sultana* material. The statoblasts were placed in deionised water at 4°C and kept in the dark for a minimum period of 4 mo. They were then transferred to Petri dishes containing artificial pond water (Wood 1996) to induce hatching under

ambient laboratory conditions (exposure to normal daylight period, 20°C). The small colonies that emerged from statoblasts were grown in the culture system to provide material for trials. In keeping with the absence of *Tetracapsuloides bryosalmonae* during long-term monitoring of the source population (Tops & Okamura 2003), infections were never observed in any *F. sultana* material in the culture system. In addition, PCR using *T. bryosalmonae*- and *Buddenbrockia plumatellae*-specific primers (see below) on portions of *F. sultana* colonies used as source material in transmission trials consistently provided negative results. This approach assumes that *T. bryosalmonae* is distributed throughout a colony and hence should be detected by screening portions of colonies by PCR, but there are no data to corroborate this assumption. Source material was only used during Trials 1a and 1b. For the remainder of the trials, statoblast-hatched *F. sultana* colonies were employed.

***Plumatella fungosa*:** The source of all *P. fungosa* colonies was a single colony collected from Blenheim Palace Lake (51°51'N, 01°23'W) in July 2000. The material used in the transmission trials was the result of several generations of statoblast-derived colonies produced from this original colony, as described for *Fredericella sultana* above. No malacosporeans have ever been observed in this *P. fungosa* material in culture; however, as a further check that the material was uninfected, small portions of live bryozoans were cut from the stock material and subjected to PCR using *Tetracapsuloides bryosalmonae*- and *Buddenbrockia plumatellae*-specific primers (see below). Since none of these were positive for malacosporean infection, the material was deemed to be suitable for transmission studies. Severed branches were induced to attach to surfaces for transmission studies as described above.

***Plumatella emarginata*:** *P. emarginata* colonies collected from Burghfield Lake, Berkshire (51°26'N, 01°04'W) in August 2002 released larvae in the laboratory. The larvae metamorphosed into small colonies on plastic discs which were placed in culture for onward growth, thus providing bryozoan colonies of larval origin for transmission studies. As a check that the material was uninfected by malacosporeans, small portions of colonies (n = 6 colonies) were subject to PCR using *Tetracapsuloides bryosalmonae*- and *Buddenbrockia plumatellae*-specific primers (see below). Negative PCR results provided evidence that the material was suitable for transmission studies.

**Sources of potential infection.** Fish used for exposing bryozoans to infected kidney included rainbow trout *Oncorhynchus mykiss* obtained from a fish farm on the Kennet and Avon Canal, Berkshire, UK (Table 1: Trial 1a), wild brown trout *Salmo trutta* collected during routine survey work of trout populations

conducted by the Environment Agency on the River Test, Hampshire, UK (Table 1: Trial 1b), and rainbow trout from a hatchery on the American River, California, USA (Table 1: Trial 1c). Fish used in cohabitation studies were naturally infected with *Tetracapsuloides*

*bryosalmonae* and included rainbow trout from the American River Hatchery, California (Table 1: Trial 2b), and Chinook salmon *Oncorhynchus tshawytscha* from the Merced River Hatchery, California (Table 1: Trials 2a and 2c). Brown trout from Mount Shasta

Table 1. Summary of transmission trials, including identity of host and target species used in each trial, details of transmission including the exposure time of bryozoans to potential infection, the number (No.) of experimental (Exp) and control (Ctl) bryozoan colonies used in the trials (and subsequently subject to PCR), and the post-exposure culture period. Trials 1a to 3d involved transmission with *Tetracapsuloides bryosalmonae*; Trials 4a and 4b involved transmission with *Buddenbrockia plumatellae*

Trial (Date)	Host species	Target species	Transmission details	No. experimental and control colonies in trial			Culture period
				<i>Fredericella sultana</i>	<i>Plumatella fungosa</i>	<i>Plumatella emarginata</i>	
1a (Oct 2000)	<i>Oncorhynchus mykiss</i>	<i>Fredericella sultana</i>	Exposure to macerated kidney for 24 h	Exp: 6 (0) Ctl: 7 (0)			54 d
1b (Jan 2001)	<i>Salmo trutta</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Exposure to macerated kidney for 24 h	Exp: 4 (4) Ctl: 4 (0)	Exp: 10 (10) Ctl: 10 (10)		37 d
1c (Jul 2002)	<i>Oncorhynchus mykiss</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Exposure to macerated kidney for 8 h	Exp: 6 (4)	Exp: 1 (1)		14 d
2a (Apr 2002)	<i>Oncorhynchus tshawytscha</i>	<i>Fredericella sultana</i>	Cohabitation with fish for 80 h	Exp: 20 (9) Ctl: 20 (14)			22 d
2b (Apr 2002)	<i>Oncorhynchus mykiss</i>	<i>Fredericella sultana</i>	Cohabitation with fish for 80 h	Exp: 20 (12)			22 d
2c (Jul 2002)	<i>Oncorhynchus tshawytscha</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Cohabitation with fish for 19.5 h	Exp: 10 (8) Ctl: 6 (6)	Exp: 1 (0) Ctl: 1 (1)		14 d
2d (Jul 2002)	<i>Salmo trutta</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Cohabitation with fish for 19.5 h	Exp: 10 (9)	Exp: 1 (0)		14 d
3a (Jun 2002)	<i>Fredericella sultana</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Spores directed into lophophores (4 h); retention in spore suspension (20 h)	Exp: 5 (5)	Exp: 2 (2)		30 d
3b (Oct 2002)	<i>Fredericella sultana</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i> <i>Plumatella emarginata</i>	Spores directed into lophophores (4 h); retention of spore suspension (16 h)	Exp: 3 (3)	Exp: 1 (1)	Exp: 3 (3)	33 d
3c (Jun 2002)	<i>Fredericella sultana</i>	<i>Fredericella sultana</i>	Exposure to shredded bryozoans for 135 min	Exp: 3 (2) Ctl: 2 (2)			30 d
3d (Aug 2003)	<i>Fredericella sultana</i>	<i>Plumatella fungosa</i>	Exposure to naturally released spores for 21 d		Exp: 10 (10)		21 d
4a (Aug 2001)	<i>Plumatella fungosa</i>	<i>Plumatella fungosa</i>	Spores directed into lophophores (4 h); retention in spore suspension (12 h)		Exp: 2 (0) Ctl: 2 (0)		52 d
4b(i) (Aug 2001)	<i>Cristatella mucedo</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Spores directed into lophophores (4 h); retention in spore suspension (20 h)	Exp: 5 (4) Ctl: 5 (5)	Exp: 5 (4)		60 d
4b(ii) (Aug 2001)	<i>Cristatella mucedo</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Exposure to immature sacs for 24 h	Exp: 2 (2)	Exp: 3 (2)		60 d
4b(iii) (Aug 2001)	<i>Cristatella mucedo</i>	<i>Fredericella sultana</i> <i>Plumatella fungosa</i>	Spores directed into lophophores (4 h); retention in spore suspension (20 h)	Exp: 2 (1)	Exp: 1 (1)		60 d

Hatchery, California, were hatched from eggs and maintained in well water at 15°C until intraperitoneally injected as in previous studies (Feist et al. 2001) with homogenised kidneys of Chinook salmon infected with extrasporogonic and sporogonic stages of *T. bryosalmonae* 6 wk prior to the transmission trials (Table 1: Trial 2d).

Colonies of *Fredericella sultana* infected with *Tetracapsuloides bryosalmonae* collected from the River Cerne, Dorset, UK, provided material for bryozoan to bryozoan transmission (Table 1: Trials 3a to 3d). The infection of a *Plumatella fungosa* colony by vermiform stages of *Buddenbrockia plumatellae* during a field experiment (Tops & Okamura 2003) provided material for Trial 4a (Table 1), while infected colonies of *Cristatella mucedo* from Bonnerweiher, Bavaria, Germany, provided sac-like stages of *B. plumatellae* for Trial 4b(i to iii) (Table 1).

**Transmissions.** Table 1 summarises details of the following transmission trials and should be referred to for information on the number of experimental and control bryozoan colonies used in trials. Note that PCR of portions of experimental colonies prior to exposure in trials served as an additional control unless otherwise specified. As previously mentioned, this method assumes that *Tetracapsuloides bryosalmonae* is distributed throughout a colony and hence can be detected by screening portions of colonies by PCR. Other controls are described below for individual trials.

**(1) Fish to bryozoan. Exposure of bryozoans to infected fish kidney:** Fish were collected from sites enzootic for PKD (Trials 1a to 1c; Table 1) and transported to the laboratory. They were killed by a blow to the head, followed by severance of the spinal cord. For Trials 1a and 1b, approximately 27 mm<sup>3</sup> of tissue was dissected from the posterior portion of the kidney and imprints were made for histological analysis to confirm infection by *Tetracapsuloides bryosalmonae*. For Trial 1b, kidney tissue was fixed in absolute ethanol and stored prior to PCR at -20°C. For Trial 1c, wet mounts of the kidney material were prepared and examined according to Kent & Hedrick (1986) to confirm the presence of sporogonic stages of *T. bryosalmonae* in kidney tubules.

For Trials 1a and 1b, the remaining kidney tissue was removed and placed in a minimal amount of 1 M phosphate buffered saline (PBS) (Oxoid™ tablets; pH 7.3) on ice. Kidney material from all fish collected for each Trial was combined in a dish with a minimal amount of PBS, and disrupted using forceps. This macerated kidney material was added to an aerated tank containing colonies of *Fredericella sultana* and 16 l of water from the bryozoan culture system. After 24 h the bryozoans were transferred to the culture system

(unless otherwise specified, see below) and were then visually monitored every 2 to 3 d, using a dissection microscope, for signs of infection. For Trial 1c, the kidney tissue was disrupted by passing it through a 150 µm sieve with a minimal amount of PBS. Kidney homogenates were then combined as above and added to an aerated tank containing colonies of *F. sultana* and *Plumatella fungosa*, and 9.5 l of well water. After 8 h the bryozoans were transferred to aquaria with water from a local pond at 15°C and visually monitored for signs of infection every 2 to 3 d. Controls for Trial 1a consisted of *F. sultana* colonies similarly treated but not exposed to kidney homogenate, although PBS was added. Trials 1b and 1c were constrained by availability of bryozoan material, and no separate colonies were available to act as controls. Consequently, for Trial 1b, portions of colonies collected prior to exposure to homogenate were subjected to PCR and served as controls. At the conclusion of the culture period, colonies in Trials 1b and 1c were saved for PCR to assess potential cryptic infection (Tops & Okamura 2003). Results of Trial 1a are based on visual inspection only.

**(2) Fish to bryozoan. Cohabitation of bryozoans with infected fish:** Bryozoans were exposed to fish recovering from PKD in cohabitation trials (see Trials 2a to d; Table 1) to determine whether sporogonic stages of *Tetracapsuloides bryosalmonae* that pass through the urinary tract of recovering fish are infective to bryozoans. Four exposures were carried out using naturally infected rainbow trout and Chinook salmon and artificially infected brown trout (see Table 1). The fish were maintained in flow-through tanks (133 l) supplied with well water at 15°C.

Plastic discs (5.4 cm diameter) with attached bryozoan colonies were randomly allocated positions within cylindrical plastic cages (Fig. 1). Twenty *Fredericella sultana* colonies were available for Trials 2a and 2b, and 10 colonies for Trials 2c and 2d. Only 2 *Plumatella fungosa* colonies were available for Trials 2c and 2d (see Table 1). Each cage could hold a maximum of 10 discs with colonies. Bryozoans were exposed to fish for 16 h d<sup>-1</sup> (Trial 2a and 2b) and 6.5 h d<sup>-1</sup> (Trial 2c and 2d) over a period of 5 and 3 d respectively. In between fish exposure periods, bryozoans were returned to pond water to feed. This was deemed to be essential since no food was available in well water and it was important to ensure colonies did not degenerate before parasites might develop. For Trials 2a and 2b, bryozoans were cohabited with 50 fish per treatment in flow-through tanks (133 l) supplied with well water. For Trials 2c and 2d, bryozoans were cohabited with fish in buckets filled with aerated well water (19 l) in order to increase the likelihood of exposure to potential infective stages, since these would have been continuously lost in the flow-through

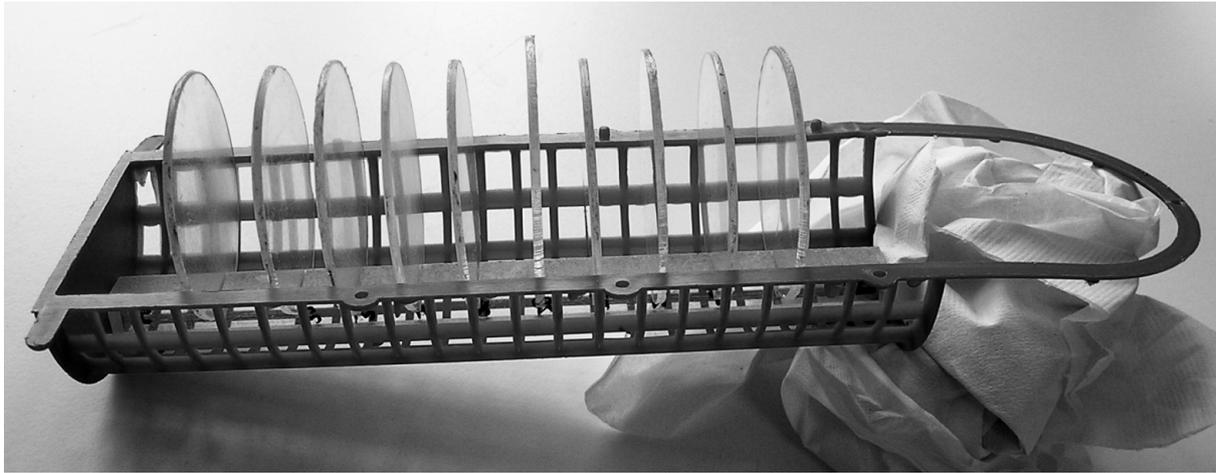


Fig. 1. Discs (5.4 cm diameter) and one half of the cage apparatus used in exposing bryozoans in Trials 2a to d and 3c. During trials bryozoans were enclosed within the fully assembled cage using cable ties, and were submersed in a downward facing position in experimental containers

arrangement used in Trials 2a and 2b. For Trials 2c and 2d, 8 fish were caught at random each day from a large group of infected individuals and were cohoused with the bryozoans in the bucket. At the end of each cohousing period the fish were returned to the flow-through system. Hence, different sets of fish were likely involved in the transmission study on different days. Bryozoans cohoused with a group of Chinook salmon hatched from eggs and reared in well water acted as controls for all trials.

After completion of exposure, the bryozoans were returned to the laboratory culture system and monitored for the presence of *Tetracapsuloides bryosalmonae* stages every 2 to 3 d using a dissection microscope. Regular observations of colonies allowed us to fix moribund bryozoans in absolute ethanol prior to their death to check for possible transmission of *T. bryosalmonae* by PCR. Nonetheless, some mortality occurred.

The presence of spores and sporogonic stages in the kidney tubules of fish was assessed post-exposure through histological examination of wet mounts as described earlier. Examination of a subset of fish in Trials 2a and 2b indicated that extrasporogonic stages were present but sporogonic stages were rare. This prompted us to undertake Trials 2c and 2d in hopes of including more mature stages. Examination of a subset of fish indicated that sporogonic stages were present in 3/10 fish in Trial 2c and 2/10 fish in Trial 2d at low to moderate levels (based on subjective scoring by R. P. Hedrick, assessing the relative abundance of parasite stages as a result of familiarity gained during previous studies: Kent & Hedrick 1986, Foott & Hedrick 1987).

**(3) Bryozoan to bryozoan. Exposure of bryozoans to *Tetracapsuloides bryosalmonae* infected *Fredericella sultana*:** For Trial 3a, colonies of *F. sultana* were torn apart with forceps ('shredded') and observed with a dissection microscope. Sacs of *T. bryosalmonae* released by shredding were transferred to a small petri dish (with 10 µl disposable microcapillary tubes) and burst open using a sterile, mounted needle to release spores. Concentrated spores were pipetted up into clean microcapillary tubes and wafted into the open lophophores of *F. sultana* and *Plumatella fungosa* colonies (see Table 1) from our laboratory stock, in separate petri dishes containing 20 ml of culture water. The bryozoans were exposed to potential infection by spores in the petri dish for 24 h. The water was agitated every 30 min for 4 h to re-suspend any viable spores which may have settled to the bottom of the dish and were thus unavailable for infection. After 24 h, the bryozoans were transferred to the culture system and checked every 2 to 3 d for visual signs of infection. Colonies which survived culture for 30 d were saved for PCR as a further check on infection status. Colonies that appeared to be moribund were fixed for PCR at earlier dates prior to death.

A single colony of *Fredericella sultana* provided material for Trial 3b when it developed infection in culture 10 wk after collection from the River Cerne, Dorset. Sacs were collected as for Trial 3a. Target colonies of *F. sultana*, *Plumatella fungosa* and *P. emarginata* (see Table 1) were exposed to spores for 20 h, which were wafted into the lophophore every 30 min for 4 h. Colonies were subsequently checked for visual signs of infection every d. After 33 d of culture the colonies were saved for PCR.

For Trial 3c, *Fredericella sultana* colonies collected from the River Cerne were shredded (see above) and placed in a container with approximately 30 l of water containing naïve rainbow trout and cage-enclosed target colonies of *F. sultana* for 135 min, as part of a larger study (data not presented) on the transmission of *Tetracapsuloides bryosalmonae* to fish. Target bryozoans were then returned to culture and checked for visual signs of infection every 2 d. After 30 d of laboratory culture the remaining live colonies were saved for PCR.

The development of persistent *Tetracapsuloides bryosalmonae* infections in colonies of *Fredericella sultana* (originally collected from the River Cerne) in 1 culture system and the coincident development of new *Plumatella fungosa* colonies that hatched from statoblasts in the same system allowed our fortuitous, unplanned Trial 3d. The *P. fungosa* colonies became established adjacent to heavily infected *F. sultana* colonies on the same petri dishes. The latter produced sacs over a prolonged period, and released mature spores and possibly sacs at a variety of developmental stages into the recirculating culture system. This should have resulted in the relatively continuous exposure to infection of the *P. fungosa* colonies during the period of spore viability. The *P. fungosa* colonies were maintained for 21 d alongside the infected *F. sultana* colonies, were checked for visual signs of infection every 2 to 3 d, and were then fixed in absolute ethanol for PCR. There were no controls for this trial.

**(4) Bryozoan to bryozoan. Exposure of bryozoans to *Buddenbrockia plumatellae* infected *Plumatella fungosa* and *Cristatella mucedo*:** For Trial 4a, spore-filled vermiform stages of *B. plumatellae* infecting a colony of *P. fungosa* were collected by applying gentle external pressure to the infected portion of the bryozoan colony. Spores were collected and wafted into the open lophophores of uninfected *P. fungosa* colonies (see Table 1) in petri dishes as described above, and were resuspended once per hour for 4 h. The *P. fungosa* colonies were exposed to spores for 16 h and then transferred to an aerated 2 l aquarium with water from the culture system (from a system never exposed to infected bryozoan colonies) for 5 d prior to transfer to culture. Fifty percent of the volume in the 2 l tank was replaced daily with fresh water from the culture system. The 2 colonies were maintained in culture for 52 d, and were checked for visual signs of infection every 2 to 5 d. Two uninfected, laboratory-reared *P. fungosa* colonies were used as controls. These were initially maintained in petri dishes filled with culture-system water. Deionised water was wafted into the lophophores and the water was agitated as described above. The colonies were then transferred to an aerated 2 l aquarium containing water from the culture

system never exposed to infected bryozoan colonies. After 5 d in the 2 l aquarium, colonies were transferred to a culture system.

For Trial 4b(i) (see Table 1), spore-filled sacs of *Buddenbrockia plumatellae* were obtained by applying gentle external pressure to infected *Cristatella mucedo* colonies. Colonies of *Fredericella sultana* and *Plumatella fungosa* were exposed to the spores in an aerated 2 l aquarium containing water from the culture system, as described above for *Tetracapsuloides bryosalmonae*. Spores were again resuspended every hour for 4 h, and the total exposure time was 24 h. Colonies were then placed into culture. Controls for this experiment were uninfected, laboratory-reared *F. sultana* colonies, which were maintained in 2 l of aerated, culture-system water. Deionised water was wafted into the lophophores and the water was agitated as described above. In addition, 20 *C. mucedo* colonies with immature sacs (lacking spores) were disrupted (Trial 4b[ii]). The disrupted material was added to 2 l of aerated culture water containing uninfected colonies of *F. sultana* and *P. fungosa*. The water was stirred every hour for 4 h to maintain the material in suspension. After 24 h the colonies were transferred to culture. Finally, 1 colony of *C. mucedo* with mature, spore-filled sacs of *B. plumatellae* was placed in 500 ml culture water along with uninfected colonies of *F. sultana* and *P. fungosa* (Trial 4b[iii]). The water was gently agitated every hour for 4 h to suspend any spores which may have been released by the infected *C. mucedo* colony. The passive release of sacs has been observed for both *B. plumatellae* (B. Okamura pers. obs.) and *T. bryosalmonae* (M. Longshaw pers. comm.). After 24 h of exposure, the bryozoans were transferred to culture. The bryozoans exposed to potential infection in Trials 4b (i to iii) were maintained in culture for 60 d, and were checked for visual signs of infection every 2 to 3 d. Appropriate culture conditions have not been identified for *C. mucedo*, so uninfected colonies of this species were not included in the study.

**DNA extractions and PCR.** DNA extraction of both bryozoan and fish material, that had been saved in absolute ethanol and stored at  $-20^{\circ}\text{C}$ , was conducted using a standard CTAB (1% N-cetyl N,N,N-trimethylammonium bromide)-Proteinase K method as described by Winnepenninckx et al. (1993), with the following modifications: (1) samples were stored at  $-20^{\circ}\text{C}$ , (2) the samples were not ground in liquid nitrogen prior to extraction, (3) samples were suspended in 675  $\mu\text{l}$  preheated ( $60^{\circ}\text{C}$ ) CTAB buffer, and (4) all steps were carried out using 1.5 ml plastic eppendorf tubes.

PCR amplification was conducted using either *Tetracapsuloides bryosalmonae*-specific primers (514F and 776R, from D. C. Morris et al. 2002b) or primers we designed to specifically amplify *Buddenbrockia plumatel-*

*lae* (BZ1 F: 5'-CGTAAGGCTTCAAGCGAAAG-3' and BZ2R: 5'-CGAGCGTTTTAAATGCAACA-3', product size 397 base pairs). The primers were not cross-reactive between the 2 species of parasite. Conditions for amplification were according to D. C. Morris et al. (2002b). Negative (adding ultrapure water as a template) and positive controls were included in all amplifications. An internal, competitive-standard mimic molecule developed by D. C. Morris et al. (2002b) was included as a means of checking the absence of PCR inhibition in assessing *T. bryosalmonae* infection. No internal competitive-standard was available for the assessment of *B. plumatellae* infections.

## RESULTS

We obtained no evidence that the stages of *Tetracapsuloides bryosalmonae* that develop in fish and bryozoans caused infections in target bryozoan colonies, based on visual inspection of exposed colonies or by subsequent PCR. No infections were evident in any controls. The conclusion that the *T. bryosalmonae* stages found in bryozoans in our transmission trials are incapable of infecting bryozoans is bolstered by the finding that the naïve rainbow trout exposed to the parasite (Trial 3c) developed PKD; recipient bryozoans present during the same exposure period did not develop infections. Similarly, we obtained no evidence that either the sac-like or vermiform stages of *Buddenbrockia plumatellae* that develop in bryozoans are capable of producing new infections in bryozoans. The identity of the parasites used in transmission trials was confirmed by PCR using species-specific primers.

## DISCUSSION

### Lack of transmissions

Our studies provide evidence that the life cycles of the 2 malacosporean parasites known to date do not involve direct transmission from bryozoan to bryozoan, either by mature spores or by earlier sac stages. No infections resulted from exposing bryozoans to concentrated spores (Trials 3a to d, 4a, 4b[i and iii]), to immature sacs (Trial 4b[ii]) or to long-term cohabitation with infected colonies releasing spores and possibly sacs at different stages of development (Trial 3d). The lack of direct transmission from bryozoan to bryozoan by exposure to immature sac stages suggests that there is no parallel in the life cycle of *Tetracapsuloides bryosalmonae* to the direct fish to fish transmission indicated for immature trophozoites of some

myxosporeans (Diamant 1997, Redondo et al. 2002, Yasuda et al. 2002). The consistent absence of evidence for bryozoan to bryozoan transmissions for both *T. bryosalmonae* and *Buddenbrockia plumatellae* strongly indicates that such transmission is precluded in malacosporean life cycles.

Our trials also provided no evidence for transmission to bryozoans of *Tetracapsuloides bryosalmonae* stages that develop in fish. This result supports the suggestion that fish are abnormal hosts, a view supported by the intensity of the immune response mounted by fish to the presence of *T. bryosalmonae* (Kent & Hedrick 1986) and by recent phylogeographic studies (Henderson & Okamura 2004). The latter provide no indication that *T. bryosalmonae* was transferred to Europe during the introduction of rainbow trout through fisheries activities nor subsequently within Europe, a result that strikingly contrasts with the introductions of pathogens and parasites typically associated with aquaculture. For instance, such transfer is evident in the case of *Myxobolus cerebralis*, where introduction to North America from Europe in the 1950s rapidly established this myxozoan parasite in native populations of the obligate alternate host, the oligochaete worm *Tubifex tubifex* (Hoffman 1990). However, despite these caveats, the possibility of fish to bryozoan transmission requires further attention, as discussed below.

While the availability of material and variation in protocols meant that each study lacked replication, our consistent results are compelling. Indeed, a strength of the work is the incorporation of both a broad range of parasite stages and of potential new hosts. Thus, 3 bryozoan species were assayed as potential malacosporean hosts (*Fredericella sultana*, *Plumatella fungosa* and *P. emarginata*) and we utilised parasite stages from rainbow trout, brown trout and Chinook salmon, and the bryozoans *F. sultana*, *P. fungosa* and *Cristatella mucedo*. Also included were wild and farmed sources of fish, naturally and artificially infected fish, and fish from the UK and North America. In addition, we tested exposure to artificially and naturally released spores, and to parasitic sacs from infected bryozoan colonies. In total we assessed infection in 97 bryozoan colonies (a conservative estimate based on number of colonies subjected to PCR following post-exposure culture), and each bryozoan species tested as potential host material was sourced from different water bodies. Furthermore, transmission studies indicate that our stock material of *F. sultana* was susceptible to infection by *Tetracapsuloides bryosalmonae* in the field: colonies derived from our laboratory stock became infected when transplanted into the River Itchen, Hampshire, UK, while stock colonies in culture did not develop infections (Tops & Okamura 2003). Stock material of *P. fungosa* was similarly sus-

ceptible to infection by *T. bryosalmonae* in the River Itchen and was also infected by *Buddenbrockia plumatellae* in the Kennet and Avon Canal, Berkshire, UK (Canning et al. 2002, Tops & Okamura 2003). These observations, together with the range of source localities for infected material, suggest that lack of infection in our transmission trials is not a result of using resistant host strains. It should also be pointed out that infection of bryozoans does not appear to be influenced by bryozoan maturity since *T. bryosalmonae* infections have been encountered in, e.g., small (young) colonies of *Pectinatella magnifica* (Okamura et al. 2001) and in established colonies of *F. sultana* which are producing statoblasts (S. Tops & B. Okamura unpubl. data). Nonetheless, we offered a range of stages of potential new hosts, including branches of colonies derived from statoblasts (most trials), colonies derived from statoblasts (Trial 3d), and colonies derived from larvae (*P. emarginata* in Trial 3b).

### Constraints of study

Negative results such as ours are justifiably subject to criticism on methodological grounds. Thus, failure to close the life cycle of either malacosporean could be attributable to inappropriate conditions for transmission or for subsequent development of malacosporeans. The latter is unlikely since the development and persistence of infections in field-collected *Fredericella sultana* from the River Cerne in our laboratory culture system (Tops & Okamura 2003, current study) demonstrate that the post-exposure culture conditions were appropriate for the development of *Tetracapsuloides bryosalmonae* in bryozoan hosts.

Constraints on transmission include the possibility that spore concentrations may have been too low to be effective or that mature infective stages were not present in the material used. We attempted to control for such contingencies given the limitations of each transmission trial. Thus, in our bryozoan to bryozoan transmission studies, we ensured physical contact between bryozoans and spores by wafting spores directly into bryozoan lophophores, and we exposed bryozoans to spores in small volumes of water to maximise spore concentrations. The inclusion of both bryozoans and fish in Trial 3c demonstrated that functional spores were indeed present at least in this trial, since the fish subsequently developed PKD (M. Longshaw & S. Feist unpubl. data).

There is a greater possibility that transmissions of *Tetracapsuloides bryosalmonae* from fish to bryozoans were hampered by lack of mature infective stages, low concentrations of infective spores released in fish urine and short periods of exposure to allow bryozoans to

feed. Unfortunately, it was not feasible to predict before the transmission trials whether fish were releasing spores. Post-transmission microscopic examination of kidney indicated that the fish used in Trials 2a and 2b had not yet reached the stage of recovery from PKD dominated by release of spores from sporogonic stages, although some sporogonic stages were detected. Conducting these trials in flow-through systems may also have influenced transmission by greatly reducing spore concentrations. Microscopic examination indicated that a proportion of the fish used in Trials 2c and 2d had sporogonic stages, and the conduction of these trials in buckets increased the possibility of spore contact with bryozoans. However, random selection of fish for trials and lack of knowledge about actual spore release in urine during trials hamper the interpretation of results. Similarly, infective stages may have been present in low concentration when bryozoans were exposed to macerated fish kidney. Filtering of the water used in transmission studies to determine concentrations of infective stages was not conducted. This would not have been feasible in the trials conducted in the flow-through systems. In addition, the soft, unprotected spores are likely to adhere and disintegrate on contact with filters, making their identification and quantification exceedingly difficult.

Despite the above caveats, there is some evidence that low concentrations of *Tetracapsuloides bryosalmonae* spores must be effective at achieving infection. Concentrations of spores released from bryozoans should be exceedingly low for much of the year given the generally low prevalence of infections in bryozoan populations (Okamura et al. 2001, Okamura & Wood 2002, S. Tops & B. Okamura unpubl. data). In addition, fish are routinely immunised by developing subclinical infections through exposure to waters enzootic for PKD in autumn, when water temperatures are decreasing (Longshaw et al. 2002) and when prevalence of infection in bryozoan populations is low (S. Tops & B. Okamura unpubl. data). Furthermore, Feist et al. (2001) obtained transmission of *T. bryosalmonae* by cohabitation of fish with infected bryozoan material in a flow-through system where the continuous loss of spores should have resulted in low concentration. Hedrick et al. (2004) estimated a maximum concentration of 120 *T. bryosalmonae* spores ml<sup>-1</sup> fish urine, thus spores should become highly dilute upon release from fish in natural populations. These observations suggest that spores from both bryozoans and fish would have to be effective in contacting and infecting hosts at very low concentrations. An ability to achieve transmission at low concentration should be highly adaptive for dilute water-borne spores, and we would expect this to be true for all infective stages in the life cycle of *T. bryosalmonae*, be they stages achieving transmission

from fish to bryozoans, from bryozoans to fish or from any other potential host.

Despite the expected effectiveness of spores at low concentrations, given the uncertainties associated with our fish to bryozoan transmissions, more work is required to assess this route of infection, particularly given the possible infection of bryozoans by malaco-sporeans through exposure to macerated fish kidney (D. C. Morris et al. 2002a) and to stages released from fish with PKD (D. J. Morris et al. 2002). However, even if such transmission is eventually demonstrated, it may only be facultative since infected bryozoans occur in sites lacking salmonids in Ohio and Michigan (Okamura et al. 2001). Furthermore, even if spores released by fish are demonstrated to be infective for bryozoans, it is difficult to envision how such spores could regularly infect bryozoan populations upstream from fish farms and hatcheries to cause the annual and massive PKD outbreaks that occur in many sites. For instance, very high prevalences of *Tetracapsuloides bryosalmonae* infections coincide with early seasonal growth of the bryozoan population just upstream from a fish farm which suffers PKD outbreaks on the River Cerne, Dorset (Longshaw et al. 1999, S. Tops & B. Okamura unpubl. data). The bryozoan population and fish farm are no more than 6 km from the headwaters of the Cerne, and the only salmonids present in this stretch of water are a few farm escapees and a small population of brown trout (M. Longshaw pers. comm.). It seems unlikely that the high prevalences of infections (over 40%; S. Tops & B. Okamura unpubl. data) in dense stands of bryozoans on many willow roots could directly result from spores released by only a few fish. Another informative site is a salmonid hatchery in California (Hot Creek Hatchery) which experiences PKD outbreaks. Fish from the hatchery have rarely been collected upstream (probably due to difficulties in swimming through pipes) and rare cyprinids are the only resident fish in the short reach of stream between the spring source and the diversion of flow-through pipes to the hatchery (Okamura & Wood 2002).

### Implications of study

Our results imply that there may be another host incorporated in the life cycles of *Tetracapsuloides bryosalmonae* and *Buddenbrockia plumatellae*. Prior to the discovery of phylactolaemate bryozoans as hosts of the 'PKX' parasite, researchers screened many aquatic invertebrates and other fish species from sites enzootic for PKD in an attempt to locate the source of the infection. For instance, Morris et al. (1999) surveyed 35 000 oligochaete worms. In another study, 60 to 80 species of aquatic invertebrates were investi-

gated, including *Gammarus* sp., caddis flies, leeches and snails (Longshaw & Feist 2000, M. Longshaw & S. Feist unpubl. data cited in Feist et al. 2001). None of these invertebrates tested positive for the parasite, as determined by PCR investigation using specific primers. In addition, our culture systems contain many of the most cosmopolitan aquatic invertebrates, including chironomids, cladocerans, rotifers, oligochaetes, flatworms and snails, which have been introduced along with bryozoans brought in from the field. If any of these organisms were acting as hosts for *T. bryosalmonae*, we would anticipate that infections would eventually have developed within the long-term cultured bryozoans that were cohabited with infected bryozoans introduced from the field. Such infection has never been observed. However, it is possible that our culture conditions are unsuited to development in another host or that such a host was lacking.

So far there is no evidence that fish, other than salmonids and pike, act as hosts of *Tetracapsuloides bryosalmonae* (Hedrick et al. 1993, Adams & Morris 1999). A number of other potential fish hosts have been investigated, but none have been identified (Adams & Morris 1999). Furthermore, the broad diversity of habitats associated with PKD outbreaks and *T. bryosalmonae* infections in bryozoans provide evidence that no single fish can be an obligate host. *T. bryosalmonae* has been recorded from sites ranging from clear, cool streams to eutrophic lakes in North America, and the number of resident non-salmonids range from single (e.g. the rare cyprinid at Hot Creek Hatchery) to many species (Okamura et al. 2001). If fish hosts are obligate in the life cycle, this broad distribution of *T. bryosalmonae* suggests an unusual capacity to exploit a diversity of obligate fish hosts, although, as discussed below, the life cycle of bryozoans may complicate the interpretation of distributional data.

We anticipate that if another invertebrate or non-salmonid fish host is utilised in the life cycle of *Tetracapsuloides bryosalmonae*, infection is likely to be patchy, as it is in bryozoan populations (Anderson et al. 1999, S. Tops & B. Okamura unpubl. data), and that this has led to the lack of detection. The best chance of identifying what such a host might be rests in focusing on simple systems, as was done in the elucidation of the life cycle of the oyster parasite *Marteilia refringens* (Audemard et al. 2002). The simplest system for *T. bryosalmonae* would be the recirculating system in France (Gay et al. 2001), but PCR and transmission-based investigations have so far failed to determine whether resident invertebrates in this system are incorporated in the life cycle of *T. bryosalmonae* (de Kinkelin et al. 1999, P. de Kinkelin & D. J. Morris unpubl. data cited in Gay et al. 2001). If our transmission results are correct in demonstrating that transmis-

sion is not direct from bryozoan to bryozoan, and further work confirms that salmonids are dead-end hosts, the search for an unknown host might once again become akin to looking for a needle in a haystack (Audemard et al. 2002).

### Bryozoan life cycles and cryptic malacosporean stages

It is now clear that malacosporeans can persist for at least 2 to 3 mo as cryptic stages in the body wall of bryozoans prior to proliferating as sacs in the body cavity (Canning et al. 2002, Tops & Okamura 2003). Myxosporeans appear to behave in a similar manner. *Myxobolus cerebralis* maintains persistent infections in oligochaetes characterised by periods of parasite dormancy and intermittent shedding of actinospores (Gilbert & Granath 2001), while at low temperatures cryptic immature forms undergo cycles of replication but do not produce actinospores (El-Matbouli et al. 1999). These recent studies suggest that persistent cryptic infection is a general myxozoan feature and that cryptic infections of malacosporeans may be maintained indefinitely within bryozoan populations.

We predict that the life history of bryozoans plays an important role in the maintenance and spread of cryptic malacosporean infections without the necessity of regular transmissions from any other hosts. If cryptic stages proliferate as colonies grow, this would ensure that all regions of colonies become infected. Also, colonies of *Fredericella sultana* fragment as they become larger, and branches that drift downstream reattach (Wood 1973, S. Tops & B. Okamura pers. obs.). Such fragmentation and reattachment could increase local prevalence of infection and introduce infected colonies elsewhere. Furthermore, as *F. sultana* itself overwinters as live colonies (Wood 1973, Raddum & Johnsen 1983), cryptic infections can be maintained year-round in this species (Gay et al. 2001). Finally, if cryptic stages are present in dormant bryozoan statoblasts, infections could be passed to new colonies through hatching from statoblasts. All of these processes could promote long-term infections in bryozoan populations without the necessity of regular transmissions from any other hosts, thus explaining the high prevalences of mature infections in bryozoan populations early in the growing season (Longshaw et al. 1999, S. T.ops & B. Okamura unpubl. data).

If cryptic stages of *Tetracapsuloides bryosalmonae* are capable of undergoing dispersal in bryozoan statoblasts, then the persistence of infections in some sites may simply be explained by colonisation through infected statoblasts and the subsequent spread of parasitism in the population through proliferation of para-

sites within clonally reproducing bryozoan hosts. The growing body of evidence that waterfowl act as dispersal vectors of bryozoan statoblasts lends credence to this possibility. Such evidence includes ongoing gene flow amongst sites traversed by migratory waterfowl (Freeland et al. 2000), the presence of intact statoblasts in waterfowl digestive tracts and faeces (Figuerola et al. 2003), and the viability of some statoblasts following ingestion and excretion by waterfowl (Charalambidou et al. 2003). Such waterfowl-mediated dispersal of infected statoblasts, followed by the subsequent spread of parasitism in the bryozoan population, provides one explanation for the presence of infected bryozoan populations in sites lacking salmonids. Unravelling the ecology of early cryptic stages of malacosporeans, identification of the cues that promote proliferation of spore-producing sacs from these cryptic stages and investigating the clonal structure of bryozoan host populations are important areas for further research.

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