

## ORIGINAL ARTICLE

# Infection dynamics of *Bonamia exitiosa* on intertidal *Ostrea angasi* farms

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## Abstract

*Bonamia* spp. cause epizootics in oysters worldwide. In southern Australia, *Bonamia exitiosa* Hine, Cochennac and Berthe, 2001 threatens aquaculture of *Ostrea angasi* Sowerby, 1871. *Bonamia* spp. infections can display strong seasonality, but seasonal dynamics of *B. exitiosa*–*O. angasi* are unknown. *Ostrea angasi* naïve to *B. exitiosa* infection were stocked onto farms in three growing regions, and *B. exitiosa* was monitored seasonally for one year. Environmental parameters we measured did not correlate with *B. exitiosa* prevalence or infection intensities. Extreme temperatures suggest *O. angasi* culture systems need development. *Bonamia exitiosa* prevalence increased over time. After three months, *O. angasi* had *B. exitiosa* prevalence of 0.08–0.4, and after one year, the prevalence was 0.57–0.88. At some sites, *O. angasi* had >0.5 *B. exitiosa* prevalence in >6 months, but at other sites, >9 months passed before prevalence was >0.5. *Bonamia exitiosa* infection intensities were low with no seasonal pattern but were affected by the interaction of site, season and oyster meat:shell ratio. Understanding infection and initiating a breeding programme for resistance would provide benefits for *O. angasi* industry expansion.

## KEYWORDS

*Bonamia exitiosa*, farm trial, *Ostrea angasi*, South Australia

## 1 | INTRODUCTION

Mollusc aquaculture provides a substantial supply of food; global mollusc production in 2016 was 17.1 million tonnes (FAO, 2018). Since the 1970s, diseases caused by the intracellular protozoa *Bonamia* spp. have negatively affected oyster industries in Europe, North America, South America, North Africa, New Zealand and Australia (Arzul et al., 2006; Hill et al., 2014).

A *Bonamia* sp. was identified in Australia in the early 1990s in farmed and wild populations of the Native Oyster (*Ostrea angasi*, Sowerby, 1871) (see Hine & Jones, 1994). Extensive investigations including genome sequencing have identified all southern Australian *Bonamia* isolates as *Bonamia exitiosa* Hine, Cochennac and Berthe, 2001 (see Bradley, 2019; Buss, Harris, Tanner, Wiltshire, & Deveney,

2019a). Handlinger et al. (1999) reported that histological surveys of wild *O. angasi* across SA in 1992–93 did not detect *Bonamia*. Surveys of farmed Pacific oysters (*Crassostrea gigas* Thunberg, 1793) in SA in 2002, however, reported *Bonamia*-like cells (Diggle, 2003), but no confirmatory diagnosis was made. If these cells were *B. exitiosa*, these data support that *C. gigas* is a host of *Bonamia* spp. as proposed by Lynch et al. (2010). It is therefore likely that *B. exitiosa* was introduced or became widespread in SA between 1993 and 2002.

*Bonamia* spp. transmit directly between hosts (Arzul & Carnegie, 2015). Large farmed oyster populations may have facilitated the proliferation of *B. exitiosa* in SA, leading to the high prevalence observed by Buss, Wiltshire, Prowse, Harris, and Deveney (2019b). Although the origin of SA *B. exitiosa* is unknown, it likely arrived via an anthropogenic route. Transmission of *Bonamia ostreae* Pichot, Comps, Tigé, Grizel and

Rabouin, 1980 is linked to vessel biofouling (Howard, 1994), translocation of stock (Hudson & Hill, 1991; Peeler, Oidtmann, Midtlyng, Miossec, & Gozlan, 2011) and the seafood trade (Feng et al., 2013). Given that *B. exitiosa* is now established across southern Australia, aquaculture operators and reef restoration proponents need to understand and take the pathogen into account in their planning.

Environmental conditions affect infection dynamics of *Bonamia* spp.–host systems. High salinity (>20 psu) is associated with higher *B. exitiosa* prevalence and/or infection intensities in *Crassostrea ariakensis* Fujita, 1913 (see Audemard, Carnegie, Bishop, Peterson, & Burrenson, 2008; Audemard, Carnegie, Hill, Peterson, & Burrenson, 2014; Audemard, Carnegie, Stokes, et al., 2008; Bishop, Carnegie, Stokes, Peterson, & Burrenson, 2006); higher *B. exitiosa* prevalence and infection intensities in *Ostrea chilensis* Philippi, 1844 (see Hine, Diggles, Parsons, Pringle, & Bull, 2002); and increased survival of *B. ostreae* cells (Arzul et al., 2009). *Bonamia* spp. in different hosts have a range of seasonal patterns of infection. In *B. exitiosa*–*C. ariakensis*, higher *B. exitiosa*-associated prevalence and mortality occurs in spring, summer and early autumn, but *B. exitiosa* was undetectable in *C. ariakensis* at <25°C in late autumn and winter (Carnegie et al., 2008). For *B. exitiosa*–*O. chilensis*, the prevalence peaks in autumn and winter (Hine, 1991). In *B. ostreae*–*Ostrea edulis* Linnaeus, 1758 transmission occurs throughout the year, but higher *B. ostreae* prevalence is observed in winter (Arzul et al., 2006; Culloty & Mulcahy, 1996) and spring (Culloty & Mulcahy, 1996; Engelsma et al., 2010).

Field infection dynamics and *B. exitiosa* seasonality are undescribed in *O. angasi*. This study aimed to assess seasonal changes in *B. exitiosa* prevalence and infection intensities in intertidally cultured *O. angasi* over a year. Understanding *B. exitiosa*–*O. angasi* infection dynamics will inform farm management decisions and aid development of the *O. angasi* aquaculture industry.

## 2 | METHODS

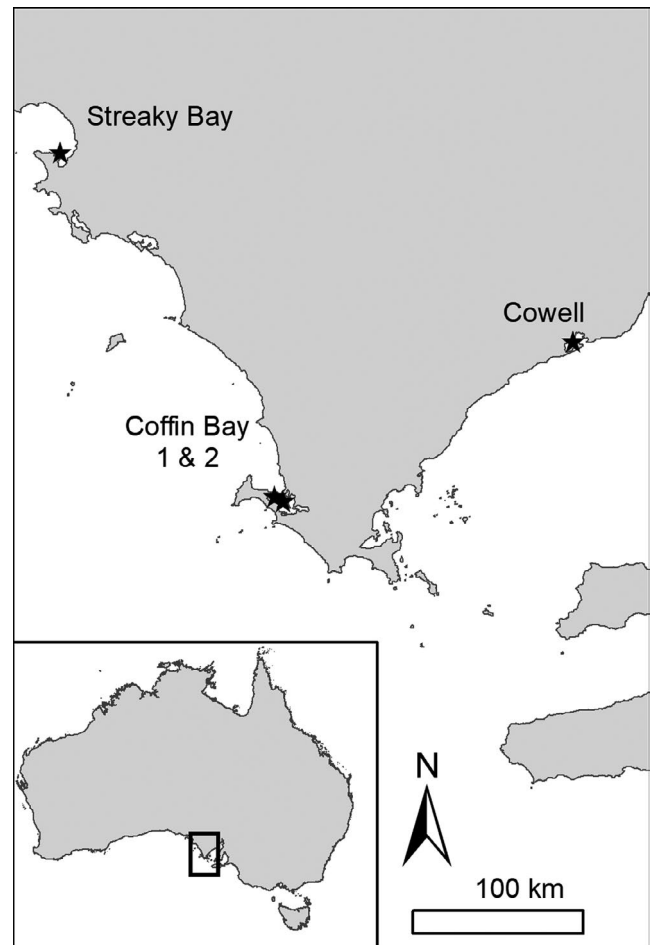
### 2.1 | Experimental animals

Common names used are consistent with Australian Fish Names Standard AS 5300-2015 (Standards Australia, 2015).

Fourteen-month-old (7–10 mm) *O. angasi* that had not been used for any experiments were sourced from the South Australian Research and Development Institute (SARDI) Aquatic Sciences Centre (SAASC) Mollusc Hatchery (West Beach, Adelaide, South Australia). Testing by real-time PCR (Corbeil et al., 2006) ( $n = 150$ ) and histology ( $n = 150$ ) did not detect *Bonamia* (mean Bayesian estimated prevalence, credible intervals: 0.017, 0.000–0.05; Buss et al., 2019b).

### 2.2 | Farm stocking and location

Four oyster farms on the Eyre Peninsula (South Australia) were chosen as experimental sites: Coffin Bay 1, Coffin Bay 2, Cowell and Streaky Bay (Figure 1). Approximately 600 juvenile *O. angasi* (total  $n = 1,780$ )



**FIGURE 1** *Ostrea angasi* farm stocking locations in the Eyre Peninsula, South Australia. Each star shape represents one farm

were stocked in three replicate baskets (15 L, SEAPA) and deployed at each of the four sites. At Coffin Bay and Cowell, oysters were deployed in 3-mm mesh size baskets, and at Streaky Bay, oysters were deployed in 6-mm mesh size baskets. All oysters were chosen from the same cohort and transported to farms within 48 hr of leaving the hatchery.

Coffin Bay 1 is in western Coffin Bay. It has a moderate-energy environment, is sheltered from prevailing winds and is characterized by a sandy seafloor and high biofouling, comprising mostly barnacles and mussels. Coffin Bay 2 is in the central channel of Coffin Bay and experiences large tidal fluctuations. It is a high energy, deep-water site, with a sandy substrate and high biofouling, comprising barnacles and mussels. Cowell is in southern Franklin Harbour, adjacent to a mangrove (*Avicennia marina* (Forssk.) Vierh.) forest. It is a shallow and low-energy environment with a fine sediment substrate and dense *Posidonia australis* Hook. f. beds. The Cowell site has high turbidity and extreme barnacle biofouling with settlement predominantly in summer. Streaky Bay is in southern Streaky Bay and is a shallow, exposed site with high wave and wind energy. This site is characterized by a sandy benthos with *P. australis* and *Amphibolis antarctica* (Labill.) Asch. beds and a *Pinna bicolor* Gmelin, 1791 population. All sites are intertidal; oysters were periodically exposed to air and then submerged under water. In summer during tidal maxima,

**TABLE 1** Sampling timeline of *Ostrea angasi* from four South Australian farms over four seasons<sup>a</sup>

	Jan	Feb	Mar	Apr	May	Jun	Jul	Aug	Sep	Oct	Nov	Dec
Year	S	S	A	A	A	W	W	W	Sp	Sp	Sp	S
2017		St			X1			X2			X3	
2018		X4										

<sup>a</sup>"St" denotes date of stocking, "X" denotes sampling date, and the number corresponds to the number of sampling trips since stocking. "S" is summer, "A" is autumn, "W" is winter, and "Sp" is spring.

the substrate at Cowell and Streaky Bay sites was completely exposed. Sites were stocked in the Austral summer in late February 2017 (Table 1).

### 2.3 | Experimental design: oyster sampling

To assess the dynamics of *B. exitiosa* in *O. angasi*, the cohort of oysters was monitored seasonally over a year (Table 1). Every three months at the end of each Austral season, all baskets were inspected and a sample of 20 oysters per replicate basket ( $n = 60$ ) were removed from the baskets following a randomization plan, placed in an ice box, transported ashore and sampled. Oysters were weighed (OHAUS, Scout General (SPX) Portable Balance Scales, Model # SPX223), and shell length was measured with a digital calliper (Craftright 150-mm Stainless Steel Digital Vernier Caliper) along the longest shell axis (hinge to top of shell). Oyster meat was removed, the empty shell was weighed, and the meat-to-shell ratio was calculated (meat weight (g)/shell weight (g)  $\times 100$ ). At the end of the trial, all oysters were removed and survival was estimated.

### 2.4 | Diagnostic sampling: heart smears and terminology

Heart smears were used to determine the prevalence and infection intensities of *B. exitiosa* in *O. angasi*. Within 24 hr of sampling, heart smears from sample oysters were made on slides, dried, fixed and stained as per Buss et al. (2019b). Heart smears were viewed under a compound light microscope (Brightfield Olympus BX53), and *B. exitiosa* infection intensities were graded using a semi-quantitative score following Buss et al. (2019b). Parasitology terminology used for *B. exitiosa* prevalence and infection intensity is consistent with Bush, Lafferty, Lotz, and Shostak (1997).

### 2.5 | Experimental design: environmental parameters

Two temperature loggers (HOBO Pendant® Temperature/Light Data Logger 64K-UA-002-64, accuracy  $\pm 0.47^\circ\text{C}$  at  $25^\circ\text{C}$ ) were used per site, with one placed at the same height as the oyster baskets (the above probe) to measure the temperature oysters were experiencing, and one placed directly below (the below probe) on the

benthos to monitor the water temperature. A conductivity logger (Odyssey Conductivity and Temperature Logger, 80 mS/cm, accuracy 3% of reading) was also used at each site to assess conductivity. Each conductivity logger was placed on the benthos below the oysters. Every 3 months, environmental data were retrieved and collated. Phytoplankton data per season and site were provided by the South Australian Shellfish Quality Assurance Program (SASQAP).

### 2.6 | Statistical analyses

Infection intensity data were compared using the Quantitative Parasitology 3.0 (Reiczigel, Marozzi, Fábíán, & Rózsa, 2019), using Sterne's exact confidence intervals for 95% bootstrap confidence intervals for mean infection intensities (with 5,000 replications). Mean infection intensities were considered significantly different when confidence intervals did not overlap.

A Bayesian Latent Class Model (LCM) was built and used to calculate estimated prevalence with credible intervals using JAGS code modified from the prevalence R package (Devleeschauwer et al., 2015). This model used uninformative beta (1, 1) priors for prevalence estimates. Priors for DSe (diagnostic sensitivity) and DSp (diagnostic specificity) of heart smear were informed by outputs from the analysis of Buss et al. (2019b). These beta priors reflected 95% confidence that heart smear DSe and DSp fall within the credible interval with the mean specified in Table 2. The Markov chain Monte Carlo (MCMC) simulations were obtained by running the model in JAGS v. 4.3.0 (Plummer, 2017) using three chains for 10,000 iterations, thinned at a rate of 10, following 2,000 iterations for adaptation and 10,000 iterations for burn-in. JAGS was run using the R2jags package (Su & Yajima, 2015) in R (R Core Team, 2017). The convergence was assessed using the Gelman–Rubin convergence statistic and confirmed by visual inspection of trace, density and autocorrelation plots generated using the MCMCvis package (Youngflesh, 2018). Estimated prevalence values were assessed as being different when credible intervals did not overlap.

A generalized linear model (GLM) was used to assess *B. exitiosa* infection intensities in oysters across site and season. A negative binomial distribution was used for infection intensity analysis due to the overdispersion of data relative to a Poisson distribution. Negative binomial GLMs were conducted in R using the MASS package (Venables & Ripley, 2002). *Bonamia exitiosa* infection intensity plots for each site and season were created using the R package ggplot2 (Wickham, 2016).

**TABLE 2** The beta priors, associated mean and 95% credible intervals for diagnostic sensitivity (DSe) and diagnostic specificity (DSp) for heart smear, used to calculate estimated prevalence, plus the posterior predictions for DSe and DSp of heart smear<sup>a</sup>

Test	DSp or DSe	Beta priors	Mean, 95% credible intervals	Posterior predication (mean, credible intervals)
Heart smear	DSe	(113, 72)	0.61 (0.54–0.68)	0.66 (0.61–0.70)
Heart smear	DSp	(27, 18)	0.60 (0.45–0.73)	0.71 (0.64–0.78)

<sup>a</sup>Heart smear priors were derived from Buss et al. (2019b).

### 3 | RESULTS

#### 3.1 | Prevalence

Estimated *B. exitiosa* prevalence from heart smears increased over time; estimated prevalence ranged from 0.08–0.40 after 3 months to 0.57–0.88 after 12 months (Table 3). *Ostrea angasi* from Coffin Bay sites 1 and 2 exceeded 0.5 *B. exitiosa* prevalence after six months, whereas those from Cowell and Streaky Bay took nine months for *B. exitiosa* prevalence to exceed 0.5. *Ostrea angasi* collected at the end of winter from Coffin Bay 2 that had higher estimated *B. exitiosa* prevalence than *O. angasi* from Cowell, but credible intervals for other sites overlapped (Table 3). At the end of spring, there were no differences in estimated *B. exitiosa* prevalence in *O. angasi* (Table 3).

The posterior predicted means of DSe and DSp for heart smear were higher than the prior means from Buss et al. (2019b) (Table 2), but the 95% credible intervals of the posterior predictions overlapped those of the priors.

#### 3.2 | Infection intensity

Mean *B. exitiosa* infection intensities did not differ at each site between seasons (Table 3). There were no differences between sites within each season except for winter, when *O. angasi* from Cowell and Streaky Bay had lower mean infection intensities than *O. angasi* from Coffin Bay 2 (Table 3).

GLM showed a significant three-way interaction between site, season and meat:shell ratio (LRT:  $\chi^2(9) = 26.12$ ,  $p = 1.95e^{-3}$ ). *Ostrea angasi* from Coffin Bay 2 and Streaky Bay had higher infection intensities associated with higher meat:shell ratio in summer and spring, but in autumn and winter, higher infection intensities were associated with lower meat:shell ratio (Figure 2). This pattern was not consistent between sites (Figure 2).

#### 3.3 | Survival and size data

The final proportion of survivors at all sites varied between 17.01% and 32.77% (Table 4). The lowest proportion of surviving oysters was at Streaky Bay.

After 1 year, Cowell had larger oysters (20 g mean weight) than other sites (<9 g mean weight (Table 3).

#### 3.4 | Environmental parameters

Initial data exploration using a correlation matrix showed infection intensities, and the prevalence did not correlate with any measured environmental parameter. Cowell experienced the greatest temperature difference: maximum temperature: 48.7°C, and minimum temperature: 5.8°C (Table 5).

In summer, temperature at Coffin Bay 1 had a maximum of 35°C, but only December data were available because the probe malfunctioned. At Coffin Bay 2, temperature exceeded 40°C once. At Cowell, the temperature exceeded 35°C 19 times and five of these events exceeded 40°C. Streaky Bay exceeded 35°C 11 times, and five of these events exceeded 40°C.

### 4 | DISCUSSION

We observed the clear patterns of prevalence for *B. exitiosa* in farmed *O. angasi*. *Bonamia exitiosa* was detected in *O. angasi* at all four farm sites within three months of stocking. *Bonamia exitiosa* prevalence increased in *O. angasi* at all sites over time, with highest prevalence (>0.57) after one year, but *B. exitiosa* infection intensities in *O. angasi* were not different between autumn and summer. Our findings are consistent with Buss et al. (2019b) who found *B. exitiosa* estimated prevalence of >0.59 and infection intensities of <4.36 in *O. angasi* on SA farms. At Cowell and Streaky Bay, *B. exitiosa* prevalence in *O. angasi* took nine months to exceed 0.5, whereas Coffin Bay sites exceeded 0.5 infection prevalence within six months of deployment. Sites with slower increase in prevalence may be more suitable for farming *O. angasi*. In warmer water, *O. angasi* grows more rapidly (Dix, 1980), and higher temperatures at Cowell could explain why oysters stocked in that area were larger than oysters from other sites. These data indicate that *B. exitiosa*–*O. angasi* infections are site-dependent as described for *B. ostreae* in *O. edulis* in Europe (Culloty, Cronin, & Mulcahy, 2004). These data therefore can inform decisions about the suitability of sites for oyster aquaculture (Culloty et al., 2004).

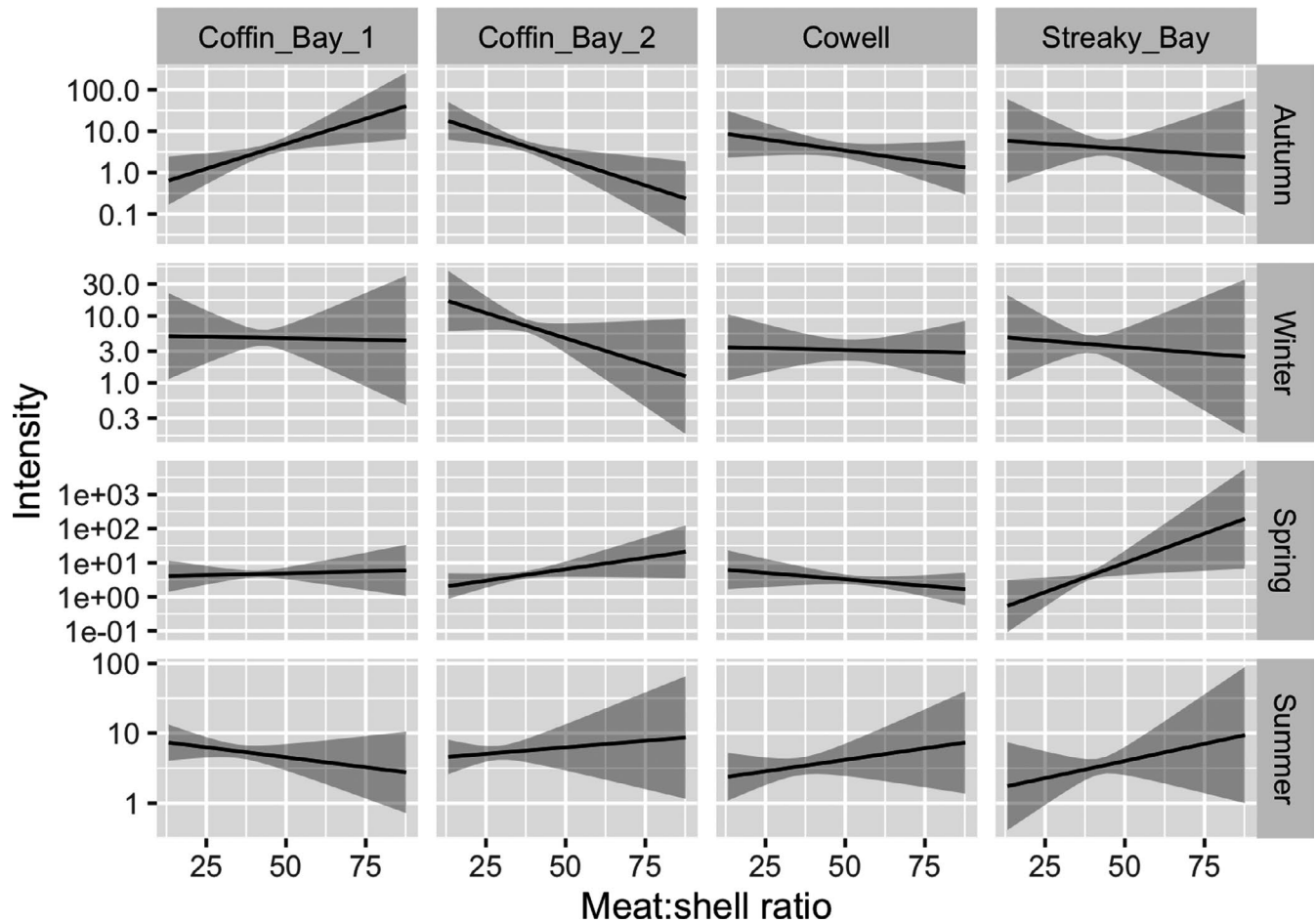
Diagnostic tests and their accuracy are well described for *B. exitiosa* in farmed *O. angasi* populations (Buss et al., 2019b). Understanding DSe and DSp of each test means that cost- and time-effective tests such as heart smear (Diggles, Cochennec-Laureau, & Hine, 2003) can be used and provide more accurate prevalence estimates than for less well-characterized tests and provide better interpretation of infection dynamics (Buss et al., 2019b; McDonald & Hodgson, 2018). The posterior predictions of

**TABLE 3** Size data (mean  $\pm$  standard deviation), Bayesian estimated prevalence and mean infection intensity (calculated through quantitative parasitology) of *Bonamia exitiosa* in *Ostrea angasi* from heart smears over four seasons in South Australia<sup>†,‡</sup>

Season	Site	Oyster data			Bonamia exitiosa data		
		Weight (g) (mean $\pm$ SD)	Shell length (mm) (mean $\pm$ SD)	Meat:shell ratio (%) (mean $\pm$ SD)	n	Estimated prevalence (Bayesian credible intervals)	Mean infection intensity, cell count (confidence intervals)
Autumn	Coffin Bay 1	6.02 $\pm$ 2.04	39.77 $\pm$ 4.78	45.20 $\pm$ 8.07	60	0.40 (0.07–0.76) <sup>ab</sup>	3.65 (2.62–5.04) <sup>b</sup>
	Coffin Bay 2	6.66 $\pm$ 2.13	38.91 $\pm$ 4.28	38.63 $\pm$ 7.39	60	0.33 (0.03–0.65) <sup>ab</sup>	4.38 (2.88–7.74) <sup>ab</sup>
	Cowell	10.17 $\pm$ 6.46	45.31 $\pm$ 13.52	46.10 $\pm$ 11.47	60	0.09 (0.00–0.31) <sup>b</sup>	3.64 (2.07–5.64) <sup>ab</sup>
	Streaky Bay	3.05 $\pm$ 1.15	29.70 $\pm$ 4.12	44.65 $\pm$ 6.27	60	0.08 (0.00–0.26) <sup>b</sup>	4.00 (2.50–7.15) <sup>ab</sup>
Winter	Coffin Bay 1	7.79 $\pm$ 4.36	40.74 $\pm$ 6.66	41.92 $\pm$ 5.84	60	0.53 (0.16–0.88) <sup>ab</sup>	4.61 (3.56–5.82) <sup>ab</sup>
	Coffin Bay 2	5.22 $\pm$ 2.78	32.76 $\pm$ 5.98	37.84 $\pm$ 5.90	59	0.88 (0.62–1.00) <sup>a</sup>	7.02 (5.28–9.67) <sup>a</sup>
	Cowell	9.62 $\pm$ 4.42	44.46 $\pm$ 7.52	49.42 $\pm$ 10.85	60	0.20 (0.01–0.50) <sup>b</sup>	3.10 (2.30–4.50) <sup>b</sup>
	Streaky Bay	3.80 $\pm$ 1.44	29.61 $\pm$ 4.08	40.48 $\pm$ 5.07	60	0.44 (0.10–0.79) <sup>ab</sup>	3.78 (2.74–5.22) <sup>b</sup>
Spring	Coffin Bay 1	6.36 $\pm$ 2.91	38.14 $\pm$ 5.51	40.51 $\pm$ 5.89	60	0.89 (0.66–1.00) <sup>a</sup>	4.67 (3.64–6.93) <sup>ab</sup>
	Coffin Bay 2	6.39 $\pm$ 4.17	33.05 $\pm$ 6.91	35.86 $\pm$ 7.13	59	0.80 (0.49–0.99) <sup>a</sup>	4.31 (3.50–5.29) <sup>ab</sup>
	Cowell	9.34 $\pm$ 5.44	45.63 $\pm$ 8.93	53.06 $\pm$ 10.25	60	0.53 (0.16–0.87) <sup>ab</sup>	3.03 (2.34–3.83) <sup>b</sup>
	Streaky Bay	5.37 $\pm$ 2.68	33.71 $\pm$ 7.18	38.67 $\pm$ 4.58	59	0.63 (0.26–0.94) <sup>ab</sup>	4.03 (2.87–7.56) <sup>ab</sup>
Summer	Coffin Bay 1	7.70 $\pm$ 4.21	38.07 $\pm$ 5.71	35.92 $\pm$ 8.38	60	0.88 (0.65–1.00) <sup>a</sup>	5.51 (4.12–8.36) <sup>ab</sup>
	Coffin Bay 2	8.44 $\pm$ 3.79	37.97 $\pm$ 7.57	28.94 $\pm$ 6.99	60	0.87 (0.61–1.00) <sup>a</sup>	5.22 (4.12–7.48) <sup>ab</sup>
	Cowell	20.69 $\pm$ 10.64	53.90 $\pm$ 10.56	36.34 $\pm$ 7.94	60	0.57 (0.20–0.91) <sup>ab</sup>	3.37 (2.77–4.20) <sup>b</sup>
	Streaky Bay	9.24 $\pm$ 3.28	42.49 $\pm$ 4.98	41.91 $\pm$ 4.96	60	0.78 (0.46–0.98) <sup>ab</sup>	3.36 (2.73–4.11) <sup>b</sup>

<sup>†</sup>Different superscripts denote differences at a 5% level, with <sup>a</sup> representing the highest value.

<sup>‡</sup>SD: standard deviation; n: sample size per site.



**FIGURE 2** *Bonamia exitiosa* mean infection intensity in *Ostrea angasi* per site (Coffin Bay sites 1 and 2, Cowell and Streaky Bay), per season (autumn, winter, spring and summer) and the relation to meat:shell ratio

prevalence from heart smears had overlapping credible intervals with the heart smear priors, indicating that the data in this study are consistent with priors in Buss et al. (2019b). This similarity increases confidence in the estimated prevalence in this study. Understanding test accuracy improves the certainty of results from field trials and surveillance programmes.

Sites with different environments were included in this study, but the environmental parameters we measured did not influence the prevalence or infection intensities of *B. exitiosa*. It is likely that parameters we did not measure, including proximity and density of oysters (Arzul & Carnegie, 2015; Lallias et al., 2008) such as farmed and wild populations of *O. angasi* and possibly *C. gigas*,

had greater influence than environmental parameters. *Crassostrea gigas* is the predominant oyster farmed in SA (Nell, 2001) and a likely host of *Bonamia* spp. (see Diggles, 2003; Lynch et al., 2010). Coffin Bay is the best known and largest oyster farming region in SA with 145 leases (184 ha), followed by Streaky Bay (38 leases, 172 ha) and Cowell (37 leases, 115 ha) (PIRSA, 2017). Populations of farmed *C. gigas* and *O. angasi* in Coffin Bay are larger than in the other regions, which may drive the higher *B. exitiosa* prevalence in *O. angasi*, but estimates of total farmed oyster population sizes were not available to us. The first record of wild *C. gigas* in SA was in 1990 (Hone, 1993), and feral oyster numbers in growing regions can be substantial but are subject to control (EPA, 2005; Wear,

**TABLE 4** Count data of *Ostrea angasi* from four South Australia farms after four seasons

Site	Initial count at stocking	Final total counts (dead and alive) <sup>a</sup>	Final counts (live)	Proportion live oysters remaining at end of trial (%)
Coffin Bay 1	1,785	685	585	32.77
Coffin Bay 2	1,785	399	386	21.62
Cowell	1,785	438	414	23.19
Streaky Bay	1,775	181	302	17.01

<sup>a</sup>Final live counts included oysters that were removed for sampling per season.



**TABLE 5** Mean seasonal (Au = autumn, Wi = winter, Sp = spring, Su = summer) water conductivity and temperature data from four South Australian oyster farms (CB1 = Coffin Bay 1, CB2 = Coffin Bay 2, CW = Cowell, SB = Streaky Bay)

	Water conductivity (mS/cm) (mean $\pm$ SE) <sup>a</sup>				Water temperature: top probe (°C) (mean $\pm$ SE) <sup>a</sup>				Water temperature: bottom probe (°C) (mean $\pm$ SE) <sup>a</sup>			
	CB1	CB2	CW	SB	CB1	CB2	CW	SB	CB1	CB2	CW	SB
Au	54.03 $\pm$ 0.16	51.04 $\pm$ 0.52	55.65 $\pm$ 0.11	54.53 $\pm$ 0.16	18.63 $\pm$ 0.03 (13.85–25.71)	18.75 $\pm$ 0.03 (14.90–31.17)	19.34 $\pm$ 0.04 (10.94–31.47)	19.19 $\pm$ 0.04 (7.88–40.07)	20.77 $\pm$ 0.03 (16.71–25.22)	18.67 $\pm$ 0.02 (14.71–23.97)	19.28 $\pm$ 0.03 (11.82–29.75)	19.34 $\pm$ 0.03 (11.14–27.27)
Wi	58.49 $\pm$ 0.09	52.44 $\pm$ 0.79	55.13 $\pm$ 0.20	43.01 $\pm$ 0.91	13.18 $\pm$ 0.01 (9.47–18.52)	13.53 $\pm$ 0.07 (7.98–25.03)	12.82 $\pm$ 0.01 (5.76–16.62)	13.49 $\pm$ 0.01 (7.58–22.05)	13.20 $\pm$ 0.01 (9.47–17.00)	13.35 $\pm$ 0.01 (10.75–15.76)	12.76 $\pm$ 0.01 (6.06–15.95)	13.53 $\pm$ 0.01 (9.87–18.71)
Sp	48.81 $\pm$ 0.54	48.06 $\pm$ 0.65	60.33 $\pm$ 0.16	42.48 $\pm$ 0.71	17.68 $\pm$ 0.21 (12.21–35.0)	17.34 $\pm$ 0.04 (10.75–41.23)	17.88 $\pm$ 0.04 (9.37–35.01)	15.74 $\pm$ 0.07 (9.77–28.56)	17.28 $\pm$ 0.03 (12.11–24.93)	17.07 $\pm$ 0.03 (11.43–23.20)	17.98 $\pm$ 0.03 (10.26–27.17)	18.23 $\pm$ 0.03 (10.16–34.80)
Su	42.84 $\pm$ 1.40	59.14 $\pm$ 0.26	63.13 $\pm$ 0.15	44.23 $\pm$ 2.30	21.64 $\pm$ 0.07 (14.80–32.60)	22.74 $\pm$ 0.02 (15.28–40.07)	24.21 $\pm$ 0.04 (15.66–48.69)	23.73 $\pm$ 0.03 (14.33–48.16)	22.69 $\pm$ 0.02 (17.19–26.68)	22.64 $\pm$ 0.02 (17.38–28.26)	23.58 $\pm$ 0.02 (17.57–32.81)	23.29 $\pm$ 0.02 (14.71–34.90)

Note: Numbers in parentheses list the minimum and maximum temperature for that site and season.

Abbreviation: SE, standard error.

<sup>a</sup>Conductivity probes were positioned at the bottom of the oyster farm post. The temperature probes included one at the top and one at the bottom of the oyster farm post.

Theil, Bryars, Tanner, & de Jong, 2004). *Ostrea angasi* populations are not monitored. The size of wild populations of *O. angasi* and *C. gigas* at each site is therefore unknown. Other bivalve species could also be hosts; Coffin Bay has large and growing populations of blue mussel (*Mytilus galloprovincialis* Lamarck, 1819) (unpublished data). Northern Hemisphere *M. galloprovincialis* held on a *B. ostreae*-affected oyster farm for 2 years were not infected when assessed by histology (Figueras & Robledo, 1994), but it is unknown whether *M. galloprovincialis* can become infected with *B. exitiosa*. Coffin Bay also has a large, commercially fished population of Vongoles (*Katelysia* spp.; see Dent, Mayfield, & Carroll, 2016). Streaky Bay has large populations of *P. bicolor* on the substrate of the oyster leases, and *P. bicolor* shells are a favoured habitat for *O. angasi* settlement (Crawford, 2016). Further investigation of the capacity of other bivalves to act as hosts or reservoirs for *B. exitiosa* could better inform the risk profile for farms.

Contaminants are associated with increased prevalence, mortality and disease in molluscs for a variety of pathogens (Morley, 2010). Mangrove systems capture and retain wastewater-borne contaminants and pollutants (Maiti & Chowdhury, 2013; Tam & Wong, 1995). Environmental contaminants were not measured in this study, but lower prevalence near mangroves in Cowell may be linked to lower environmental contamination at this site.

There were no differences in *B. exitiosa* infection intensities in *O. angasi* over one year; infection intensities of *O. angasi* varied seasonally but with the pattern differing between sites and meat:shell ratio. Higher infection intensities in *O. angasi* at Coffin Bay 2 and Streaky Bay were associated with higher meat:shell ratios in spring and summer, but with lower meat:shell ratios in autumn and winter. Hine (1991) found that *O. chilensis* in New Zealand had light *B. exitiosa* infections in spring and summer and heavy *B. exitiosa* infections in autumn. Buss et al. (2019b) found higher *B. exitiosa* infection intensities were associated with lower meat:shell ratios for harvest size 20- to 22-month-old *O. angasi* in winter. Meat:shell ratio is driven by complex oceanographic and climatic factors that vary seasonally between sites (Grangeré, Ménesguen, Lefebvre, Bacher, & Pouvreau, 2009; Rahman, Henderson, Miller-Ezzy, Li, & Qin, 2020). Oyster populations undergoing mortality caused by *Bonamia* spp. have poor condition including *O. angasi* (see Corbeil, Handler, & Crane, 2009) and *O. edulis* (see Rogan, Culloty, Cross, & Mulcahy, 1991). Meat:shell ratio may therefore be an indicator of *B. exitiosa* infection severity; poor condition may occur when oysters have advanced infection, and better condition may occur when oysters have earlier stages of *B. exitiosa* infection. Data from this study show oysters from some sites in autumn and winter were prone to lower condition and more advanced stages of *B. exitiosa* infection, but in spring and summer, oysters in better condition were associated with earlier stages of *B. exitiosa* infection. A longer study or a study with more sites may clarify these differences and reveal whether the variability is driven more by season or site.

Hine (1991) also found that prevalence and infection intensity of *B. exitiosa* infection in the gonad increased after spawning in

summer. *Ostrea angasi* spawn between mid-spring and early autumn (O'Sullivan, 1980) when they are over two years of age (Crawford, 2016) and brood larvae only when they are over 68-mm shell length (O'Sullivan, 1980). Oysters in this study were not old enough to be sexually mature and were not incubating larvae. Seasonal patterns in *B. exitiosa* infections may be more pronounced and consistent in mature *O. angasi* than in the juvenile animals in our experiment.

We found that temperature did not influence *B. exitiosa* prevalence or infection intensities in *O. angasi*. In other *Bonamia* sp.-host systems, the effect of temperature varies; higher water temperature (>20°C) is associated with increased *B. exitiosa* prevalence and infection intensities in *C. ariakensis* (see Audemard et al., 2014; Carnegie et al., 2008), but *B. ostreae* prevalence increased in *O. edulis* at lower water temperatures (10°C) (see Cochenne-Laureau & Auffret, 2002). The environment, parasite and host influence infection (Scholthof, 2007), but observational understanding of *B. exitiosa*-*O. angasi* infection dynamics is more useful for management than understanding the effects of individual environmental parameters. Infection dynamics could be better understood by monitoring *B. exitiosa* infection in *O. angasi* in several year classes over multiple years and would aid farm management decisions including timing of stocking, stocking density, site management and husbandry and timing harvest of *O. angasi*.

Survival assessment at the end of the experiment showed that survival after one year was low (17%–33%). *Bonamia exitiosa* infection intensities were lower (3.01–7.02 cells/heart smear) than described for *O. angasi* experiencing clinical *B. exitiosa* disease and mortality (15.40–24.20 cells/heart smear) by Buss et al. (2019a). It is therefore unlikely that oysters were dying from clinical *B. exitiosa* infection in this study. Buss et al. (2019b) surveyed farmed *O. angasi* and also found low infection intensity (heart smear intensity <4.36 cells/slide). Low *O. angasi* survival was not associated with *B. exitiosa* prevalence or infection intensity and was probably caused by sub-optimal management of the oysters. Streaky Bay had lower *O. angasi* survival than other sites, but oysters at this site were cultured in baskets with larger mesh size (6 mm vs. 3 mm), and the grower noted loss of oysters within the first season. *Ostrea angasi* can have high survival (52%–92%) when grown subtidally (Mitchell, Crawford, & Rushton, 2000) and also intertidally (>99%) (Li, Miller-Ezzy, Crawford, Gardner, & Deveney, 2019), but optimized approaches to maximizing survival in farmed *O. angasi* have not been identified.

In SA, *O. angasi* are mostly cultivated on intertidal leases in adjustable long-line systems designed for *C. gigas* (Nell, 2001; Wear et al., 2004). These systems are different to benthic, rack or hanging rope systems used to farm *O. edulis* in Europe (Héral & Deslous-Paoli, 1991). Our temperature data demonstrate that oysters experience extreme conditions in intertidal systems. *Crassostrea gigas* is tolerant to elevated water temperature (Rahman, Henderson, Miller-Ezzy, Li, & Qin, 2019) and can survive acute exposures of up to 34°C (Holliday, 1995), but 42°C water temperature for 1 hr caused 100% acute mortality (Rajagopal et al., 2005). Information on the thermal tolerance of *O. angasi* is limited; temperatures above 31°C increase mortality in *O. angasi* larvae (O'Connor, 2015), but

optimal and lethal temperatures for other *O. angasi* life stages have not been assessed. *Ostrea angasi* is a subtidal species (Edgar, 2008) and is likely to have a lower upper temperature tolerance than intertidal adapted *C. gigas*. Intertidal oyster farms are more prone to impacts of climate change, and a breeding programme could select traits that facilitate better production in challenging climates (Leith & Haward, 2010).

*Ostrea angasi* reefs are being restored across southern Australia (Gillies, Crawford, & Hancock, 2017) and are also at risk of *B. exitiosa* infection. *Bonamia exitiosa* transmission between oyster farms and restored reefs could create mutual disease risk. To date, however, Australian sites for reef restoration have been approved based on oceanographic modelling and ecosystem-based development principles to avoid increasing pathogen and biosecurity risks. *Ostrea angasi* farming and reef restoration could also co-expand if careful management is applied. Susceptible-infected parasite models (Bidegain et al., 2017) simulating *Perkinsus marinus* (Mackin, Owen, & Collier) Levine, 1978 infection dynamics between wild and farmed *Crassostrea virginica* Gmelin, 1791 show that farmed *C. virginica* could act as a pathogen sink by removing parasites if the farmed population was harvested before onset of clinical disease and shedding of parasites (Ben-Horin et al., 2018). Timing harvests may therefore provide an alternate management strategy and facilitate expansion of both *O. angasi* aquaculture and reef restoration, but an understanding of parasite uptake and shedding prior to onset of clinical signs and optimal harvest times would need to be established to develop practically implementable farm management.

Aquaculture and reef restoration could benefit from a programme to breed for resistance against *B. exitiosa*. In Rossmore, Ireland, an *O. edulis* breeding programme produces *B. ostreae*-resistant *O. edulis*. *Bonamia ostreae* prevalence decreased from 90% to <15% with negligible mortality in selected stock. This programme, however, was based on mass selection and breeding from survivors and took 30 years to achieve substantial *B. ostreae* resistance (Lynch, Flannery, Hugh-Jones, Hugh-Jones, & Culloty, 2014). If the Australian oyster industry chose to expand *O. angasi* production, a breeding programme incorporating an infection model to test susceptibility and molecular methods to identify markers of resistance could decrease time to achieving resistant stock. Such a programme would be expensive, and cost-benefits would need to be examined to justify the expense.

This is the first analysis of *B. exitiosa* infection dynamics in *O. angasi*. *Bonamia exitiosa* occurred at all our study sites, and although *O. angasi* from some sites had slower increases in *B. exitiosa* prevalence and may be more suited to *O. angasi* cultivation, *O. angasi* at all sites had >0.5 prevalence after one year. Intensification of *O. angasi* farming at Cowell and Streaky Bay may, however, change infection dynamics. Reef restoration projects also need to continue to be well planned to ensure that they are not developed where they can negatively interact with oyster farms. Pairing site-specific information with a programme to breed for resistance against *B. exitiosa* would aid expansion of *O. angasi* aquaculture.



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## CONFLICT OF INTEREST

The authors have no conflict of interest.

## DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available from the corresponding author upon reasonable request.

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