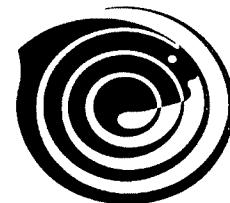


**TROPICAL AUSTRALIAN MICROALGAE AS LIVE FOOD  
FOR PEARL OYSTER (*Pinctada maxima*) SPAT**

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**F I S H E R I E S  
R E S E A R C H &  
D E V E L O P M E N T  
C O R P O R A T I O N**

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# 1. NON-TECHNICAL SUMMARY

95/131	<b>Tropical Australian microalgae as live food for pearl oyster, <i>Pinctada maxima</i>, spat</b>
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## OBJECTIVES:

1. To assess the suitability of Australian microalgae as live food source for young pearl oyster spat.
2. To prove that Australian tropical microalgae species are superior to the northern hemisphere species in the conditioning of spat of pearl oyster.

## NON TECHNICAL SUMMARY

Currently the Australian pearl oyster industry is using live food consisting of mixed microalgae originating from overseas. These species include *Pavlova lutheri*, *Isochrysis galbana* (clone T. Iso), *Chaetoceros mulleri*, *Tetraselmis* sp. and *Cryptomonas* sp. Since these species were isolated from overseas, they may not perform optimally under Australian tropical conditions. A suite of tropical Australian microalgae are available from the Northern Territory University (NTU) and CSIRO Microalgae Collections, so this is an opportune time to test the above hypothesis.

Pearl oyster (*Pinctada maxima* (Jameson)) spat were fed on 10 species of tropical Australian micro-algae, *Chaetoceros* sp. (CS256), *Cryptomonas* sp. (CRFI01), *Fragilaria pinnata* (PS5), *Fragilaria pinnata* (GOC01), *Isochrysis* sp. (NT14), *Isochrysis* sp. (PS11), *Nephroselmis* sp. (GOC52), *Nitzschia paleacea* (NT7), *Tetraselmis* sp. (TEQL01), *Rhodosorus* sp. (CS249), either as unialgal, 100 % rations or in 50:50 combinations. The Tahitian *Isochrysis galbana* (strain T. Iso.) was included for comparison.

Spat fed on a 100 % ration of either CS256, CRFI01, T. Iso, PS 11, TEQL01, or NT7 for 4-6 weeks showed 700 - 2800 per cent and 360 - 460 per cent better growth and survival, respectively, than those of the unfed control. Spat fed CRFI01, PS11, CS256 or T.Iso had significantly higher organic weight, carbohydrate, lipid, protein and polyunsaturated fatty acid

(PUFA) contents at harvest than unfed controls. Spat fed CS256, PS11 or T.Iso. had higher amounts of the PUFA docosahexaenoic acid (DHA) than the unfed control, while spat fed CS256 or CRFI01 had higher amounts of eicosapentaenoic acid (EPA) than the unfed control. Spat fed 50:50 combinations of CRFI01 + T. Iso, PS11 + T. Iso, CS256 + T. Iso, CRFI01 + PS11, or CS256 + PS11 all showed significantly better growth than unfed control in terms of shell length, dry weight and organic weight. In terms of shell length and dry weight, the combination T.Iso/CS256 showed the best results. There were positive correlations between spat carbohydrate content and spat shell length ( $r^2 = 0.717$ ), dry weight ( $r^2 = 0.874$ ) and organic weight ( $r^2 = 0.796$ ). There were also medium correlations between spat DHA and spat shell length ( $r^2 = 0.595$ ). There were no correlations between spat physical parameters and microalgal ash, carbohydrate, lipid, protein or individual fatty acids.

The Australian microalgae *Chaetoceros* sp. (CS256), *Cryptomonas* sp. (CFRI01) and *Isochrysis* sp. (PS11) showed consistent performance comparable to that of T.Iso, and therefore could be used in the rearing of pearl oyster spat instead of the northern hemisphere *Isochrysis* (T. Iso). This means that it is possible to use these tropical Australian isolates in combination with overseas species in the rearing of Pearl Oyster spat. The reduction in use of overseas microalgal isolates could also reduce the possibility that the microalgae may enter the environment.

Further work needs to be done on

- feeding trials involving combinations of different Australian microalgae at different densities;
- feeding trials on industrial scales;
- growout performance of spat fed tropical Australian microalgae; and
- supplementary feeding trials at growout sites.

**KEYWORDS:** Tropical microalgae, pearl oysters, *Pinctada maxima*, aquaculture, nutrition

## 2. BACKGROUND

Since 1990, the Pearl Oyster Industry has routinely spawned the silver-lip (or gold-lip) pearl oyster, *Pinctada maxima* (Jameson), on a commercial basis (Gervis and Sims, 1992; O'Sullivan, 1994; Rose, 1994). The hatchery-spawned and -reared spat are then placed in the sea for the grow-out production phase. In 1996, the price for hatchery produced spat was high at approximately \$US 0.1-0.14 per millimetre hinge length (Rose, R.A, unpublished data 1996, cited in Taylor et al., 1997a). Although hatchery methods for *P. maxima* are now well established (Rose and Baker, 1994), sporadic occurrence of high, unexplained mortalities experienced with spat recently entering ocean nursery facilities in the Northern Territory is still a problem (Rose, R.A. pers.comm.).

There are many hypotheses and complex reasons for the above mentioned mortalities. The high mortality of spat may be due to poor water quality or nutritional status of the spat, for example. Nutritionally, the high mortalities may be the result of two separated but interconnected events. Firstly, the high mortality may be due to the weak spat resulting from the union of poor quality gametes produced by the broodstock. Secondly, it may be due to the sudden switch from a hatchery diet (which consists of northern hemisphere phytoplankton) to a diet of local species present in the waters of the grow-out areas. In this work, we carried out preliminary investigation on the use of tropical Australian microalgae with the objective to test the second hypothesis.

The northern hemisphere microalgae that are currently being used as live food for pearl oysters and pearl oyster spat are: *Pavlova lutheri* and Tahitian *Isochrysis* sp. (T.Iso) for feeding *P. fucata* (Krishnan and Alagarswami, 1993; Numaguchi, 1994) and T.Iso, *Chaetoceros mulleri*, *Tetraselmis* and *Cryptomonas* for feeding *P. maxima* spat (Bellanger, 1995; Mills, 1997; Taylor et al., 1997b). A suite of tropical Australian microalgae are available from the NTU and CSIRO Microalgae Culture Collections, and it is an opportune time to test the above hypothesis. Moreover, the possible replacement of overseas microalgae species with Australian ones will simplify quarantine procedures for the disposal of the microalgae.

## 3. NEED

A bank of tropical microalgae isolated from the northern Australian waters by the microalgae group at the Northern Territory University (NTU) in Darwin and by the personnel at the CSIRO Marine laboratories in Hobart, Tasmania, is available at NTU and CSIRO. These microalgae species needed to be trialed to establish whether they could be used in the pre-conditioning of pearl oyster spat in the hatchery, so that the northern hemisphere phytoplankton could be replaced.

## 4. OBJECTIVES

- a. To assess the suitability of Australian microalgae as live food source for young pearl oyster spat.
- b. To prove that Australian tropical microalgae species are superior to the northern hemisphere species in the conditioning of spat of pearl oyster.

## 5. METHODS

### 5.1 Micro-algae

The following tropical Australian micro-algae were used in the feeding trials. They were chosen on the basis of size, ease of culturing and the taxonomic characteristics similar to the overseas species currently used by the industry (see background):

#### Cryptomonad:

*Cryptomonas* sp. (Isolated from Fitzroy Island, CRFI 01)

#### Diatoms:

*Chaetoceros* sp. (Isolated from Innisfail bay, CS 256)

*Fragilaria pinnata* Ehrenb.(Isolated from Port Steven, PS5)

*Fragilaria pinnata* Ehrenb.(Gulf of Carpentaria, GOC 01)

#### Green flagellates:

*Nephroselmis* sp. (Gulf of Carpentaria, GOC 52)

*Tetraselmis* sp. (MBPF Queensland, TEQL 01)

#### Rhodophyte;

*Rhodosorus* sp. (Isolated from Beaver Cay, CS 249)

#### Controls

*Isochrysis galbana* Parke (strain T. Iso.)

Unfed control.

Micro-algae were subject to a streak plate purifying procedure. Axenic conditions of the cultures was not achieved. However, the population of bacteria of the cultures has been kept to a low number as observed under phase contrast microscopy. The harmful bacteria, *Vibrio* sp., was not present in any cultures as tested using TCBS agar. Microalgae grown in 2 L batches until the end of the log phase, as determined by cell counts at intervals, were fed to the spat.

All microalgae were grown in 1.5 L sterile medium f in 2L flasks or 5L sterile medium f in 10L carboys at 27 °C, and 80-100  $\mu\text{moles/photons m}^{-2} \text{ s}^{-1}$  under 14h:10h light/dark cycles. 0.5 - 1% CO<sub>2</sub> was bubbled in the cultures at a rate of 0.4 L min<sup>-1</sup> as required to maintain pH of the cultures at 7.8- 8.0.

## 5.2 Experimental design of feeding trials

### 5.2.1 Feeding trials with single species of microalgae

The following methodology and experimental design were employed for the feeding trials:

Duration:	4-6 weeks
Location:	The Darwin Hatchery Project, Stokes Hill Wharf, Darwin
Water temperature:	Ambient (25 - 28 °C, minimum and maximum)
Water treatment:	The water used in feeding trials was pumped directly from the sea outside the hatchery to a settling reservoir. Once large particulates were settled out of the water, the water was then pumped through 10 µm sand filter, 5 µm cartridge filter and finally 1 µm cartridge filter. The 1µm filtered water was UV sterilised before use.
Water salinity:	35 +/- 1 ‰ as measured with a hand held refractometer.
Containers:	Spat were placed in 1.5L plastic containers containing 1L of above water, which was aerated with air stones. The containers were placed in a 10 tonne water bath to keep their water temperature constant.
Spat size:	6 - 15 mm
Number of spat per replicate:	15 - 20
Number of replicates:	3 - 6 replicates per treatment
Treatment:	10 new species of microalgae isolated from the Australian tropical waters (see species list above). These can be considered fixed factors which were chosen from a community of microalgae.
Controls:	Unfed and fed with <i>Isochrysis</i> Tahitian (T.Iso). Unfed control is used to identify new microalgae that cannot be used as feed, and T. Iso control is used to identify new microalgae that are superior or equivalent quality to those currently used as feed for Pearl Oyster spat.
Feeding history of spat:	Prior to experimentation, spat were fed a ration of T. Iso and <i>Chaetoceros mulleri</i> . After spat were placed in containers (see above), they were fed T.Iso for two days, then starved for two days before treatments were imposed. Unfed control ensures that the carry-over effect, if any, is accounted for.



At the beginning of each feeding trial, a sub population of spat of roughly equal shell length (range 0.610 - 0.770 mm, for all experiments, except the pilot experiment where spat range 1.520 - 1.555 mm were used) was selected from a population of recently spawned broodstock. For each experiment, spat were selected and were randomly assigned to each replicate (15 spat per replicate for growth experiments, 20 spat per replicate for experiments involving biochemical analysis). Homogeneity amongst the replicates was statistically tested before each feeding trial. Replicated buckets were floated in a 10,000 L circulated water bath kept in an outdoor shade house. The position of the replicated buckets in the water bath was also selected by a randomised design. Spat in 3 replicate buckets were harvested at time zero for initial biochemical analysis (dry weight, ash weight, organic weight).

Algal density was adjusted to 100 000 cells per mL (see 4.2.2) at the beginning of each day and dead spat were removed. The water in each bucket was changed every 2 days. To monitor growth, spat shell length was measured weekly. The biochemical composition (carbohydrate, lipid, protein and fatty acid composition) of 3-5 replicate buckets of spat and 2 L of each microalgal feed, was analysed at the end of the experimental period.

### 5.2.2 Effect of microalgal concentration on spat growth rate

To establish the optimum feeding density of microalgae for pearl oyster spat, the species *Isochrysis* PS11 was fed to spat at three concentrations ( $0.3 \times 10^5$ ,  $1 \times 10^5$  and  $3 \times 10^5$  cells per mL). The controls were unfed spat and spat fed T.Iso. There were five replicates per treatment.

The following parameters were measured at the beginning and the end of the experiments: shell length, dry weight, ash weight and organic weight (ash-free dry weight).

Survival and specific growth rate (SGR) were calculated using the following equations.

$$\text{Survival \%} = (N_t - N_{t_0})/N_{t_0} \times 100$$

$$\text{SGR} = \frac{\ln SL_t - \ln SL_{t_0}}{t - t_0} \times 100$$

Where  $SL_t$  is the shell length at time  $t$ ,  $SL_{t_0}$  is the shell length at time  $t_0$ ,  $N_t$  is the number of spat at time  $t$  and  $N_{t_0}$  is the number of spat at time  $t_0$ .

Specific growth rate was used so that results can be normalised between experiments.

### 5.2.3 Supplement feeding

The objective of this part of the study was to test the effect of combinations of promising species of Australian microalgae with or without *Isochrysis* (T.Iso.) on the growth and biochemistry of the spat. The following combinations (50:50 ratio based on organic weight of the algae) were used:

*Cryptomonas* CRFI01 + *Isochrysis* T. Iso  
*Isochrysis* PS11 + *Isochrysis* T. Iso  
*Chaetoceros* CS256 + *Isochrysis* T. Iso  
*Cryptomonas* CRFI01 + *Isochrysis* PS11  
*Chaetoceros* CS256 + *Isochrysis* PS11

Together with the above combinations, there were controls of unfed spat and spat fed T.Iso only. There were 8 replicates per treatment, with the duration of the experiment 28 days.

Parameters measured included shell length, dry weight, ash weight, ash-free dry weight, carbohydrate, lipid, protein and fatty acid composition. The first four parameters were measured at the beginning and the end of the experiment; the last four parameters, the end of the experiment only.

### **5.3 Biochemical analysis**

For chemical analysis, harvested microalgae and spat were rinsed with 0.5 M ammonium formate, freeze-dried and stored at -75 °C prior to chemical analysis.

Duplicate samples of microalgal cells or spat were analysed for ash (inorganic matter) total carbohydrate, total lipid, total protein, and fatty acid composition. Total carbohydrates were determined by the colorimetric method of Dubois et al., (1956) and total lipid was analysed gravimetrically after extraction with chloroform-methanol (2:1) by the method of Bligh and Dyer (1959). Total inorganic matter (ash) was determined gravimetrically after heating at 550 °C. Total nitrogen content was determined by Flow Injection Analysis after Kjeldahl digestion and total protein calculated from total nitrogen (%) x 6.25. (Renaud et al., 1994).

Lipids extracted from microalgae and spat were saponified then acidified, the fatty acids extracted into hexane and then converted to fatty acid methyl esters (FAME) by digestion with 14% BF<sub>3</sub>-methanol complex. The lipids from microalgal species were trans-methylated directly. FAME samples were reduced to 1 mL under a stream of nitrogen and analysed immediately or stored at -75 °C. Fatty acid compositions were determined using gas chromatography with a BP225, polar capillary column (50% cyanopropyl 50% phenyldimethylsiloxane, 25m x 0.2 mm id) and GC-MS using a non-polar column (DB-5MS, 30 m x 0.25 mm id) (Renaud et al., 1994). The shorthand notation used in fatty acid nomenclature is L:Bn-x where L = chain length, B = number of double bonds and (n-x) = position of the ultimate double bond from the terminal methyl group.

### **5.4 Statistical analysis**

Significance at 0.05 level between the results were tested by ANOVA (SuperANOVA), followed by paired comparisons using Scheffe's or by Student-Newman-Kuels (S-N-K) test. Equality of variance was checked using Barlett's test. Correlation analyses of spat growth parameters (shell length, dry weight, organic weight) and biochemical composition of both microalgal food species and spat (ash, carbohydrate, lipid, protein, individual fatty acids) were carried out.

## 6. RESULTS

### 6.1 Spat fed single species of microalgae

#### 6.1.1 *Effect of different algae food on growth and survival of pearl oyster spat*

The results described here (Table 1) are the summary of four separate feeding trials, extending from January 1995 to July 1996. At the end of the experimental period (4-6 weeks), spat fed CRFI01, CS256 showed better specific growth rates than the northern hemisphere microalga T. Iso. PS 11 performed at about 95% of the T. Iso control, whilst TEQL01 and CS249 showed 47% and 40%, respectively, below the T. Iso. control (Table 1). Those fed NT7, PS5, NT14, GOC01 showed poorer specific growth rates than that of unfed control (Table 1). It is interesting to note that GOC52 showed the best specific growth rate of all microalgae used, but also had the highest mortality of 93.3%.

#### 6.1.2 *Effect of different algal diets on the biochemistry of pearl oyster spat*

Spat fed microalgal species CRFI01, PS11, CS256 and T.Iso control had organic weights at harvest that were significantly higher than those of the unfed controls (Table 2). The shell weights (ash) at harvest of spat fed CRFI01, PS11 and T.Iso were also significantly higher than the control (Table 2).

Further biochemical analysis of spat found that animals fed the microalgal species CRFI01, PS11, CS256 and T.Iso control, had higher carbohydrate, lipid and protein contents at harvest than unfed controls (Table 3).

Spat fed CRFI01, PS11, CS256 or T.Iso also had significantly higher polyunsaturated fatty acid (PUFA) contents at harvest (33.7-36 % total fatty acids) than unfed controls (Table 4). Spat fed CS256, PS11 or T.Iso. had higher amounts of the PUFA docosahexaenoic acid (DHA) than the unfed control (Table 4), while spat fed CS256, CRFI01 or NT7 had higher amounts of eicosapentaenoic acid (EPA) than the unfed control (Table 4).

#### 6.1.3 *Effect of microalgal concentration on spat growth rate and proximate chemical composition*

Under the experimental conditions stated in section 4.2.1, the optimum microalgal concentration with respect to specific growth rate of spat was  $3 \times 10^5$  cells per mL. Spat fed T.Iso or PS11 at concentration  $3 \times 10^5$  cells per mL were found to have a significantly higher specific growth rate than spat fed either  $0.3 \times 10^5$  or  $1 \times 10^5$  cells per mL (Table 5).

Spat fed PS11 at concentration  $3 \times 10^5$  cells per mL had significantly lower carbohydrate and lipid contents than all other treatments. There were no significant differences in protein or ash contents.

**Table 1:** Specific growth rate (SGR) and survival of pearl oyster spat fed on different microalgae over a 4-6 weeks period. Figures in brackets = 1 standard deviation. \* = SD within replicates of the one experiment. 3-5 replicates per experiment. Each replicate had 15 spat. \*\* = calculated on basis of mean of all replicates. \*\*<sup>1</sup> = value during first 2 weeks only, all spat died after this period.

<b>Treatment</b>	<b>Mean SGR of spat (% increase /day)</b>	<b>Mortality (%)</b>	<b>No. of experiment</b>
<i>Nephroselmis</i> sp. (GOC52)	0.810** <sup>1</sup>	93.3 (0)*	1
<i>Cryptomonas</i> sp. (CRFI01)	0.734 (0.308)	9.39 (7.98)	3
<i>Chaetoceros</i> sp. (CS256)	0.710 (0.231)	22.07 (34.81)	3
<i>Isochrysis</i> Tahitian (T. Iso)	0.685 (0.181)	18.56 (25.64)	4
<i>Isochrysis</i> sp. (PS11)	0.650 (0.141)	0.67 (0.95)	2
<i>Tetraselmis</i> sp. (TEQL01)	0.322 (0.168)	26.4 (3.39)	2
<i>Rhodorus</i> sp. (CS249)	0.273**	84.47 (16.76)*	1
Unfed control	0.189 (0.098)	56.3 (36.58)	3
<i>Nitzschia paleacea</i> (NT7)	0.126 (0.048)	1.34 (1.90)	2
<i>Fragilaria pinnata</i> (PS5)	0.113**	39.97 (29.06)*	1
<i>Isochrysis</i> sp. (NT14)	0.05**	17.34 (12.09)	1
<i>Fragilaria pinnata</i> (GOC01)	-0.377**	77.7 (10.19)*	1

**Table 2:** The dry weight, organic weight and ash weight of spat at harvest.  
Data given as mean of 3 replicates with 15 animals per replicate.  
Standard deviation in parenthesis.

Algal species	Code	Dry weight (mg)	Organic weight (mg)	Ash weight (mg)
Unfed control		10.0 (3.1)	0.4 (0.2)	9.4 (2.9)
<i>Isochrysis</i> sp.	T.Iso	19.4 (5.1)	1.8 (0.5) *	17.5 (4.6) *
<i>Chrytomonas</i> sp.	CRFI01	20.6 (8.3)	1.9 (0.8) *	18.7 (7.5) *
<i>Isochrysis</i> sp.	PS11	16.7 (6.3)	1.5 (0.6) *	15.2 (5.8) *
<i>Chaetoceros</i> sp.	CS256	14.3 (3.6)	1.4 (0.4) *	12.9 (3.2)
<i>Nitzschia paleacea</i>	NT7	11.0 (2.5)	0.9 (0.3)	10.1 (2.3)
<i>Tetraselmis</i> sp.	TEQL01	11.0 (0.5)	0.9 (1.0)	10.0 (5.0)
<i>Isochrysis</i> sp.	NT14	8.9 (2.4)	0.9 (0.3)	8.0 (2.2)

\* Significantly higher than unfed control, at 95% level, but not significantly different from one another. ANOVA and Scheffe's test were used.

**Table 3:** Proximate biochemical composition of **(A)** tropical Australian microalgal feed species, Day 28 (n = 2), and **(B)** pearl oyster spat fed new microalgae (data given as mean; n = 3; pooled samples, each consisting of 60 spat).

Algal species	Code	Proximate chemical analysis (% dry wt)			
		Ash	Carbohydrate	Lipid	Protein
<b>(A) Microalgal food species</b>					
<i>Isochrysis</i> sp. control	T.Iso	10.6	10.9	25.3	35.8
<i>Cryptomonas</i> sp.	CRFI01	14.8	4.0	22.3	48.1
<i>Isochrysis</i> sp.	PS11	11.8	14.6	22.7	24.3
<i>Chaetoceros</i> sp.	CS256	20.9	6.1	16.6	40.1
<i>Nitzschia paleacea</i>	NT7	21.1	6.8	24.3	38.2
<b>(B) Pearl oyster spat</b>					
Unfed control		94.0	0.22	0.21	5.79
<i>Isochrysis</i> sp. control	T.Iso	90.2	0.41 *	0.42 *	7.23
<i>Cryptomonas</i> sp.	CRFI01	90.8	0.38 *	0.45 *	7.41 *
<i>Isochrysis</i> sp.	PS11	91.0	0.40 *	0.50 *	7.46 *
<i>Chaetoceros</i> sp.	CS256	90.2	0.37 *	0.36 *	7.00
<i>Nitzschia paleacea</i>	NT7	91.8	0.27	0.24	6.29

\* Significantly higher than unfed control, at 95% level, but not significantly different from one another. ANOVA and Scheffe's test were used.

**Table 4:** Fatty acid composition (% total fatty acids) of pearl oyster *Pinctada maxima* spat fed tropical Australian microalgae for 28 days (mean; n=2; pooled samples of 50 spat). Average coefficients of variation:  $\pm 9.0\%$  for FAME < 1 % total fatty acids;  $\pm 5.0\%$  for FAME > 2 % total fatty acids.

Fatty acid	Fatty acid composition of spat (% total fatty acids)						
	Unfed	Microalgae food species					
		TISO	CRFI01	PS11	CS256	NT7	NT14
<b>Saturated fatty acids (SFA)</b>							
14:0	2.8	3.5	2.5	3.8	1.1	2.3	3.3
16:0	17.4	18.1	20.6 *	18.8	18.6	22.8 *	26.7*
17:0	2.9	1.3	0.7	1.7	0.6	1.3	1.5
18:0	11.6	9.1	9.3	8.4	9.6	12.7	8.5
20:0	6.8	5.3	6.1	5.1	5.9	7.5	4.9
<b>Total SFA</b>	<b>41.5</b>	<b>37.3</b>	<b>39.2</b>	<b>37.8</b>	<b>35.8</b>	<b>46.5</b>	<b>44.9</b>
<b>Monounsaturated fatty acids (MUFA)</b>							
16:1n-7	3.4	2.1	5.1	1.1	1.1	1.7	1.8
18:1n-9	4.6	6.2	2.6	6.8 *	3.5	4.5	5.9
18:1n-7	3.6	3.8	5.1	3.1	2.8	4.2	2.3
20:1	1.5	2.4	1.0	2.3	1.8	1.1	2.1
22:1	7.8	9.1	5.3	8.7	10.3 *	7.0	6.9
<b>Total MFA</b>	<b>20.9</b>	<b>23.6</b>	<b>19.1</b>	<b>22.0</b>	<b>19.5</b>	<b>18.5</b>	<b>19.0</b>
<b>Polyunsaturated fatty acids (PUFA)</b>							
16:2n-7	2.2	1.1	1.7	0.9	1.1	2.5	1.8
16:3	-	-	-	-	-	-	-
18:2n-6	1.9	0.8	1.1	2.1	0.5	1.7	2.5
18:3n-3	-	2.1	0.9	0.9	-	-	0.6
18:4n-3	1.4	4.9 *	0.9	4.4 *	2.6	-	1.8
18:5n-3	-	-	-	1.0	-	-	-
20:2	-	-	-	-	-	-	-
20:4n-6	6.8	4.5	7.4	4.2	3.6	6.1	4.4
20:5n-3	1.5	1.0	6.3 *	0.9	3.8 *	4.7 *	1.6
22:2	3.2	1.7	4.4	1.7	2.3	3.7	0.9
22:4	1.5	1.3	1.0	1.1	0.9	1.3	1.2
22:5	3.8	4.8 *	2.1	4.8 *	5.4 *	3.2	5.6 *
22:6n-3	8.4	11.3 *	7.0	10.7 *	10.8 *	7.8	7.1
<b>Total PUFA</b>	<b>30.7</b>	<b>34.7 *</b>	<b>32.8</b>	<b>33.7 *</b>	<b>36.0 *</b>	<b>31.0</b>	<b>28.3</b>

\* = Significantly higher than the unfed control  
 - = Less than 0.1 % total fatty acids

**Table 5:** Specific growth rate (SGR) and proximate chemical analysis of pearl oyster spat fed different concentrations of microalgae (mean, n = 3; standard deviation in brackets).

Species	Concentration (cells per mL)	SGR (%)	Proximate chemical analysis (% dry wt)			
			Ash	Carbohydrate	Lipid	Protein
TISO	0.3 x 10 <sup>5</sup>	0.77 (0.05)*	89.3	0.94	0.51	4.61
	1 x 10 <sup>5</sup>	1.17 (0.19)	89.7	0.72	0.45	4.49
	3 x 10 <sup>5</sup>	1.32 (0.45)	88.0	0.76	0.43	4.59
PS11	0.3 x 10 <sup>5</sup>	0.63 (0.22)*	90.0	0.80	0.51	4.98
	1 x 10 <sup>5</sup>	1.13 (0.42)	89.7	0.81	0.56	6.96
	3 x 10 <sup>5</sup>	1.73 (0.12)	87.9	0.66 *	0.3 *	5.57
Unfed control: All died						

\* = significantly lower than other treatments. ANOVA and Scheffe's test were used.

## 6.2 Spat fed a combination of two microalgae species (supplementary feeding)

### 6.2.1 Effect of supplementary feeding on mortality

The spat showed very little mortality over the 28 days of experimentation. The unfed control showed the highest mortality (Mean = 8%, Standard deviation (SD) = 4%). The three treatments TISO, TISO/PS11, and TISO/CRFI01 showed no mortality, whilst the next three treatments TISO/CS256, PS11/CRFI01, and PS11/CS256 had a mean of 1% mortality (SD = 2%) each. These mortality figures are in contrast with those in the experiment conducted in 1996 when there was greater than 50% mortality for all treatments. The difference in mortality may be due to the difference in water quality of the two seasons attributed to the cyclonic activities in 1996 season.

### 6.2.2 Effects of treatments on shell length, dry weight and organic weight (ash-free dry weight, AFDW) of spat

Table 6 shows the results of supplementary feeding experiments. As compared to the shell length at the beginning of the experiment, unfed control did not show any growth while all other treatments did. Dry weight and organic weight showed the same trend. In terms of shell length and dry weight, the combination TISO/CS256 showed the best results (significance at 5% level).

In terms of organic weight, the three treatments TISO, TISO/CRFI01 and TISO/CS256 were the best compared to all other treatments. However, there is no significant difference in organic weight between the four treatments TISO, TISO/CRFI01, TISO/CS256 and



**Table 6:** Effects of supplementary feeding on shell length, dry weight and organic weight of spat. Data given as mean, with standard deviation in parenthesis; n = 3 replicates with 15 animals per replicate.

Treatment	Shell length (cm)		Dry weight (mg)		Organic weight (mg)	
Unfed control	0.731	(0.049)	8	(1)	0.6	(0.2)
TIso	0.942	(0.072)	17	(3)	1.8	(0.5)
TIso/PS11	0.928	(0.086)	15	(3)	1.6	(0.5)
TIso/CRFI01	0.941	(0.093)	17	(4)	1.8	(0.4)
TIso/CS256	0.975	(0.079)	18	(3)	1.8	(0.4)
PS11/CRFI01	0.920	(0.066)	15	(3)	1.6	(0.4)
PS11/CS256	0.915	(0.085)	16	(6)	1.7	(0.4)

**Table 7:** Proximate biochemical composition of (A) tropical Australian microalgal feed species (mean, Day 0 and Day 28, n=3), and (B) pearl oyster *Pinctada maxima* spat fed tropical Australian microalgae for 28 days (mean, n=3, pooled samples of 50 spat). Average coefficients of variation: ash  $\pm$  7.8 %; carbohydrate  $\pm$  8.0 %; lipid  $\pm$  7.5 %; protein  $\pm$  7.5%.

	Proximate analysis (% dry weight)			
	Ash	Carbohydrate	Lipid	Protein
<b>(A) Microalgae food species</b>				
TISO	11.3	15.2	27.9	44.2
TISO + PS11	13.1	12.8	26.9	43.0
TISO + CRFI01	13.9	10.2	22.1	54.2
TISO + CS256	13.6	13.1	24.5	43.7
PS11 + CRFI01	15.9	9.2	22.2	50.2
PS11 + CS256	15.6	12.1	24.6	39.7
<b>(B) Pearl oyster spat</b>				
Unfed control	92.4	0.28	0.89	6.5
TISO	88.4	0.44	0.77	7.7
TISO+ PS11	89.4	0.39	0.79	8.3
TISO + CRFI01	89.3	0.52*	0.56*	9.1
TISO + CS256	90.3	0.45	0.57*	8.4
PS11+ CRFI01	88.2	0.47*	0.55*	8.9
PS11 + CS256	87.6*	0.45	0.56*	9.0

\* Significantly higher than unfed control, at 95% level, but not significantly different from one another. ANOVA and Scheffe's test were used.

PS11/CS256. The ranking of the supplement feeding results in terms of organic weight is as follows:

$T_{Iso} = T_{Iso/CRFI01} = T_{Iso/CS256} = PS11/CS256 > PS11/CRFI01 = T_{Iso/PS11} > \text{unfed control}$

Based on shell length, dry weight and organic weight, it would appear that T<sub>Iso</sub>/CS256 may have the best potential as a live feed for pearl oyster spat.

### ***6.2.3 Effect of different combined algal diets on the biochemistry of pearl oyster spat***

Spat fed the combined microalgal species T<sub>Iso</sub>/CRFI01 or PS11/CRFI01 had significantly higher carbohydrate and lipid contents at harvest than unfed controls (Table 7). Significantly higher lipid content was also found in spat fed T<sub>Iso</sub>/CS256 or PS1/CS256 (Table 7). There was no significant difference in protein contents of all treatments.

Spat fed combinations T<sub>Iso</sub>/PS11, T<sub>Iso</sub>/CRFI01, T<sub>Iso</sub>/CS256 or PS11/CRFI01 had significantly higher percentages of polyunsaturated fatty acid (PUFA) at harvest (33.8-38.6 % total fatty acids), than the unfed control (Table 8). Spat fed the same four combinations or PS11/CS256 had significantly higher percentages of the PUFA docosahexaenoic acid (DHA) than the unfed control (Table 8). However, it is important to note that the absolute amount of fatty acid in the unfed control was very low.

### ***6.2.4 Correlation between spat parameters, spat biochemistry or microalgal biochemistry***

In feeding trials with single species of microalgae, there were positive correlations between spat carbohydrate content and spat shell length ( $r^2 = 0.717$ ), dry weight ( $r^2 = 0.874$ ) and organic weight ( $r^2 = 0.796$ ). There were also medium correlations between spat DHA and spat shell length ( $r^2 = 0.595$ ). In experiments investigating the effect of different combined algal diets, there were no correlations between spat physical parameters and microalgal ash, carbohydrate, lipid, protein (Table 7) or individual fatty acids (Table 9).

## **7. DISCUSSION**

The Australian Pearl Oyster Industry has been very successful in the spawning of broodstock and the rearing of larvae and spat in hatcheries. It has been able to meet the demand of farmers for spat used in the growout phase. Currently, the industry is using live food consisting of mixed microalgae originating from overseas. This work is a preliminary investigation of the possible use of Australian tropical microalgae as a replacement for overseas species. The significance of the work is two fold. It may provide a better diet resulting in better growth and survival of spat; and if overseas microalgae species are replaced by Australian species, the quarantine procedures could be simpler.

The response of Pearl Oyster spat to different treatments shows that the pretreatment period for the experiments was adequate, and the 4-6 weeks duration was similar to other published work (Mills 1997). The positive specific growth rates and the high standard deviation of the mortality of unfed control (Table 1) could be due to the variation of the carry-over effect of the prehistory of the spat. The density of spat in containers was adequate as shown by the

ratio of shell height (SH) to shell length (SL) during growth (0.69-0.75). Taylor et al. (1997) reported abnormal growth in *P. maxima* with SH:SL of 0.9-0.93.

The results show clearly that Australian tropical microalgae have mixed quality as far as usage as live feed for Pearl Oyster spat is concerned. *Cryptomonas* sp. (CRFI01), *Chaetoceros* sp. (CS256) and *Isochrysis* sp. (PS11) have the potential of being able to replace the northern hemisphere *Isochrysis* sp. (T. Iso) in the rearing of Pearl Oyster spat. Specific growth rates and mortality rates of Pearl Oyster spat reported here for the above microalgae were comparable to that reported by Mills (1997) who fed spat with T. Iso and *Chaetoceros muelleri*. However, *Nitzschia paleacea* (NT7), *Fragilaria pinnata* (PS5 & GOC51) and *Isochrysis* sp. (NT14), and to certain extent *Rhodosorus* sp. (CS249), were not suitable as a source of live food for Pearl Oyster spat.

Although the above results are preliminary, it is interesting to note that CRFI01, CS256 and GOC52 are better food for *Artemia salina* than T. Iso, and that NT14 performs worse than unfed control (Luong-Van Thinh et al. 1999). One of the reasons for the poor performance of NT14 may be due to volatile chemical compounds secreted by the microalga, such as an unidentified long-chain alkanone, which was detected by gas chromatography but not further characterized (Luong-Van Thinh et al. 1999).

It is clear from these results that spat fed Australian tropical microalgae CRFI01 and CS256 had higher concentration of 20:5n-3 than T. Iso and unfed control. Supplementary feeding experiments involving mixed diets of microalgae showed that spat constantly had higher concentrations of 22:6n-3 than those fed T. Iso alone and unfed control. These polyunsaturated fatty acids are essential to the growth and development of many aquaculture animals (Brown et al. 1989), and they could influence survival in the growout situation. On the other hand, gross energy and digestibility (see below) may play an important role. This hypothesis may be tested in the next phase of the project. It is worth mentioning here that gross energy content of dietary ingredients does not account for the ability of an organism to digest that energy (Fleming et al. 1995).

There were no clear correlations between the gross chemical composition of mixed feed of microalgae and growth of spat. The same results were found in supplementary feeding of juvenile Pacific Oysters, *Crassostrea gigas* (Brown & McCausland 1999). It is possible that mixed diet induces digestive enzymes formation in the gut of spat. This prepares spat to digest other live food present in the growout environment, which influences survival of the spat. Mixed diet also stimulates the formation of a balanced gut flora of spat, and enhances the digestive physiological ontological changes. This may lead to increased filtering rate, digestibility and physical quality of the spat, resulting in higher growth and survival rates.

In conclusion, this preliminary work shows the potential of several Australian tropical microalgae in the rearing of Pearl Oyster spat. Extensive further work is needed to confirm that the Australian microalgae are superior to the existing ones used by the industry.

## 8. BENEFITS

It is clear from the above results that the Australian microalgae *Chaetoceros* sp. (CS256), *Cryptomonas* sp. (CFRI01) and *Isochrysis* sp. (PS11) showed consistent performance and therefore could be used in the rearing of pearl oyster spat instead of the northern hemisphere *Isochrysis* (T. Iso). These results could be further refined by future research and implementation. The use of Australian microalgae would benefit the industry in terms of possible environmental problems created by imported microalgae species.

## 9. FURTHER DEVELOPMENT

It is clear from the above results that further work on the Australian microalgae *Chaetoceros* CS256 and *Cryptomonas* CRFI01 is warranted. Further supplementary feeding experiments can be conducted with these two Australian microalgae used as the base to which other microalgae can be combined. Different microalgal combinations (e.g., other than the 50:50 ratio used in this experiment) can also be used. Additional parameters should be included such as filtering rate, digestibility, and quality of the shell (form, shape, toughness), and the nutritive value (especially the energy content) of individual and mixed microalga feed (van Barneveld 1995). Larger scale feeding trials should then be carried out.

Monitoring of spat growth and survival in the growout phase should be planned to confirm the performance of spat reared on Australian microalgae. Supplementary feeding of spat under grow out conditions should also be planned.

## 10. CONCLUSION

The Australian microalgae *Chaetoceros* sp. (CS256), *Cryptomonas* sp. (CFRI01) and *Isochrysis* sp. (PS11) showed consistent performance comparable to that of T.Iso, and therefore could be used in the rearing of pearl oyster spat instead of the northern hemisphere *Isochrysis* (T. Iso). On the basis of the growth of spat, the Australian microalgal species were superior to T. Iso. However, on the basis of nutrition, further work is required to form a firm conclusion. The outcomes above are consistent with the objectives of the project, which state

1. To assess the suitability of Australian microalgae as live food source for young pearl oyster spat.
2. To prove that Australian tropical microalgae species are superior to the northern hemisphere species in the conditioning of spat of pearl oyster.

**Table 8:** Fatty acid composition (% total fatty acids) of pearl oyster *Pinctada maxima* spat fed tropical Australian microalgae for 28 days (mean; n=2; pooled samples of 50 spat). Average coefficients of variation:  $\pm 9.0$  % for FAME < 1 % total fatty acids;  $\pm 5.0$  % for FAME > 2 % total fatty acids.

Fatty acid	Fatty acid composition of spat (% total fatty acids)						
	Microalgal food species						
	Unfed	TISO	TISO/PS11	TISO/CRFI01	TISO/CS256	PS11/CRFI01	PS11/CS256
<b>Saturated fatty acids (SFA)</b>							
14:0	1.6	4.9 *	6.1 *	3.0	3.0	2.5	2.5
16:0	16.7	17.2	15.1	16.1	15.8	15.6	16.0
17:0	2.3	2.1	1.8	2.6	2.4	2.4	2.3
18:0	10.2	7.4	6.1	6.8	6.1	6.5	6.1
20:0	5.0	3.5	4.3	5.1	5.0	4.9	5.3
<b>Total SFA</b>	<b>35.8</b>	<b>35.1</b>	<b>33.4</b>	<b>33.6</b>	<b>32.3</b>	<b>31.9</b>	<b>32.5</b>
<b>Monounsaturated fatty acids (MUFA)</b>							
16:1n-7	2.6	3.5	2.1	2.6	5.8	2.5	5.6
18:1n-9	5.9	8.0 *	9.5 *	4.9	4.8	5.2	5.3
18:1n-7	2.5	3.3	3.2	2.5	3.1	2.2	3.0
20:1	4.4	2.7	3.6	2.9	2.5	3.1	2.6
22:1	9.5	8.5	8.3	8.1	6.7	8.7	7.3
<b>Total MFA</b>	<b>24.9</b>	<b>26.0</b>	<b>26.7</b>	<b>21.0</b>	<b>22.9</b>	<b>21.7</b>	<b>23.8</b>
<b>Polyunsaturated fatty acids (PUFA)</b>							
16:2n-7	1.1	2.6	2.0	3.9	3.7	5.4	4.0
16:3	-	1.9	1.7	4.3	3.2	2.0	2.8
18:2n-6	1.9	3.0	3.1	2.2	2.1	1.8	1.9
18:3n-3	0.1	1.5	1.5	3.5	0.2	3.6	0.3
18:4n-3	0.1	5.3 *	6.4 *	4.5	2.7	3.8	1.8
18:5n-3	-	-	0.9	0.8	1.0	0.2	0.2
20:2	2.1	1.2	1.1	0.1	1.1	1.0	1.1
20:4n-6	5.0	3.3	3.3	3.5	4.1	3.4	4.0
20:5n-3	3.1	0.9	0.6	1.6	2.1	1.9	2.4
22:2	3.2	0.1	1.2	1.5	2.4	1.2	3.0
22:4	1.4	1.3	0.9	0.3	0.3	1.4	0.6
22:5	3.4	3.5	3.7	4.4	3.2	4.3	2.9
22:6n-3	6.4	7.2	8.2 *	8.0 *	7.7 *	8.3 *	7.8 *
<b>Total PUFA</b>	<b>27.8</b>	<b>31.8</b>	<b>34.6*</b>	<b>38.6*</b>	<b>33.8*</b>	<b>38.3*</b>	<b>32.8</b>

\* = significantly higher than the unfed control

Other abbreviations as in Table 5

**Table 9:** Fatty acid composition (% total fatty acids) of mixed Australian tropical microalgal species fed to pearl oyster *Pinctada maxima* spat (mean, Day 0 and Day 28, n = 4). Average coefficients of variation:  $\pm 9.0\%$  for FAME < 1 % total fatty acids;  $\pm 5.0\%$  for FAME > 2 % total fatty acids.

Fatty acid	Fatty acid composition of microalgae (% total fatty acids)					
	TISO	TISO/PS11	TISO/CRFI01	TISO/CS256	PS11/CRFI01	PS11/CS256
<b>Saturated fatty acids (SFA)</b>						
14:0	29.9	28.1	16.0	21.7	14.5	20.3
16:0	11.3	10.6	13.3	18.4	12.7	17.8
18:0	0.5	0.6	1.2	0.7	1.2	0.7
<b>Total SFA</b>	<b>41.7</b>	<b>39.3</b>	<b>30.5</b>	<b>40.8</b>	<b>28.4</b>	<b>38.8</b>
<b>Monounsaturated fatty acids (MUFA)</b>						
16:1n-7	5.8	5.3	4.4	20.7	3.9	20.2
18:1n-9	10.8	11.2	6.6	5.8	7.0	6.2
18:1n-7	0.7	1.2	1.0	0.7	1.5	1.2
20:1	-	-	-	-	-	-
<b>Total MFA</b>	<b>17.3</b>	<b>17.7</b>	<b>12.0</b>	<b>27.2</b>	<b>12.4</b>	<b>27.6</b>
<b>Polyunsaturated fatty acids (PUFA)</b>						
<b>Polyunsaturated fatty acids (PUFA)</b>						
18:2n-6	3.7	4.3	2.4	2.1	3.0	2.7
18:3n-3	4.2	3.7	12.7	2.1	12.8	2.1
18:4n-3	16.1	15.7	20.1	8.1	20.3	8.2
18:5n-3	4.6	4.8	4.5	4.5	5.2	5.1
20:4n-6	-	-	0.7	0.4	0.7	0.4
20:5n-3	0.1	0.2	3.6	2.1	3.7	2.2
22:6n-3	3.9	2.9	3.2	1.3	3.7	1.8
<b>Total PUFA</b>	<b>32.6</b>	<b>31.6</b>	<b>47.2</b>	<b>20.6</b>	<b>49.3</b>	<b>22.5</b>

Abbreviations as in Table 5

## 11. RECOMMENDATIONS

Based on the results of this work, it is recommended that pearl hatchery operators could use the Australian microalgae *Chaetoceros* sp. (CS256), *Cryptomonas* sp. (CFRI01) and *Isochrysis* sp. (PS11) to supplement T. Iso in the rearing of small spat in the range of 0.600 - 1.500 cm shell length.

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## **Appendix 1**

### **Intellectual property**

No intellectual property can be placed on these microalgae since they are available at the NTU and CSIRO microalgal collections.

## **Appendix 2**

### **Staff**

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