

## The selection of an ideal diet for *Ostrea edulis* (L.) broodstock conditioning (part B)

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### Abstract:

Four microalgae species (*Rhodomonas salina*, *Thalassiosira weissflogii*, *Thalassiosira pseudonana* and *Pavlova lutheri*) were evaluated to estimate their potential as food for *Ostrea edulis* (L.) reproductive conditioning. Best ingestion and absorption were observed with *R. salina* (3.44 and 1.59 mg g<sup>-1</sup> h<sup>-1</sup>, respectively), followed by *T. pseudonana* (2.75 and 0.98 mg g<sup>-1</sup> h<sup>-1</sup>) and *P. lutheri* (2.40 and 0.91 mg g<sup>-1</sup> h<sup>-1</sup>). Oysters fed *T. weissflogii* exhibited the lowest ingestion and absorption values (1.40 and 0.68 mg g<sup>-1</sup> h<sup>-1</sup>). Proximate composition (proteins and carbohydrates) and lipid content (fatty acids and sterols) analysed in four main tissues (gonad, digestive gland, muscle and gills) also differed significantly with diet. Protein ranged from 355 mg g<sup>-1</sup> in the gonad of oysters fed *P. lutheri* to 837 mg g<sup>-1</sup> in gills of oysters fed *T. weissflogii*; whereas carbohydrates ranged from 17.5 mg g<sup>-1</sup> in gills of oysters fed *P. lutheri* to 271 mg g<sup>-1</sup> in gonads of oysters fed *R. salina*. An overall poor enrichment in total PUFAs across all diets masked some of their potential impact on nutrition. In gonad, however, the major polyunsaturated fatty acids (polar lipid fraction) were EPA (≈ 19% for oysters fed *T. weissflogii* and 14% for those fed *P. lutheri*) and DHA (17% for oysters fed *P. lutheri* and 15% for those fed *R. salina*). Sterol contents showed a clear transfer from food to oyster tissues except with *P. lutheri*, from which neither methylpavlovol nor ethylpavlovol (characteristic of Pavlophyceae) were detected in oyster tissues. Histological analysis showed that gametogenesis was active in oysters fed *R. salina* and *T. weissflogii*, whereas only low gonadic development occurred in unfed oysters or those fed *P. lutheri*. *R. salina* is accordingly highly recommended for *O. edulis* broodstock conditioning whereas *P. lutheri* should be excluded.

### Highlights

► We study the feeding assessment of four single microalgal species by flat oyster ► Best ingestion and absorption were observed with *R. salina* or *T. weissflogii* ► In gonad, the major polyunsaturated fatty acids (polar lipids) were EPA and DHA ► A transfer of sterols from food to oyster tissues occurred except with *P. lutheri* ► Gametogenesis was very active in oysters fed *R. salina* or *T. weissflogii*

**Keywords:** *Ostrea edulis*; Conditioning; Algal diets; Ingestion; Absorption; Biochemical composition

## 1. Introduction

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Broodstock conditioning is a key step in the process of rearing bivalves under standardized conditions. Its success has often been estimated in terms of the quality of bivalve eggs and larvae produced. Thus, initial egg lipid contents have been found to be positively correlated with either larval survival (e.g., *Mercenaria mercenaria* and *Crassostrea virginica*: Gallager and Mann, 1986; *Pecten maximus*: Le Pennec et al., 1991) or larval growth (*O. edulis*: Helm et al., 1973). Although temperature has been considered to be the main environmental factor regulating bivalve reproduction (e.g., *C. gigas*: Fabioux et al., 2005; *Mytilus galloprovincialis*: Fearman and Moltscchaniwskyj, 2010); feeding (i.e., the amount of food supplied) also seems to be an important factor for increasing fecundity (e.g., *C. gigas*: Chavez-Villalba et al., 2003; *Argopecten purpuratus*: Martinez et al., 2000). In contrast, the influence of the relative food value of different phytoplankton species (nutritional quality) on mollusc gonadic development has been very little explored, especially in *O. edulis*. The pioneer works of Frolov and Pankov (1992) and Millican and Helm (1994) provided relevant information in this field but only a few studies (Berntsson et al., 1997) have been carried out since. Indeed the appearance in France in the 1970-1980s of epizooties of *Marteilia refringens* (Comps, 1970) and *Bonamia ostreae* (Comps et al., 1980) and their progressive extension throughout Europe (see review in Laing et al., 2005) led to the collapse of *O. edulis* culture and research then focused more effort on recently introduced species (e.g., *C. gigas*: Helm and Millican, 1977; Robert et al., 1982, and *R. philippinarum*: Helm, 1990; Utting and Spencer, 1991). In Europe, particularly France and Spain, *O. edulis* remains an emblematic species and attempts to develop "resistant strains" have been made in both countries (Naciri-Graven et al., 1988; Montes et al., 2003; Lallias et al., 2010). Moreover, such interest in flat oyster cultivation is now increasing in France due to high *C. gigas* juvenile mortalities (Samain and McCombie, 2007; Pernet et al., 2010).

To allow *O. edulis* genetic improvement through selection, the reliability of hatchery methods for this species needs to be improved. As already pointed out, conditioning is an important step in hatchery production of molluscs and particular attention needs to be paid to flat oyster feeding during this stage because hatchery-conditioned broodstock has been found to have lower fecundity than wild stock (Helm et al., 1991). We had already made an initial study to look for an ideal diet for *O. edulis* (Gonzalez-Araya et al., 2010). This work compared four monospecific microalgal diets based on ecophysiological and biochemical approaches, and assumed that the best microalgae should be those that were highly ingested, digested, assimilated and efficiently allocated to the reproductive compartment. The present study was designed to provide complementary information by testing the influence of four more microalgae on *O. edulis* consumption, ingestion, assimilation and reproduction.

## 2. Material and methods

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The techniques used in this study were previously detailed in González-Araya et al. (2010), so only a brief outline will be given here.

### Experimental design

In August 2008, *O. edulis* aged 18 months ( $\approx$  5 cm length and 0.5 g flesh dry weight), originating from Bay of Quiberon (South Brittany, France) were submersed, at 5 m depth, for 1 month, in mesh bags tied to trestles in the Bay of Brest. They were then returned to the quarantine area of the Argenton hatchery, where they were maintained at 14 °C for an additional month, during which they were treated for a week with chloramphenicol at 8 mg l<sup>-1</sup> to limit any development of vibrios. Thereafter, seawater temperature was increased by 1 °C weekly and, at beginning of October 2008, the flat oysters were transferred to translucent 50-l tanks where they were distributed homogeneously (30 oysters per tank, corresponding to

an equivalent biomass of  $\approx 1$  kg total weight and 16 g dry flesh weight). During this pre-conditioning period oysters were fed a mixed diet of *T. Iso* and *Chaetoceros gracilis* used routinely in Argenton to feed most of mollusc at different stages of development (Ben Kheder et al., 2010). Triplicate tanks were set up for each of the four single diet species tested here. Oysters were maintained in a flow-through system at 19 °C and fed constantly at  $900 \mu\text{m}^3 \mu\text{l}^{-1}$  or unfed (only receiving continuously  $1 \mu\text{m}$ -filtered-seawater). Four different microalgae were tested as mono-specific diets at the same biovolume (measured daily and accordingly including variation in cell volume) : *Rhodomonas salina* (mean volumetric size  $160 \mu\text{m}^3$ , mean dry weight  $130 \text{ pg cell}^{-1}$ , strain CCAP 978/24), *Thalassiosira weissflogii* ( $900 \mu\text{m}^3$ ,  $250 \text{ pg cell}^{-1}$ , strain CCAP 1085/1), *Thalassiosira pseudonana* ( $40 \mu\text{m}^3$ ,  $35 \text{ pg cell}^{-1}$ , strain CCAP 1085/3) and *Pavlova lutheri* ( $40 \mu\text{m}^3$ ,  $20 \text{ pg cell}^{-1}$ , strain CCAP 931/1). The choice of these species was based on their frequency of utilization in different mollusc commercial hatcheries worldwide (Robert and Trintignac, 1997; Borowitzka, 1997). Ingestion and absorption of the different microalgae were studied according to Beiras et al. (1994) over six consecutive weeks. It was hypothesized that the microalgae species that was best absorbed by oysters represented the best potential diet; hence, we tested this hypothesis by examining nutrient biochemical allocation in the gonads in the polar fraction compared with other tissues to control potential reproductive specificity.

Microalgae were grown in standard batch culture, based on the step-by-step method (Robert et al., 2004). Cultures grown in 300-l cylinders were used as feed for oysters when they attained the late logarithmic phase after 3-5 days. Conway medium (Walne, 1966) was used at  $1 \text{ ml l}^{-1}$  except for *R. salina*, which was cultivated on a double dose of nutrients to improve growth.

### Ecophysiological measurements

Ingestion was estimated by measuring algal concentration twice a day using an electronic coulter counter (Multisizer 3) at the inlet ( $C_i$ ) and outlet ( $C_o$ ) of each tank with consumption  $C = C_i - C_o$ . For all diets, pseudofaeces production (PF) was  $< 10\%$  and considered as nil so that consumption  $\approx$  ingestion ( $C \approx I$ ).

Tanks were drained and oysters cleaned three times a week (Monday, Wednesday and Friday) and daily faeces production, established over precise 24-h periods, was accordingly measured on the two other days: Tuesday for *R. salina* and *T. weissflogii* and Thursday for the other two diets. Faeces samples were collected onto a  $450 \text{ }^\circ\text{C}$  pre-combusted GF/C (Glass Filter, type C : grade  $1.2 \mu\text{m}$ ) filter using a vacuum pump and washed with ammonium formate solution. Faeces total weight was measured after drying at  $75 \text{ }^\circ\text{C}$  and the Organic Matter fraction (% OM) calculated by the difference after combustion at  $450 \text{ }^\circ\text{C}$  for 4 h. The procedure was similar for microalgae, for which 25 ml of culture were filtered for each sample.

For each diet, the coefficient of variation was  $< 10\%$  and the data were accordingly pooled to express the mean faeces production over the entire experimental period. Under such conditions, absorption (A) was defined as  $A = I$  (Ingestion)  $\times$  ae (absorption efficiency) and ae was defined as :  $ae = 100 \times (\text{OM}_A - \text{OM}_F) / [(1 - \text{OM}_F) \times \text{OM}_A]$ , where  $\text{OM}_A$  is the microalgae relative organic content and  $\text{OM}_F$  the faeces relative organic content (Conover, 1966).

### Biochemical analysis

At the beginning and end of the experimental period, 15 oysters per feeding condition were dissected to sample four different organs separately: gonads (Gn), digestive gland (Dg), adductor muscle (Am) and gills (G). For each diet, three pools were prepared of each of the four organs, each pool containing the tissues of five oysters; these were then stored at  $-80 \text{ }^\circ\text{C}$  for a period of up to 6 months prior to analysis.

Aliquots of the homogenate were analysed separately for protein (Lowry et al., 1957), carbohydrate (Dubois et al., 1956), fatty acids and sterols (Folch et al., 1957). Fatty acids were analysed after transesterification with BF<sub>3</sub> according to Marty et al. (1992) and

Delaporte et al. (2006), whereas sterols were evaluated after transesterification with sodium methoxide (Eder et al., 1992; Soudant et al., 2000). Fatty acids and sterols were identified by comparing their retention time with standards (23:0 for FA and cholestane for sterols). In the present study, we initially only planned to report the fatty acids of the polar lipid fraction, as we did in our sister work (Gonzalez et al., 2010), because they correspond to real assimilation. However, we extended our analysis to some of the main fatty acids of the neutral lipid fraction because this additional data was useful to improve our understanding of the fate of these compounds, as mentioned in the discussion.

### Gametogenesis survey

At the beginning and end of the experimental period, 15 oysters per diet were frozen and stored at -80 °C until gonad histological analysis. Gamete activity was assessed using Mann's maturity index (Mann, 1979) to classify oysters into five categories on a scale of 0-4 ranging from inactivity (0) to spent (4).

### Statistical analyses

After logarithmic transformation ( $\log_{10} [x_i]$ ) of ingestion and absorption data, and angular transformation of percentage data by the function  $[\arcsin(\sqrt{x_i/100})]$  for biochemical composition, statistical analyses were carried out using SIGMAPLOT software (version 11.0). Significant differences were detected between the means at the 5% threshold using ANOVA and a *a posteriori* multiple comparison test of the means (Pairwise Holm-Sidak Method).

## 3. Results

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### Effects of food on physiological parameters

Cumulative mortality remained low in all diets (2%) meaning that they could be objectively compared.

The highest ingestion rate reached by *O. edulis* was  $3.44 \pm 0.22 \text{ mg g}^{-1} \text{ h}^{-1}$ , achieved when they were fed *Rhodomonas salina*. In contrast, *Thalassiosira weissflogii* led to low ingestion, with a mean value of  $1.40 \pm 0.17 \text{ mg g}^{-1} \text{ h}^{-1}$ . Oysters fed *T. pseudonana* or *Pavlova lutheri* showed intermediate positions, with mean ingestion rates of  $2.75 \pm 0.22$  and  $2.41 \pm 0.30 \text{ mg g}^{-1} \text{ h}^{-1}$ , respectively.

Similarly, flat oysters fed *R. salina* exhibited a significant higher absorption value ( $A = 1.62 \pm 0.04 \text{ mg g}^{-1} \text{ h}^{-1}$ ) than those fed the other three diets ( $p < 0.001$ ). Despite *R. salina* and *T. weissflogii* showed similar absorption efficiencies ( $ae = 47\%$  and  $51\%$  respectively), *R. salina* absorption was twice as high as that of *T. weissflogii* ( $A = 0.68 \pm 0.02 \text{ mg g}^{-1} \text{ h}^{-1}$ ). In contrast oysters fed *T. pseudonana* or *P. lutheri* showed similar absorption values and absorption efficiencies ( $A = 0.98 \pm 0.10$  and  $0.91 \pm 0.11 \text{ mg g}^{-1} \text{ h}^{-1}$ ,  $ae = 33-34\%$ , respectively). When flat oysters were supplied with feed composed of these two last microalgae, no significant differences were recorded in ingestion, absorption and assimilation values ( $p \gg 0.05$ ).

### Diet composition

Diatoms and flagellates differed in their fatty acid and sterol contents (Table 1). Whereas *R. salina* and *P. lutheri* were rich in 22:6(n-3), which represented 8 to 11% of total fatty acids, respectively (vs. 3.5 to 4.5% for the *Thalassiosira*). Diatoms were characterized by high EPA concentration (14.5 to 20.5%), but 20:5(n-3) content was surprisingly the highest in *P. lutheri*

(23.5% vs. 9.5% in *R. salina*). *R. salina* was also particularly rich in 18:2(n-6) (18% vs. 0.5 to 2.5% for the three other microalgae;  $p = 0.011$ ; Table 1). In contrast, *R. salina* was poor in 16:1(n-7), with only 0.7% vs. 16 to 20% for the other three diets and accordingly such overall differences were significant ( $p = 0.036$ ). However, no mean row differences were found by use of a multiple posteriori test. Arachidonic acid (AA; 20:4 (n-6)) was highly variable between species without any overall trend. Thus, with only 0.2% AA *T. weissflogii* was the microalga with the poorest level, contrasting with the other *Thalassiosira (pseudonana)*, which had the highest content (9.4%). AA represented 2.4% in *R. salina* but was a fifth of this amount in *P. lutheri* ( $\approx 0.5\%$ ). Lastly, with values ranging from 0.2 to 0.9, the DHA/EPA ratio did not fluctuate widely (Table 1).

With around 79% of total sterols, *T. weissflogii* and *T. pseudonana* were rich in 24-methylene-cholesterol compared with the two flagellates (vs. 0 to 2%). Despite lower contents (4.4 to 6%), cholesterol was also characteristic of the diatoms (vs. 0.4% for both flagellates) (Table 1). *R. salina* was characterized by a high brassicasterol content (97%); whereas, this sterol was inexistent in the other diets tested here. *P. lutheri* contained specific sterols, including methylpavlovol (36% of total sterols), ethylpavlovol (16%), methylporifera (14%) and  $\beta$ -sitosterol (12%), that were not detected in the other diets (Table 1).

### Effect of food on oyster biochemical composition

Mean gonad protein content increased from 335 mg g<sup>-1</sup> at the beginning of the experiment to 470 to 501 mg g<sup>-1</sup> at the end in oysters fed all diets except *P. lutheri*, which led to no significant rise (355 mg g<sup>-1</sup>). After 6 weeks, the highest gonad protein content was observed in oysters fed *R. salina* 501.5  $\pm$  178 mg g<sup>-1</sup>, but differences with the other diets were not significant ( $p = 0.43$ ); (Fig. 1a). Protein increase was also independent of diet in flat oyster muscle (802-815 mg g<sup>-1</sup> vs. 718 mg g<sup>-1</sup> initially;  $p = 0.63$ ), gills (692-837 mg g<sup>-1</sup> vs. 734 mg g<sup>-1</sup> initially;  $p = 0.32$ ) and digestive gland (476-608 mg g<sup>-1</sup> vs. 449 mg g<sup>-1</sup> initially;  $p = 0.18$ ) (Fig. 1a).

There was no enrichment in carbohydrate in oyster gonad, gills or digestive gland with any of the diets ( $0.59 > p > 0.13$ ) (Fig. 1b). Oysters did, however, show carbohydrate accumulation in muscle ( $p < 0.001$ ) with all the diets except *P. lutheri*: levels increased from an initial 13.9  $\pm$  2 mg g<sup>-1</sup> to 38.9  $\pm$  5 mg g<sup>-1</sup> with *R. salina*; whereas, they only reached 17.5  $\pm$  7 mg g<sup>-1</sup> with *P. lutheri* (Fig. 1).

Whatever tissues were analyzed, the main polar fatty acids found over all diets were 16:0, 18:0, 20:4 (n-6), 20:5(n-3), 22:2j and 22:6(n-3) which represented  $\approx$  50-65% of total fatty acids (gonad = 62%: Table 2), (digestive gland = 64%: Table 3), (muscle = 55%: Table 4) gills = 52%: Table 5). These specific fatty acids represented initially 65-67% of total FA regardless oyster tissues.

Neither total fa nor specific fa enrichment of the gonad was found (Table 2). Thus, no significant differences between diets were found for 16:0 ( $p = 0.99$ ), with values ranging from 18.3 to 20.2% after six weeks of conditioning, compared with 20.7% at the beginning of the experiment (Table 2). Similarly no significant 18:0, 20:4 (n-6), 20:5(n-3), 22:2j, 22:6(n-3), values fluctuations were noted over time ( $p = 0.83$ ;  $p = 0.10$ ;  $p = 0.88$ ;  $p = 0.89$ ;  $p = 0.98$  respectively: Table 2). In contrast, significant 20:2 (n-6) enrichment occurred when oysters were fed *R. salina* (1.3%) ( $p = 0.03$ ) (Table 2). It was the only fa exhibiting significant differences over time in the gonad when values were expressed as relative content. When considering absolute content and beyond the main fatty acids only a significant 20:4 (n-6) increment was noted in oysters fed *T. pseudonana*.

There was no specific fatty acid transfer to the gonad because the trends found for the main fatty acids in the gonad were similar to those in the digestive gland except for 20:4(n-6), which accumulated when oysters were fed *T. pseudonana* (Table 3). When considering absolute content in minor fatty acids, an enrichment in 20:2(n-6) occurred when oysters were fed *R. salina* (Table 3); whereas, a decrease in 22:5(n-6) was noted for those fed *T. weissflogii* and *T. pseudonana* contrasting with the enrichment found with *P. lutheri* (Table 3). Most of the other minor fatty acids in the digestive gland showed differences when considering absolute values but no clear trend can be seen (Table 3). A similar overall

situation was also found for the main fatty acids in muscle except for 16:0, whose absolute values decreased ( $p < 0.001$ ) with all diets (Tables 4), 20:4(n-6) and 22:2j which increased in oysters fed *T. pseudonana* and *T. weissflogii* respectively (Table 4). As for digestive gland some specific trends can be seen in minor fatty acids when considering absolute values. Thus an increase in 16:1(n-7), 18:1(n-7), 18:2(n-6) was noted in muscle when oysters were fed *T. weissflogii*, and *R. salina* contrasting with the decrease observed in 16:3(n-3) for all diets (Table 4). A similar overall situation was also found for the main fatty acids in gills except for 20:4(n-6) whose absolute value increased when oysters were fed *T. pseudonana* (Table 5). Similar trends as those reported in muscle can be seen in minor fatty acids when considering absolute values. Thus an increase in 16:1(n-7), 18:1(n-7), 18:2(n-6) was also noted in gills when oysters were fed *T. weissflogii*, and *R. salina*. (Table 5). In contrast a specific increase in 22:5(n-6) occurred with *P. lutheri*.

In the unfed oysters, no significant differences ( $p > 0.05$ ) were found in main polar fatty acid composition except for 16:0 in most of tissues at the end of the 6-week experiment (Table 6). Because no gonad development occurred in unfed oysters, it was impossible to separate this organ from the mantle and no data were therefore available on this tissue at the end of conditioning (Table 6).

When fed different microalgae, neutral fatty acid contents followed similar trends to polar fatty acids in all oyster tissues including gonad (Fig. 2). When oysters were fed *T. weissflogii* or *T. pseudonana*, an accumulation of 20:5(n-3) was noted but depletion of 22:6(n-3) was observed in oysters fed the same microalga. An accumulation of 20:4(n-6) was only observed in oysters fed *T. pseudonana* and, to a lesser extent, in oysters fed *R. salina*. No accumulation of 22:6(n-3) in gonad was observed in oysters fed *R. salina*, whatever fraction (neutral or polar) of lipids was analysed (Fig. 2).

Cholesterol allocation in all oyster tissues were not related to diet species because its content remained globally constant (Table 7) even with *T. weissflogii* and *T. pseudonana* supply, however rich in this sterol (Table 1). In contrast, in oysters fed *T. pseudonana*, 24-Methylen-cholesterol enrichment occurred in all tissues except digestive gland (Table 7). Such pattern was also found with *T. weissflogii*, however, to a lesser extent (Table 7). There was accordingly a clear food imprint for 24-Methylen-cholesterol that is the main sterol in both diatoms (80%: Table 1). When oysters were fed *R. salina*, brassicasterol enrichment seemed to occur in all tissues, doubling the concentrations, from  $\approx 20\%$  to 35-45% (Table 7) but such differences were not significant due to high variability in tissues samples of oysters fed *R. salina* (CV = 10-15%). On the other hand, campesterol allocation seemed to be more erratic, with a significant increase in gonad and gills of oysters fed *T. weissflogii* but an enhancement in the digestive gland of oysters fed *T. pseudonana* (Table 7). Lastly, when oysters were fed *P. lutheri*, no assimilation or incorporation of specific *P. lutheri* sterols (methylpavlovol and ethylpavlovol) was observed in any of the tissues which contrasted with the enrichment of  $\beta$ -sisterol in the gonad and gills (Table 7).

## Gonad development

At the beginning of experiment, 33% of oysters had recently spawned (stage 4: 33%) or were sexually inactive (stage 0: 50%), while only 17% were at the initiation stage of gametogenesis (stage 1) (Table 8). At the end of experiment, a clear effect of the diets on gametogenesis evolution was visible. Gametogenesis had occurred more rapidly when oysters were fed *R. salina* and *T. weissflogii*, than when they were fed *T. pseudonana* and *P. lutheri*. When oysters were fed *R. salina* at least 50% of the population became ripe, while 40% spawned (Table 8). When fed *T. weissflogii*, 61% of the population became ripe during the experiment, while 20% were still in development (Table 8); thus, on week 6, a total of 1.5 million larvae were collected from oysters fed *R. salina* and 1 million from oysters fed *T. weissflogii*. Among oysters fed *T. pseudonana* 48% were ripe, but 25% of the population were still inactive (Table 8). In contrast, among oysters fed *P. lutheri*, only 7% of individuals were ripe, 43% were developing, and inactive individuals represented 40% of the population (Table 8). No larval release was recorded during the experimental period in either of these last two diets.

## 4. Discussion

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### Impact of diets on physiological responses

The four microalgae differed significantly in their value for *Ostrea edulis* broodstock conditioning. First, they had a differential influence on response of flat oyster nutritional physiology. Ingestion of oysters fed *R. salina* ( $3.4 \text{ mg g}^{-1} \text{ h}^{-1}$ ) was double that of oysters fed *T. weissflogii* ( $1.4 \text{ mg g}^{-1} \text{ h}^{-1}$ ). This discrepancy could be explained by the difference in microalgae cell size between these species, which have a six fold volume difference:  $900 \mu\text{m}^3$  vs.  $160 \mu\text{m}^3$ . It may also explain the intermediate position occupied by *T. pseudonana* and *P. lutheri* ( $2.4$  to  $2.8 \text{ mg g}^{-1} \text{ h}^{-1}$ ), both of which have a lower cell size ( $40 \mu\text{m}^3$ ). Nevertheless, it has been reported that, among food particles of similar size, *Crassostrea gigas* larvae showed a target preference for *Chaetoceros calcitrans forma pumilum* (Rico-Villa et al., 2006). Moreover, it has been reported that external organic components of the diatom *Coscinodiscus perforatus* act as a major selection cue for *Pecten maximus* (Beninger and Decottignies, 2005). Lastly, epicellular molecules, such as lectins, have been showed to play a significant role in food selection of *Mytilus edulis* (Espinosa et al., 2010). Food selection seems to be an active process and, in the present case, *O. edulis* clearly shows a greater affinity preference for *R. salina*.

A similar trend was observed for absorption, but there was also high absorption efficiency with *T. weissflogii*. (51%). Although the oysters fed *R. salina* showed the best ingestion and absorption of the study, the values remained lower than those previously reported with *C. gracilis* and *S. marinoi* ( $\approx 5 \text{ mg g}^{-1} \text{ h}^{-1}$ ; González-Araya et al., 2010). Absorption efficiency (ae%) ranged from 33 to 51%, while those obtained with the four other microalgae tested in our sister study ranged from 19 to 46% (González-Araya et al., 2010). It is interesting to note that, whichever of the eight different species the microalgae was supplied, *O. edulis* absorption efficiency on single diets was lower than that achieved with a mixed diet, as reported by Savina and Pouvreau (2004) who observed 80-90% ae with *Glycymeris glycymeris* and *Phaphia romboïdes*. It is also notable that high ae has been reported in tropical mollusc species fed single-species diets: 70-90% ae in *Pinctada maxima* fed T- Iso or *Dunaliella primolecta* and 80-90% ae in *Pinctada margaritifera* fed the same diets (Yukihira et al., 1998).

### Impact of diets on biochemical responses

In bivalves, mantle, digestive gland and adductor muscle tissues are considered to serve for storage and translocation of protein and glycogen, lipids and proteins, respectively (Barber and Blake, 1985; Saucedo et al., 2002). Thus, the reproductive cycle of a bivalve can be divided into two processes: storage accumulation (generally glycogen) and gametogenesis (Pouvreau et al., 2006), which is achieved by using previously accumulated reserves and/or available food. The seasonal changes in energy storage and depletion in relation to gametogenesis are well documented (Bayne et al., 1982, Barber and Blake, 1985). Indeed, the adductor muscle of pectinids is known to be an important energy reserve site, and its utilization is associated with reproductive effort (Soudant et al., 1996). The utilization of carbohydrate from the *O. edulis* adductor muscle as source of energy was not clearly shown in the present study. Nevertheless, at the end of experiment, higher carbohydrate concentrations were recorded in gonads of oysters fed *R. salina*, which were also the oysters with the best gonadic development. For the other tissues analyzed, there were no statistical differences in either protein and carbohydrate contents during conditioning ( $p > 0.05$ ) regardless of diet. In the present study, conditioning was carried out from September 2008 to December 2008, and these results could be explained by specific environmental effects prior to oyster collection. In our experiment, despite a pre-conditioning period in which the oysters

were exposed to low temperature and food conditions just after harvest, it is likely that reserves from summer storage remained in most tissues and that these contributed to the initial proximate values measured. This idea is in agreement with results already reported in *O. edulis* (Gabbot and Walker, 1971) *C. gigas* (Delaporte et al., 2006) and *Pinctada mazatlanica* (Saucedo et al., 2002), for which the apparent glycogen (80% of carbohydrate contents) and protein increase were observed from January to July, with a second minor peak from August to October.

Polyunsaturated fatty acids (PUFAs) EPA (20:5(n-3)) and DHA (22:6(n-3)) have been shown to be essential for a wide variety of molluscs, as well as prawns and fish larvae (Volkman et al., 1989). Thus, phytoplankton species deficient in EPA and DHA have been reported to be poor food value for *C. gigas* spat (Langdon and Waldock, 1981). Moreover, it is likely that DHA plays a major structural and functional role in the cell membranes involved in oogenesis and embryogenesis (Soudant et al., 1996) in *P. maximus* broodstock. The specific role of 20:5(n-3) has been found to be related to energetic functions during embryogenesis of *Crassodoma gigantea* (Whyte et al., 1992) and *P. maximus* larvae (Delaunay et al., 1993; Soudant et al., 1998).

For *O. edulis* broodstock fed diets rich in PUFAs, a transfer of these fatty acids to their larvae was reported by Frolov and Pankov (1992). Lastly, many studies have documented relations between the fatty acid profile of the diet and that of the gonad or larvae contents (Knauer and Southgate, 1997; Caers et al., 2000; Flores-Vergara et al., 2004). The results of the present study differ in that the fatty acid profiles of main tissues did not show a clear correlation with the diets as it has been shown in our previous work with four other microalgae (Fig. 3; Gonzalez et al., 2010). These results were unexpected, because fatty acid composition in microalgae is generally specific. Here *R. salina* and *P. lutheri* exhibited high DHA contents, whereas the two *Thalassiosira* species showed high EPA values. Despite this specific composition, no accumulation of DHA above the initial contents was recorded in any organs of oysters fed *R. salina* or *P. lutheri*. Initial DHA contents seemed accordingly to be sufficient to cover the needs of the oysters, or had already reached a maximum level of accumulation in the gonads. Broodstock for this experiment were collected at the end of summer, and it is possible that the second peak of phytoplankton in the natural environment contributed to the PUFAs already present in oyster gonads in their initial state in our experiment. This idea is supported by data from the phytoplankton survey network (REPHY, 2011), which show a second peak of phytoplankton at the end of summer 2008 in Bay of Quiberon (Brittany, France).

Indeed it has been reported that proportions of 22:6(n-3) and 20:5(n-3) from neutral and polar lipids of artificially conditioned oysters are generally lower than those developing in the natural environment (Soudant et al., 1999). In our study, it is noteworthy that initial DHA contents at the beginning of autumn conditioning (16% : Fig. 2b) were similar to DHA values recorded at the end of spring conditioning in oysters fed T-Iso (16% vs 13.5% for spring initial value: Fig. 3b) known to be particularly rich in 22:6(n-3). Gonad needs apparently a determined DHA content which should be achieved during preconditioning period in the natural environment depending on this season at collection.

In contrast, an increase in EPA was only observed in gonads of oysters fed *T. weissflogii* when polar lipids were analyzed. When the neutral lipids were examined, however, EPA storage was found in all gonads except those of oysters fed *R. salina*. Such EPA accumulation during conditioning could be related to an insufficient initial 20:5(n-3) content (12%: Fig. 2b), which differs from the final values (17%: Fig. 3b) in all diets tested in this previous study (González-Araya et al., 2010).

Fatty acid 20:4(n-6) was incorporated in both neutral and polar lipids of the gonad, but this incorporation was only noted in oysters fed *T. pseudonana*. This fatty acid is a major precursor of prostaglandins, which may influence the reproduction process in molluscs (Osada et al., 1989). Maturation was however more active in oysters fed *R. salina* and *T. weissflogii*.

Sterols are known to play a variety of roles in living organisms, acting as structural components of cell membranes, steroid hormones and vitamin D precursors (Soudant et al., 2000). In our study, the relative sterol composition of the organs was significantly influenced by diet. Thus, for oysters fed *T. weissflogii* and *T. pseudonana*, 24-Methylen-cholesterol, was

efficiently transferred to the gonad. Similarly, brassicasterol was only accumulated in gonads of oysters fed *R. salina*.

In the present work, the sterols characteristic of *P. lutheri* (methylpavlovol and ethylpavlovol) did not show any accumulation in oyster gonads. Because the same trend was also observed in other tissues, we consider *P. lutheri* to be a poor microalgae species for *O. edulis* broodstock conditioning. This verdict is also based on the moderate physiological performances observed (moderate ingestion and absorption) and the poor gametogenesis development noted in oysters fed this species.

Despite the differing fatty acid and sterol composition of the diets, gametogenesis of *O. edulis* was globally successful, except when oysters were fed *P. lutheri* (as already mentioned) or not fed at all. This last point means that, although flat oyster tissue storage was clearly insufficient to support gametogenesis from an energetic point of view, initial stored reserves could have been crucial for essential requirements poorly covered by some of the diets. Our present study found the highest *O. edulis* reproductive performances in oysters fed *R. salina*, while our sister work recommended *Chaetoceros gracilis* or *Skeletonema marinoi* as the best from a comparison of four other species (González-Araya et al., 2010). T-Iso also showed some potential for *O. edulis* reproductive conditioning, although this was counterbalanced by low physiological performances with poor ingestion and absorption (González-Araya et al., 2010). *R. salina* offers similar biochemical characteristics as T-Iso but with higher physiological performances, which make it a better candidate. To balance diet and enhance flat oyster fecundity a mixed diet has been recommended (González-Araya et al., 2012). To improve reproductive performances a mixture based on *C. gracilis* (or *S. marinoi*) plus *R. salina* is highly recommended for *O. edulis* broodstock conditioning. The potential benefits of such microalgal assemblages for *O. edulis* broodstock fecundity and the quality of the larvae produced will be examined in a forthcoming paper.

## Conclusions

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*Rhodomonas salina* and *Thalassiosira weissflogii* are both as efficient feeds for *Ostrea edulis* conditioning.

1. Despite moderate physiological performances, *Pavlova lutheri* has no value for *O. edulis* conditioning due to low or inexistent transfer of dietary components.
2. The analysis of physiological and biochemical performances of flat oysters fed eight different microalgae species tested in separate experiments led us to recommend a mixed diet for *O. edulis* conditioning based on the association of *Chaetoceros gracilis* (or *Skeletonema marinoi*) plus *R. salina*.

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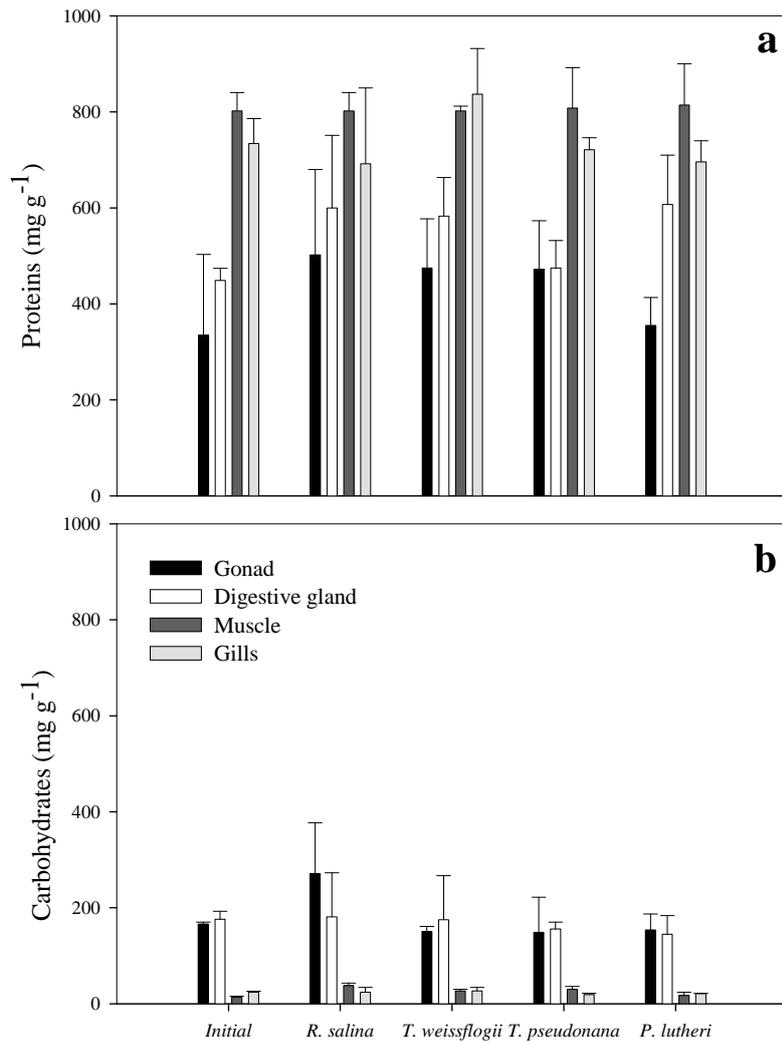
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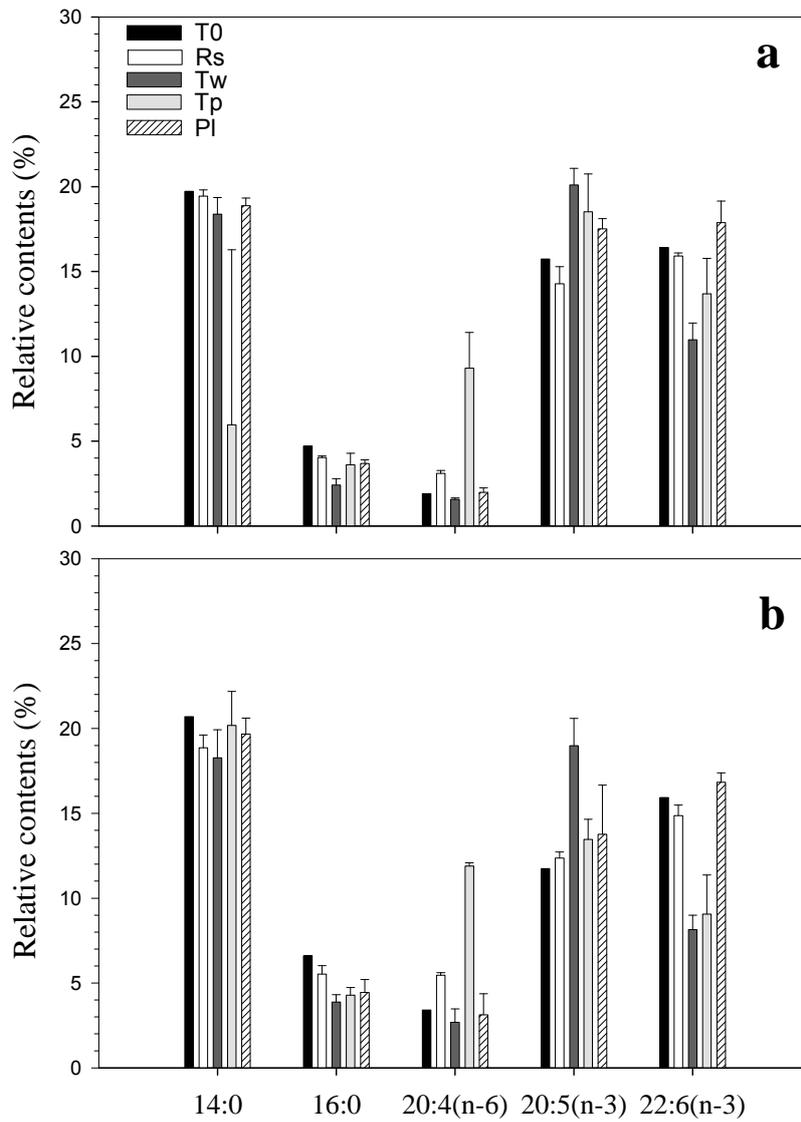
### **Figures**

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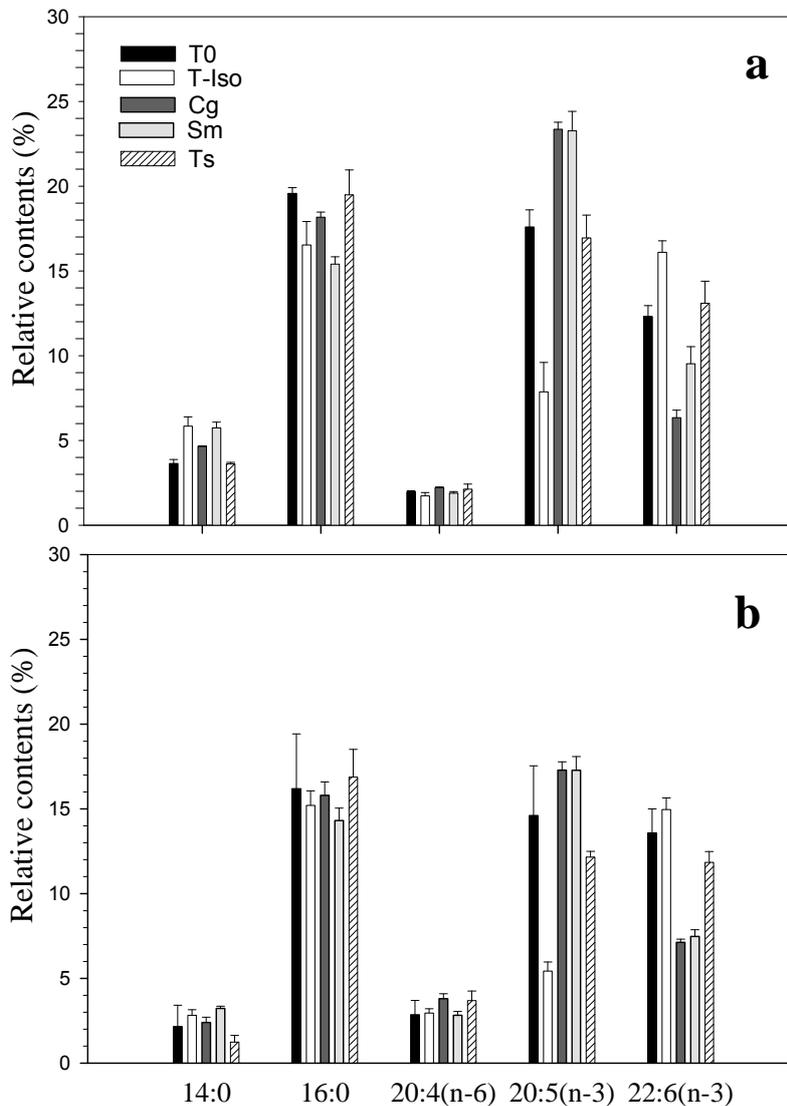
**Fig. 1**

**Figure 1.** Protein (a) and carbohydrate (b) contents of gonad, digestive gland, muscle and gills in European flat oyster *Ostrea edulis* (L.) broodstock, fed 4 microalgae species (values expressed in dw g<sup>-1</sup> tissue ± S.D.; n=3).



**Fig. 2**

**Figure 2.** Main fatty acid composition of neutral (a) and polar lipids (b) in gonad of *Ostrea edulis* broodstock according to diet: **T0**: Initial contents; **Rs**: *Rhodomonas salina*; **Tw**: *Thalassiosira weissflogii*; **Tp**: *Thalassiosira pseudonana* and **Pl**: *Pavlova lutheri*, expressed in relative contents (mean weight % of fatty acids  $\pm$  S.D.).



**Fig. 3**

**Figure 3.** Main fatty acid composition of neutral (a) and polar lipids (b) in gonad of *Ostrea edulis* broodstock according to diet: **T0**: Initial contents; **T-Iso**: *Isochrysis affinis galbana*; **Cg**: *Chaetoceros gracilis*; **Sm**: *Skeletonema marinoi*; **Ts**: *Tetraselmis suecica*, expressed in relative contents (mean weight % of fatty acids  $\pm$  S.D.).

## Tables

**Table 1.** Fatty acid and sterol composition of total lipids of *Rhodomonas salina*, *Thalassiosira weissflogii*, *Thalassiosira pseudonana* and *Pavlova lutheri* expressed in mean relative content (weight % of total polar fatty acids  $\pm$  S.D., n = 3).

**Table 2.** Fatty acid composition of the polar fraction in gonad of flat oysters fed mono-specific diets (weight % of total acids  $\pm$  S.D.). Values within the same line with a common

superscript letter, in the corresponding column (x' for absolute value, x for relative value), are not significantly different at  $p = 0.05$ .

**Table 3.** Fatty acid composition of the polar fraction in digestive gland of flat oysters fed mono-specific diets (weight % of total acids  $\pm$  S.D.). Values within the same line with a common superscript letter, in the corresponding column (x' for absolute value, x for relative value), are not significantly different at  $p = 0.05$ .

**Table 4.** Fatty acid composition of the polar fraction in muscle of flat oysters fed mono-specific diets (weight % of total acids  $\pm$  S.D.). Values within the same line with a common superscript letter, in the corresponding column (x' for absolute value, x for relative value), are not significantly different at  $p = 0.05$ .

**Table 5.** Fatty acid composition of the polar fraction in gill of flat oysters fed mono-specific diets (weight % of total acids  $\pm$  S.D.). Values within the same line with a common superscript letter, in the corresponding column (x' for absolute value, x for relative value), are not significantly different at  $p = 0.05$ .

**Table 6.** Main fatty acid composition of the polar fraction of the main tissues of unfed flat oysters (weight % of total acids  $\pm$  S.D.). \* indicate significant differences at  $p = 0.05$ .

**Table 7** Sterol composition of different tissues of *Ostrea edulis* fed mono-specific diets (weight %  $\pm$  S.D.). Values within the same line with a common superscript letter are not significantly different at  $p = 0.05$ .

**Table 8** Gonad development (%) in *Ostrea edulis* broodstock fed *R. salina*, *T. weissflogii*, *T. pseudonana* or *P. lutheri* for 6-weeks. Stages: 0, inactive; 1, early active; 2, late active; 3 ripe; 4, spent (see text for explanation: section Gonad development).

**Table 2**

Fatty acid	Oyster diets									
	Initial		<i>R. salina</i>		<i>T. weissflogii</i>		<i>T. pseudonana</i>		<i>P. lutheri</i>	
	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)								
14:0	0.16 ± 0.00	1.16 ± 0.88	0.24 ± 0.06	1.63 ± 0.16	0.30 ± 0.11	2.23 ± 0.36	0.33 ± 0.03	3.21 ± 1.39	1.13 ± 1.56	3.28 ± 1.59
16:0	2.81 ± 0.00	20.70 ± 1.25	2.79 ± 0.65	18.86 ± 0.75	2.60 ± 1.20	18.27 ± 1.66	2.22 ± 0.54	20.18 ± 2.00	4.98 ± 5.33	19.68 ± 0.94
18:0	0.89 ± 0.00	6.59 ± 0.45	0.80 ± 0.09	5.52 ± 0.50	0.53 ± 0.16	3.87 ± 0.44	0.49 ± 0.19	4.28 ± 0.45	1.00 ± 0.93	4.44 ± 0.76
16:1(n-7)	0.32 ± 0.00	2.35 ± 1.13	0.14 ± 0.01	0.99 ± 0.11	0.98 ± 0.49	6.82 ± 0.69	0.56 ± 0.06	5.43 ± 2.35	1.16 ± 1.69	3.10 ± 2.02
18:1(n-9)	0.31 ± 0.00	2.27 ± 1.25	0.21 ± 0.03	1.41 ± 0.12	0.24 ± 0.15	1.62 ± 0.48	0.15 ± 0.04	1.35 ± 0.07	0.74 ± 0.94	2.43 ± 0.65
18:1(n-7)	0.23 ± 0.00	1.72 ± 1.69	0.15 ± 0.03	1.00 ± 0.02	0.58 ± 0.35	3.93 ± 0.77	0.38 ± 0.05	3.52 ± 0.82	0.95 ± 1.23	3.02 ± 0.99
16:3(n-3)	0.37 ± 0.05	2.72 ± 1.06	0.40 ± 0.08	2.86 ± 1.02	0.38 ± 0.25	2.65 ± 1.73	0.27 ± 0.19	2.20 ± 1.13	0.41 ± 0.05	2.96 ± 1.90
18:2(n-6)	0.17 ± 0.03	1.22 ± 0.29	0.95 ± 0.37	6.33 ± 1.27	0.13 ± 0.08	0.83 ± 0.25	0.15 ± 0.02	1.39 ± 0.40	0.46 ± 0.63	1.34 ± 0.63
18:4(n-3)	0.16 ± 0.14	1.19 ± 0.45	0.27 ± 0.09	1.80 ± 0.30	0.08 ± 0.03	0.59 ± 0.05	0.18 ± 0.02	1.71 ± 0.49	0.48 ± 0.69	1.33 ± 0.78
20 :2i	0.05 ± 0.03	0.33 ± 0.00	0.02 ± 0.00	0.12 ± 0.02	0.05 ± 0.02	0.37 ± 0.13	0.03 ± 0.01	0.26 ± 0.02	0.06 ± 0.03	0.33 ± 0.14
20 :2j	0.03 ± 0.00	0.22 ± 0.01	0.01 ± 0.00	0.07 ± 0.02	0.04 ± 0.02	0.29 ± 0.03	0.03 ± 0.02	0.26 ± 0.13	0.04 ± 0.02	0.21 ± 0.09
20:2(n-6)	a' 0.03 ± 0.02	ab 0.22 ± 0.01	b' 0.19 ± 0.08	a 1.26 ± 0.27	a' 0.02 ± 0.01	ab 0.13 ± 0.06	a' 0.01 ± 0.00	b 0.09 ± 0.00	a'b' 0.07 ± 0.08	ab 0.25 ± 0.03
20:4(n-6)	a' 0.46 ± 0.09	3.41 ± 1.01	a'b' 0.80 ± 0.14	5.46 ± 0.15	a' 0.35 ± 0.08	2.68 ± 0.80	b' 1.33 ± 0.41	11.89 ± 0.19	a' 0.58 ± 0.37	3.13 ± 1.24
20:5(n-3)	1.59 ± 0.97	11.74 ± 2.33	1.82 ± 0.41	12.36 ± 0.36	2.71 ± 1.32	18.98 ± 1.62	1.48 ± 0.36	13.46 ± 1.19	4.05 ± 5.00	13.77 ± 2.89
22 :2i	a' 0.17 ± 0.02	1.22 ± 0.08	b' 0.10 ± 0.00	0.67 ± 0.12	b' 0.08 ± 0.02	0.56 ± 0.07	b' 0.06 ± 0.04	0.48 ± 0.26	a'b' 0.13 ± 0.02	0.92 ± 0.58
22 :2j	0.88 ± 0.31	6.52 ± 0.69	0.45 ± 0.02	3.15 ± 0.48	0.97 ± 0.38	6.94 ± 0.90	0.55 ± 0.34	4.52 ± 2.03	0.93 ± 0.34	5.86 ± 3.10
22:4(n-6)	0.09 ± 0.06	0.69 ± 0.09	0.08 ± 0.07	0.58 ± 0.50	0.05 ± 0.01	0.40 ± 0.17	0.11 ± 0.08	0.93 ± 0.44	0.07 ± 0.02	0.44 ± 0.24
22:5(n-6)	a'b' 0.15 ± 0.04	1.10 ± 0.09	a' 0.07 ± 0.05	0.54 ± 0.41	a' 0.05 ± 0.02	0.38 ± 0.05	a' 0.06 ± 0.04	0.47 ± 0.19	b' 0.45 ± 0.28	2.49 ± 1.06
22:5(n-3)	0.16 ± 0.10	1.15 ± 0.11	0.16 ± 0.01	1.11 ± 0.17	0.16 ± 0.03	1.22 ± 0.29	0.09 ± 0.05	0.73 ± 0.23	0.18 ± 0.15	0.85 ± 0.22
22:6(n-3)	2.16 ± 0.99	15.93 ± 2.89	2.18 ± 0.37	14.87 ± 0.62	1.15 ± 0.50	8.15 ± 0.85	1.07 ± 0.52	9.06 ± 2.30	4.27 ± 4.60	16.83 ± 0.56
<b>TO.MONO</b>	2.23 ± 0.57	16.45 ± 1.58	1.85 ± 0.34	12.58 ± 0.20	3.18 ± 1.51	22.25 ± 2.03	2.12 ± 0.55	19.19 ± 1.50	4.70 ± 5.61	16.62 ± 2.51
TO.(n-9)	0.46 ± 0.02	3.37 ± 1.36	0.35 ± 0.12	2.40 ± 0.71	0.33 ± 0.13	2.39 ± 0.00	0.24 ± 0.03	2.23 ± 0.56	0.96 ± 1.22	3.17 ± 0.83
TO.(n-7)	1.25 ± 0.09	9.23 ± 1.64	0.81 ± 0.12	5.54 ± 0.31	2.41 ± 1.16	16.85 ± 1.58	1.53 ± 0.32	14.00 ± 2.06	3.16 ± 3.87	10.85 ± 2.20
<b>TO.POLY</b>	6.87 ± 1.36	50.69 ± 3.58	8.49 ± 1.64	57.72 ± 0.71	6.67 ± 2.84	47.29 ± 2.87	5.74 ± 2.13	50.34 ± 4.06	13.11 ± 13.55	53.56 ± 3.35
TO.(n-4)	0.05 ± 0.00	0.37 ± 0.01	0.03 ± 0.01	0.18 ± 0.05	0.18 ± 0.12	1.23 ± 0.33	0.08 ± 0.00	0.72 ± 0.22	0.15 ± 0.21	0.45 ± 0.20
TO.(n-6)	0.93 ± 0.09	6.87 ± 0.99	2.26 ± 0.55	15.27 ± 0.74	0.65 ± 0.20	4.83 ± 0.91	1.77 ± 0.54	15.77 ± 0.38	1.82 ± 1.64	8.20 ± 1.50
TO.(n-3)	4.77 ± 1.25	35.14 ± 3.47	5.62 ± 1.06	38.25 ± 0.77	4.70 ± 2.11	33.04 ± 2.95	3.22 ± 1.17	28.29 ± 2.21	9.96 ± 11.24	37.54 ± 2.45
TO. NMI	1.13 ± 0.08	8.30 ± 1.87	0.58 ± 0.03	4.01 ± 0.63	1.14 ± 0.44	8.16 ± 1.07	0.68 ± 0.41	5.52 ± 2.42	1.15 ± 0.41	7.32 ± 3.89
<b>(n-3)/(n-6)</b>	a'b' 5.11 ± 2.33	5.11 ± 0.78	a' 2.51 ± 0.15	2.51 ± 0.15	b' 7.05 ± 1.76	7.05 ± 1.76	a' 1.80 ± 0.18	1.80 ± 0.18	a'b' 4.73 ± 1.26	4.73 ± 1.26
<b>22:6/20:5</b>	1.36 ± 0.78	1.36 ± 0.01	1.20 ± 0.08	1.20 ± 0.08	0.43 ± 0.05	0.43 ± 0.05	0.68 ± 0.21	0.68 ± 0.21	1.26 ± 0.24	1.26 ± 0.24
<b>22:5/20:4</b>	a' 0.32 ± 0.05	0.32 ± 0.08	b' 0.10 ± 0.07	0.10 ± 0.07	a'b' 0.15 ± 0.02	0.15 ± 0.02	b' 0.04 ± 0.02	0.04 ± 0.02	c' 0.80 ± 0.16	0.80 ± 0.16
<b>TOTAL</b>	13.56 ± 8.49	100.00 ± 0.00	14.71 ± 2.92	100.00 ± 0.00	13.99 ± 5.61	100.00 ± 0.00	11.23 ± 3.52	100.00 ± 0.00	25.53 ± 27.63	100.00 ± 0.00

**Table 3**

Fatty acid	Oyster diets									
	Initial		<i>R. salina</i>		<i>T. weissflogii</i>		<i>T. pseudonana</i>		<i>P. lutheri</i>	
	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)								
14:0	0.20 ± 0.03	1.37 ± 0.16	0.22 ± 0.11	1.06 ± 0.16	0.23 ± 0.04	1.39 ± 0.15	0.20 ± 0.02	1.64 ± 0.05	0.25 ± 0.05	1.61 ± 0.24
16:0	2.70 ± 0.13	18.12 ± 0.61	3.53 ± 1.26	17.50 ± 0.24	2.84 ± 0.12	17.51 ± 0.36	2.27 ± 0.18	18.17 ± 0.35	3.07 ± 0.62	19.53 ± 1.65
18:0	1.01 ± 0.06	6.75 ± 0.23	1.49 ± 0.55	7.37 ± 0.25	0.85 ± 0.03	5.24 ± 0.14	0.73 ± 0.04	5.84 ± 0.20	0.87 ± 0.19	5.50 ± 0.27
16:1(n-7)	a <sup>c</sup> 0.24 ± 0.02	b 1.64 ± 0.08	a <sup>r</sup> 0.11 ± 0.05	a 0.54 ± 0.08	b <sup>r</sup> 0.67 ± 0.11	b 4.12 ± 0.47	c <sup>r</sup> 0.33 ± 0.03	ab 2.66 ± 0.12	a <sup>c</sup> 0.22 ± 0.04	ab 1.41 ± 0.08
18:1(n-9)	a <sup>r</sup> 0.32 ± 0.01	2.16 ± 0.04	a <sup>b</sup> 0.22 ± 0.08	1.06 ± 0.06	a <sup>b</sup> 0.24 ± 0.02	1.50 ± 0.03	b <sup>r</sup> 0.15 ± 0.02	1.22 ± 0.04	a <sup>r</sup> 0.28 ± 0.05	1.79 ± 0.01
18:1(n-7)	a <sup>r</sup> 0.24 ± 0.01	ab 1.63 ± 0.02	b <sup>r</sup> 0.15 ± 0.06	b 0.72 ± 0.04	c <sup>r</sup> 0.53 ± 0.07	a 3.28 ± 0.25	a <sup>d</sup> 0.33 ± 0.04	a 2.62 ± 0.09	e <sup>r</sup> 0.00 ± 0.00	c 0.00 ± 0.00
16:3(n-3)	0.37 ± 0.02	2.50 ± 0.07	0.77 ± 0.33	3.75 ± 0.33	0.53 ± 0.20	3.32 ± 1.50	0.35 ± 0.04	2.80 ± 0.19	0.51 ± 0.08	3.24 ± 0.19
18:2(n-6)	a <sup>r</sup> 0.15 ± 0.01	a 1.03 ± 0.03	b <sup>r</sup> 0.91 ± 0.32	b 4.49 ± 0.08	a <sup>r</sup> 0.09 ± 0.02	a 0.58 ± 0.07	a <sup>r</sup> 0.10 ± 0.01	a 0.82 ± 0.09	a <sup>r</sup> 0.14 ± 0.02	a 0.91 ± 0.06
18:4(n-3)	a <sup>r</sup> 0.13 ± 0.02	0.85 ± 0.13	b <sup>r</sup> 0.27 ± 0.09	1.35 ± 0.05	a <sup>r</sup> 0.08 ± 0.01	0.50 ± 0.11	a <sup>r</sup> 0.14 ± 0.01	1.09 ± 0.04	a <sup>r</sup> 0.11 ± 0.02	0.68 ± 0.06
20:2i	a <sup>r</sup> 0.08 ± 0.01	0.51 ± 0.03	b <sup>r</sup> 0.03 ± 0.01	0.16 ± 0.01	a <sup>c</sup> 0.08 ± 0.01	0.51 ± 0.08	b <sup>r</sup> 0.04 ± 0.00	0.32 ± 0.02	a <sup>r</sup> 0.08 ± 0.02	0.52 ± 0.06
20:2j	a <sup>r</sup> 0.04 ± 0.00	0.26 ± 0.01	b <sup>r</sup> 0.01 ± 0.00	0.07 ± 0.01	a <sup>r</sup> 0.06 ± 0.01	0.35 ± 0.07	a <sup>r</sup> 0.05 ± 0.01	0.38 ± 0.03	a <sup>r</sup> 0.05 ± 0.01	0.30 ± 0.02
20:2(n-6)	a <sup>r</sup> 0.03 ± 0.00	a 0.21 ± 0.00	b <sup>r</sup> 0.23 ± 0.06	b 1.17 ± 0.11	a <sup>r</sup> 0.03 ± 0.01	a 0.18 ± 0.05	a <sup>r</sup> 0.01 ± 0.00	a 0.09 ± 0.01	a <sup>r</sup> 0.04 ± 0.01	a 0.25 ± 0.00
20:4(n-6)	a <sup>b</sup> 0.72 ± 0.01	b 4.86 ± 0.11	a <sup>c</sup> 1.33 ± 0.50	ab 6.58 ± 0.26	b <sup>r</sup> 0.44 ± 0.03	b 2.73 ± 0.10	c <sup>r</sup> 1.64 ± 0.20	a 13.07 ± 0.54	a <sup>b</sup> 0.77 ± 0.17	b 4.87 ± 0.20
20:5(n-3)	a <sup>b</sup> 1.71 ± 0.08	11.45 ± 0.36	a <sup>b</sup> 2.37 ± 0.82	11.76 ± 0.04	a <sup>r</sup> 2.86 ± 0.32	17.59 ± 0.92	b <sup>r</sup> 1.30 ± 0.16	10.36 ± 0.47	a <sup>b</sup> 1.74 ± 0.39	11.04 ± 0.51
22:2i	a <sup>r</sup> 0.19 ± 0.00	a 1.27 ± 0.00	b <sup>r</sup> 0.13 ± 0.04	ab 0.64 ± 0.03	c <sup>r</sup> 0.08 ± 0.00	b 0.47 ± 0.02	c <sup>r</sup> 0.07 ± 0.00	ab 0.56 ± 0.02	a <sup>r</sup> 0.17 ± 0.04	ab 1.09 ± 0.03
22:2j	a <sup>b</sup> 1.18 ± 0.02	7.93 ± 0.18	a <sup>r</sup> 0.75 ± 0.27	3.69 ± 0.15	b <sup>r</sup> 1.63 ± 0.02	10.12 ± 0.73	a <sup>r</sup> 0.99 ± 0.11	7.94 ± 0.33	b <sup>r</sup> 1.63 ± 0.36	10.33 ± 0.21
22:4(n-6)	a <sup>b</sup> 0.10 ± 0.01	0.66 ± 0.05	a <sup>r</sup> 0.13 ± 0.05	0.64 ± 0.03	b <sup>r</sup> 0.04 ± 0.00	0.24 ± 0.03	a <sup>r</sup> 0.14 ± 0.02	1.14 ± 0.11	a <sup>b</sup> 0.08 ± 0.02	0.48 ± 0.01
22:5(n-6)	a <sup>r</sup> 0.21 ± 0.00	ab 1.43 ± 0.06	a <sup>b</sup> 0.14 ± 0.06	a 0.71 ± 0.02	b <sup>r</sup> 0.05 ± 0.00	a 0.33 ± 0.01	b <sup>r</sup> 0.08 ± 0.01	a 0.62 ± 0.05	c <sup>r</sup> 0.60 ± 0.07	b 3.87 ± 0.29
22:5(n-3)	a <sup>b</sup> 0.17 ± 0.00	1.12 ± 0.04	a <sup>r</sup> 0.21 ± 0.08	1.01 ± 0.03	a <sup>b</sup> 0.16 ± 0.01	0.98 ± 0.02	b <sup>r</sup> 0.10 ± 0.01	0.81 ± 0.04	a <sup>b</sup> 0.11 ± 0.02	0.70 ± 0.06
22:6(n-3)	a <sup>b</sup> 2.40 ± 0.06	16.07 ± 0.46	b <sup>r</sup> 3.17 ± 1.07	15.75 ± 0.15	a <sup>r</sup> 1.48 ± 0.14	9.13 ± 0.35	a <sup>r</sup> 1.30 ± 0.09	10.40 ± 0.25	a <sup>b</sup> 2.45 ± 0.44	15.60 ± 0.74
<b>TO.MONO</b>	2.43 ± 0.08	16.32 ± 0.11	2.29 ± 0.83	11.31 ± 0.22	3.07 ± 0.26	18.93 ± 0.46	2.07 ± 0.17	16.58 ± 0.32	2.05 ± 0.40	13.04 ± 0.15
TO.(n-9)	a <sup>c</sup> 0.51 ± 0.03	3.41 ± 0.20	a <sup>c</sup> 0.55 ± 0.20	2.72 ± 0.06	a <sup>r</sup> 0.68 ± 0.05	4.23 ± 0.15	b <sup>r</sup> 0.21 ± 0.02	1.69 ± 0.03	b <sup>c</sup> 0.36 ± 0.06	2.30 ± 0.13
TO.(n-7)	a <sup>r</sup> 1.29 ± 0.04	8.64 ± 0.12	a <sup>r</sup> 0.94 ± 0.37	4.64 ± 0.22	b <sup>r</sup> 2.19 ± 0.21	13.47 ± 0.48	a <sup>r</sup> 1.42 ± 0.14	11.34 ± 0.15	a <sup>r</sup> 1.08 ± 0.21	6.88 ± 0.23
<b>TO.POLY</b>	7.90 ± 0.20	52.95 ± 1.02	11.74 ± 4.09	58.25 ± 0.75	8.03 ± 0.34	49.59 ± 1.00	6.58 ± 0.68	52.66 ± 1.15	9.04 ± 1.76	57.50 ± 1.59
TO.(n-4)	a <sup>r</sup> 0.06 ± 0.01	ab 0.40 ± 0.03	a <sup>r</sup> 0.03 ± 0.01	a 0.12 ± 0.02	b <sup>r</sup> 0.15 ± 0.02	b 0.92 ± 0.05	a <sup>r</sup> 0.06 ± 0.01	ab 0.46 ± 0.02	a <sup>r</sup> 0.04 ± 0.01	ab 0.27 ± 0.02
TO.(n-6)	a <sup>r</sup> 1.27 ± 0.02	ab 8.50 ± 0.13	b <sup>r</sup> 2.91 ± 1.02	a 14.43 ± 0.24	a <sup>r</sup> 0.69 ± 0.06	b 4.27 ± 0.09	a <sup>b</sup> 2.05 ± 0.24	a 16.41 ± 0.65	a <sup>b</sup> 1.69 ± 0.30	ab 10.75 ± 0.20
TO.(n-3)	a <sup>b</sup> 5.07 ± 0.16	34.01 ± 0.84	a <sup>r</sup> 7.88 ± 2.72	39.12 ± 0.32	a <sup>b</sup> 5.33 ± 0.27	32.90 ± 0.39	b <sup>r</sup> 3.32 ± 0.32	26.58 ± 0.59	a <sup>b</sup> 5.38 ± 1.04	34.23 ± 1.40
TO. NMI	a <sup>b</sup> 1.49 ± 0.03	9.97 ± 0.19	a <sup>r</sup> 0.92 ± 0.33	4.56 ± 0.18	b <sup>r</sup> 1.85 ± 0.02	11.45 ± 0.74	a <sup>r</sup> 1.15 ± 0.12	9.20 ± 0.33	b <sup>r</sup> 1.93 ± 0.42	12.24 ± 0.23
<b>(n-3)/(n-6)</b>	a <sup>r</sup> 4.00 ± 0.13	ab 4.00 ± 0.13	b <sup>r</sup> 2.71 ± 0.02	a 2.71 ± 0.02	c <sup>r</sup> 7.71 ± 0.25	b 7.71 ± 0.25	d <sup>r</sup> 1.62 ± 0.04	aa 1.62 ± 0.04	e <sup>r</sup> 3.18 ± 0.11	a 3.18 ± 0.11
<b>22:6/20:5</b>	a <sup>r</sup> 1.40 ± 0.03	1.40 ± 0.03	a <sup>r</sup> 1.34 ± 0.02	1.34 ± 0.02	b <sup>r</sup> 0.52 ± 0.02	0.52 ± 0.02	c <sup>r</sup> 1.01 ± 0.07	1.01 ± 0.07	a <sup>r</sup> 1.41 ± 0.06	1.41 ± 0.06
<b>22:5/20:4</b>	a <sup>r</sup> 0.29 ± 0.01	ab 0.29 ± 0.01	b <sup>r</sup> 0.11 ± 0.00	b 0.11 ± 0.00	b <sup>r</sup> 0.12 ± 0.00	b 0.12 ± 0.00	b <sup>r</sup> 0.05 ± 0.00	b 0.05 ± 0.00	c <sup>r</sup> 0.80 ± 0.08	a 0.80 ± 0.08
<b>TOTAL</b>	14.91 ± 0.40	100.00 ± 0.00	20.15 ± 6.95	100.00 ± 0.00	16.20 ± 1.00	100.00 ± 0.00	12.48 ± 1.07	100.00 ± 0.00	15.73 ± 3.10	100.00 ± 0.00

**Table 4**

Fatty acid	Oyster diets									
	Initial		<i>R. salina</i>		<i>T. weissflogii</i>		<i>T. pseudonana</i>		<i>P. lutheri</i>	
	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)
14:0	<sup>a,b</sup> 0.08 ± 0.01	1.02 ± 0.04	<sup>a</sup> 0.07 ± 0.01	0.70 ± 0.07	<sup>a,b</sup> 0.11 ± 0.02	1.14 ± 0.18	<sup>b</sup> 0.12 ± 0.02	1.21 ± 0.11	<sup>a,b</sup> 0.08 ± 0.02	0.94 ± 0.13
16:0	<sup>a</sup> 1.36 ± 0.05	16.30 ± 0.11	<sup>b</sup> 0.89 ± 0.01	9.45 ± 0.23	<sup>b</sup> 0.96 ± 0.04	10.04 ± 0.25	<sup>b</sup> 0.99 ± 0.14	10.32 ± 0.56	<sup>b</sup> 0.84 ± 0.12	9.89 ± 0.60
18:0	0.52 ± 0.00	6.22 ± 0.24	0.56 ± 0.02	5.96 ± 0.19	0.52 ± 0.06	5.43 ± 0.54	0.58 ± 0.08	5.99 ± 0.45	0.52 ± 0.06	6.22 ± 1.31
16:1(n-7)	<sup>a</sup> 0.07 ± 0.01	0.89 ± 0.10	<sup>a</sup> 0.07 ± 0.01	0.71 ± 0.13	<sup>b</sup> 0.28 ± 0.03	2.89 ± 0.28	<sup>c</sup> 0.17 ± 0.01	1.76 ± 0.08	<sup>a</sup> 0.08 ± 0.02	0.97 ± 0.13
18:1(n-9)	0.14 ± 0.01	1.64 ± 0.10	0.12 ± 0.01	1.29 ± 0.10	0.13 ± 0.02	1.35 ± 0.22	0.12 ± 0.00	1.24 ± 0.06	0.12 ± 0.01	1.39 ± 0.03
18:1(n-7)	<sup>a,b</sup> 0.09 ± 0.01	1.10 ± 0.06	<sup>b</sup> 0.06 ± 0.01	0.68 ± 0.12	<sup>c</sup> 0.21 ± 0.01	<sup>c</sup> 2.16 ± 0.13	<sup>c</sup> 0.17 ± 0.03	1.80 ± 0.13	<sup>a,b</sup> 0.10 ± 0.02	1.22 ± 0.18
16:3(n-3)	<sup>a</sup> 0.25 ± 0.13	3.03 ± 1.40	<sup>b</sup> 0.05 ± 0.02	0.55 ± 0.25	<sup>a</sup> 0.08 ± 0.02	0.87 ± 0.19	<sup>a</sup> 0.08 ± 0.02	0.86 ± 0.19	<sup>a</sup> 0.06 ± 0.01	0.75 ± 0.10
18:2(n-6)	<sup>b</sup> 0.06 ± 0.01	<sup>ab</sup> 0.72 ± 0.04	<sup>a</sup> 0.24 ± 0.03	<sup>a</sup> 2.55 ± 0.33	<sup>c</sup> 0.00 ± 0.00	<sup>b</sup> 0.00 ± 0.00	<sup>b</sup> 0.06 ± 0.01	<sup>ab</sup> 0.64 ± 0.06	<sup>b</sup> 0.04 ± 0.01	<sup>ab</sup> 0.51 ± 0.09
18:4(n-3)	<sup>a</sup> 0.05 ± 0.01	0.55 ± 0.14	<sup>a</sup> 0.05 ± 0.01	0.58 ± 0.09	<sup>b</sup> 0.02 ± 0.00	0.21 ± 0.00	<sup>a,c</sup> 0.06 ± 0.01	0.61 ± 0.04	<sup>b</sup> 0.02 ± 0.01	0.22 ± 0.08
20 :2i	<sup>a</sup> 0.02 ± 0.00	0.25 ± 0.01	<sup>a</sup> 0.02 ± 0.00	0.18 ± 0.02	<sup>b</sup> 0.04 ± 0.00	0.37 ± 0.04	<sup>c</sup> 0.03 ± 0.00	0.31 ± 0.04	<sup>c</sup> 0.03 ± 0.00	0.30 ± 0.01
20 :2j	<sup>a</sup> 0.02 ± 0.00	0.28 ± 0.00	<sup>b</sup> 0.03 ± 0.00	0.30 ± 0.03	<sup>c</sup> 0.05 ± 0.00	0.48 ± 0.04	<sup>c</sup> 0.05 ± 0.00	0.49 ± 0.01	<sup>b</sup> 0.03 ± 0.00	0.40 ± 0.01
20:2(n-6)	<sup>a</sup> 0.04 ± 0.00	<sup>ab</sup> 0.46 ± 0.00	<sup>b</sup> 0.13 ± 0.01	<sup>a</sup> 1.35 ± 0.11	<sup>a</sup> 0.03 ± 0.00	<sup>b</sup> 0.34 ± 0.02	<sup>a</sup> 0.03 ± 0.00	<sup>b</sup> 0.35 ± 0.02	<sup>a</sup> 0.03 ± 0.01	<sup>ab</sup> 0.39 ± 0.04
20:4(n-6)	<sup>a</sup> 0.29 ± 0.01	3.53 ± 0.04	<sup>a</sup> 0.40 ± 0.03	4.27 ± 0.37	<sup>a</sup> 0.29 ± 0.01	3.04 ± 0.05	<sup>b</sup> 0.71 ± 0.11	7.36 ± 0.53	<sup>a</sup> 0.30 ± 0.04	3.59 ± 0.15
20:5(n-3)	<sup>a,c</sup> 1.21 ± 0.06	14.54 ± 0.08	<sup>a,b</sup> 1.09 ± 0.04	11.47 ± 0.32	<sup>c</sup> 1.39 ± 0.01	14.50 ± 0.30	<sup>a,b</sup> 1.08 ± 0.08	11.30 ± 0.35	<sup>b</sup> 0.96 ± 0.15	11.31 ± 0.75
22 :2i	<sup>a</sup> 0.07 ± 0.00	0.89 ± 0.02	<sup>a</sup> 0.07 ± 0.00	0.72 ± 0.03	<sup>a</sup> 0.07 ± 0.00	0.68 ± 0.01	<sup>b</sup> 0.06 ± 0.00	0.68 ± 0.05	<sup>a</sup> 0.07 ± 0.00	0.88 ± 0.04
22 :2j	<sup>a,b</sup> 0.49 ± 0.00	5.90 ± 0.20	<sup>b</sup> 0.45 ± 0.02	4.79 ± 0.17	<sup>c,d</sup> 0.66 ± 0.02	6.89 ± 0.29	<sup>d,e</sup> 0.60 ± 0.04	6.31 ± 0.35	<sup>a,e</sup> 0.54 ± 0.04	6.42 ± 0.09
22:4(n-6)	<sup>a</sup> 0.06 ± 0.00	0.76 ± 0.03	<sup>b</sup> 0.08 ± 0.00	0.84 ± 0.03	<sup>a</sup> 0.05 ± 0.00	0.57 ± 0.03	<sup>b</sup> 0.09 ± 0.01	0.91 ± 0.15	<sup>a</sup> 0.06 ± 0.01	0.73 ± 0.05
22:5(n-6)	<sup>a</sup> 0.09 ± 0.00	1.09 ± 0.00	<sup>a</sup> 0.09 ± 0.00	0.99 ± 0.03	<sup>a</sup> 0.08 ± 0.00	0.81 ± 0.03	<sup>a</sup> 0.08 ± 0.01	0.86 ± 0.05	<sup>b</sup> 0.15 ± 0.02	1.77 ± 0.14
22:5(n-3)	<sup>a</sup> 0.16 ± 0.01	1.94 ± 0.07	<sup>a,b</sup> 0.15 ± 0.01	1.57 ± 0.12	<sup>a,b</sup> 0.14 ± 0.00	1.45 ± 0.02	<sup>a,b</sup> 0.14 ± 0.01	1.48 ± 0.06	<sup>b</sup> 0.13 ± 0.01	1.49 ± 0.03
22:6(n-3)	1.73 ± 0.07	20.78 ± 0.08	1.64 ± 0.05	17.34 ± 0.40	1.40 ± 0.05	14.64 ± 0.61	1.42 ± 0.15	14.82 ± 0.37	1.55 ± 0.20	18.24 ± 0.68
<b>TO.MONO</b>	<sup>a</sup> 1.36 ± 0.04	16.37 ± 0.17	<sup>a</sup> 1.36 ± 0.09	14.31 ± 0.82	<sup>b</sup> 1.85 ± 0.05	19.27 ± 0.25	<sup>c</sup> 1.63 ± 0.08	17.05 ± 1.06	<sup>a</sup> 1.30 ± 0.11	15.37 ± 0.34
TO.(n-9)	0.29 ± 0.01	3.50 ± 0.03	0.33 ± 0.02	3.53 ± 0.16	0.29 ± 0.06	3.02 ± 0.55	0.28 ± 0.01	2.89 ± 0.29	0.28 ± 0.01	3.33 ± 0.18
TO.(n-7)	<sup>a</sup> 0.82 ± 0.05	9.82 ± 0.20	<sup>a</sup> 0.70 ± 0.08	7.41 ± 0.80	<sup>b</sup> 1.23 ± 0.01	12.84 ± 0.24	<sup>c</sup> 1.06 ± 0.08	11.05 ± 0.67	<sup>a</sup> 0.77 ± 0.08	9.14 ± 0.14
<b>TO.POLY</b>	4.75 ± 0.32	57.09 ± 1.34	4.85 ± 0.06	51.23 ± 0.52	4.52 ± 0.06	47.16 ± 1.13	4.70 ± 0.42	48.95 ± 0.27	4.18 ± 0.55	49.35 ± 1.91
TO.(n-4)	<sup>a</sup> 0.02 ± 0.00	0.24 ± 0.04	<sup>a</sup> 0.01 ± 0.00	0.14 ± 0.02	<sup>b</sup> 0.05 ± 0.01	0.54 ± 0.09	<sup>c</sup> 0.03 ± 0.00	0.28 ± 0.04	<sup>a</sup> 0.01 ± 0.00	0.15 ± 0.03
TO.(n-6)	<sup>a</sup> 0.56 ± 0.02	6.79 ± 0.05	<sup>b</sup> 1.00 ± 0.06	10.57 ± 0.76	<sup>a</sup> 0.49 ± 0.01	5.10 ± 0.08	<sup>b</sup> 1.01 ± 0.13	10.53 ± 0.61	<sup>a</sup> 0.61 ± 0.09	7.22 ± 0.45
TO.(n-3)	3.56 ± 0.29	42.73 ± 1.65	<sup>b</sup> 3.27 ± 0.12	34.53 ± 1.03	<sup>c</sup> 3.17 ± 0.06	33.07 ± 0.96	<sup>d,e</sup> 2.91 ± 0.26	30.35 ± 0.30	<sup>a,e</sup> 2.88 ± 0.40	33.99 ± 1.60
TO. NMI	<sup>a,b</sup> 0.61 ± 0.01	7.32 ± 0.21	<sup>b</sup> 0.57 ± 0.02	5.99 ± 0.18	<sup>c,d</sup> 0.81 ± 0.02	8.41 ± 0.30	<sup>d,e</sup> 0.75 ± 0.05	7.79 ± 0.40	<sup>a,e</sup> 0.68 ± 0.06	8.00 ± 0.12
<b>(n-3)/(n-6)</b>	<sup>a,b</sup> 6.29 ± 0.29	6.29 ± 0.29	<sup>c</sup> 3.28 ± 0.33	3.28 ± 0.33	<sup>b</sup> 6.49 ± 0.28	6.49 ± 0.28	<sup>c</sup> 2.89 ± 0.19	2.89 ± 0.19	<sup>d</sup> 4.71 ± 0.15	4.71 ± 0.15
<b>22:6/20:5</b>	<sup>a</sup> 1.43 ± 0.01	1.43 ± 0.01	<sup>a</sup> 1.51 ± 0.02	1.51 ± 0.02	<sup>b</sup> 1.01 ± 0.03	1.01 ± 0.03	<sup>c</sup> 1.31 ± 0.07	1.31 ± 0.07	<sup>d</sup> 1.61 ± 0.06	1.61 ± 0.06
<b>22:5/20:4</b>	<sup>a</sup> 0.31 ± 0.00	0.31 ± 0.00	<sup>b</sup> 0.23 ± 0.03	0.23 ± 0.03	<sup>b</sup> 0.27 ± 0.01	0.27 ± 0.01	<sup>c</sup> 0.12 ± 0.01	0.12 ± 0.01	<sup>d</sup> 0.49 ± 0.03	0.49 ± 0.03
<b>TOTAL</b>	8.32 ± 0.36	100.00 ± 0.00	9.47 ± 0.11	100.00 ± 0.00	9.59 ± 0.14	100.00 ± 0.00	9.59 ± 0.82	100.00 ± 0.00	8.46 ± 0.80	100.00 ± 0.00

**Table 5:**

Fatty acid	Oyster diets									
	Initial		<i>R. salina</i>		<i>T. weissflogii</i>		<i>T. pseudonana</i>		<i>P. lutheri</i>	
	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)	Contents ( $\mu\text{g mg}^{-1}$ )	Rel. Contents (%)
14:0	0.19 ± 0.07	0.85 ± 0.04	0.13 ± 0.02	0.54 ± 0.05	0.24 ± 0.04	0.82 ± 0.04	0.22 ± 0.02	0.92 ± 0.06	0.20 ± 0.02	0.84 ± 0.06
16:0	3.79 ± 1.58	16.72 ± 0.15	1.84 ± 0.28	7.50 ± 0.51	2.34 ± 0.40	8.00 ± 0.19	2.02 ± 0.12	8.39 ± 0.24	1.90 ± 0.14	8.18 ± 0.49
18:0	1.41 ± 0.61	6.20 ± 0.04	1.28 ± 0.17	5.22 ± 0.33	1.22 ± 0.14	4.18 ± 0.10	1.07 ± 0.05	4.43 ± 0.08	0.97 ± 0.06	4.18 ± 0.32
16:1(n-7)	a <sup>i</sup> 0.19 ± 0.07	ab 0.87 ± 0.06	a <sup>i</sup> 0.12 ± 0.01	a 0.49 ± 0.02	b <sup>i</sup> 0.62 ± 0.09	b 2.13 ± 0.04	c <sup>i</sup> 0.37 ± 0.02	ab 1.54 ± 0.05	a <sup>i</sup> d <sup>i</sup> 0.23 ± 0.02	ab 0.98 ± 0.07
18:1(n-9)	0.39 ± 0.18	1.69 ± 0.05	0.27 ± 0.03	1.11 ± 0.01	0.33 ± 0.03	1.13 ± 0.06	0.25 ± 0.02	1.05 ± 0.11	0.31 ± 0.01	1.33 ± 0.05
18:1(n-7)	a <sup>i</sup> d <sup>i</sup> 0.26 ± 0.10	1.13 ± 0.03	a <sup>i</sup> 0.12 ± 0.02	0.50 ± 0.05	b <sup>i</sup> 0.65 ± 0.14	2.23 ± 0.13	c <sup>i</sup> d <sup>i</sup> 0.40 ± 0.03	1.68 ± 0.09	d <sup>i</sup> 0.33 ± 0.03	1.43 ± 0.09
16:3(n-3)	a <sup>i</sup> 0.80 ± 0.02	3.90 ± 1.73	b <sup>i</sup> 0.12 ± 0.01	0.47 ± 0.05	b <sup>i</sup> 0.23 ± 0.11	0.82 ± 0.44	b <sup>i</sup> 0.23 ± 0.04	0.94 ± 0.13	b <sup>i</sup> 0.15 ± 0.01	0.67 ± 0.04
18:2(n-6)	a <sup>i</sup> 0.14 ± 0.06	a 0.63 ± 0.02	b <sup>i</sup> 0.59 ± 0.12	b 2.40 ± 0.28	a <sup>i</sup> 0.00 ± 0.00	c 0.00 ± 0.00	a <sup>i</sup> 0.09 ± 0.01	ac 0.37 ± 0.01	a <sup>i</sup> 0.08 ± 0.01	ac <sup>i</sup> 0.36 ± 0.04
18:4(n-3)	a <sup>i</sup> 0.07 ± 0.02	0.35 ± 0.07	a <sup>i</sup> 0.07 ± 0.02	0.28 ± 0.06	b <sup>i</sup> 0.02 ± 0.00	0.08 ± 0.01	a <sup>i</sup> c <sup>i</sup> 0.07 ± 0.01	0.29 ± 0.03	b <sup>i</sup> d <sup>i</sup> 0.02 ± 0.00	0.10 ± 0.02
20 :2i	a <sup>i</sup> 0.12 ± 0.05	0.53 ± 0.01	a <sup>i</sup> 0.06 ± 0.00	0.24 ± 0.02	b <sup>i</sup> 0.23 ± 0.05	0.77 ± 0.09	a <sup>i</sup> 0.13 ± 0.01	0.56 ± 0.04	a <sup>i</sup> 0.14 ± 0.01	0.60 ± 0.07
20 :2j	a <sup>i</sup> 0.06 ± 0.02	abc 0.26 ± 0.00	a <sup>i</sup> b <sup>i</sup> 0.04 ± 0.01	a 0.16 ± 0.02	c <sup>i</sup> 0.17 ± 0.03	b 0.58 ± 0.02	c <sup>i</sup> d <sup>i</sup> 0.15 ± 0.00	bc 0.60 ± 0.03	e <sup>i</sup> 0.10 ± 0.00	abc 0.43 ± 0.01
20:2(n-6)	a <sup>i</sup> 0.05 ± 0.02	a 0.20 ± 0.00	b <sup>i</sup> 0.24 ± 0.04	b 0.97 ± 0.13	a <sup>i</sup> 0.06 ± 0.01	a 0.20 ± 0.01	a <sup>i</sup> 0.02 ± 0.00	a 0.10 ± 0.01	a <sup>i</sup> 0.04 ± 0.00	a 0.18 ± 0.02
20:4(n-6)	a <sup>i</sup> 1.47 ± 0.64	ab 6.49 ± 0.05	a <sup>i</sup> 1.98 ± 0.13	a 8.11 ± 0.24	a <sup>i</sup> 1.25 ± 0.18	b 4.29 ± 0.07	b <sup>i</sup> 3.34 ± 0.24	c 13.86 ± 0.53	a <sup>i</sup> 1.32 ± 0.07	a b 5.69 ± 0.18
20:5(n-3)	a <sup>i</sup> 2.22 ± 0.96	9.75 ± 0.07	a <sup>i</sup> 2.08 ± 0.26	8.50 ± 0.35	b <sup>i</sup> 4.21 ± 0.71	14.40 ± 0.55	a <sup>i</sup> 1.49 ± 0.10	6.18 ± 0.22	a <sup>i</sup> 1.71 ± 0.12	7.37 ± 0.36
22 :2i	a <sup>i</sup> b <sup>i</sup> 0.39 ± 0.17	1.72 ± 0.03	b <sup>i</sup> c <sup>i</sup> 0.22 ± 0.02	0.89 ± 0.08	b <sup>i</sup> c <sup>i</sup> 0.17 ± 0.02	0.58 ± 0.03	c <sup>i</sup> 0.16 ± 0.01	0.68 ± 0.03	b <sup>i</sup> c <sup>i</sup> 0.24 ± 0.01	1.04 ± 0.04
22 :2j	b <sup>i</sup> c <sup>i</sup> 2.28 ± 1.01	10.03 ± 0.16	a <sup>i</sup> b <sup>i</sup> 1.28 ± 0.13	5.25 ± 0.63	c <sup>i</sup> 3.37 ± 0.47	11.57 ± 0.55	b <sup>i</sup> c <sup>i</sup> 2.33 ± 0.02	9.66 ± 0.25	b <sup>i</sup> c <sup>i</sup> 2.35 ± 0.09	10.15 ± 0.40
22:4(n-6)	a <sup>i</sup> 0.20 ± 0.08	ab 0.89 ± 0.02	b <sup>i</sup> 0.27 ± 0.02	ab 1.09 ± 0.03	a <sup>i</sup> 0.14 ± 0.01	a 0.49 ± 0.03	c <sup>i</sup> 0.47 ± 0.03	b 1.95 ± 0.05	a <sup>i</sup> b <sup>i</sup> 0.17 ± 0.01	ac 0.73 ± 0.02
22:5(n-6)	a <sup>i</sup> 0.37 ± 0.15	ab 1.64 ± 0.03	a <sup>i</sup> 0.22 ± 0.02	a 0.91 ± 0.07	a <sup>i</sup> c <sup>i</sup> 0.17 ± 0.02	a 0.59 ± 0.03	a <sup>i</sup> 0.21 ± 0.00	a 0.86 ± 0.03	b <sup>i</sup> 0.85 ± 0.05	b 3.68 ± 0.29
22:5(n-3)	0.29 ± 0.12	1.28 ± 0.01	0.26 ± 0.02	1.07 ± 0.00	0.31 ± 0.05	1.06 ± 0.06	0.21 ± 0.01	0.85 ± 0.02	0.16 ± 0.01	0.68 ± 0.05
22:6(n-3)	3.70 ± 1.60	16.26 ± 0.13	3.79 ± 0.40	15.49 ± 0.36	3.24 ± 0.55	11.07 ± 0.61	2.61 ± 0.20	10.82 ± 0.48	3.23 ± 0.11	13.91 ± 0.18
<b>TO.MONO</b>	a <sup>i</sup> b <sup>i</sup> 3.25 ± 1.23	14.50 ± 0.77	a <sup>i</sup> 2.70 ± 0.31	11.04 ± 0.36	b <sup>i</sup> 4.67 ± 0.63	16.01 ± 0.24	a <sup>i</sup> b <sup>i</sup> 3.57 ± 0.07	14.81 ± 0.22	a <sup>i</sup> b <sup>i</sup> 3.02 ± 0.10	13.04 ± 0.25
TO.(n-9)	0.50 ± 0.14	2.30 ± 0.38	0.59 ± 0.07	2.42 ± 0.09	0.52 ± 0.04	1.79 ± 0.12	0.42 ± 0.03	1.75 ± 0.20	0.49 ± 0.01	2.10 ± 0.03
TO.(n-7)	a <sup>i</sup> 1.77 ± 0.75	ab 7.79 ± 0.03	a <sup>i</sup> 1.12 ± 0.08	a 4.58 ± 0.12	b <sup>i</sup> 2.99 ± 0.45	b 10.23 ± 0.13	a <sup>i</sup> b <sup>i</sup> 2.17 ± 0.09	ab 9.02 ± 0.14	a <sup>i</sup> 1.65 ± 0.09	ab 7.10 ± 0.27
<b>TO.POLY</b>	12.70 ± 5.15	56.25 ± 1.28	12.48 ± 1.11	51.12 ± 0.27	14.05 ± 2.06	48.14 ± 0.14	a <sup>i</sup> b <sup>i</sup> c <sup>i</sup> d <sup>i</sup> 11.87 ± 0.68	49.22 ± 1.16	11.57 ± 0.32	49.91 ± 0.34
TO.(n-4)	a <sup>i</sup> 0.08 ± 0.02	abc 0.39 ± 0.08	b <sup>i</sup> d <sup>i</sup> 0.02 ± 0.00	0.09 ± 0.01	a <sup>i</sup> c <sup>i</sup> 0.08 ± 0.03	0.29 ± 0.09	a <sup>i</sup> b <sup>i</sup> c <sup>i</sup> d <sup>i</sup> 0.05 ± 0.01	ac 0.21 ± 0.04	d <sup>i</sup> 0.03 ± 0.00	0.14 ± 0.01
TO.(n-6)	a <sup>i</sup> b <sup>i</sup> 2.28 ± 0.95	10.05 ± 0.08	a <sup>i</sup> c <sup>i</sup> 3.38 ± 0.31	a 13.85 ± 0.15	b <sup>i</sup> 1.69 ± 0.24	b 5.80 ± 0.09	c <sup>i</sup> 4.20 ± 0.27	ac 17.41 ± 0.54	a <sup>i</sup> b <sup>i</sup> 2.51 ± 0.06	ab 10.82 ± 0.18
TO.(n-3)	7.49 ± 2.92	33.28 ± 1.31	7.48 ± 0.79	30.63 ± 0.77	8.31 ± 1.28	28.48 ± 0.75	4.83 ± 0.37	20.03 ± 0.86	6.20 ± 0.24	26.71 ± 0.45
TO. NMI	a <sup>i</sup> b <sup>i</sup> 2.85 ± 1.26	12.54 ± 0.19	a <sup>i</sup> 1.59 ± 0.16	6.54 ± 0.75	b <sup>i</sup> 3.94 ± 0.56	13.51 ± 0.64	a <sup>i</sup> b <sup>i</sup> 2.77 ± 0.02	11.50 ± 0.33	a <sup>i</sup> b <sup>i</sup> 2.84 ± 0.09	12.23 ± 0.37
<b>(n-3)/(n-6)</b>	a <sup>i</sup> 3.31 ± 0.10	ab 3.31 ± 0.10	b <sup>i</sup> 2.21 ± 0.04	ac 2.21 ± 0.04	c <sup>i</sup> 4.91 ± 0.06	b 4.91 ± 0.06	d <sup>i</sup> 1.15 ± 0.02	c 1.15 ± 0.02	e <sup>i</sup> 2.47 ± 0.07	abc 2.47 ± 0.07
<b>22:6/20:5</b>	a <sup>i</sup> 1.67 ± 0.00	1.67 ± 0.00	e <sup>i</sup> b <sup>i</sup> 1.82 ± 0.03	1.82 ± 0.03	c <sup>i</sup> d <sup>i</sup> 0.77 ± 0.02	0.77 ± 0.02	a <sup>i</sup> b <sup>i</sup> d <sup>i</sup> 1.75 ± 0.07	1.75 ± 0.07	e <sup>i</sup> 1.89 ± 0.07	1.89 ± 0.07
<b>22:5/20:4</b>	a <sup>i</sup> 0.25 ± 0.01	ab 0.25 ± 0.01	b <sup>i</sup> c <sup>i</sup> d <sup>i</sup> 0.11 ± 0.01	a 0.11 ± 0.01	c <sup>i</sup> 0.14 ± 0.01	a 0.14 ± 0.01	d <sup>i</sup> 0.06 ± 0.00	a 0.06 ± 0.00	e <sup>i</sup> 0.65 ± 0.07	a 0.65 ± 0.07
<b>TOTAL</b>	22.69 ± 9.67	100.00 ± 0.00	24.41 ± 2.21	100.00 ± 0.00	29.17 ± 4.23	100.00 ± 0.00	24.10 ± 0.83	100.00 ± 0.00	23.19 ± 0.51	100.00 ± 0.00

**Table 6** les sanalysys

Fatty acid	Oyster organs									
	Gonad		Digestive gland				Muscle		Gills	
	Initial Contents (%)	Final Contents (%)	Initial Contents (%)	Final Contents (%)	Initial Contents (%)	Final Contents (%)	Initial Contents (%)	Final Contents (%)		
14:0	1.16 ± 0.88	n.d.	n.d.	1.37 ± 0.16	1.01 ± 0.22	1.02 ± 0.04	0.81 ± 0.06	0.85 ± 0.04	0.56 ± 0.07	
16:0	20.70 ± 1.25	n.d.	n.d.	18.12 ± 0.61	16.37 ± 0.90	16.30 ± 0.11	* 9.89 ± 0.22	16.72 ± 0.15	* 7.42 ± 0.25	
18:0	6.59 ± 0.45	n.d.	n.d.	6.75 ± 0.23	6.43 ± 0.06	6.22 ± 0.24	6.20 ± 0.23	6.20 ± 0.04	5.27 ± 0.42	
20:4(n-6)	3.41 ± 1.01	n.d.	n.d.	4.86 ± 0.11	6.72 ± 0.78	3.53 ± 0.04	4.05 ± 0.25	6.49 ± 0.05	6.78 ± 0.16	
20:5(n-3)	11.74 ± 2.33	n.d.	n.d.	11.45 ± 0.36	11.98 ± 0.60	14.54 ± 0.08	12.04 ± 0.60	9.75 ± 0.07	8.77 ± 0.42	
22:6(n-3)	15.93 ± 2.89	n.d.	n.d.	16.07 ± 0.46	17.41 ± 1.02	20.78 ± 0.08	19.00 ± 1.24	16.26 ± 0.13	15.84 ± 0.65	
<b>Σ PUFA</b>	<b>50.69 ± 3.58</b>	<b>n.d.</b>	<b>n.d.</b>	<b>52.95 ± 1.02</b>	<b>58.53 ± 2.06</b>	<b>57.09 ± 1.34</b>	<b>50.77 ± 0.07</b>	<b>56.25 ± 1.28</b>	<b>51.38 ± 0.39</b>	

Table 7

	Oyster diets				
	Initial	<i>R. salina</i>	<i>T. weissflogii</i>	<i>T. pseudonana</i>	<i>P. lutheri</i>
<b>Gonad</b>					
Brassicasterol					ab
	ab 19.7 ± 1.2	a 44.2 ± 5.0	b 8.7 ± 1.0	b 8.8 ± 0.9	12.7 ± 1.0
Cholesterol	36.3 ± 1.1	24.5 ± 3.6	26.1 ± 0.5	23.0 ± 0.3	29.3 ± 2.5
Campesterol	a 2.2 ± 0.2	a 1.7 ± 0.3	b 13.5 ± 0.8	ab 5.2 ± 0.3	ab 5.2 ± 0.7
24-Methylen-cholesterol	a 14.0 ± 0.8	a 11.1 ± 0.6	29.9 ± 0.9	b 48.0 ± 2.4	a 8.9 ± 2.1
Stigmastérol	2.6 ± 0.0	5.9 ± 0.8	2.4 ± 0.5	0.3 ± 0.5	6.1 ± 1.0
Isofucosterol	2.5 ± 0.2	1.0 ± 0.2	2.5 ± 0.3	1.6 ± 1.4	2.1 ± 0.7
MethylPorifera	1.8 ± 0.2	1.1 ± 0.2	1.2 ± 0.1	1.4 ± 0.8	1.4 ± 0.1
β-sitosterol	ab 3.2 ± 0.4	a 1.8 ± 0.2	ab 3.5 ± 0.1	a 2.2 ± 0.7	b 22.7 ± 5.8
Fucosterol	3.6 ± 0.2	1.2 ± 0.2	3.8 ± 0.1	2.7 ± 0.3	2.2 ± 0.2
Desmosterol	3.7 ± 0.1	1.2 ± 0.1	2.3 ± 0.5	1.3 ± 0.4	1.1 ± 0.1
MethylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
EthylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
<b>Digestive gland</b>					
Brassicasterol	19.3 ± 0.2	45.0 ± 7.7	8.6 ± 0.9	11.3 ± 9.5	16.1 ± 0.5
Cholesterol	33.8 ± 0.3	26.3 ± 3.4	49.2 ± 1.7	25.5 ± 3.2	32.0 ± 1.1
Campesterol	a 2.4 ± 0.0	a 2.4 ± 0.2	a 1.6 ± 0.1	b 22.4 ± 2.2	a 6.0 ± 0.5
24-Methylen-cholesterol	14.1 ± 0.5	9.5 ± 0.9	10.4 ± 0.4	16.7 ± 1.1	16.8 ± 0.7
Stigmastérol	4.1 ± 0.5	4.0 ± 1.0	2.1 ± 0.3	0.0 ± 0.0	2.9 ± 1.0
Isofucosterol	2.5 ± 0.2	2.7 ± 0.6	4.1 ± 0.4	2.5 ± 0.6	6.8 ± 0.8
MethylPorifera	1.9 ± 0.1	0.4 ± 0.1	0.0 ± 0.0	5.2 ± 6.2	0.6 ± 0.3
β-sitosterol	ab 4.3 ± 0.1	a 1.8 ± 0.6	ab 2.0 ± 0.1	b 9.5 ± 0.8	ab 4.7 ± 0.2
Fucosterol	a 3.7 ± 0.1	a 0.5 ± 0.1	b 15.0 ± 1.0	a 0.9 ± 0.5	a 1.0 ± 0.1
Desmosterol	a 5.0 ± 0.2	b 0.9 ± 0.2	b 0.8 ± 0.1	b 1.0 ± 0.1	b 0.9 ± 0.2
MethylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
EthylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
<b>Muscle</b>					
Brassicasterol	20.3 ± 0.4	32.9 ± 2.9	15.0 ± 0.4	13.5 ± 1.0	18.5 ± 0.5
Cholesterol	35.3 ± 0.8	33.2 ± 1.4	31.8 ± 0.3	30.1 ± 0.8	35.2 ± 1.0
Campesterol	2.5 ± 0.0	1.9 ± 0.0	7.6 ± 0.3	3.5 ± 0.2	2.9 ± 0.9
24-Methylen-cholesterol	ab 16.6 ± 0.1	b 13.3 ± 1.0	26.1 ± 0.4	c 35.9 ± 0.9	b 13.2 ± 0.8
Stigmastérol	3.1 ± 0.2	3.4 ± 0.9	3.8 ± 0.8	3.9 ± 0.7	4.6 ± 0.1
Isofucosterol	3.0 ± 0.3	2.7 ± 0.1	2.9 ± 0.2	2.5 ± 1.2	2.9 ± 0.9
MethylPorifera	1.5 ± 0.2	0.8 ± 0.7	0.9 ± 0.1	0.8 ± 0.1	1.1 ± 0.1
β-sitosterol	3.6 ± 0.3	2.8 ± 0.6	3.6 ± 0.3	2.2 ± 0.3	10.7 ± 1.7
Fucosterol	1.8 ± 0.1	1.0 ± 0.0	2.1 ± 0.2	1.3 ± 0.6	1.6 ± 0.1
Desmosterol	2.2 ± 0.2	0.9 ± 0.2	0.7 ± 0.2	0.7 ± 0.1	1.1 ± 0.2
MethylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
EthylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
<b>Gills</b>					
Brassicasterol	ab 19.2 ± 0.2	b 37.3 ± 3.2	a 9.3 ± 0.2	a 9.3 ± 0.4	a 12.6 ± 0.4
Cholesterol	33.4 ± 0.3	27.3 ± 3.8	27.6 ± 0.4	26.5 ± 0.6	27.4 ± 1.0
Campesterol	a 2.3 ± 0.2	a 1.7 ± 0.1	b 14.0 ± 0.5	a 5.3 ± 0.3	a 5.4 ± 0.2
24-Methylen-cholesterol	bc 13.8 ± 0.2	b 11.8 ± 0.7	31.8 ± 1.0	a 43.0 ± 0.9	b 9.2 ± 0.9
Stigmastérol	a 4.5 ± 0.1	a 7.7 ± 1.8	b 0.0 ± 0.0	ab 3.0 ± 0.3	a 7.5 ± 0.2
Isofucosterol	2.6 ± 0.0	1.1 ± 0.2	1.2 ± 0.5	1.5 ± 0.0	1.5 ± 0.2
MethylPorifera	2.0 ± 0.1	1.3 ± 0.2	1.5 ± 0.1	0.9 ± 0.2	1.3 ± 0.0
β-sitosterol	a 4.0 ± 0.2	a 1.9 ± 0.5	a 4.6 ± 0.2	a 2.5 ± 0.4	b 24.3 ± 2.5
Fucosterol	3.4 ± 0.0	1.1 ± 0.2	3.1 ± 0.1	2.4 ± 0.1	2.0 ± 0.1
Desmosterol	6.0 ± 0.1	4.7 ± 0.5	3.6 ± 0.2	2.2 ± 1.6	3.2 ± 0.1
MethylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
EthylPavlovol	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0

**Table 8:**

Broodstock diet	Stages of gonad maturation				
	0	1	2	3	4
Initial	50.0	17.0	0.0	0.0	33.0
<i>R. salina</i>	0.0	3.5	3.5	51.7	41.3
<i>T. weissflogii</i>	6.5	9.7	16.1	61.3	6.4
<i>T. pseudonana</i>	24.2	18.2	6.1	48.5	3.0
<i>P. lutheri</i>	40.0	23.1	20.0	6.9	10.0