Production Performance of *Moina macrocopa* (Straus 1820) (Crustacea, Cladocera) Cultured in Different Salinities: The Effect on Growth, Survival, Reproduction, and Fatty Acid Composition of the Neonates

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Abstract: Salinity is a known factor in shaping population dynamics and community structure through direct and indirect effects on aquatic ecosystems. Salinity changes further influence food webs through competition and predation. The responses of *Moina macrocopa* (Cladocera) collected from Setiu Wetland lagoon (Terengganu) was evaluated through manipulative laboratory experiments to understand the ability of *M. macrocopa* to tolerate high salinity stress. Specifically, the fatty acid composition, growth, survival, and reproduction of this cladocerans species was examined. Sodium chloride (NaCl) as used in the treatments water with the concentration 0, 4, 6, 8, 12, and 15 salinity. Fatty acid levels were determined using Gas Chromatography and Mass Spectrophotometry (GC-MS). The results indicated that optimal conditions produced the highest fatty acid content, especially the polyunsaturated fatty acid content, such as EPA (eicosapentaenoic acid), ALA (alpha-linoleic acid), ARA (arachidonic acid), and DHA (docosahexaenoic acid). Furthermore, *M. macrocopa* survival was best at salinity 0, with a percentage of 98%, whereas the opposite occurred at salinity 15, with approximately 20% of viable animals surviving. Besides, *M. macrocopa* also showed the highest reproduction rate at salinity 0 (e.g., average initial age of reproduction, 4.33 ± 0.58 days) compared with other salinities level. Interestingly, the difference in growth at different salinities was not evident, an unusual finding when considering adverse effects such as osmoregulation pressure on the organism. Based on the results, we conclude that *M. macrocopa* can only tolerate salinity below salinity 8 and cannot withstand stressful environmental conditions associated with salinities above 8.

Keywords: freshwater zooplankton; *Moina*; salinity tolerance; NaCl; fatty acid

1. Introduction

The global climate is changing rapidly in recent years. Industrialization, sea level rise, and urbanization often cause a variety of negative impacts on the aquatic ecosystems. Global warming and climate change have been described as the key factors contributing to rising sea levels. The increase in sea level rise has one of the most severe consequences. Consequently, coastal areas such as saltwater intrusion, coastal flooding, and coastal erosion can affect the inland ecosystem. They can also cause changes in the severity and frequency of severe events due to the combined effects of high spring tides, storm surges, surface...
waves, and river flooding. Zooplankton populations address environmental fluctuation conditions ranging from seasonal and predictable to unusual and unpredictable occurrences [1]. Salinity typically fluctuates temporarily and is a vital influence on zooplankton’s composition and dynamics in inland coastal systems, as even marginally increased salinity adversely affects species. Salinity rises particularly as a result of marine intrusions in these aquatic systems. In contrast, salinity decreases after heavy rain and the main river inflow [2,3].

Furthermore, salt concentrations in freshwater habitats have increased because of industrial activities and urbanization, with resultant impacts on freshwater animals [4]. Agricultural operations, urbanization, and other coastal growth projects are rapidly taking place in Southeast Asia that can drastically alter the geomorphological structures [5]. The Setiu Wetland lagoon is considered a major location for brackish water mariculture activities, including finfish cage, pond culture, and shellfish farming [5]. This anthropogenic interference has both directly and indirectly degraded the natural environment of Setiu Wetland lagoon. Increased seawater levels and coastal erosion have significantly led to increases in salinity [4], affecting aquatic ecosystems and biodiversity [5]. Aquatic organisms are sensitive to changes in water quality. Small fish are more opportunistic in nature in that they can adapt and survive in stressful conditions found in shallow lagoons [6]. Zooplankton is an important community in the aquatic ecosystem for energy transfer from primary producers to fishes [6,7]. Factors such as zooplankton and ecosystem species type heavily influence the feeding of small fish [8]. The ecosystem’s condition affects both foraging efficiency and predation risk and may beneficially increase the former and decrease the latter in fish [9].

Zooplankton species distribution and abundance are influenced by environmental factors such as water transparency, climate change, and nutritional food content [10–12]. The abundance and diversity [13] of zooplankton and phytoplankton [4] are affected by salinity and available diets in the environment. Therefore, water salinity shifts also can alter the original taxa composition and ecological processes, such as primary productivity, decomposition, nutrient cycles, and food web function [14]. Salinity increase in freshwater can reduce zooplankton richness, especially in the cladoceran community, and thus, change the adaptation of the species to a more salt-tolerant species [5]. Research on cladocerans has been widely conducted due to their ecological importance, sensitivity towards environmental changes, and ease of handling [15]. Cladocerans are an important group in the zooplankton community with most of the species living in freshwater environments with salinities of <1 [4]. This is due to osmoregulation adaptations that only allow them to tolerate lower salinity levels [16]. This represents rapid changes in osmotic hemolymph concentration during salinity acclimation. Rapid osmotic adjustment and the relatively wide tolerance range of cladocera can encourage the colonization of contrasting environments whenever other ecological constraints are less important [17]. A few cladoceran species have shown an ability to adapt to salinity changes. However, this adaptation can negatively impact the growth and reproduction of the cladoceran species compared to their original habitat. Cladocerans are reported to be abundant in freshwater ecosystems as compared to brackish water environments [18]. However, a few cladoceran species can live in a saline environment, with brackish water species, tolerating beyond salinity 13 [16]. The dynamics and abundance of cladocerans are influenced by increased salt levels in freshwater environments [19]. A previous study by Ismail [20] showed that the highest salinity at the study area was 17.76 while the normal in-situ parameter was supposed to be in the range of 28–29 °C, 3.8–4.5 mg/L, 7.5pH and salinity 2–6, respectively.

*Moina* sp., commonly referred to as water fleas, are crustaceans within the family of Moinidae, which inhibit freshwater and thrives in both brackish and marine environment [21]. The reproductive cycle of *Moina* sp. has both a sexual and asexual phase. Normally, the population consists of all females that reproduce asexually. Under optimum conditions, *Moina* sp. reproduce at only 4 to 7 days of age, with a brood size of 4 to 22 per female [22]. Broods are produced every 1.5 to 2.0 days [22], with most females producing 2
to 6 broods during their lifetime [22]. Under adverse environmental conditions, males are produced, and sexual reproduction occurs, resulting in resting egg (ephippia) production. This case is similar to brine shrimp eggs production. The stimuli for the switch from asexual to sexual reproduction in populations of *Moina* sp. are an abrupt reduction in the food supply and significant change in environmental conditions. The density of *Moina* sp. cultures routinely reaches densities of 5000 individuals per Litre (19,000/gallon) and therefore, they are well adapted to an intensive culture [22]. *M. macrocopa* are rich in protein and nutrients. They are excellent live foods for fish and prawn larvae when compared to other live feeds such as rotifer and *Artemia* [9] However, the fatty acid content of *Moina* sp. and other live food species such as copepods varies when they are cultivated with different culture media and different levels of environmental stress [23,24].

Studies have shown that lowering ambient temperature tends to increase C20–22 polyunsaturated fatty acid production in planktonic crustaceans [25]. However, the study on the impact of other environmental parameters on freshwater cladocerans, such as salinity change, is still limited and some quantitative data are only available on the fatty acid profiles of zooplankton at different temperatures but salinity-induced effects on the fatty acid composition of *M. macrocopa* have not been evaluated [26]. Lipids and fatty acids are structural components of cell membranes that play an essential role in growth, survival, and reproduction of aquatic organisms, especially in the early life stages of larvae [26]. Salinity changes in the freshwater habitat can impact the distribution of, community structure of water fleas, and can affect feeding behavior and also have an indirect impact on nutritional composition. The current study investigated the tolerance of *M. macrocopa* to salinity changes. This study’s objectives were to evaluate the ability of *M. macrocopa* to adapt to a saline environment up to salinity 15 [27] and, to investigate the potential of *M. macrocopa* to saline tolerance [16]. Specifically, this study aimed to assess the impact of salinity on fatty acid composition, growth, survival, and reproduction of *M. macrocopa* when exposed to different salinity levels. *M. macrocopa* is also considered a useful indicating animal to evaluate the impacts of increasing salinity on the aquatic environment.

2. Materials and Methods

2.1. Cultivation of *Moina Macrocopa*

Live *M. macrocopa* were collected from the lagoon of Setiu Wetland (Figure 1) at the beginning of the month (March, April, May), located in the state of Terengganu (N 5°68′ E 102°70′), and cultured in the Hatchery of University Malaysia Terengganu before use in the experiments. A zooplankton net (sizes from 50 to 200 µm) was used to sample zooplankton in the sampling area. Salinity, dissolved oxygen, and water temperature at the sampling site were measured using a YSI Multi-probe instrument (YSI Model 33) (Table 1). The plankton were collected by pulling the plankton net through the water horizontally and the plankton were further filtered and rinsed before placed into the bottle.

*Moina* sp. were isolated and cultured in a 100 mL flask (Figure 2) filled with freshwater (salinity 0) and, were periodically upscaled to 2000 L tanks for mass culture once the density reached 10 individuals per milliliter (ind/mL). Water parameters were maintained at 26 °C to 30 °C, salinity 0 with a light regime of 12 h light: 12 h dark [23]. In the current study, individuals in the neonates’ stage >24 h old were used [4] and isolation of neonates was carried out using nets with different mesh sizes (50 µm to 200 µm) into each container with 10 mL of culture medium. A single neonates per replicate and 20 replicates of each treatment with salinity 2, 4, 6, 8, 12, and 15 were studied. Hence, at least 20 egg-carrying females were required for each treatment. The test animals were observed twice daily until egg being carried were noticed in their brood chamber. Following this, animals were monitor every hour, the time of first reproduction was recorded when the first neonate was released from the brood chamber and the experiment was then ceased. The time was recorded in hour in the reproduction period between salinity treatments. During the experiment, *M. macrocopa* were fed fresh microalgae at a concentration of $14 \times 10^7$ cells/mL [28]. Microalgae *Nannochloropsis* sp. were used as feed to maintain
similarity with the original habitat food resource. Pure microalgae stock was cultured at room temperature (25 °C to 28 °C) with 24 h of light. The algal strain *Nannochloropsis* sp. was obtained from the UMT hatchery algal laboratory and, the algae were cultured as feed for *M. macrocopa*. Microalgae were harvested when the culture reached the exponential phase [29] or before the decline phase [30]. To maintain *M. macrocopa* cultivation’s salinity, the *Nannochloropsis* sp. was cultured at a lower salinity by gradually decreasing the salinity until it reached the same range of *M. macrocopa* cultivation.

![Figure 1](https://www.google.com/maps/search/setiu+wetland/@4.2777739,101.7082448,7.36z) via Google map.

**Table 1.** The in-situ parameters of the sampling area.

| Month     | Temperature (°C) | Dissolved Oxygen (mg/L) | pH   | Salinity |
|-----------|------------------|-------------------------|------|----------|
| 1 March 2020 | 28.32            | 4.21                    | 7.16 | 6.23     |
| 1 April 2020 | 29.67            | 4.45                    | 7.50 | 2.11     |
| 2 May 2020  | 29.33            | 3.89                    | 7.24 | 2.08     |

Changes in growth, survival, and reproduction of *M. macrocopa* were examined at six different salinities. The salinity level used was within the range of those able to occur within the original habitat of Setiu Lagoon. According to [5,31], the water salinity in Setiu Wetland ranges from salinity 0 to 35.0. In the laboratory, *M. macrocopa* was tested at salinity 0, 4, 6, 8, 12, and 15 [13], for the similarity of the range of salinity in Setiu wetland. The effect of salinity treatments was conducted by using sodium chloride (NaCl) solution as test waters. Every treatment ran in parallel with control in three replicates, each replicate contained 10 neonates (average number were taken from each replicate for all treatments) in 100 mL test water in 250 mL glass beakers. The artificial test water was diluted with the synthetic freshwater media to the respective test salinity. Test media were prepared by diluting saline water with synthetic freshwater media until the required salinities were recorded with a salinity conductivity- temperature Meter (YSI Model 33). The culture medium (salinity 0) was used as a control to mimic the original habitat of freshwater *M. macrocopa* in nature with no salinity level.
The average percentage survival of *M. macrocopa* was determined every alternate day by using a zooplankton counting chamber. The initial density was known, the final density of *M. macrocopa* was then recorded to generate a survival estimate (final density divided by initial counted density). The growth of *M. macrocopa* was determined with 10 individuals daily by measuring body length (from the base of the caudal spine to the anterior edge of the head) with a projection microscope. The experiment was carried out for 15 days until the entire cohort died. This experiment was conduct by using the equation by Pianka 1988 [32] to measure the reproductive capacity of *M. macrocopa*, and life tables were used to record and analyze reproduction data. The life table variables included the initial age at maturation, longevity (days), survival/survivorship, average longevity, gross reproduction rate, net reproduction rate, generation time, and life expectancy. These variables were estimated as follows:

\[
\text{Age of first maturation (day) = age at which the first brood appears from a female}
\]

\[
\text{Longevity (day) = the average number of days of survival of the female}
\]

\[
\text{Survival/Survivorship} = \sum l_x
\]

\[
\text{Average longevity} = \sum \frac{n_x}{n}
\]

\[
\text{Gross reproduction rate} = \sum m_x
\]
Net reproduction rate \( R_0 \) = \[ \sum l_x m_x \] (6)

Generation time \( T \) = \[ \sum \frac{l_x m_x x}{R_0} \] (7)

Life expectancy \( e_x \) = \[ \frac{T_x}{n_x} \] (8)

where, \( l_x \): Proportion individual surviving to age \( x \); \( n_x \): Number of individual alive for each age class; \( n \): Number of animals; \( m_x \): Age-specific fecundity (number of neonates produced per surviving female at age \( x \)); \( T_x \): Generation time at age \( x \).

The intrinsic rate of population increases \( r \) was calculated at the end of the experiment, and all the data were recorded at 15 to 16 days which was equivalent to the average lifespan (cycle) of \( M. macrocopa \).

2.2. Analysis of Fatty Acid

The zooplankton samples from each treatment were collected and freeze-dried before fatty acid analysis. Fatty acids (FAs) were extracted from freeze-dried \( M. macrocopa \) samples using a technique of \[33\] for both qualitative and quantitative examination. The extracted FAs were transesterified into fatty acid methyl esters (FAME) using a strong acid at 80 °C to 85 °C for about an hour. After this treatment, purified water and hexane were added and the upper organic layer transferred to a vial \[34\]. This step was performed several times to achieve complete extraction of FAME. Samples were then dried and dissolved again in 20 \( \mu \)L hexane to get 50 times concentration, removing all solvent peaks (i.e., toluene). The concentrated samples were then injected into gas chromatography-mass spectrometry (GC-MS) to read the spectra using caprylic acid (\( CH_3(CH_2)_6COOH \)) as an internal standard.

2.3. Data Analysis

This study’s data were expressed as Mean ± SD and were analyzed using one-way analysis of variance (ANOVA) to test for differences among salinity treatments. Post hoc Tukey’s multiple comparison tests were used to determine the significant differences of means between treatments. The level of significant difference was set at \( p < 0.05 \).

3. Results

3.1. Survival and Growth of \( M. macrocopa \) in Different Salinities

The survival and growth of \( M. macrocopa \) at different salinity levels are shown in Tables 2 and 3 below. The highest survival of \( M. macrocopa \) occurred at salinity 0 (91.16 ± 1.67%, \( p < 0.05 \), refer to Table 2, Figure 3) while the lowest survival (approximately 13%) occurred at the salinity of 15. The survival rate at salinity 4 (81.89 ± 8.08%) and salinity 6 (75.31 ± 7.93%) were lower than in the control at salinity 0 (\( p < 0.05 \), refer to Table 2, Figure 3). The growth of \( M. macrocopa \) showed a positive relationship with the lowering of salinities at 0 (1.35 ± 0.28 mm), 4 (1.32 ± 0.25 mm), and 6 (1.30 ± 0.25 mm) with growth not being significantly affected. When the salinity reached 8 (0.72 ± 0.61 mm), 12 (0.39 ± 0.54 mm) and 15 (0.23 ± 0.43 mm), the growth of \( M. macrocopa \) became slower (Figure 4 and Table S4). There was no significant difference between salinity ranges on the growth of \( M. macrocopa \) (Table 3).
Table 2. The survival rate of *M. macrocopa* on different ranges of salinity. All values are mean ± standard deviation (*n* = 3). The different small letters indicate significant differences between different salinity (*p* < 0.05).

| Salinity | Survival Rate (%) (Mean ± SD) |
|----------|-------------------------------|
| 0        | 91.16 ± 1.67<sup>a</sup>      |
| 4        | 81.89 ± 8.08<sup>b</sup>      |
| 6        | 75.31 ± 7.93<sup>c</sup>      |
| 8        | 43.69 ± 2.20<sup>d</sup>      |
| 12       | 21.04 ± 1.13<sup>e</sup>      |
| 15       | 13.35 ± 1.25<sup>e</sup>      |

Table 3. Growth rate of *M. macrocopa* on different ranges of salinity. All values are mean ± standard deviation (*n* = 3). The different small letters indicate a significant difference between different salinity levels (*p* < 0.05).

| Salinity | Growth Rate (mm) (Mean ± SD) |
|----------|------------------------------|
| 0        | 1.35 ± 0.28<sup>a</sup>      |
| 4        | 1.32 ± 0.25<sup>a</sup>      |
| 6        | 1.30 ± 0.25<sup>a</sup>      |
| 8        | 0.72 ± 0.61<sup>b</sup>      |
| 12       | 0.39 ± 0.54<sup>c</sup>      |
| 15       | 0.23 ± 0.43<sup>c</sup>      |

**Figure 3.** The survival rate of *M. macrocopa* for different salinity treatments.

**Figure 4.** The growth rate of *M. macrocopa* for different salinity treatments. The different small letters indicate significant differences between different salinity treatments (*p* < 0.05).
3.2. Age of First Maturation and Average Longevity

The life table parameters of *M. macrocopa* are shown in Table 4. Salinity significantly influenced the initial age of reproduction (*p* < 0.05, refer to Figure 5). Females began to reproduce after 4.33 ± 0.58 days at salinity 0, compared longer delay for salinity 4 (5.33 ± 0.58 days), 6 (46.67 ± 1.15 days), and 8 (11.67 ± 0.58 days). However, there was no difference in reproduction between salinity 0 and 4 (*p* > 0.05, refer to Figure 5). The average longevity of *M. macrocopa* was positively correlated with salinity (*p* < 0.05, refer to Figure 6), where average longevity was reduced with increased salinity. Female of *M. macrocopa* survived longer at salinity 0 (12.67 ± 0.58 days), compared with other treatment (Figure 6). Furthermore, no life table data can be recorded at salinity 12 and 15, since all the *M. macrocopa* were found dead before each cohort was able to reproduce (Figure 6).

**Table 4.** Life table parameters for *M. macrocopa*. All values are mean ± standard deviation (n = 3). The different small letters indicate significant differences between different salinity (*p* < 0.05).

| Salinity | Average Initial Age of Reproduction (Days) | Average Longevity (Days) | Net Reproduction Rate | Gross Reproduction Rate | Generation Time | Intrinsic Rate of Population Increases |
|----------|-----------------------------------------|--------------------------|-----------------------|-------------------------|-----------------|----------------------------------------|
| 0        | 4.33 ± 0.58 a                           | 12.67 ± 0.58 a           | 29.58 ± 7.27 a        | 7.33 ± 1.53 a           | 6.78 ± 2.18 a   | 0.14 ± 0.04 a                          |
| 4        | 5.33 ± 0.58 b                           | 9.67 ± 0.58 b            | 16.00 ± 4.23 b        | 5.67 ± 0.57 b           | 9.11 ± 0.84 b   | 0.087 ± 0.023 b                        |
| 6        | 6.67 ± 1.15 c                           | 7.33 ± 1.53 b            | 8.17 ± 1.78 c         | 5.00 ± 1.00 c           | 10.56 ± 3.29 a | 0.060 ± 0.01 c                        |
| 8        | 11.67 ± 0.58 d                          | 3.00 ± 0.01 c            | 3.52 ± 1.04 d         | 3.00 ± 1.00 d           | 11.67 ± 0.577 a| 0.012 ± 0.006 d                       |
| 12 d     | -                                       | -                        | -                     | -                       | -               | -                                      |
| 15 d     | -                                       | -                        | -                     | -                       | -               | -                                      |

* * Life table parameters for salinity 12 and 15 cannot be measured and recorded because all the treatments reached mortality before being able to reproduce.

**Figure 5.** The average initial age of reproduction of *M. macrocopa* for different salinity treatments. The different small letters indicate significant differences between different salinity treatments (*p* < 0.05).

**Figure 6.** The average longevity of *M. macrocopa* for different salinity treatments. The different small letters indicate significant differences between different salinity treatments (*p* < 0.05).
3.3. Net Reproduction Rate and Gross Reproduction Rate

The net reproduction rate is the average number of females in a population produced during a female’s lifetime (Figure 7). The net reproduction rates of *M. macrocopa* ranged with rates for salinity 0 (29.58 ± 7.27 offspring/female), 4 (16.001 ± 4.23 offspring/female), 6 (8.17 ± 1.78 offspring/female), and 8 (3.52 ± 1.04 offspring/female) that revealed significant differences in reproductive rates among salinity treatments (*p* < 0.05, refer to Figure 8). The increase of salinity showed a statistical difference in gross reproduction rates of *M. macrocopa* (*p* < 0.05, refer to Figure 8). The highest gross reproduction rate is at salinity 0 (7.33 ± 1.53 offspring/female) followed by salinity 4 (5.67 ± 0.57 offspring/female), 6 (5.00 ± 1.00 offspring/female) and 8 (3.00 ± 1.00 offspring/female) (Figure 8). The net reproduction rates and gross reproduction rate of *M. macrocopa* decreased with salinity increases (Figures 7 and 8). There was a significant difference between the salinity 0 as control and salinity 4 (*p* < 0.05, refer to Table 4), for net reproduction or gross reproduction rate (Figure 8).

![Figure 7](image_url) **Figure 7.** The net reproduction rate of *M. macrocopa* for different salinity treatments. The different small letters indicate significant difference between different salinity treatments (*p* < 0.05).

![Figure 8](image_url) **Figure 8.** The gross reproduction rate of *M. macrocopa* for different salinity treatments. The different small letters indicate significant differences between different salinity treatments (*p* < 0.05).
3.4. Generation Time and Intrinsic Rate of Population Increase

The time from the laying of eggs to when an individual reaches sexual maturity is known as generation time (Figure 9). The generation time for *M. macrocopa* was shorter at salinity 0 (6.78 ± 2.18 days) and longer at salinity 8 (11.67 ± 0.57 days). There was no significant difference in generation time between all the treatments (*p* > 0.05, refer to Figure 9). Based on the results, salinity showed an inverse relationship with the intrinsic rate (*p* < 0.05, refer to Figure 10). The *M. macrocopa* showed a high intrinsic rate at salinity 0 (0.14 ± 0.04) compared with the rate at salinity 4 (0.087 ± 0.023), 6 (0.060 ± 0.01) and 8 (0.012 ± 0.006) as seen in Figure 10.

![Figure 9](image-url)  
**Figure 9.** The generation time of *M. macrocopa* for different salinity treatments. The different small letters indicate significant differences between different salinity treatments (*p* > 0.05).

![Figure 10](image-url)  
**Figure 10.** The intrinsic rate of population increases of *M. macrocopa* for different salinity treatments. The different small letters indicate significant differences between different salinity treatments (*p* < 0.05).

3.5. Fatty Acid Compositions of *M. Macrocopa*

The fatty acid compositions of *M. macrocopa* grown at different salinities are shown in Tables 5 and 6. The fatty acid contents were significantly affected by salinity levels that cohorts were exposed to (*p* < 0.05). The highest level of fatty acids content considering saturated and polyunsaturated fatty acids in *M. macrocopa* was recorded for the salinity of 0 (*p* < 0.05). The highest level of fatty acid as the sum of the saturated, monosaturated, and...
polyunsaturated, as a percentage of total fatty acids, was recorded in the control treatment at the lowest salinity level. The amount of EPA (C20:5) in *M. macrocopa* were higher at the lowest salinity levels (3.407 ± 0.189%) than at higher salinity levels (p < 0.05). The increase of salinity from 0 (8.867 ± 0.252%) to 15 (0.474 ± 0.047%) significantly abated the C18:1 monosaturated fatty acids in *M. macrocopa* (p < 0.05, refer to Tables 5 and 6). The content of polyunsaturated fatty acids (PUFA) was significantly higher for a salinity of 0 (0.725 ± 0.014%) than at higher salinity levels in *M. macrocopa* (p < 0.05). The highest polyunsaturated fatty acids were recorded for eicosapentaenoic acid (EPA) (0.725 ± 0.047) at the salinity of 0 (p < 0.05). The highest alpha-linoleic acid (ALA) content in *M. macrocopa* was also found at salinity 0 and 4 (0.474 ± 0.047%, 0.135 ± 0.047%) respectively, p < 0.05, refer to Tables 5 and 6).

**Table 5.** Fatty acid composition (% total fatty acids) of *Moina macrocopa* grown at different salinities. All values are mean ± standard deviation (n = 3). The different small letters indicate significant differences between; different salinity (p < 0.05).

The bold fatty acid species are described in the results.

| Fatty Acid | 0          | 4          | 6          | 8          | 12         | 15         |
|------------|------------|------------|------------|------------|------------|------------|
| C14:0      | 3.407 ± 0.189 \(^a\) | 2.977 ± 0.157 \(^a\) | 1.789 ± 0.65 \(^b\) | 0.511 ± 0.168 \(^c\) | 0.183 ± 0.002 \(^c\) | 0.008 ± 0.006 \(^c\) |
| C16:0      | 27.746 ± 2.585 \(^b\) | 26.220 ± 1.926 \(^b\) | 19.451 ± 1.534 \(^b\) | 4.427 ± 1.536 \(^c\) | 0.017 ± 0.001 \(^d\) | 0.007 ± 0.004 \(^d\) |
| C16:1      | 1.200 ± 0.290 \(^a\) | 0.563 ± 0.235 \(^b\) | 0.037 ± 0.023 \(^c\) | 0.012 ± 0.002 \(^c\) | -           | -           |
| C18:1 (n9) | 8.867 ± 0.252 \(^b\) | 5.923 ± 0.484 \(^b\) | 3.380 ± 1.124 \(^c\) | 0.920 ± 0.459 \(^d\) | 0.103 ± 0.031 \(^d\) | 0.047 ± 0.014 \(^d\) |
| C22:1 (n9) | 0.229 ± 0.0325 \(^a\) | 0.062 ± 0.099 \(^b\) | 0.007 ± 0.002 \(^c\) | 0.002 ± 0.002 \(^c\) | -           | -           |
| C18:2 (n6) | 24.033 ± 1.952 \(^a\) | 13.269 ± 2.376 \(^b\) | 5.070 ± 0.354 \(^c\) | 1.116 ± 0.703 \(^d\) | 0.368 ± 0.319 \(^d\) | 0.055 ± 0.044 \(^d\) |
| C18:3 (n3)(ALA) | 0.474 ± 0.047 \(^a\) | 0.135 ± 0.047 \(^b\) | 0.014 ± 0.020 \(^c\) | 0.001 ± 0.001 \(^c\) | -           | -           |
| C20:4 (n6)(ARA) | 0.616 ± 0.082 \(^a\) | 0.145 ± 0.364 \(^b\) | 0.041 ± 0.442 \(^c\) | 0.006 ± 0.001 \(^d\) | 0.002 ± 0.002 \(^d\) | 0.001 ± 0.001 \(^d\) |
| C20:5 (n3)(EPA) | 0.725 ± 0.047 \(^a\) | 0.420 ± 0.318 \(^b\) | 0.004 ± 0.004 \(^b\) | -           | -           | -           |
| C22:6 (n3)(DHA) | 1.151 ± 0.133 \(^a\) | 1.173 ± 0.356 \(^a\) | 0.427 ± 0.300 \(^b\) | 0.002 ± 0.002 \(^b\) | 0.001 ± 0.001 \(^b\) | -           |

ALa: Alpha-linolenic acid; ARA: Arachidonic acid; EPA: Eicosapentaenoic acid; DHA: Docosahexaenoic acid.

**Table 6.** Fatty acid composition (% total fatty acids) of *Nannochloropsis* sp. The bold fatty acid species are described in the results.

| Fatty Acid | *Nannochloropsis* sp. |
|------------|-------------------------|
| C14:0      | 3.77 ± 1.60             |
| C16:0      | 28.14 ± 2.15            |
| C16:1      | 18.17 ± 2.68            |
| C18:1 (n9) | 8.02 ± 1.63             |
| C18:2 (n6) | 4.25 ± 0.73             |
| C18:3 (n3)(ALA) | 0.84 ± 0.17         |
| C20:4 (n6)(ARA) | 5.28 ± 1.05         |
| C20:5 (n3)(EPA) | 25.88 ± 3.79         |
| C22:6 (n3)(DHA) | 0.49 ± 0.25         |

ALa: Alpha-linolenic acid; ARA: Arachidonic acid; EPA: Eicosapentaenoic acid; DHA: Docosahexaenoic acid.

**4. Discussion**

Freshwater ecosystems are becoming increasingly threatened, partly due to the rise in the salinity of groundwater (i.e., from seawater intrusion) and water regime modifications, which reduce the frequency of high flow flushing events [25,35,36]. These changes will significantly impact both the survival rate [37] and reproduction [37,38] of *M. macrocopa*. This study demonstrated how the increase of salinity, especially sodium chloride (NaCl) affects survival and life history traits of a freshwater cladoceran species through an examination of effects on *M. macrocopa*, which may be representative of the impact to other members of zooplankton communities. The present study results indicated that the increase in salinity influenced the survival, growth, and reproduction of *M. macrocopa*.

*M. macrocopa* achieved the highest recorded survival rate at treatment salinity of 0, compared to other salinity range treatments. As indicated by [39] for another species of cladoceran, *Latonopsis australis*, *L. australis* had the highest survival rate recorded for an
environmental salinity of 0, which also decreased, with the increase of the salinity level in the natural environment. Moreover, according to the study of [40], the survival rate of Moina eugenie decreased with increased salinity and [4], further supported this where the survival rate of Daphnia magna was shown to decrease with increasing salt concentration. It is concluded that the survival rate of M. macrocopa is undoubtedly, significantly affected by the increase of salinity levels.

The current study showed that the length of M. macrocopa does not experience a significant reduction in environments with high salinities, up to a point. The results showed the positive growth of M. macrocopa in salinity levels from 0 until 6. However, when the salinity was increased to 8, 12, and 15, the growth of M. macrocopa became slower. The slowest growth was observed at salinity 15, which might have resulted from some regulatory stress that occurred in the culture [41]. Increased salinity stress can also add additional energy costs for metabolism and osmoregulation to redirect energy from somatic growth. In a stressful environment, animals which spend more energy in reproduction are an adaptation to increase mortality [12]. This conclusion is also aligned with the findings of [27] investigating the growth rate of exotic cladoceran, Daphnia exilis which also did not show any changes under a salinity regime of 6 but started to decline in growth when salinity reached 8 and above.

Female reproduction was affected by salinity and the maturation of M. macrocopa was delayed at the highest salinity. These results are comparable to those obtained by [42], who noted that the sexual maturation of M. macrocopa was delayed by 1 day at a salinity level of 5.5 when compared to 0. Additionally, [4], noted that the reproduction age of D. magna was delayed from 7 days until 9 days at salinity levels of 0 and 2.66, respectively.

The current study results showed that the average lifespan of M. macrocopa was 12.3 days of salinity 0 and, 9.6 days at salinity 6. These are similar results to a study by [43], where the average lifespan of D. magna was also significantly affected by salt concentration. The results showed that maximum average lifespan (57.7 days) occurred at the salinity of 0. In contrast, the shortest lifespan (25.2 days) occurred at salinity 7. Furthermore, in studies of Simocephalus vetulus, the results also showed that this cladoceran’s lifespan decreased drastically when the salinity level was increased apart from food impacts that positively influence the lifespan of copepods [10,11,44].

The offspring net and gross reproduction rates of M. macrocopa were also affected when salinity increased. The previous study also showed that the number of offspring produce was the most affected parameters on the life table as caused by an increase of salinity for D. magna and D. longispina [45]. Furthermore, in an observation-based study by [39], the findings also showed that the offspring of Latonopsis australis were significantly influenced by high salinity. It is concluded that salinity clearly impacts reproduction of offspring of this cladoceran.

Salinity levels impact the intrinsic rate (births minus deaths) for M. macrocopa population. This claim was supported in similar research done by [39] where the intrinsic rate of Latonopsis australis decreased when the salinity levels of treatments increased. In the current study, life table parameters at salinity more than 8 also cannot be reliably acquired because salinity higher than 8 is beyond the tolerance of M. macrocopa. [13] D. longispina reproduced and lived well in salinity below salinity 5 but then started to decline when salinity levels increased. Thus, the present study concludes that M. macrocopa can achieve the optimum growth population when salinity is below 6. Zooplankton is sensitive to change in the aquatic environment that can significantly change ambient conditions within the aquatic ecosystem [46]. The increase of salt concentration can cause the disappearance of species that cannot tolerate highly saline conditions. Meanwhile, salinity is a strong mechanism that can change aquatic communities [47].

Furthermore, no life table data can be recorded for salinities of 12 and 15 since the M. macrocopa are dead before being able to reproduce. Therefore, various freshwater invertebrate species (including cladocerans) are more sensitive to NaCl salinity than to the effect produced by the array of chemical compounds present in sea salt [41]. Higher
The content of salinity level might be fatal to freshwater cladocerans especially *M. macrocopa*. The highest salinity at which life table could be recorded was only at salinity 8. It is strongly supported by [4], the highest salinity will cause cladoceran stress and mortality. Apart from that, *M. macrocopa* that spending most of their life stages under good conditions (food and environment) may adapt their salinity tolerance only with considerable time-lags.

The observations in the current study showed that fatty acid expression is higher at the optimum culture conditions (salinity 0 and 4) compared to other salinity level treatments. This effect is probably due to the minimal stress undergone by the cultured *M. macrocopa* at these optimum conditions, which is finally reflected in obtaining a rich fatty acid profile. Similar observations were also reported [48–50] where live feed underwent minimal stress during the culture period produced superior survival, growth and reproduction. High polyunsaturated fatty acids achieved by *M. macrocopa* might be due to *Nannochloropsis* sp. as food sources, which were conducted to imitate feed in the natural environment. The amount of EPA and DHA in algae varies greatly among different algal species and environmental conditions. EPA content in *Nannochloropsis* sp. is 23.4 mg g$^{-1}$ [24]. Microalgae such as *Nannochloropsis* sp. have received increasing interest as a target live feed for aquatic animals because of the high contents of EPA [7].

5. Conclusions

In conclusion, the survival, growth, reproduction rate, and fatty acid profile of *M. macrocopa* were affected by different salinity treatment ranges. *M. macrocopa* produced better growth rate, survival rate, and productivity at the salinity of 0 which is a normal freshwater environment. Salinity at 4 also provided relatively good growth for *M. macrocopa*. Most of the present study results showed no significant difference between salinity treatments of 0 and 4. The salinity above 8 was intolerable for *M. macrocopa* and to be above the critical salinity maximum (CSMax) or acute phase for cladoceran. Apart from that, fatty acid content also significantly abated with increased salinity levels. The highest content of fatty acid produced was with *M. macrocopa* being cultured in optimal salinity levels of 0 and 4, which are considered to be normal in their natural habitat. Some cladocera species’ ability to withstand stressful physiological environments, such as increased salinity, leads to a shelter from severe predation and competition typically found in more complex populations. It also can be expected that cladocera species capable of surviving under these conditions are those competitively disadvantaged or more susceptible to predation. That might be the case for *Moina* sp., which can grow at Setiu Wetland during times of significantly increased salinity [51]. Furthermore, salinity can directly and indirectly affect aquatic organism population dynamics. Small changes in salinity tend to be beneficial under natural conditions, where salinity can directly and indirectly mediate multiple biotic and abiotic processes [52]. Overall, the result of direct and indirect effects of minor salinity changes under natural conditions, advantaging cladocera’s population growth.

Comparing results in the whole NaCl concentration range tested, we can conclude that the impairment effects on fatty acid content, growth, reproduction, and survival under the current treatments were provoked by salinity stress. Although the result of the present study shows that the survival rate, life history and fatty acids of *M. macrocopa* were significantly affected by the increase in salinity, future detailed studies are required to understand how freshwater species, including other species of zooplankton which can adapt to a higher saline stressed environment. The present results shown for the responses of neonates towards different salinity can further indicated the similar physiological reactions on the adult’s stages of *Moina macrocopa*. Thus, future researchers can use these findings for further study and research on the development of species characterization and on the factors affecting the abundance and zooplankton composition within the freshwater community, thus providing more ecologically relevant information on primary productivity in the ecosystem.
Supplementary Materials: The following are available online at https://www.mdpi.com/1424-2818/13/3/105/s1, Figure S1: The hypothesis test summary of the survival in M. macrocopa, Figure S2: The hypothesis test summary of the survival in M. macrocopa. Figure S3: The hypothesis test summary of the survival data, Table S1: The detailed SPSS data on the growth of M. macrocopa., Table S2: The ANOVA results on the growth of M. macrocopa in responds to different salinity, Table S3: The survival of M. macrocopa in responds to different salinity, Table S4: Population density data of M. macrocopa, Table S5: The mean, standard deviation, sum of squares data on the effect of salinity on survival and reproductive capacity of M. macrocopa in response to different salinity; Table S6: The ANOVA data on fatty acid effects in response to different salinity.

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