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Juliana Ferreira dos Santos

Influência dos sistemas de cultivo intensivo e semi-intensivo e dietas nos parâmetros zootécnicos e enzimas digestivas da tilápia do Nilo (*Oreochromis niloticus* L.)

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Co-orientador: Prof. Dr. Daniel Lemos (USP)

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BANCA EXAMINADORA

Prof. Dr. Ranilson de Souza Bezerra- Orientador (Presidente)
Centro de Ciências Biológicas – UFPE

Profa. Dra. Maria Tereza dos Santos Correia (Membro Interno)
Centro de Ciências Biológicas – UFPE

Profa. Dra. Mércia Andréa da Silva Lino (Membro Externo)
Unidade Acadêmica de Serra Talhada – UFRPE

Dra. Carolina Nunes Costa Bonfim (Membro Externo)
Departamento de Pesca e Aquicultura – UFRPE

Dra. Helane Maria Silva da Costa (Membro Interno)
Departamento de Bioquímica – UFPE

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Onde leva? Não perguntas, segue-o!

Nietzsche

RESUMO

A tilápia do Nilo pode ser produzida em diferentes sistemas de cultivo e escalas, dentre eles os sistemas semi-intensivo em viveiros (SSI), e o intensivo em tanques-rede (SI). Seus manejos são completamente distintos, por isso o gerenciamento se faz necessário a fim de não desestabilizar parâmetros de crescimento e afetar os custos de produção, principalmente a ração, que pode chegar a representar até 80% destes custos. Para um menor gasto, ingredientes alternativos podem ser usados na elaboração das dietas, no entanto, eles devem apresentar resultados semelhantes ou melhores, quando comparados às dietas convencionais. Sendo assim, pretende-se avaliar se a inclusão de hidrolisado proteico de camarão (HPC) na dieta pode influenciar as atividades enzimáticas da tilápia do Nilo e avaliar a influência do SSI e do SI nos parâmetros de crescimento e nas enzimas digestivas deste peixe. Para avaliar a influência do HPC nas enzimas digestivas de juvenis de tilápia do Nilo, os cultivos foram realizados em aquários. O HPC foi incluído nas dietas nas concentrações de 0, 15, 30 e 60 g.kg⁻¹ (HPC 0, 15, 30 e 60, respectivamente). Ao término dos ensaios (45 dias), o estômago e o intestino dos espécimes experimentais foram removidos para realização das análises. Para avaliação da influência do SSI e do SI sob os parâmetros de crescimento e as enzimas digestivas da tilápia do Nilo, os espécimes experimentais foram cultivados durante 90 dias. Foram adotados dois planos nutricionais: trifásico para SSI (ração com 35, 32 e 28% de proteína bruta – PB) e bifásico para SI (35 e 32% PB). Semanalmente foram realizadas coletas de fitoplâncton e zooplâncton, diariamente os principais parâmetros de qualidade de água foram mensurados e mensalmente foram realizadas biometrias para análise dos parâmetros de crescimento, ajustes na alimentação e remoção dos tecidos (intestino) para análises enzimáticas. A atividade de protease alcalina total foi significativamente maior nos tratamentos HPC15 e HPC60, no entanto, o efeito não foi dose-dependente. Zimograma revelou 12 bandas proteolíticas, oito delas responderam a incorporação do HPC. Já o zimograma de inibição indicou uma redução na atividade de três enzimas, uma diminuição na atividade da tripsina com o aumento da concentração de HPC, e o inverso com duas aminopeptidases. Com relação aos sistemas de criação o alimento natural foi mais abundante no SSI. Sobrevivência, eficiência proteica, eficiência alimentar e fator de condição com 31 dias de cultivo foram melhores no SSI. A relação peso-comprimento não foi alterada pelos sistemas. E quando comparado o peso relativo dos peixes do SSI e do SI com tilápias de outros ambientes ou cultivos, eles foram maiores em nove das 14 comparações. As enzimas digestivas também foram avaliadas no decorrer dos cultivos (31, 63 e 94 dias). Protease total, amilase e lipase não foram estatisticamente diferentes entre os tratamentos ao longo dos períodos analisados. No entanto, atividade de tripsina e quimotripsina foram maiores, com 31 e 63 dias de cultivo em peixes do SSI, sugerindo que o alimento natural pode ter influenciado essas atividades. Uma correlação positiva foi observada entre a concentração recomendada de aminoácidos essenciais para a tilápia do Nilo e atividade específica de aminopeptidase nos peixes do SI. Zimograma também revelou 12 bandas proteolíticas ativas em ambos os sistemas. Sendo que os valores de densidade integrada (*DI*) foram maiores nas bandas do SSI. Espécimes de ambos os sistemas exibiram cinco bandas com atividade amilolítica, com os maiores valores de *DI* aos 31 dias de cultivo. Os menores valores de *DI* foram observados com peixes do SSI, com 63 e 94 dias de cultivo. A comparação da eletroforese bidimensional do intestino de animais cultivados em SSI e SI mostraram um total de 1.233 spots, sendo 731 diferentes. Tilápia do Nilo demonstrou adaptação às dietas utilizadas e aos sistemas de cultivo empregados.

Palavras-chave: tilápia do Nilo, hidrolisado proteico de camarão, enzimas digestivas, sistema intensivo, sistema semi-intensivo.

ABSTRACT

The Nile tilapia can be produced in different culture systems and scales, including the semi-intensive systems in ponds (SIS), and the intensive in cages (IS). Their managements are completely different, so it is required management to stabilize the growth parameters and production costs, mainly feed, which can represent up to 80% of these costs. For lower expenses, alternative ingredients can be used in the preparation of diets, however, they should deliver results similar or better, when compared to conventional diets. Therefore, intends to evaluate whether the inclusion of shrimp protein hydrolyzate (SPH) in the diet can influence the enzymatic activity of juvenile Nile tilapia and evaluate the influence of SIS and IS on growth parameters and digestive enzymes of this fish. To evaluate the effect of SPH on digestive enzymes of juvenile Nile tilapia, the cultures were performed in aquaria. The SPH was included in the diet at concentrations of 0, 15, 30 and 60 g.kg⁻¹ (SPH0, 15, 30 and 60, respectively). At the end of the assay (45 days) the stomach and intestine of experimental specimens were removed for analyzes. To assess the influence of SIS and IS on growth parameters and digestive enzymes of Nile tilapia experimental specimens were cultured for 90 days. Were adopted two nutritional plans: triphasic for SIS (diet with 35, 32 and 28% crude protein - CP) for SI and biphasic (35 and 32% CP). Weekly samples were taken for phytoplankton and zooplankton, the main daily water quality parameters were measured and biometrics were performed monthly for analysis of growth parameters, adjustments in diet and removal of tissues (intestine) for enzymatic analyzes. The activity of total alkaline protease was significantly higher in treatments SPH15 and SPH60, however, the effect was not dose-dependent. Zymogram revealed 12 proteolytic bands, eight of which responded to SPH incorporation. Inhibition zymogram indicated a decrease in the activity of three enzymes, with trypsin activity decreasing with the increase of SPH concentration, whereas occurred the opposite for two aminopeptidases. With respect to culture systems natural food was more abundant in SIS. Survival, protein efficiency, feed efficiency and condition factor with 31 days of cultivation were better in SIS. The weight-length relationship was not altered by the systems. And comparing the relative weight of the fish from SIS and SI systems with tilapia from other environments, they were higher in 9 of 14 comparisons. Digestive enzymes also were evaluated during the cultivation (31, 63 and 94 days). Total proteolytic, amylase and lipase activity were not statistically different between the treatments along of the periods analyzed. However, trypsin and chymotrypsin activity were higher with 31 and 63 days of culture in fish from SIS, suggesting that natural food may have influenced these activities. A positive correlation was observed between the recommended concentration of essential amino acids for Nile tilapia and specific aminopeptidases activity in fish IS. Zymogram also revealed 12 active proteolytic bands in both systems. Since the integrated density (*ID*) values were higher in the bands of SIS. Specimens of both systems exhibited five bands with amylase activity, with higher values of *ID* to 31 days of cultivation. The lowest values of *ID* were observed with fish from SIS with 63 and 94 days of cultivation. The comparison of two-dimensional electrophoresis of the intestine of cultured animals in SIS and IS showed a total of 1,233 spots, among them 731 different. Nile tilapia showed adaptation to the diets and employed culture systems.

Keyword: Nile tilapia, intensive system, semi-intensive system, digestive enzymes, shrimp protein hydrolyzate.

LISTA DE FIGURAS

REVISÃO BIBLIOGRÁFICA

Figura 1: Exemplar de tilápia do Nilo (<i>Oreochromis niloticus</i> L.).....	24
Figura 2: Cultivo semi-intensivo de tilápia do Nilo.....	25
Figura 3: Desenho esquemático da relação entre produção, alimento natural e alimentação suplementar em viveiros escavados.	26
Figura 4: Cultivo intensivo de tilápia do Nilo.....	27
Figura 5: Resíduos da indústria de camarão.	37
Figura 6: Desenho esquemático de camarão identificando os resíduos da indústria pesqueira passíveis de utilização na produção de proteína hidrolisada.....	38

ARTIGOS

CAPÍTULO I

Figure 1: Hydrolysis profile of shrimp protein hydrolysate (using autolysis) in sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) using stacking gel at 4 % (w/v) and separation gel at 17 %. Lanes correspond to molecular mass marker (MM) and incubation times of 0, 30, 60, 90, 120, 150 and 180 min of enzymatic hydrolysis. SPH was produced through autolysis of <i>Litopenaeus vannamei</i> cephalothorax(heads).....	64
--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	----

Figure 2: Zymogram of digestive proteases (3 % casein as substrate) of intestine enzyme extracts from juvenile Nile tilapia <i>O. niloticus</i> fed on diets containing different concentrations of shrimp protein hydrolysate (SPH). Lanes correspond to dietary SPH inclusion (%): SPH 0, SPH 1.5, SPH 3 and SPH 6.....	65
---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	----

Figure 3: Inhibition zymogram of digestive proteases (3 % casein as substrate) of intestine enzyme extracts from juvenile Nile tilapia <i>O. niloticus</i> fed the SPH 0 diet. Lanes correspond to treatments: control without inhibitors, TPCK, PMSF, Benzamidine, TLCK and Bestatin.....	66
--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	----

CAPÍTULO II

Figure 1: Mean \pm SE of mean of the evolution of the abundance of phytoplankton (cel.mL $^{-1}$ \times 10 2) intensive (Figure 1A) and semi-intensive cultivation (Figure 1B) of Nile tilapia (<i>O. niloticus</i>)....	115
------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	-----

Figure 2: Mean \pm SE of mean of the evolution of the abundance of zooplankton (Ind.L $^{-1}$) intensive (Figure 2A) and semi-intensive cultivation (Figure 2B) of Nile tilapia (<i>O. niloticus</i>).	115
------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	-----

Figure 3: Parameters of the mathematical models of Nile tilapia ($W_t = a L_t^b$) adjusted to total weight-length data from fish under different culture systems.	116
--------------------------------------------------------------------------------------------------------------------------------------------------------------------------	-----

Figure 4: Mean of the evolution of Fulton's condition factor (K) of fish subjected to different cultures in comparison with the corresponding population K _{mean}	116
----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	-----

Figure 5: Boxplot of the evolution in the values of K _{rel} (%) from fish under different cultures.....	117
------------------------------------------------------------------------------------------------------------------	-----

Figure 6: Boxplot of the evolution of Total length (cm) (Figure 6A) and Weight (g) (Figure 6B) of fish under different cultures..... 118

CAPÍTULO III

Figure 1 : Alkaline proteolytic activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD, using azocasein (A), BApNA (Na -benzoyl-DL-arginine-*p*-nitroanilide) (B), and SApNA (Suc-Ala-Ala-Pro-Phe-*p*-nitroanilide) (C) as substrate, expressed as U mgProtein $^{-1}$. For the same times, means with different superscript letters differ significantly ($P < 0.05$) by Student's t test ($n = 3$).. 158

Figure 2: Aminopeptidase activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD, using aminoacyl- β -naphthylamide as substrates with Leu (A), Arg (B), Tyr (C) Gly (D), Ser (E), Ala (F) His (G) Iso (H) and Pro (I) as specific substituents and expressed as mU mgProtein $^{-1}$. For the same times, means with different superscript letters differ significantly ($P < 0.05$) by Wilcoxon-Mann-Whitney test and Student's t test ($n = 3$). 160

Figure 3: Amylase activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD, using starch as substrate, expressed as μg of maltose. min^{-1} .mg Protein $^{-1}$. For the same times, means with different superscript letters differ significantly ($P < 0.05$) by Student's t test ($n = 3$)..... 161

Figure 4: Lipase activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD, using *p*-nitrophenyl palmitate (*p*-NPP), as substrate, expressed as U mgProtein $^{-1}$. For the same times, means with different superscript letters differ significantly ($P < 0.05$) by Wilcoxon-Mann-Whitney test ($n = 3$). 161

Figure 5: Correlation between aminopeptidase activity of Nile tilapia (*Oreochromis niloticus*) using aminoacyl- β -naphthylamide as substrates and recommended concentration of essential amino acids for Nile tilapia assessed by the Ideal Protein Concept. Values are shown as mean \pm SD of three crude extracts obtained from intensive culture. Concentration of amino acids expressed in % of dietary protein: arginine (4.1), leucine (4.3), isoleucine (2.6) and histidine (1.5) *Celik (2012), based on Fagbenro (2000)..... 162

Figure 6: (A) Zymogram of digestive proteases of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system, using casein as substrate. Lanes 1, 2 and 3 correspond to intensive culture in the times 31, 63 and 94 days, respectively; Lanes 4, 5 and 6 correspond to semi-intensive culture in the times 31, 63 and 94 days, respectively. (B) Evolution of the integrated density of the bands of proteases zymograms from the intestines of Nile tilapia cultivated in the intensive and semi-intensive system. (C) Pixel intensity and respective means values of integrated density (*ID*) for each lane of protease bands (lanes 1-6), subfigures 1, 2 and 3 correspond to the intensive system and 4, 5 and 6 semi-intensive system, means with different superscript letters differ significantly ($P < 0.05$) using One-way analysis of variance (ANOVA) followed by Tukey test ($n = 3$)..... 163

Figure 7: (A) Zymogram of amylase activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system, using starch as substrate. Lanes 1, 2 and 3 correspond to intensive culture in the times 31, 63 and 94 days , respectively; Lanes 4, 5 and 6 correspond to semi-intensive culture in the times 31, 63 and 94 days, respectively. (B) Evolution of the integrated density of the bands of amylase zymograms from the intestines of Nile tilapia cultivated in the intensive and semi-intensive system. (C) Pixel intensity and respective means values of integrated density (*ID*) for each lane of amylase bands (lanes 1-6), subfigures 1, 2 and 3 correspond to the intensive system and 4, 5 and 6 semi-intensive system, means with different superscript letters differ significantly ($P < 0.05$) using One-way analysis of variance (ANOVA) followed by Tukey test (n=3)..... 164

Figure 8: Two-dimensional gel electrophoresis profile of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive system. First proteins were separated by charge in the first dimension (immobiline Drystrip pH 3–10) and then by molecular weight in the second dimension (12% SDS-PAGE)..... 165

Figure 9: Two-dimensional gel electrophoresis profile of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in semi-intensive system. First proteins were separated by charge in the first dimension (immobiline Drystrip pH 3–10) and then by molecular weight in the second dimension (12% SDS-PAGE)..... 166

LISTA DE TABELAS

REVISÃO BIBLIOGRÁFICA

Tabela 1: Classificação das enzimas segundo IUBMB 32

Tabela 2: Classificação e divisão das proteases 33

ARTIGOS

CAPÍTULO I

Table 1: Proximate composition (%) and energy of lyophilized shrimp protein hydrolysate SPH, from *Litopenaeus vannamei* 58

Table 2: Amino acid composition of the shrimp protein hydrolysate SPH from *Litopenaeus vannamei* 59

Table 3: Composition and approximate analysis of the experimental diets 60

Table 4: Growth performance and nutrient utilization in Nile tilapia fed on diets containing shrimp protein hydrolysate instead of fish meal replacement 61

Table 5: Digestive enzyme activity of crude extracts recovered from stomach (acid) and intestine of juvenile Nile tilapia *O. niloticus* fed on experimental diets containing increasing shrimp protein hydrolysate levels (0–60 g kg⁻¹) 65

CAPÍTULO II

Table 1: Proximate analysis of the commercial diets used in the experiment 110

Table 2: Nutritional plan used as feed for Nile tilapia (*O. niloticus*) subjected to different culture systems 110

Table 3: Quality indicators of water in intensive and semi-intensive cultured Nile tilapia (*O. niloticus*) 111

Table 4: Abundance of phytoplankton (cel.ml⁻¹) in water of intensive and semi-intensive systems Nile tilapia (*O. niloticus*) in time "0" (zero), 31, 63 and 94 days 112

Table 5: Abundance of zooplankton (Ind.L⁻¹) in water of intensive and semi-intensive systems Nile tilapia (*O. niloticus*) in time "0" (zero), 31, 63 and 94 days 112

Table 6: Growth performance and nutrient utilization of Nile tilapia (*O. niloticus*) cultivated in intensive and semi-intensive system in 94 days 113

Table 7: Fulton's condition factor of Nile tilapia cultivated in intensive and semi-intensive systems at times "0" (zero), 31, 63 and 94 days 113

Table 8: Relative weight (W_{rm}) of fish with 94 days of cultivation in the intensive and semi-intensive systems compared with total weight-length relationships of fish obtained in other

environments (natural and cultivated) of the experimental specimens in the analyzed environments.....114

CAPÍTULO III

Table 1: Proximate analysis of the commercial diets used in the experiment 155

Table 2: Nutritional plan used as feed for Nile tilapia (*O. niloticus*) subjected to different culture systems 155

Table 3: Growth performance and nutrient utilization of Nile tilapia (*O. niloticus*) cultivated in intensive and semi-intensive system 156

Table 4: Fulton's condition factor of Nile tilapia cultivated in intensive and semi-intensive systems at times "0" (zero), 31, 63 and 94 days 156

Table 5: Integrated density (*ID*) of the bands of protease zymograms from the intestine of Nile tilapia (*O. niloticus*) cultured in intensive and semi-intensive systems 157

Table 6: Integrated density (*ID*) of the bands of amylase zymograms from the intestine of Nile tilapia (*O. niloticus*) cultured in intensive and semi-intensive systems 157

LISTA DE ABREVIATURAS

a: Coefficient of the arithmetic weight-length relationship
Coeficiente da relação aritmética peso-comprimento

ADG: Average daily gain
ganho médio diário

Ala: Alanine
Alanina

am: coefficient of the weight-length relationship (geometric mean)
Coeficiente da relação peso-comprimento (média geométrica)

ANPU: Apparent net protein utilization
Utilização da proteína líquida aparente

AOAC: Official methods of analysis
Métodos oficiais de análise

Arg: Arginine
Arginina

b: Exponent of the arithmetic form of the weight–length relationship
Expoente da forma aritmética da relação peso-comprimento

BApNA: N α -benzoyl-DL-arginine-*p*-nitroanilide
N α -benzoil-DL-arginina-*p*-nitroanilida

BCP_f: Final body crude protein
Proteína bruta corporal final

BCP_i: Initial body crude protein
Proteína bruta corporal inicial

BHT: Butylated hydroxytoluene
Hidroxitolueno butilado

bm: Exponent of the weight–length relationship (geometric mean)
Expoente da relação peso-comprimento (média geométrica)

BW_f: Average final body weight_(g) of fish
Peso corporal médio final (g) do peixe

BW_i: Average initial body weight_(g) of fish
Peso corporal médio inicial (g) do peixe

CONAMA: Brazilian Council for Environment
Conselho Nacional do Meio Ambiente

DMSO: Dimethyl sulfoxide
Dimetil sulfóxido

DNSA: 3,5-Dinitro salicylic acid
3,5-Ácido dinitro salicílico

DTT: Dithiothreitol
Ditiotreitol

EC: Enzyme Commission
Comissão de enzimas

EMPAF: Empresa de Armazenagem Frigorífica

FAO: Food and Agriculture Organization
Organização para a Alimentação e Agricultura

FB: final biomass
Biomassa final

FCR: Feed conversion ratio
Taxa de conversão alimentar

FE: Feed efficiency
Eficiência alimentar

GIFT: Genetically Improved Farmed Tilapia
Tilapia cultivada melhoradas geneticamente

Gly: Glycine
Glicina

Hb: Hemoglobin
Hemoglobina

His: Histidine
Histidina

IBAMA: Instituto Brasileiro do Meio Ambiente e dos Recursos Naturais Renováveis

Iso: Isoleucine
Isoleucina

IUBMB: União Internacional de Bioquímica e Biologia Molecular

K: Condition factor
Fator de condição

kDa: Kilodaltons
Quilodaltons

Kmean: Mean condition factor
Fator de condição médio

Krel: Relative condition factor
Fator de condição relativo

L: Total length of fish
Comprimento total do peixe

L:D: Light:dark
Luz:escuro

Leu: Leucine
Leucina

MM: Mass marker
Marcador de massa

mM: milimolar

µM: micromolar

µS: microsiemens

MPA: Ministério da Pesca e Aquicultura

Ns: No statistical differences
Sem diferença estatística

PBS: Phosphate buffered saline
Tampão fosfato-salino

P/GE: Protein/ gross energy
Proteína / energia bruta

PepT1: Peptide transport
Peptídio de transporte

PER: Protein efficiency ratio
Taxa de eficiência proteica

PMSF: Phenyl-methylsulfonyl-fluoride
Fluoreto de fenilmetilsulfônilo

p-NPP: *p*-nitrophenyl palmitate
p-nitrofenil palmitato

p-NP: *p*-nitrophenol
p-nitrofenol

Pro: Proline
Prolina

S: Survival
Sobrevivência

SApNA: Suc-Ala-Ala-Pro-Phe-*p*-nitroanilide
Suc-Ala-Ala-Pro-Fe-*p*-nitroanilida

SDS-PAGE: Dodecyl sulfate polyacrylamide gel electrophoresis
Dodecil sulfato eletroforese em gel de poliacrilamida

Ser: Serine
Serina

SGR: Specific growth rate
Taxa de crescimento específico

SPH: Shrimp protein hydrolysate
Hidrolisado proteico de camarão

TCA: Trichloroacetic acid
Ácido tricloroacético

TLCK: Tosyl-lysine chloromethyl ketone
Tosil-lisina clorometil cetona

TPCK: Tosyl phenylalanine chloromethyl ketone
Tosil fenilalanina clorometil cetona

Tris: Tris-hydroxymethyl aminomethane
Tris-hidróximetil aminometano

Trp: Tryptophan
Triptofano

Tyr: Tyrosine
Tirosina

U: Unit of enzyme activity
Unidade de atividade enzimática

UI: International unit
Unidade internacional

W: Eeighth of fish in grams
Peso do peixe em gramas

WG: Weight gain rate
Taxa de ganho de peso

WLR: Weight-length relationships
Relação peso-comprimento

Wr_m: Relative weight
Peso relativo

SUMÁRIO

1. INTRODUÇÃO	21
2. REVISÃO BIBLIOGRÁFICA	23
3. OBJETIVOS	40
4. REFERÊNCIAS	41
5. CAPÍTULO I.....	55
Digestive enzyme activity in juvenile Nile tilapia (<i>Oreochromis niloticus</i> , L) submitted to different dietary levels of shrimp protein hydrolysate.....	58
6. CAPÍTULO II	71
Use of condition factor and its derivatives to evaluate the effect of different culture systems on the growth of Nile tilapia (<i>Oreochromis niloticus</i> L.)	72
7. CAPÍTULO III.....	119
Enzymatic changes in the intestine of Nile tilapia (<i>Oreochromis niloticus</i> L.) under intensive and semi-intensive culture systems.....	120
8. CONCLUSÃO	167
9. ANEXOS	168

1. INTRODUÇÃO

O aumento na demanda por recursos pesqueiros para consumo humano tem gerado um crescimento na produção aquícola mundial. De acordo com a Organização das Nações Unidas para Agricultura e Alimentação (FAO, 2012) foram produzidos 154 milhões de toneladas de pescado em 2011 (90,4 milhões provenientes da pesca e 63,6 milhões da aquicultura). O extrativismo ainda é responsável por cerca de 60% do total de pescado produzido.

No entanto, as pressões antrópicas exercidas sobre os ecossistemas têm levado a uma estabilização na obtenção de recursos pesqueiros nos últimos anos, sendo improvável que ocorra um aumento substancial nas capturas totais (JABLONSKI, 2005). Desta forma, o suprimento de organismos aquáticos para atender a demanda global inevitavelmente terá que vir através da aquicultura.

Nas últimas três décadas (1980-2010), a produção mundial da aquicultura se expandiu por quase 12 vezes mais, com uma taxa média de crescimento anual de 8,8 %. Com destaque para produção em águas interiores, que além de representar cerca de 60 % do total cultivado, apresentou um crescimento de mais de 40 % no período de 2006 a 2011, passando de 31,3 para 44,3 milhões de toneladas (FAO, 2012).

Espécies exóticas têm sido amplamente introduzidas e utilizadas para a produção em massa. Bem sucedidas internacionalmente, espécies como as tilápias têm merecido destaque na aquicultura mundial, especialmente a tilápia do Nilo (*Oreochromis niloticus* L.).

No Brasil, o seu cultivo apresenta-se em expansão, principalmente nas regiões Nordeste e Sudeste (IBAMA, 2008), devido as suas características biológicas e mercadológicas relevantes (FARIA et al., 2001).

O aumento da produtividade tem trazido benefícios ao setor aquícola em todos os elos da cadeia produtiva do pescado. Entretanto, este crescimento deve ser direcionado com o intuito de dar subsídios para a prática sustentável da atividade (BUENO et al., 2011). Levando em consideração o conceito de sustentabilidade dos recursos naturais no contexto da exploração florestal e pesqueira, e de acordo com a definição da Comissão Mundial para o Desenvolvimento Econômico: “*O desenvolvimento sustentável é aquele que atende às necessidades do presente, sem comprometer a capacidade das gerações futuras em satisfazer as suas próprias*” (CASTELLO, 2007).

Corroborando com o conceito de sustentabilidade, Abery et al. (2005) e Henry-Silva et al. (2006) demonstraram que se não há um planejamento da produção, impactos negativos podem ser gerados não só sobre o sistema, mas também sobre os ecossistemas aquáticos, tornando a produção insustentável.

Uma forma de manter a sustentabilidade de um sistema de produção é a aplicação de um manejo viável tanto para os aquicultores como para os mananciais. Manejo inadequado além de promover uma produção ecologicamente insustentável pode desestabilizar parâmetros de crescimento, afetando diretamente os custos de produção. No entanto, os animais podem demonstrar algumas compensações fisiológicas, adequando-se quando necessário às mudanças ocorridas no meio.

A aquicultura brasileira têm se profissionalizado bastante nos últimos anos, e os modelos produtivos adotados estão baseados em sistemas intensivos de criação, onde são utilizadas rações como principal fonte de nutrientes para peixes (BUENO et al., 2011). Isto pode levar os produtores a optarem por dietas menos onerosas gerando uma falta de controle nutricional e um aumento nos custos e nos resíduos depositados no meio.

Saldaña e Lopez (1988) mencionam que o valor nutricional de uma dieta não se baseia simplesmente na sua composição química, mas também na capacidade fisiológica do peixe em digerir e absorver, de acordo com seus hábitos alimentares. Esta digestão é executada no trato digestório com o auxílio de enzimas (NIELSEN -SCHIMIDT, 1996).

As enzimas digestivas são estudadas como uma maneira de compreender as exigências nutricionais e o efeito dos constituintes da dieta. Sabendo que a alimentação e o manejo podem influenciar a atividade e a expressão gênica de algumas enzimas, como também em outras proteínas, refletindo no crescimento dos peixes e nos impactos ambientais, entender seu modo de regulação é importante para o uso racional do tipo e quantidade de nutrientes empregados nas dietas (MUHLIA-ALMAZÁN et al., 2003), podendo também fornecer informações sobre a forma de adaptação dos animais a diferentes sistemas de cultivo.

2. REVISÃO BIBLIOGRÁFICA

2.1 Tilápia do Nilo

A tilápia é um peixe de água doce pertencente à família Cichlidae. Elas são nativas da África, mas foram introduzidas em muitas regiões tropicais, subtropicais e temperadas do mundo durante a segunda metade do século 20 (EL-SAYED, 2006).

As espécies de importância comercial estão divididas em três gêneros de acordo com seu comportamento reprodutivo. O gênero *Tilapia* incuba seus ovos no substrato; *Sarotherodon*, tanto os machos como as fêmeas incubam os ovos na boca e *Oreochromis*, as fêmeas são responsáveis pela incubação dos ovos (TREWAVAS, 1982).

Elas habitam uma grande variedade de ecossistemas, como rios permanentes e temporários, rios com corredeiras, grandes rios equatoriais, tropicais e subtropicais, lagos pantanosos, lagos rasos e profundos, lagos artificiais, alcalinos, ácidos e salinos, fontes termais, lagos em cratera, lagos com baixo conteúdo de minerais, estuários abertos e fechados, lagos de água salobra e habitat marinho. A diversidade de *habitats* mencionados promove uma fácil adaptação às mudanças ambientais, refletindo em sua tolerância a parâmetros físicos (variações de temperatura, fotoperíodo, correnteza, profundidade, turbidez, etc), químicos (pH, salinidade, minerais, oxigênio dissolvido, amônia, etc) e biológicos (concorrência, disponibilidade de alimentos, etc.) (EL-SAYED, 2006).

Durante os estágios larvais, a tilápia se alimenta inicialmente de fitoplâncton e zooplâncton, especialmente crustáceos (copépodos) (BOWEN, 1982). O período de transição do estágio planctófago para uma dieta especializada é geralmente curto (BOWEN, 1976). A alimentação de juvenis e adultos consiste de uma variedade considerável de vegetação aquática, zooplâncton, fitoplâncton, perifítón e detritos de origem vegetal. Por esta razão, a tilápia pode mudar de uma fonte de alimento para outro com poucas mudanças na composição da dieta (EL-SAYED, 2006).

O século 20 firmou o cultivo de tilápia. Este desenvolvimento se deu principalmente pela demanda do filé pelos Estados Unidos, o cultivo na África e América do Sul, o melhoramento genético e a expansão do cultivo em águas salinas nas Filipinas. China se tornou o maior produtor e grande fornecedor dos EUA (GUERRERO, 2008).

Dados da FAO (2009) demonstraram claramente que a produção mundial de tilápia aumentou de 398.066 t em 1991 para 2.326.413 t em 2006, promovendo um rendimento de US\$

559.438 milhões para mais de US\$ 2,7 bilhões. De acordo com Wing-Keong e Hanin (2007), *Oreochromis niloticus* (Figura 1) representa 80% das espécies de tilápia cultivadas no mundo, sendo considerada a mais importante.

No Brasil, sua produção em 2006 foi de aproximadamente 71.000 t, representando 37,8% da produção aquícola continental (IBAMA, 2008). Em 2010 esta produção ultrapassou 155.000 t representando 39% da aquicultura continental no país (BRASIL, 2012).

Devido as suas características biológicas e mercadológicas como rápido crescimento, rusticidade, alimentação em baixos níveis tróficos, tolerância a variações ambientais, resistência a doenças, ausência de espinhos intra-musculares, dentre outros, a tilápia do Nilo é considerada uma das principais espécies da piscicultura mundial e a principal espécie brasileira.

Figura 1: Exemplar de tilápia do Nilo (*Oreochromis niloticus* L.).



Fonte: <http://www.terradagente.com.br/fauna/0,0,2,288;5,tilapia-do-nilo.aspx>

2.2 Sistemas de cultivo

A tilápia é produzida em diferentes sistemas de cultivo e escalas, com maior ou menor dependência de insumos, principalmente ração (KÖPRÜCÜ e ÖZDEMİR, 2005). Sua produção em países em desenvolvimento ocorre primeiramente em sistema semi-intensivo, onde fertilizantes inorgânicos ou adubos são usados para produção do alimento natural (MATAKA e KANG'OMBE, 2007). Sistema semi-intensivo (Figura 2) constitui um meio de produção de baixo custo, sua

produtividade é geralmente menor do que a dos sistemas intensivos devido à menor densidade de estocagem e insumos de produção. Apesar disto, sua popularidade tem aumentado nas duas últimas décadas em vários países do mundo, especialmente entre os pequenos agricultores (EL-SAYED, 2006).

Figura 2: Cultivo semi-intensivo de tilápia do Nilo



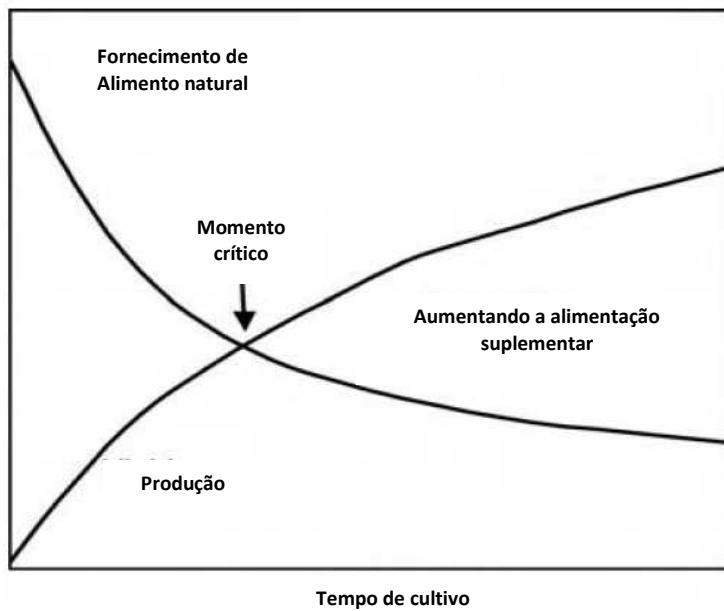
Fonte: Juliana Ferreira dos Santos

O cultivo em viveiros escavados apresenta uma menor incidência de doenças, maior sobrevivência na recria e engorda, contribuição do alimento natural e uma melhor conversão alimentar quando comparado a sistema intensivo em tanques rede (KUBITZA, 2009).

Neste sistema pode-se variar a biomassa de 0,1 a 0,5 Kg/m² dependendo do aporte de alimentação suplementar. Quando a alimentação da tilápia é suplementada com baixas concentrações de proteína em cultivo semi-intensivo, o déficit é suprido pelo alimento natural (LITI et al., 2006). No entanto, alimento natural é deficiente em energia não protéica, induzindo a utilização de parte desta proteína natural no metabolismo energético (EDWARDS et al., 2000).

Diana et al. (1994) avaliaram os efeitos da adição de alimento suplementar em viveiros fertilizados no crescimento de tilápia do Nilo, e observaram que a combinação da alimentação com fertilizantes foram mais eficientes do que um desses dois itens sozinhos. A Figura 3 demonstra a importância da alimentação suplementar em cultivos semi-intensivos. Green et al. (2002) também avaliaram o cultivo de tilápia do Nilo em viveiros fertilizados e constataram uma melhor produtividade com a alimentação suplementar.

Figura 3: Desenho esquemático da relação entre produção, alimento natural e alimentação suplementar em viveiros escavados.



Fonte: El-Sayed (2006).

Dietas convencionais podem aumentar a produtividade, no entanto também podem aumentar os custos neste tipo de sistema, tornando o cultivo mais oneroso. Desta forma, tem se focado no uso de recursos que possam tornar este cultivo mais viável (MATAKA e KANG'OMBE, 2007). Um regime alimentar adequado, baseado em taxas de alimentação, como demonstrado por Abdelghany e Ahmad (2002) pode reduzir tanto os custos como a carga de nutrientes nos viveiros.

A expansão do cultivo de tilápia no mundo juntamente com a escassez de água doce, competição com a agricultura e com as atividades urbanas, tem mudado gradualmente os tradicionais sistemas semi-intensivos de criação de tilápia para sistemas mais intensivos de produção (EL-SAYED, 2006). Sistema intensivo caracteriza-se pelas altas densidades (25 a 300 peixes/m³), dietas com altos níveis proteicos e uma alta produtividade. Este tipo de cultivo vem se difundido para tilápia com o uso de tanques-rede (MOHAMMAD, 2006). O cultivo em tanques-rede (Figura 4) é visto como um caminho para aumentar a produção, proporcionar empregos e oportunidades (ABERY et al., 2005). Uma das vantagens do cultivo de tilápia em tanques-rede é que ele pode ser aplicado em vários corpos aquáticos, incluindo rios, lagos, reservatórios, lagoas, água do mar, viveiros, etc (WATANABE et al., 1990; PERSAND e BHAKAJEE, 1997; FITZSIMMONS et al., 1999; ORACHUNWONG et al., 2001; YI e LIN, 2001).

O cultivo em gaiolas apresenta como características uma tecnologia relativamente barata, quando comparado a outros sistemas intensivos de criação; pode ser aplicado na maioria dos

ambientes aquáticos; assegura maior controle do estoque e melhor observação dos peixes quando comparado com viveiros (PEIXOTO, 2007). Além disso, ele também apresenta como características positivas um tratamento mais econômico de parasitas e doenças, facilidade na despensa, manuseio dos peixes com redução na mortalidade, melhores taxas de crescimento, redução do uso de terra para construção de viveiros, controle na reprodução, boa utilização do alimento artificial, etc. (EL-SAYED, 2006). Ostrensk et al. (2008) relatam que o investimento necessário para a produção de uma tonelada de peixe em tanque rede está na ordem de 30-40% da produção em viveiros convencionais. Este fato, aliado às altas produtividades tem sido responsável pela grande expansão do sistema intensivo no país. Além disso, enquanto o sistema semi-intensivo produz cerca de 1 kg a 3 kg de peixe/m³/ano, o sistema de tanques-rede pode produzir de 150 a 300 kg/m³ no mesmo período. No entanto, a intensificação do cultivo requer o desenvolvimento de rações adequadas (KÖPRÜCÜ e ÖZDEMİR, 2005), já que a maior parte dos nutrientes ofertados neste sistema vem por meio das dietas artificiais, diferente do sistema semi-intensivo de criação.

Figura 4: Cultivo intensivo de tilápia do Nilo.



Fonte: Juliana Ferreira dos Santos

Parâmetros como qualidade de água, densidade, nutrição e alimentação são os fatores que determinam a sucesso ou fracasso de um cultivo intensivo de tilápia do Nilo. De acordo com El-Sayed (2006), o cultivo em gaiolas também apresenta algumas limitações como risco de roubo, perda devido a danos na gaiola, surtos de doenças, baixa tolerância dos peixes a má qualidade da água, completa dependência de rações de alta qualidade, perda de alimentação pelas malhas da gaiola e acúmulo de fezes e metabólitos no ambiente causando impacto negativo. Boyd (1990) enfatiza que a alimentação influencia negativamente na qualidade da água do sistema se não for

bem gerenciada. Estudos feitos por Lin et al. (1989); McGinty (1991); Yi et al. (1996) e Liti et al. (2005) demonstraram que altas densidades de tilápia também podem influenciar negativamente o cultivo, promovendo redução no ganho de peso. De fato ambientes estressantes são adversos ao crescimento, função imune e reprodução, causando mortalidade e suscetibilidade a doenças e predadores. Diana et al. (1994) e Boyd (1990) também demonstraram que um manejo inadequado com quedas nos níveis de oxigênio dissolvido pode levar a diminuições nas taxas de crescimento e/ou eventuais mortalidades.

No entanto, os animais podem demonstrar algumas compensações fisiológicas recuperando-se da experiência estressante, além disso, a qualidade da alimentação fornecida pode influenciar nesta recuperação. Desta forma, sistemas de cultivos diferenciados, bem como manejo e alimentação utilizada, poderiam influenciar os animais promovendo alterações e adaptações na sua fisiologia.

2.3. Aparelho digestório X Preferência alimentar

A anatomia do trato gastrointestinal dos peixes segue o mesmo plano básico de outros vertebrados, sua compreensão também nos fornece um entendimento acerca de sua fisiologia digestiva e preferência alimentar. Os peixes podem ser classificados em quatro categorias principais com relação ao hábito alimentar: detritívoros, herbívoros, onívoros e carnívoros. Vale salientar que a versatilidade trófica envolve também a designação dos peixes como: generalistas (sem preferência acentuada por uma fonte alimentar, utilizando um amplo espectro de alimentos); especialistas (com dieta restrita a um número relativamente pequeno de itens e usualmente apresentando adaptações morfológicas tróficas) e oportunistas (que se alimentam de fonte não usual de sua dieta ou/e fazem uso de uma fonte alimentar abundante e incomum) (ABELHA et al., 2001).

Estes organismos podem ocupar todos os níveis tróficos de um ecossistema aquático, de forma a utilizar diferentes recursos alimentares, graças à grande diversificação morfológica quanto aos órgãos e estruturas relacionadas à procura, detecção, captura, ingestão e digestão do alimento (SAMPAIO e GOULART, 2011).

A compreensão da anatomia do sistema digestório nos fornece um entendimento sobre o contexto da organização espacial e sua relação com dados bioquímicos e fisiológicos, e também são essenciais para a compreensão da nutrição de peixes em aquicultura, bem como os mecanismos de adaptações fisiológicas para mudanças no ambiente (GROSELL et al., 2011).

Sampaio e Goulart (2011) observaram as relações entre os caracteres morfológicos (boca, dentes, rastros branquiais, estômago e intestino) e a alimentação dos ciclídeos, encontrando uma estreita relação entre os mesmos.

A tilápia como um peixe onívoro oportunista, é capaz de assimilar nutrientes de diferentes fontes, como as algas bentônicas, fitoplâncton, macrófitas, zooplâncton, pequenos invertebrados, detritos e cianofíceas. A escolha da alimentação é principalmente relacionada com a disponibilidade de alimentos no ambiente, indicando que a espécie possui alta plasticidade trófica (BOWEN, 1982; PHILIPPART e RUWET, 1982; FITZSIMMONS, 1997; BEVERIDGE e BAIRD, 2000; LOWE-MCCONNELL, 2000; SAMPAIO e GOULART, 2011).

Embora haja exceções, a anatomia digestiva entre espécies de uma mesma família é bastante similar. De acordo com Rust (2002), a digestão é uma ação coordenada de atividades físicas, químicas e enzimáticas que tem início quando alimento entra na boca e termina quando é excretado nas fezes.

Segundo Hyatt (1979), a forma e a posição da boca têm o potencial de influenciar na quantidade e qualidade da presa que pode ser ingerida. De acordo com Bond (1979), a boca pode ser inferior, subterminal, terminal ou superior. Sampaio e Goulart (2011) mencionam que existe profunda relação entre a morfologia da boca e o hábito alimentar. Câmara e Chellappa (1996) descrevem que a tilápia do Nilo apresenta uma boca terminal que favorece a captura de itens alimentares planctônicos.

De um modo geral a tilápia captura os alimentos por sucção (quando a presa é sugada para a cavidade bucofaringeal); captura visual (quando o peixe localiza e seleciona o alimento a ser ingerido); e pela mordida (utilizando as mandíbulas inferiores e superiores) (BEVERIDGE e BAIRD, 2000).

Os ciclídeos podem apresentar dentes na mandíbula, na maxila e na faringe, sendo estes importantes na taxonomia desta família (GOLDSTEIN, 1988). As tilápias possuem de um a cinco fileiras de dentes mandibulares, que juntamente com os dentes faringianos contribuem com a quebra dos alimentos em fragmentos menores facilitando a digestão (BOWEN, 1982; DEMPSTER et al., 1993). Câmara e Chellappa (1996) verificaram que existe estreita relação entre o desenvolvimento dos dentes e o nível trófico. A tilápia apresenta dentes rudimentares, sendo classificada como onívora-planctófaga, evidenciando que sua dieta baseia-se em organismos fitoplanctônicos e zooplancônicos adequados ao tipo de dente.

Zayed e Mohamed (2004) demonstraram que tilápia do Nilo apresenta rastros branquiais relativamente curtos com número variável nos diferentes arcos. Eles também estabeleceram uma

relação entre o comprimento e número de rastros branquiais com a categoria trófica, caracterizando a tilápia como uma espécie onívora pelos seus rastros curtos e pouco numerosos. De acordo com Sampaio e Goulart (2011) esta relação se fundamenta no fato dos peixes planctófagos ingerirem partículas pequenas, e seus rastros numerosos e longos atuarem como uma peneira para reter os alimentos menores. Já os onívoros ingerem partículas de diversos tamanhos, seus rastros branquiais em menor quantidade e curtos favorecem a captura de presas maiores.

O esôfago de peixes é geralmente um tubo curto e reto com paredes espessas, ligado da faringe até o estômago ou intestino em peixes sem estômago. Morfologicamente o esôfago é projetado principalmente para a passagem de alimentos. A junção do esôfago e estômago não é claramente demarcada anatomicamente em teleósteos, a não ser pela mudança das fibras musculares estriadas (esôfago) e lisas (estômago) (GROSELL et al., 2011). A mudança também pode ser observada com o aparecimento de glândulas gástricas, no entanto esta transição se dá de forma gradual, revelando a zona de transição entre estes dois órgãos (KAPOOR et al., 1975).

Ao passar pelo esôfago, o alimento entra no estômago onde apresenta três regiões distintas, a região anterior denominada de cardíaca, de paredes mais finas, a região mediana ou fúndica e a região terminal ou pilórica, de paredes grossas, que é uma estrutura muscular (esfíncter) que tem a função valvular (GROSELL et al., 2011). O estômago armazena temporariamente o alimento e desempenha funções mecânicas que misturam o conteúdo estomacal auxiliando na digestão dos alimentos. A superfície interna (mucosa) contém uma variedade de células glandulares endócrinas e secretoras exócrinas, que contém células secretoras de muco e células que secretam o pepsinogênio, ácido clorídrico e enzimas digestivas (RUST, 2002; ROTTA, 2003).

O esfíncter pilórico que tem uma função valvular, mantém o alimento no estômago até ele estar fluído para passar para o intestino. Ele contribui impedindo refluxo do bolo alimentar para o estômago e controla o tempo em que o alimento fica em contato com o suco gástrico, controlando também a quantidade de material gástrico que entra no intestino (RUST, 2002; ROTTA, 2003).

A função primária do intestino é a realização dos processos digestivos (com auxílio das enzimas) que se iniciaram no estômago e a posterior absorção de nutrientes, íons e água oriundos da dieta (ROTTA, 2003; GROSELL et al., 2011). De acordo com Horn (1997) o comprimento do intestino é usado como um indicador do nível trófico em peixes. Em herbívoros seu comprimento é geralmente maior, quando comparados aos carnívoros (CLEMENTS e RAUBENHEIMER, 2005). No entanto, Kramer e Bryant (1995) alertam que a relação entre o comprimento do intestino e o hábito alimentar deve ser aplicada apenas na identificação de grandes grupos. Já Harder (1975) diz

que não há relações claras entre morfologia intestinal e o tipo de alimentação, e que esta relação não é conclusiva.

O intestino é um tubo relativamente simples, iniciando na válvula pilórica e terminando no reto (em peixes sem estômago). Ele possui glândulas digestivas e um suprimento de vasos de sangue e linfa, onde se completa a digestão iniciada no estômago (ROTTA, 2003). Uma característica básica do intestino de peixes teleósteos é a presença de pelo menos dois segmentos intestinais. O intestino superior apresenta uma quantidade mais escassa de células de Globet do que o inferior, este por sua vez apresenta uma camada muscular mais fina do que o superior. Há uma mudança de um epitélio colunar de células secretoras e de absorção para um epitélio que secreta basicamente muco no intestino inferior. De um modo geral, as porções intestinais que apresentam células mais complexas estão envolvidas com processos absorтивos (SMITH, 1989).

Os cecos pilóricos são estruturas associadas ao intestino superior que apresentam formato digitiforme. Hossain e Dutta (1996) estimam que 60% das espécies de peixes conhecidas possuem cecos pilóricos. Buddington e Diamante (1987) demonstraram que os cecos pilóricos aumentam a área de superfície para a digestão e absorção, mas não apresentam um papel na fermentação ou no armazenamento. Rotta (2003) relata que não está bem definido se existe uma relação entre a presença de cecos pilóricos e o hábito alimentar dos peixes, pois eles ocorrem nos carnívoros, onívoros e herbívoros. Eles são responsáveis pela produção de enzimas digestivas e secreção de hormônios como insulina e glucagon. As tilápias não apresentam cecos pilóricos, mas em compensação possuem um intestino que pode variar de 7 a 14 vezes o comprimento do corpo, o qual pode compensar a ausência dos mesmos (POPMA e LOVSHIN, 1996).

De acordo com Rotta (2003) o pâncreas da maioria dos teleósteos é difuso e não pode ser facilmente observado durante uma dissecação total. Smith (1989) e Rotta (2003) afirmam que o pâncreas pode se apresentar em uma estrutura individualizada, ou de forma difusa, com nódulos pancreáticos espalhados no tecido adiposo, no mesentério, fígado, duto biliar, vesícula biliar, cecos pilóricos, intestino, etc. Nas lampreias, um pâncreas exócrino é inexistente e as células pancreáticas e zimogênios estão presentes no epitélio intestinal, sendo considerada uma condição primitiva (GROSELL et al., 2011).

2.4 Enzimas digestivas

Saldaña e Lopez (1988) mencionam que o valor nutricional de uma dieta não se baseia simplesmente na sua composição química, mas também na capacidade fisiológica do peixe em

digerir e absorver, de acordo com seus hábitos alimentares. Durante a digestão as proteínas, carboidratos e lipídios são degradados em compostos mais simples para então serem absorvidas e utilizadas pelo corpo. Esta degradação é executada no trato digestório com o auxílio de enzimas (NIELSEN-SCHIMIDT, 1996).

As enzimas são catalizadores biológicos que aumentam de forma extraordinária as reações químicas. Sua eficiência catalítica, em geral, é bem maior que a dos catalisadores sintéticos e inorgânicos, sendo fundamentais para qualquer processo bioquímico. Com exceção de alguns RNAs catalíticos, todas as enzimas conhecidas são proteínas e podem ser classificadas de acordo com a reação específica que ela catalisa (tabela 1) (NELSON e COX, 2004).

Tabela 1: Classificação das enzimas segundo IUBMB.

No.	Classe	Tipo de reação catalisada
1	Oxirreduases	Transferência de elétrons (íons hidretos ou átomos de H)
2	Transferases	Reações de transferência de grupos
3	Hidrolases	Reações de hidrólise
4	Liases	Adição de grupos as duplas ligações ou formação de duplas ligações por meio de remoção de grupos
5	Isomeraseas	Transferência de grupos dentro da mesma molécula para formar isômeros
6	Ligases	Formação de ligações do tipo C-C, C-S, C-O e C-N por meio de reações de condensação acopladas a quebra do ATP

IUBMB: União Internacional de Bioquímica e Biologia Molecular. C: Carbono; N: Nitrogênio; S: Enxofre; O: Oxigênio. (Fonte: NELSON e COX, 2004).

De acordo com a IUBMB as proteases estão inseridas no subgrupo 4 do grupo 3 (Hidrolases), pois por uma reação de hidrólise, elas clivam a proteína adicionando uma molécula de água à ligação peptídica (BERG et al., 2004). A IUBMB recomenda o termo "peptidase" como sinônimo de protease, que é designado para qualquer enzima que hidrolise ligações peptídicas. A nomenclatura das peptidases é problemática, de um modo geral sua especificidade é geralmente difícil de definir, pois apresentam uma grande variedade de estruturas e ações. No entanto, uma classificação envolvendo o mecanismo catalítico é, por conseguinte utilizada.

Dois conjuntos de sub-subclasses de peptidases são conhecidas; as exopeptidases (EC 3.4.11 e EC 3.4.13-19) e as endopeptidases (EC 3.4.21-25). As exopeptidases atuam próximas as extremidades de cadeias polipeptídicas em uma extremidade N-terminal liberando um resíduo de aminoácido (aminopeptidases; EC 3.4.11), de um dipeptídeo ou de um tripeptídeo (dipeptidase e tripeptidases, EC 3.4.14). As exopeptidases que atuam no C-terminal livre libera um único resíduo (carboxipeptidases, EC 3.4.16-18), ou um dipeptídeo (peptídeo-dipeptidases; EC 3.4.15). As carboxipeptidases são divididas em três grupos com base no mecanismo catalítico: as

serinocarboxipeptidases (EC 3.4.16), os metalocarboxipeptidases (EC 3.4.17) e as cisteíncarboxipeptidases (EC 3.4.18). Outras exopeptidases são específicas para os dipeptídeos (dipeptidases, EC 3.4.13), ou para a remoção de resíduos terminais que são substituídos ou ligados por ligações isopeptídicas (omega peptidases , CE 3.4.19).

As endopeptidases são divididas em sub-subclasses, com base no mecanismo catalítico, e a especificidade é utilizada apenas para identificar enzimas individuais dentro dos grupos. As sub-subclasses são: serina endopeptidases (serinoproteases) (EC 3.4.21), cisteína endopeptidases (cisteínpotease) (EC 3.4.22), aspartato endopeptidases (aspartatoproteases) (EC 3.4.23), metaloendopeptidases (metaloproteases) (EC 3.4.24) e treonina endopeptidases (treoninoprotease) (EC 3.4.25).

A tabela 2 baseada em Rao et al. (1998) demonstra a classificação e divisão das proteases.

Tabela 2: Classificação e divisão das proteases

Local de clivagem do substrato	Sítio ativo da enzima	Número de resíduos de aminoácidos removidos
Exopeptidases	Aminopeptidases	Aminopeptidases
		Aminodipeptidases
		Aminotripeptidases
		Sítio ativo da carboxipeptidase
Endopeptidases	Carboxipeptidases	Serinocarboxipeptidase
		Metalocarboxipeptidase
		Cisteíncarboxipeptidase
	Serinoproteases	Serinoproteases
		Aspartatoproteases
		Cisteínpoteases
		Metaloproteases

Fonte: Rao et al. (1998).

As carboidrases e lipases também estão incluídas no grupo das hidrolases. As lipases (triacilglicerol acilhidrolases) são carboxilesterases que apresentam habilidade para hidrolisar ésteres de ácido carboxílico de cadeia longa, elas catalisam a hidrólise de ligações éster na interface orgânica-aquosa. Elas são enzimas ubíquas e são encontradas na maioria dos organismos (microbianos, vegetais e animais) (ABOUSALHAM e VERGER, 2000). Lipases são fisiologicamente importantes porque digerem gorduras em ácidos graxos para a absorção através da membrana celular, e transformam triglicerídeos em mais moléculas polares (LÓPEZ-LÓPEZ et al., 2003).

De acordo com Rotta (2003), o estômago dos peixes é capaz de secretar lipase gástrica, no entanto, se comparada à pancreática, tem pouca atuação no processo digestivo das gorduras e, em geral, hidrolisa apenas as gorduras de baixo ponto de fusão e já emulsificadas. As lipases de origem pancreática secretadas para o lúmen intestinal são as de maior importância na digestão das gorduras.

As carboidrases, também conhecidas como glicosilases, são enzimas que catalisam a degradação de carboidratos, ou seja, hidrolisam as ligações glicosídicas entre monossacarídeos formadores de oligossacarídeos ou polissacarídeos. De acordo com Kuz'mina et al. (1996) estas enzimas atuam na cavidade intestinal e na borda em escova dos enterócitos da parede intestinal em peixes, no entanto, há relatos de sua atividade no estômago de algumas espécies. De acordo com Silveira et al. (2009) as carboidrases aparecem também no suco pancreático, estômago, intestino e bile, mas não necessariamente em todos locais descritos para as espécies pesquisadas.

De um modo geral, os peixes aproveitam melhor os carboidratos complexos, como o amido, do que os açucares simples (New, 1987). Dentre as carboidrases estudadas em organismos aquáticos as amilases merecem atenção especial, já que são responsáveis pela degradação do amido e glicogênio.

As amilases são enzimas que catalisam a hidrólise de ligações glicosídicas $\alpha(1 \rightarrow 4)$ de polissacarídeos presentes no amido e glicogênio. Sobre o amido, atuam liberando diversos produtos, incluindo dextrinas, maltose e glicose. Elas podem ser divididas em dois grupos: as endoamilases e exoamilases. As endoamilases atuam de forma aleatória no interior da molécula do amido, causando a formação de ramos lineares de oligossacarídeos de cadeias de vários comprimentos e dessa forma quebram as ligações glicosídicas $\alpha(1 \rightarrow 4)$ presentes nas cadeias de amilose ou amilopectina (GUPTA, 2003). Diferentemente das β -amilases (exoamilases), que são encontradas em vegetais superiores e em algumas bactérias gram-positivas (PUJADAS et al., 1996) as α -amilases podem ser encontradas em animais, plantas, fungos e bactérias.

Em vertebrados superiores, as enzimas podem se apresentar distintamente, demonstrando claramente suas áreas funcionais. Para peixes, isto não é mostrado de forma clara, uma vez que as mesmas podem estar distribuídas ao longo do trato digestório (LUNDSTEDT et al., 2004). Rotta (2003) afirma que a maior parte da digestão dos alimentos ocorre no intestino e nos cecos pilóricos, quando presentes. Segundo o autor as três maiores classes de enzimas (proteases, lipases e carboidrases) estão presente com grande atividade nas secreções intestinais.

Geralmente, os fatores externos, como hábito alimentar e quantidade consumida desempenham um papel importante na distribuição e intensidade das atividades enzimáticas (TENGJAROENKUL et al., 2000). Elas também podem ser influenciadas pela idade e a espécie do

animal, bem como pela quantidade e composição da dieta ministrada (PEREZ et al., 1998). Jun-sheng et al. (2006) associam a distribuição e a intensidade de enzimas intestinais com o hábito alimentar e a morfologia intestinal. Seixas-Filho et al. (1999) citam que a produção da enzima α -amilase ocorre estritamente no pâncreas e no intestino, principalmente entre as espécies de peixes carnívoras e onívoras.

Rotta (2003) ainda afirma que peixes carnívoros apresentam maior atividade de lipases do que peixes onívoros e herbívoros. Seixas Filho et al. (2000) avaliaram a atividade lipolítica de três peixes de água doce com hábito alimentar diferente, encontrando uma baixa atividade para a espécie onívora piracanjuba (*Brycon orbignyanus*), e alta para as espécies piau (*Leporinus friderici*) e surubim (*Pseudoplatystoma curuscans*), onívora e carnívora, respectivamente.

Reimer (1982), estudando o matrinchã (*Brycon melanopterus*), observou que a atividade lipolítica está relacionada com a quantidade de gordura presente no conteúdo intestinal, e a atividade amilolítica aumenta quando a dieta é rica em amido. Fountoulaki et al. (2005) também demonstraram que as diferenças no perfil enzimático estão relacionados com os nutrientes presentes na dieta.

Já Nagase (1964), observou que não houve aumento na atividade de lipase quando a tilápia mossambica (*Sarotherodon mossambicus*) recebeu dieta rica em gordura. Hidalgo et al. (1999) analisaram as atividades proteolíticas e de amilase em seis espécies de peixes com diferentes hábitos alimentares, e observaram que a proporção de amilase total: atividade proteolítica total foi maior em peixes onívoros.

Kuz'mina et al. (1996) encontraram uma correlação entre a atividade amilolítica e sacarose total da mucosa intestinal, a concentração de carboidratos da dieta, e a intensidade de alimentação dos peixes. Jun-Sheng et al. (2006) observaram um aumento de atividade enzimática em tilápia com um aumento no peso corporal, sugerindo uma capacidade para utilizar alimentação de acordo com sua fase de crescimento.

Kumar et al. (2007) avaliando atividade proteolítica em três carpas (*Catla catla*, *Labeo rohita* e *Hypophthalmichthys molitrix*) observou uma atividade proteolítica total mais elevada em *L. rohita* (plânctofaga) seguido de *H. molitrix* (fitoplânctofaga) e posteriormente *C. catla* (zooplânctofaga). As atividades de tripsina e quimotripsina foram superiores em *H. molitrix*. Os autores sugerem que peixes herbívoros podem compensar sua dieta com proteína baixa, aumentando a sua atividade enzimática. Alguns peixes herbívoros apresentam atividades semelhantes de tripsina ou mesmo superiores quando comparados às espécies carnívoras, possivelmente para maximizar a eficiência de digestão de proteína.

Cara et al. (2007) avaliando as atividades de quimotripsina e tripsina em larvas de *Dicentrarchus labrax*, observaram que a atividade dos animais com restrição alimentar foram superiores a das larvas com melhor qualidade na alimentação, sugerindo que o aumento nas atividades seriam para compensar as deficiências alimentares no qual as larvas foram submetidas.

Assim, um bom entendimento do modo de regulação das enzimas digestivas se torna importante para o uso racional do tipo e quantidade de nutrientes utilizados nas dietas (MUHLIA-ALMAZÁN et al., 2003). Estudos nutricionais e enzimáticos contribuem para formulação de rações mais eficientes, reduzindo de forma extraordinária os custos na produção de peixes comerciais (CARUSO, et al., 1996). Além disto, podem-se diminuir os resíduos depositados no ambiente, que é imprescindível para tornar a aquicultura uma atividade sustentável e ecologicamente viável.

Tilápias são bem conhecidas pela sua capacidade de utilizar uma grande variedade de alimentos (LOWE-MCCONNELL 1975; BOWEN, 1982; PHILIPPART e RUWET, 1982; TREWAVAS 1983). Molina et al. (2005) e Lu et al. (2006) demonstraram que a tilápia tem a capacidade de ingerir e digerir cianobactérias em lagos eutrofizados, sugerindo o amplo espectro de digestão deste peixe. Funções digestivas capazes de hidrolisar uma variedade maior de alimentos tornam os peixes onívoros com capacidade de digestão variada, comparada com os carnívoros (ALMEIDA et al., 2006). Dentre as espécies onívoras, a tilápia tem se destacado pela elevada capacidade de utilizar nutrientes de origem vegetal e animal, o que possibilita a elaboração de rações de baixo custo e alto valor nutritivo (SILVA et al., 2006).

2.5 Utilização de resíduos da indústria pesqueira para fabricação de rações de organismos aquáticos

Uma boa alimentação deve satisfazer as necessidades nutricionais das espécies cultivadas, como também ser de fácil digestão e com uma boa perspectiva custo-benefício (LEMOS et al., 2004).

A determinação de fontes proteicas de menor custo e que promovam bom crescimento, é vantajoso tanto para a indústria de rações, como também para os aquicultores (COYLE et al., 2004). Sendo assim, vários estudos têm sido realizados para avaliar novas fontes proteicas que poderiam ser usadas na fabricação de rações (JACKSON et al., 1982; GABER, 1996; OLVERA-NOVOA et al., 1997; OLVERA-NOVOA et al., 2002; PLASCENCIA-JATOMEA et al., 2002; EL - SAIDY e GABER, 2003; GABER, 2006, LEAL et al., 2009). Tais recursos devem garantir os mesmos ou melhores resultados de produção, sanidade do animal, crescimento e qualidade do produto.

O crescimento da indústria de pescado tem gerado uma grande quantidade de resíduos e subprodutos que representam um desafio para os empresários do setor e para a comunidade científica especializada, em buscar estratégias para que essa atividade seja sustentável e ecologicamente viável. Segundo Arruda (2004), cerca de 50 % do pescado mundial produzido em 2000 transformou-se em resíduo. Supondo-se que esse percentual tenha-se mantido, dos 154 milhões de toneladas de pescado produzidos em 2011 (FAO, 2012), 77 milhões teriam sido descartados ou subutilizados, constituindo-se em uma fonte significativa de desperdício de recursos e de contaminação ambiental.

Ao longo dos anos, diversos têm sido os esforços empreendidos por pesquisadores em todo o mundo para desenvolver métodos que possibilitem a transformação desses resíduos em produtos passíveis de utilização tanto na nutrição humana quanto na animal (ARMENTA e GUERRERO-LEGARRETA, 2009; CHEN et al., 2011; CAHÚ et al., 2012).

A produção de farinha, hidrolisado proteico ou silagem de camarão a partir de subprodutos das indústrias pesqueiras (Figura 5) representa uma excelente alternativa para o incremento da oferta de proteína animal (MACKIE, 1982; HAARD, 1992; KENT, 1997; CAVALHEIRO et al., 2007), já que estes subprodutos são usualmente descartados.

Figura 5: Resíduos da indústria de camarão.



Fonte: Juliana Ferreira dos Santos

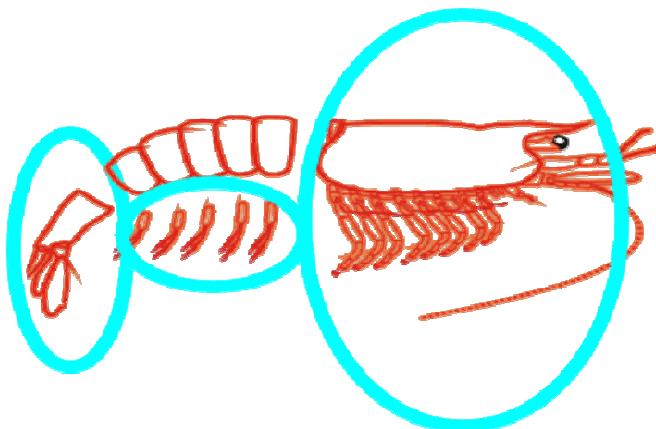
No entanto, o uso de farinha de cabeça de camarão na formulação de alimentos para organismos aquáticos não é recomendado devido a sua alta concentração de fibras e cinzas, o que resulta na formação de péletes fracos com uma baixa estabilidade em água (CAVALHEIRO et al., 2007).

Uma alternativa seria a produção de proteína hidrolisada. Vários são os métodos empregados para obtenção do hidrolisado proteico dos produtos e subprodutos pesqueiros (GILDBERG, 1993). A presença de enzimas proteolíticas no trato digestório de animais aquáticos tem uma influência significante na produção de hidrolisados, que podem ser obtidos empregando o processo de autólise ou um método de hidrólise, através da adição de enzimas (SHAHIDI et al., 1995). Gilberg e Stenberg (2001) demonstraram que a proteína dos subprodutos do camarão pode ser hidrolisada por proteases comerciais e recuperada como hidrolisado proteico com alto conteúdo de aminoácidos essenciais. A hidrólise utilizando substâncias exógenas permite um melhor controle das propriedades dos produtos resultantes. No entanto, os custos das substâncias (enzimas, ácidos e solventes) são um obstáculo para a viabilidade do processo. Plascencia-Jatomea et al. (2002) relataram que as condições ácidas em que o hidrolisado de silagem de camarão é produzido faz com que haja perda de nutrientes lábeis, como o triptofano.

A autólise é um processo enzimático que pode ser usado como uma alternativa aos processos que empregam substâncias exógenas, além de haver a possibilidade de recuperar moléculas bioativas, tais como quitina (para a produção de quitosana) e carotenóides (ARMENTA e GUERRERO-LEGARRETA, 2009; CHEN et al., 2011; CAHÚ et al., 2012).

No processamento do camarão, geralmente são removidas a cabeça, o exoesqueleto e a porção posterior (Figura 6). Estes subprodutos correspondem a aproximadamente 50% do seu peso total, o que torna importante seu aproveitamento do ponto de vista econômico, industrial e ambiental (HEU et al., 2003).

Figura 6: Desenho esquemático de camarão identificando os resíduos da indústria pesqueira passíveis de utilização na produção de proteína hidrolisada.



Designer: Lin Diniz

De acordo com Leal et al., (2009) a composição de hidrolisado proteico de camarão mostrou ser uma fonte adequada de triptofano, lisina, leucina, arginina, fenilalanina, valina e metionina.

De fato, a qualidade nutricional da proteína hidrolisada pode estar relacionada com a elevada concentração de pequenos peptídeos e aminoácidos essenciais, como também de compostos que estimulam a resposta imune nos peixes, promovendo assim crescimento e resistência a doença (GILDBERG e STENBERG, 2001). Segundo Silvestre et al. (1994), é preferível utilizar hidrolisado em dietas do que misturas de aminoácidos, uma vez que di e tripeptídeos são absorvidos pelo intestino mais rapidamente do que os aminoácidos livres. Estes peptídeos são também aceitos como atrativo alimentar devido à estimulação química que eles podem produzir em larvas e juvenis, contribuindo para o consumo de dietas artificiais nas fases iniciais de cultivo.

De acordo com Muhlia-Almazán et al. (2003) e Lundstedt et al. (2004), mudanças na origem e quantidade de nutrientes podem afetar as concentrações ou as expressões gênicas das enzimas digestivas. No entanto, Ezequerra et al. (1997); Moraes e Bidinoto (2000) sugerem que os animais aquáticos podem adaptar os níveis enzimáticos de acordo com mudanças na alimentação.

A inclusão de hidrolisado proteico na dieta de animais aquáticos tem sido amplamente estudada, principalmente avaliando seus efeitos sobre o crescimento, sobrevivência e atratividade (KOLKOVISK et al., 2000; PLASCENCIA-JATOMEA et al., 2002; SAVOIE et al., 2006). No entanto, existe pouco conhecimento sobre o efeito de hidrolisados proteicos sobre as enzimas digestivas de organismos aquáticos. Autores como Córdova-Murueta e García-Carreño (2002); Cahu et al. (2004); Zambonino-Infante e Cahu (2007) e Kotzamanis et al. (2007) realizaram estes estudos, contudo ainda são necessários mais pesquisas para fornecer a comunidade científica e aos produtores respostas mais conclusivas.

Kotzamanis et al. (2007) avaliaram o efeito da inclusão de hidrolisado proteico (dietas contendo 10 e 19% de proteína hidrolisada) sobre as atividades de enzimas digestivas em larvas de *Dicentrarchus labrax* (L.). Não foram encontradas diferenças estatísticas entre os tratamentos nas atividades de amilase e tripsina. No entanto, a atividade de aminopeptidase foi maior na concentração de 10% de hidrolisado proteico. Cahu et al. (2004) verificaram que a secreção de tripsina foi maior em larvas de *D. labrax* alimentadas com dietas contendo 14% de proteína hidrolisada, sendo reduzida a concentrações crescentes (46%).

O efeito da inclusão de proteína hidrolisada de camarão sobre as enzimas digestivas de tilápia do Nilo não estão descritos na literatura. Esta informação pode contribuir para um melhor

entendimento do modo de regulação destas enzimas e de possíveis adaptações da tilápia do Nilo na digestão de ingredientes alternativos.

3. OBJETIVOS

3.1 Objetivo geral:

Avaliar se mudanças na alimentação (inclusão de hidrolisado proteico de camarão) pode afetar as atividades enzimáticas do extrato bruto dos tecidos do trato digestório da tilápia do Nilo (*Oreochromis niloticus* L.) e avaliar a influência dos cultivos semi-intensivo e intensivo nos parâmetros zootécnicos e nas enzimas digestivas deste peixe.

3.2 Objetivos específicos:

1. Avaliar se a inclusão do hidrolisado proteico de camarão, em diferentes concentrações, pode promover mudanças nas principais atividades enzimáticas do extrato bruto dos tecidos do trato digestório da tilápia do Nilo;
2. Obter o perfil enzimático através de zimograma, de proteases dos extratos brutos do intestino dos espécimes experimentais, alimentados com diferentes concentrações de hidrolisado proteico de camarão;
3. Avaliar a diferença entre os sistemas intensivo e semi-intensivo através dos parâmetros de crescimento da tilápia do Nilo;
4. Obter o peso relativo dos peixes produzidos em sistema intensivo e semi-intensivo em comparação com tilápias provenientes de outros cultivos e de ambiente natural;
5. Determinar as atividades de enzimas digestivas nos extratos brutos dos tecidos do intestino de tilápia do Nilo nos diferentes sistemas de cultivo utilizando substratos específicos e inespecíficos;
6. Obter o perfil enzimático através de zimograma de protease e de amilase dos extratos brutos do intestino de tilápia do Nilo nos diferentes sistemas de cultivo;
7. Obter a eletroforese bidimensional dos extratos brutos do intestino de tilápia do Nilo em ambos os cultivos.

4. REFERÊNCIAS

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5. CAPÍTULO I

ARTIGO CIENTÍFICO:

Digestive enzyme activity in juvenile Nile tilapia (*Oreochromis niloticus*, L) submitted to different dietary levels of shrimp protein hydrolysate

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Digestive enzyme activity in juvenile Nile tilapia (*Oreochromis niloticus*, L) submitted to different dietary levels of shrimp protein hydrolysate

Juliana Ferreira Santos · Patrícia Fernandes Castro ·
Albino Luciani Gonçalves Leal ·
Augusto Cézar Vasconcelos de Freitas Júnior · Daniel Lemos ·
Luiz Bezerra Carvalho Jr. · Ranilson Souza Bezerra

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Abstract The effect of different dietary concentrations of shrimp protein hydrolysate (SPH) on digestive enzyme activity of Nile tilapia juveniles was evaluated. SPH concentrations in diets were 0, 15, 30 and 60 g kg⁻¹ (treatments SPH0, SPH15, SPH30 and SPH60, respectively). Hemoglobin, azocasein, BA_nNA (*N*_z-benzoyl-DL-arginine-*p*-nitroanilide), SA_nNA (Suc-Ala-Ala-Pro-Phe-*p*-nitroanilide), aminoacyl of β -naphthylamide and starch were used as substrates for enzyme activity determinations. The activity of total alkaline protease was significantly higher ($P < 0.05$) in fish under SPH15 and SPH60 treatments than in the control (SPH0). However, the effect was not dose-dependent. Substrate-SDS-PAGE was also performed to evaluate changes in the profile

J. F. Santos · A. C. V. de Freitas Júnior · L. B. Carvalho Jr. · R. S. Bezerra (✉)
Laboratório de Enzimologia (LABENZ), Departamento de Bioquímica,
Universidade Federal de Pernambuco, Cidade Universitária, Recife,
PE CEP 50670-420, Brazil
e-mail: ransoube@uol.com.br

J. F. Santos · A. C. V. de Freitas Júnior · L. B. Carvalho Jr. · R. S. Bezerra
Laboratório de Imunopatologia Keizo Asami (LIKA), Universidade Federal de Pernambuco,
Cidade Universitária, Recife, PE 50670-420, Brazil

J. F. Santos
Unidade Acadêmica de Serra Talhada, Universidade Federal Rural de Pernambuco,
Fazenda Saco, S/N, Caixa Postal 063, Serra Talhada, PE CEP 56900-000, Brazil

P. F. Castro
Embrapa Meio-Norte, Caixa Postal 341, Parnaíba, PI 64200-970, Brazil

A. L. G. Leal
Companhia Hidro Elétrica do São Francisco (CHESF), Rua Delmiro Gouveia,
333, Bongi, Recife, PE 50761-901, Brazil

D. Lemos
Laboratório de Aquicultura Marinha (LAM), Instituto Oceanográfico,
Universidade de São Paulo, Cidade Universitária, São Paulo, SP 05508-120, Brazil

of Nile tilapia digestive proteases caused by SPH. Substrate-SDS-PAGE revealed 12 active proteolytic bands, eight of which responded to SPH dietary incorporation. Inhibition substrate-SDS-PAGE indicated a decrease in the activity of three enzymes, with trypsin activity decreasing with the increase of SPH concentration, whereas the opposite occurred for two aminopeptidases. Distinct protease profiles were also found for each treatment, suggesting adaptability of digestive proteases from Nile tilapia to the different diets.

Keywords Digestive proteases and amylases · Nile tilapia · SDS-PAGE zymograms · Shrimp protein hydrolysate

Introduction

Tilapia production has increased significantly on a global scale in the last decade. This development has been followed by an increase in feed consumption, stimulating the search for new ingredients in diet formulations (Schulz et al. 2007). The growth of the aquaculture industry has generated large amounts of waste and by-products, which represent a challenge to the sustainability of the activity (Bezerra et al. 2001). Shrimp processing waste, for instance, may be disposed of in the environment, where it may potentially pollute water and land. By-products of terrestrial and aquatic farming systems may be processed into suitable ingredients and used as components in animal feeds, following biosecurity principles for feed applications. Shrimp processing waste has been identified as an animal protein source of great potential (Fanimo et al. 2000). A simple protocol for producing protein hydrolysate from white shrimp *Litopenaeus vannamei* (Boone) processing waste through autolysis has recently been developed (Cahú et al. 2012). This method renders a protein concentrate that is considered to be an excellent source of amino acids, with high levels of glutamate, aspartate, leucine, lysine, tyrosine and arginine (Leal et al. 2010). In fact, crustacean protein silage and hydrolysate have been used in fish feeds both as a new protein source (Plascencia-Jatomea et al. 2002) and, in small amounts, as flavoring to enhance the attractiveness of feeds (Kolkowski et al. 2000).

Variations in the quality and quantity of nutrients used in diet formulations may modify enzymatic profile and activity in the digestive tract of animals (Lundstedt et al. 2004). Thus, feed composition could induce biological adaptations, including an increase in nutrient absorption (Moraes and Bidinotto 2000). Digestive enzymes have been investigated as a way of understanding nutritional requirements, and the effects of diet composition on enzyme activity in order to reduce feeding costs in fish farms (Canuso et al. 1996). Most studies thus far have evaluated the effect of different concentrations of protein, carbohydrates and lipids in feed formulations, correlating these results with enzyme activity, and differences in enzyme quality profile may be related to nutrient levels in the diet (Fountoulaki et al. 2005).

The present survey focuses on the following aspects: (a) the detectable changes in the activity of the main digestive enzymes caused by different concentrations of shrimp protein hydrolysate in tilapia feeds and (b) the use of substrate-SDS-PAGE zymograms as an effective tool to improve the analysis of these changes.

Materials and methods

Materials

All reagents were of analytical grade and purchased from Sigma (St. Louis, MO, USA) and Merck (Darmstadt, Germany). The diets were prepared in Poytara LTDA (Aranquara—São Paulo—Brazil).

Shrimp protein hydrolysate production

Shrimp head processing waste, comprising the cephalothorax without the body carapace, from *L. vannamei* juveniles (total body weight about 10–12 g) was provided by a local fishery processing plant (EMPAF Ltd.). SPH used in experimental diets was prepared according to Cahú et al. (2012). In total, approximately 20 kg of shrimp cephalothoraxes was obtained. The shrimp heads were washed and stored at –20 °C in plastic bags containing 1 kg of the material. After that, the raw material was crushed in distilled water (1:1 ratio; particle size approximately 5 mm). The enzymatic autolysis (proteases from the shrimp midgut gland) occurred in a vessel placed in a water bath at 45 ± 2 °C for 180 min with constant stirring (700 rpm). The solution was then heated to 100 °C for 10 min in order to deactivate the enzymes, and the solid portion was strained through a 1.0-mm sieve (Cahú et al. 2012). The resulting material was centrifuged at 10,000 × g for 10 min, and the supernatants (SPH) was stored in plastic bottles at –20 °C until the preparation of the diets (Leal et al. 2010). The production method was similar to the industrial processes. Previously frozen-dried samples of SPH ($n = 3$) were used for proximate composition and AA analyses (Tables 1, 2) in Protein Chemical Center of the Faculdade de Medicina de Ribeirão Preto, São Paulo, Brazil (AOAC 1984).

Hydrolysis profile of SPH

Enzymatic hydrolysis of SPH (45 °C) was followed by sampling at the incubation times of 0, 30, 60, 90, 120, 150 and 180 min. The hydrolysis profile of shrimp protein hydrolysate (Aliquots of 100 µg of protein, $n = 3$) was evaluated in sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE), using stacking gel at 4 % (w/v) and separation gel at 17 % (Laemmli 1970). The gels were stained for protein overnight in 0.01 % (w/v) Coomassie Brilliant Blue. The background of the gel was destained by washing in a solution containing 10 % (v/v) acetic acid and 25 % (v/v) methanol. The molecular mass of the protein bands was estimated using the 220–10 kDa molecular mass protein standards (Sigma).

Table 1 Proximate composition (%) and energy of lyophilized shrimp protein hydrolysate

	Composition	Mean
Moisture (%)	9.7	
Crude protein (%)	43.6	
Ether extract (%)	6.2	
Ash (%)	7.3	
Carbohydrate (%)	33.1	
Energy (Kcal 100 g ⁻¹)	363.3	

SPH, from *Litopenaeus vannamei* (Leal et al. 2010)

Table 2 Amino acid composition of the shrimp protein hydrolysate

Amino acids	SPH	
	mg 100 g ⁻¹	%
<i>Essential</i>		
Arginine	3400 ± 0.043	7.3
Histidine	1060 ± 0.005	2.3
Isoleucine	2000 ± 0.021	4.3
Leucine	3490 ± 0.021	7.4
Lysine	3350 ± 0.000	7.2
Methionine	1290 ± 0.005	2.7
Phenylalanine	2370 ± 0.002	5.1
Threonine	2120 ± 0.031	4.5
TRYptophan	670 ± 0.016	1.4
Valine	2250 ± 0.012	4.8
<i>Non-essential</i>		
Tyrosine	3370 ± 0.004	7.2
Aspartic acid	4270 ± 0.031	9.1
Glutamic acid	5780 ± 0.003	12.4
Glycine	2890 ± 0.005	6.2
Serine	2030 ± 0.001	4.3
Alanine	3070 ± 0.017	6.6
Proline	2970 ± 0.024	6.3
Cystine	410 ± 0.015	0.9
SPH from <i>Litopenaeus vannamei</i> (Leal et al. 2010)	Total	100

Diets preparation

According to Leal et al. (2010), four isonitrogenous (37 % crude protein), isocaloric (total energy: 440 kcal 100 g⁻¹) experimental diets were formulated to feed *Oreochromis niloticus* juveniles (Table 3). Shrimp protein hydrolysate (SPH) was included in the diets at concentrations of 0 (control—SPH0), 15 (SPH15), 30 (SPH30) and 60 g kg⁻¹ (SPH60), as a fed basis. A 1:2 animal/plant protein ratio (fishmeal and soybean meal, respectively) in the diets was established. SPH (liquid) was mixed with soybean meal, and the dough was dried at 65 °C for 24 h. The ingredients were mixed and the diets prepared by extrusion under industrial conditions to obtain 1-mm diameter pellets.

Animals and culture conditions

Juvenile sex-reversed Nile tilapias were obtained from the Aquaculture Station of the Universidade Federal Rural de Pernambuco. The fish were stocked in fifteen 40-L glass aquaria (8 ind. per aquarium) equipped with a biological filter and continuous aeration and then submitted to a 7-day acclimatization period both for diets and experimental conditions in a completely randomized design, with four treatments and three replicates. The animals were kept in a photoperiod of 12:12 L:D. Prior to the feeding trial, fish were weighed (1.7 ± 0.4 g) and measured (4.7 ± 0.4 cm). Individuals were fed a ratio of 15, 14, 12, 10, 8 and 6 % of biomass per day, adjusted every 9 days for a period of 45 days. Feed was

Table 3 Composition and approximate analysis of the experimental diets

Ingredients (%)	Diets			
	SPH 0	SPH 15	SPH 30	SPH 60
Fish meal (57 % CP)	23.0	22.0	21.0	18.0
Shrimp protein hydrolysate (SPH)	0.0	1.5	3.0	6.0
Soybean meal (40.4 % CP)	47.0	47.5	47.5	47.5
Wheat meal	16.0	13.5	13.5	15.5
Corn starch	10.5	12.0	11.5	9.5
Soybean oil	1.0	1.0	1.0	1.0
Dicalcium phosphate	1.0	1.0	1.0	1.0
Mineral and vitamin mix ^a	1.0	1.0	1.0	1.0
Salt	0.5	0.5	0.5	0.5
Antioxidant BHT	0.02	0.02	0.02	0.02
Proximate analysis (as-fed basis, g kg ⁻¹)				
Dry matter	944.8	935.9	936.5	946.7
Crude protein	371.9	374.3	376.2	380.6
Ether extract	48.1	56.2	52.1	55.9
Crude fiber	39.7	38.8	41.1	46.6
Ash	105.7	102.9	101.6	101.9
Nitrogen-free extract	434.6	427.8	429.0	435.0
Calcium	22.2	21.7	20.0	17.2
Phosphorus	12.4	12.5	12.6	12.8
Gross energy (kcal 100 g ⁻¹) ^b	438.3	444.5	442.2	431.8
P/GE ratio (mg kcal ⁻¹)	84.9	84.1	84.6	88.0

Leal et al. 2010

CP crude protein, BHT butylated hydroxytoluene

^a Mineral and vitamin mix (quantity kg-1 premix): vitamin A (20,000 UI), vitamin D₃ (5,000 UI), vitamin E (250 mg), vitamin K₃ (25 mg), vitamin B₁ (37.5 mg), vitamin B₂ (37.5 mg), vitamin B₆ (25 mg), vitamin B₁₂ (0.053 mg), vitamin C (250 mg), niacin (200 mg), pantothenic acid (100 mg), biotin (1.25 mg), choline (1000 mg), inositol (250 mg), Fe (100 mg), Cu (12 mg), Zn (125 mg), Mn (37.5 mg), Se (0.25 mg), I (1.25 mg), Co (0.25 mg)

^b Estimate based on 5.65, 4.2 and 9.5 kcal g⁻¹ for proteins, carbohydrates and lipids, respectively

offered 4 times per day. The fish accepted and fed on all diets over the experimental period (Leal et al. 2010).

The aquaria were siphoned twice a day to remove solids, with 66 % water being replaced with clean water every day. Temperature (28.7 ± 0.59 °C), dissolved oxygen (3.5 ± 0.92 mg L⁻¹) and pH (8.1 ± 0.19) were measured twice a day ($n = 90$). Ammonia (0.14 ± 0.22 mg L⁻¹) and nitrite (0.08 ± 0.02 mg L⁻¹) were monitored once a week ($n = 6$). Values are expressed as mean \pm SE

Growth and nutrient utilization

The effects of SPH on the growth performance and nutrient utilization are shown in Table 4. Growth performance was evaluated through weight gain rate (WG), average daily gain (ADG), specific growth rate (SGR), feed conversion ratio (FCR), protein efficiency

Table 4 Growth performance and nutrient utilization in Nile tilapia fed on diets containing shrimp protein hydrolysate instead of fish meal replacement

Diet	SPH 0	SPH 15	SPH 30	SPH 60
Initial weight (g)	1.68 ± 0.14	1.72 ± 0.07	1.75 ± 0.09	1.81 ± 0.06
Final weight (g)	27.18 ± 2.43 ^{ns}	29.46 ± 1.05 ^{ns}	26.02 ± 3.07 ^{ns}	25.19 ± 2.49 ^{ns}
Survival (%)	100.0 ^{ns}	100.0 ^{ns}	100.0 ^{ns}	100.0 ^{ns}
WG (g) ^a	25.51 ± 2.57 ^{ns}	27.73 ± 1.11 ^{ns}	24.29 ± 3.04 ^{ns}	23.39 ± 2.49 ^{ns}
ADG (g day ⁻¹) ^b	0.57 ± 0.10 ^{ns}	0.62 ± 0.04 ^{ns}	0.54 ± 0.12 ^{ns}	0.52 ± 0.10 ^{ns}
SGR (% day ⁻¹) ^c	7.15 ± 0.58 ^{ns}	7.38 ± 0.10 ^{ns}	6.85 ± 0.68 ^{ns}	6.73 ± 0.57 ^{ns}
FCR ^d	1.15 ± 0.22 ^{ns}	1.09 ± 0.16 ^{ns}	1.13 ± 0.10 ^{ns}	1.17 ± 0.12 ^{ns}
PER ^e	2.26 ± 0.39 ^{ns}	2.33 ± 0.36 ^{ns}	2.20 ± 0.18 ^{ns}	2.14 ± 0.22 ^{ns}
ANPU ^f	39.31 ± 3.90 ^{ns}	40.40 ± 3.64 ^{ns}	38.59 ± 1.84 ^{ns}	34.72 ± 2.08 ^{ns}

Leal et al. 2010

Different superscripts in the same column denote statistical differences ($P < 0.05$), and "ns" denotes no statistical differences. Values are mean ± SE of five replicates

^a Weight gain rate, ^b Average daily gain, ^c Specific growth rate, ^d Feed conversion ratio, ^e Protein efficiency ratio, ^f Apparent net protein utilization

ratio (PER) and apparent net protein utilization (ANPU) based on the following formulae: $WG(g) = BW_f - BW_i$; $ADG = WG(g)/time\ (days)$; $SGR = 100 \times (\ln BW_f - \ln BW_i)/time\ (days)$; $FCR = \text{dry feed offered (g)}/\text{wet weight gain (g)}$; $PER = \text{wet weight gain (g)}/\text{protein fed (g)}$; and $ANPU = 100 \times [(BW_f \times BCP_f) - (BW_i \times BCP_i)]/(TF \times CP)$, in which BW_i and BW_f = average initial and final body weight (g) of fish, respectively; BCP_i and BCP_f = initial and final body crude protein (g 100 g⁻¹), respectively; TF = total amount of diet fed (g) and CP = crude protein of diet (g 100 g⁻¹) (Leal et al. 2010).

Extraction of digestive enzymes

At the end of trial, fish were left to fast for 24 h, and six individuals from each replicate aquarium (divided into two replicates per aquarium) were removed and killed in an ice bath for biometric measurements and tissue removal, according to standard methodology (Bezerra et al. 2001). Stomach and intestines were immediately collected and homogenized (40 mg tissue mL⁻¹) in 0.01 M Glycine-HCl pH 3.0 and 0.01 M Tris-HCl pH 8.0 buffers, respectively, containing 0.15 M NaCl, using a tissue homogenizer. The resulting preparations were centrifuged at 10,000 × g for 10 min at 4°C to remove cell debris and nuclei. The supernatants (crude enzyme extracts) were frozen at -20 °C and used in further assays (Bezerra et al. 2005). Protein concentration was determined according to Bradford (1976) using bovine serum albumin (BSA) as the standard and reported as mg of protein equivalent to BSA.

Total acid proteolytic activity

Acid protease activity was evaluated using hemoglobin (Hb) as the substrate as follows: in microcentrifuge tubes, 100 µL of 2 % Hb in 0.06 M Glycine-HCl buffer pH 3.0 was mixed with 50 µL stomach crude extract and 350 µL 0.5 M Glycine-HCl buffer pH 3.0 for 60 min at 25 °C. Five hundred microliters of 10 % trichloroacetic acid (TCA) was then added to stop the reaction. After 15 min, centrifugation was carried out at 8,000 × g for

10 min. The absorbance of supernatant (70 µL) was measured at 280 nm (Bio-Rad SmartSpec 3000, USA) versus a similarly prepared blank in which 0.01 M Glycine-HCl buffer pH 3.0 replaced the crude extract sample, based on a methodology adapted from Díaz-López et al. (1998). Previous experiments showed that, for the first 60 min, the reaction carried out under the conditions described above followed first-order kinetics. One unit (U) of enzyme activity was defined as the amount of enzyme capable of hydrolyzing hemoglobin to produce a 0.001 change in absorbance per minute per milligram of protein.

Total alkaline proteolytic activity

In microcentrifuge tubes, 1 % azocasein prepared in 0.1 M Tris-HCl buffer pH 8.0 was incubated with intestine crude enzyme extract (30 µL) for 60 min at 25 °C. Two hundred and forty µL of 10 % trichloroacetic acid (TCA) was then added to stop the reaction. After 15 min, centrifugation was carried out at 8,000×g for 5 min. The supernatant (70 µL) was added to 1 M NaOH (130 µL) in a 96-well microtiter plate, and the absorbance of this mixture was measured in a microtiter plate reader (Bio-Rad 680, Japan) at 450 nm versus a similar prepared blank in which 0.01 M Tris-HCl pH 8.0 replaced the crude extract sample. One unit (U) of enzyme activity was defined as the amount of enzyme capable of hydrolyzing azocasein to produce a 0.001 change in absorbance per minute per milligram of protein (Bezerra et al. 2005).

Trypsin and chymotrypsin activity

The activity of trypsin and chymotrypsin was determined using 8.0 mM BApNA (*N*-benzoyl-D,L-arginine-*p*-nitroanilide) and 8.0 mM SApNA (Suc-Ala-Ala-Pro-Phe *p*-nitroanilide) in DMSO (dimethyl sulfoxide), respectively. Intestine crude enzyme extract (30 µL) was incubated with 0.1 M Tris-HCl buffer pH 8.0 (140 µL) and respective substrates (30 µL) in a microtiter plate reader (Bio-Rad 680, Japan). Absorbance was measured at 405 nm versus a similar prepared blank in which 0.1 M Tris-HCl pH 8.0 replaced the crude extract sample. Enzyme activity was determined in triplicate. Trypsin and chymotrypsin units of activity were expressed as a change in absorbance per minute per milligram of protein (Bezerra et al. 2005).

Aminopeptidase activity

Aminopeptidase activity was evaluated using aminoacyl of β -naphthylamide (AA of arginine) as substrate. The procedure adapted from Oliveira et al. (1999) was carried out in triplicate, by incubating 4.2 mM substrate (50 µL), 50.0 mM sodium phosphate buffer pH 7.0 (600 µL) and deionised H₂O (50 µL) at 37 °C. After temperature equilibration, intestine crude extract (50 µL) was added, and 30 min later, the reaction was stopped by adding 1 mg mL⁻¹ fresh Garnet reagent (250 µL) in 0.2 M sodium acetate buffer pH 4.2 containing 10 % Tween 20 (v/v). After 10 min, absorbance was measured at 525 nm (Bio-Rad SmartSpec 3000, USA), and the amount of β -naphthylamine was determined using a standard β -naphthylamine curve. Activity was expressed as protease mU mg⁻¹ of protein. One unit of enzyme activity was defined as the amount of enzyme required to hydrolyze one µmol of *p*-nitroaniline per minute per milligram of protein.

Amylase activity

Amylase activity was evaluated according to Bernfeld (1955) using 2 % starch as unspecific substrate: 60 μ L intestine crude extract were incubated with 375 μ L starch solution and 375 μ L 10 mM phosphate buffer pH 8.0 containing 15 mM NaCl at 25 °C. After 20 min, 3,5-dinitro salicylic acid (DNSA) was added and the solution was heated to 100 °C for 10 min. After temperature equilibration, absorbance was measured at 570 nm (Bio-Rad SmartSpec 3000, USA) versus a similarly prepared blank in which 10 mM phosphate buffer replaced the crude extract sample. Enzyme activity was determined in triplicate. One unit of enzyme activity was defined as the amount of enzyme required to liberate 1 mg of maltose per milligram of protein per min.

Enzyme characterization in substrate-SDS-PAGE

Proteases from intestine crude extract of *O. niloticus* (Aliquots of 50 μ g of protein, $n = 3$) were studied in substrate sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) using stacking gel at 4 % (w/v) and separation gel at 12.5 % (Laemmli 1970).

Zymograms of protease activity were carried out based on Garcia-Carreño et al. (1993). After electrophoresis, the gels were immersed in 2.5 % Triton X-100 dissolved in 0.1 M Tris-HCl buffer pH 8.0 to remove the SDS and incubated with 3 % casein (w/v) in 0.1 M Tris-HCl buffer pH 8.0 for 30 min at 4 °C. The temperature was raised to 37 °C and maintained at this value for 90 min to allow the digestion of casein by the active fractions. Finally, the gel was stained overnight for protein in 0.18 % (w/v) Coomassie Brilliant Blue R250 prepared in acetic acid and methanol (10:25 % v/v), and the background of the gel was destained in acetic acid and methanol (10:25 % v/v). Clear bands in blue background denoted protease bands by digestion of casein substrate.

Inhibition assays using substrate-SDS-PAGE

The following inhibitors prepared in DMSO at a final concentration of 2 mM were used: Tosyl phenylalanine chloromethyl ketone (TPCK—chymotrypsin inhibitor); Phenyl-methyl-sulfonil-fluoride (PMSF—serine proteases inhibitor); benzamidine and tosyl-lysine chloromethyl ketone (TLCK), both trypsin inhibitors; and bestatin (leucine aminopeptidase inhibitor). For the inhibition study, only the crude extracts from animals under SPH0 treatment ($n = 3$) were used, since all caseinolytic bands were found in this treatment. Samples of enzyme extract (100 μ L) and inhibitors (100 μ L) were incubated at 25 °C for 30 min. An aliquot of 50 μ g of protein was collected and applied on each respective lane. After that, the zymogram was performed as described above. The 100 % values (control) were established using DMSO without inhibitors (Lemos et al. 2002, 2004).

Statistical analysis

All data were tested for normal distribution (Shapiro-Wilk test) and homogeneity of variances (Bartlett's test). One-way analysis of variance (ANOVA) by Tukey test was used for normally distributed data, whereas Kruskal-Wallis test was used for non-normally distributed data. Differences were reported as statistically significant when $P < 0.05$, using the SysEapro software (beta version).

Results

Figure 1 shows the hydrolysis profile of SPH. Enzymatic autolysis (proteases from the shrimp midgut gland) visibly promoted the digestion of proteins with the greatest molecular mass over time. At time 0, most proteins had molecular mass between 15 and 70 kDa, at time 150 min, most proteins were smaller than 15 kDa.

Table 5 shows the enzyme activities in the crude extracts. Statistical differences ($P < 0.05$) between animals fed the experimental diets (SPH0, SPH15, SPH30 and SPH60) were only observed for total alkaline proteases.

Substrate-SDS-PAGE zymograms revealed caseinolytic activity in crude extracts of fish from all treatments (Fig. 2). Twelve caseinolytic bands were found in animals fed on SPH0 and were named P1 to P12, showing the highest number of active bands among treatments. Proteases P1 to P4 were not significantly affected by the addition of the protein hydrolysate, but P5 to P12 underwent alterations in the caseinolytic pattern. As the concentration of hydrolysate increased, there was a decrease in the activity of proteases P6, P7 and P8. The opposite occurred for P5, P9, P10, P11 and P12, in which the caseinolytic bands proved more intense.

The effect of specific inhibitors on the caseinolytic bands from fish crude enzyme extract is shown in Fig. 3. As observed in Fig. 2, the zymogram of the crude extract (SPH0) without inhibitors (control) revealed 12 caseinolytic bands. TPCK did not significantly affect the caseinolytic activity of *O. niloticus* enzymes, and P11 was the only

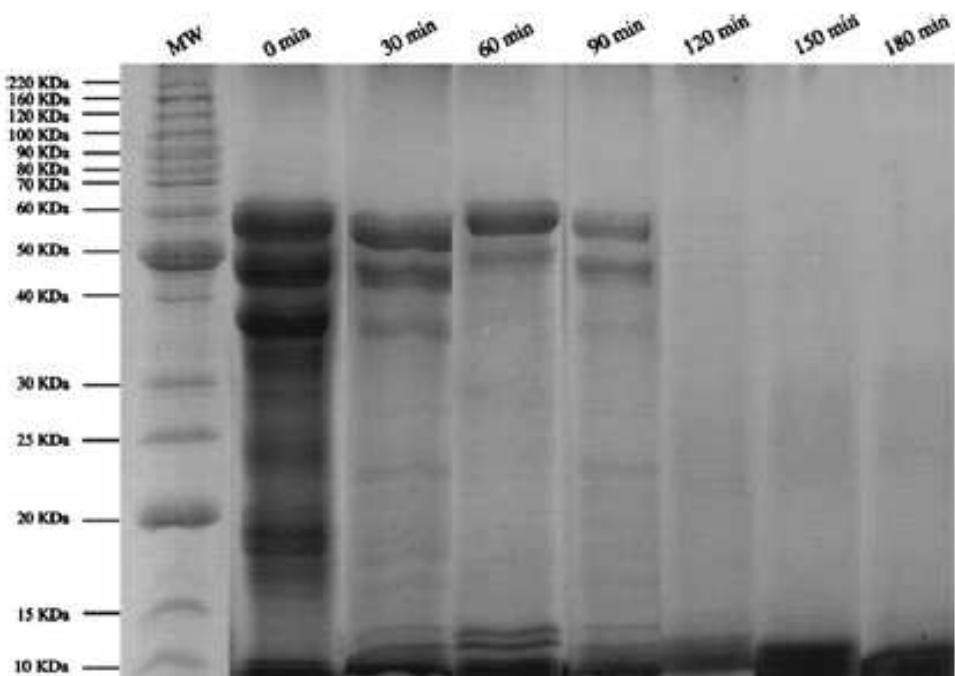


Fig. 1 Hydrolysis profile of shrimp protein hydrolysate (using autolysis) in sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) using stacking gel at 4 % (w/v) and separation gel at 17 %. Lanes correspond to molecular mass marker (MM) and incubation times of 0, 30, 60, 90, 120, 150 and 180 min of enzymatic hydrolysis. SPH was produced through autolysis of *Litopenaeus vannamei* cephalothorax (heads)

Table 5 Digestive enzyme activity of crude extracts recovered from stomach (acid) and intestine of juvenile Nile tilapia *O. niloticus* fed on experimental diets containing increasing shrimp protein hydrolysate levels (0–60 g kg⁻¹)

Diets	Acid proteolytic activity	Alkaline proteolytic activity	Trypsin	Chymotrypsin	Aminopeptidase	Amylase
SPH 0	155.84 ± 14.22 ^a	11.80 ± 0.55 ^b	2.58 ± 0.09 ^{ns}	22.70 ± 1.26 ^{ns}	41.68 ± 3.32 ^{ns}	45.47 ± 3.92 ^{ns}
SPH 15	111.38 ± 22.83 ^{ns}	15.35 ± 0.76 ^a	3.38 ± 0.17 ^{ns}	39.74 ± 3.94 ^{ns}	45.34 ± 1.63 ^{ns}	53.50 ± 4.65 ^{ns}
SPH 30	162.49 ± 16.69 ^{ns}	13.60 ± 0.73 ^{ns}	2.82 ± 0.20 ^{ns}	30.89 ± 2.42 ^{ns}	47.66 ± 2.08 ^{ns}	48.79 ± 4.82 ^{ns}
SPH 60	132.19 ± 13.14 ^{ns}	15.23 ± 0.88 ^a	2.62 ± 0.15 ^{ns}	36.66 ± 3.60 ^{ns}	39.85 ± 2.61 ^{ns}	56.77 ± 4.31 ^{ns}

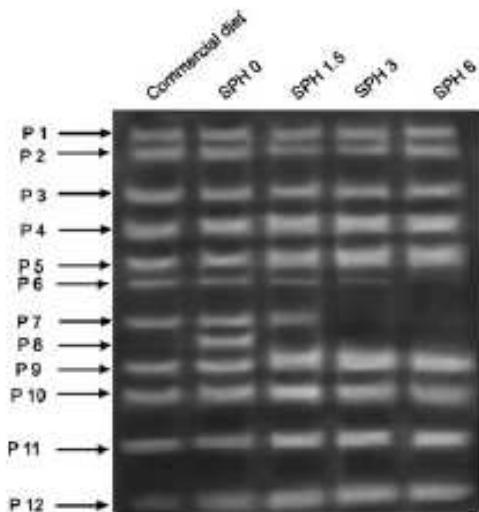
Acid and alkaline proteolytic activity expressed as U mg of Protein (P)⁻¹; trypsin, chymotrypsin and aminopeptidase activity expressed as mU mg⁻¹ of P; and amylase activity expressed as mg of maltose min⁻¹ mg of P⁻¹. Different superscripts in the same column denote statistical differences ($P < 0.05$), and "ns" denotes no statistical differences. Values are mean ± SE of five replicates

enzyme inhibited by TPCK. PMSF revealed high inhibition of P9 and P11 caseinolytic bands. Benzamidine completely inhibited P6 and P8 bands. One band between P5 and P6 enzymes (white arrow) revealed activity in the presence of benzamidine, but it did not appear in the control. TLCK strongly inhibited P3, P4, P6, P7, P8 and P11. Bestatin totally inhibited seven digestive enzyme bands of *O. niloticus* (P3, P4, P5, P6, P7, P8 and P9). Inhibitors had no effect on P1, P2, P10 and P12 caseinolytic bands.

Discussion

As can be observed in Fig. 1, autolysis efficiently solubilized and hydrolyzed protein from shrimp processing waste (cephalothorax section) and produced protein hydrolysate, which was shown to be a promising protein feedstuff for tilapia.

Fig. 2 Zymogram of digestive proteases (3 % casein as substrate) of intestine enzyme extracts from juvenile Nile tilapia *O. niloticus* fed on diets containing different concentrations of shrimp protein hydrolysate (SPH). Lanes correspond to dietary SPH inclusion (%): SPH 0, SPH 1.5, SPH 3 and SPH 6. Further details in "Materials and methods"

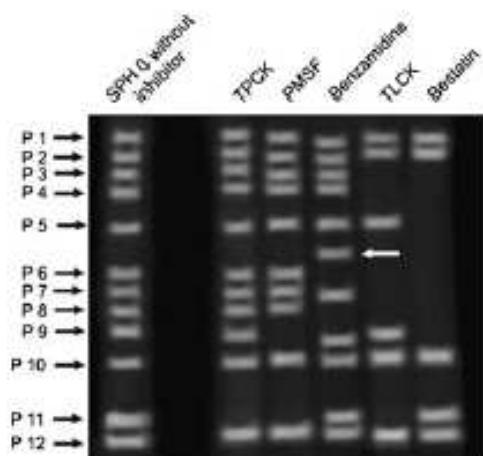


Hydrolysis using exogenous substances allows better control of the properties of the resulting products. However, the cost of the substances (enzymes, acids and solvents) is an obstacle to the viability of the process. Autolysis is an enzymatic process that can be used as an alternative to processes that employ exogenous substances to recover bioactive molecules, such as chitin (for chitosan production) and carotenoids (Armenta and Guerero-Legarreta 2009; Chen et al. 2011; Cahú et al. 2012). Moreover, autolysis is a common method used to treat fish and shrimp waste, since it does not use any exogenous chemical or biological additive, for example silage. The degree of hydrolysis affects certain product properties, like viscosity, solubility and the partition of proteins. These, in turn, influence the absorption capacity and rate of passage of the diet through the gastrointestinal tract. The solubility of the hydrolysates depends on the nature of the raw material, method, temperature and hydrolysis time (Kotzamanis et al. 2007). Plascencia-Jatomea et al. (2002) reported that the acidic conditions in which fermentative shrimp silage hydrolysate is produced cause the loss of labile nutrients, such as tryptophan. Although the two products have a very similar essential amino acid composition, the SPH produced here proved to be an adequate source of tryptophan, as well as lysine, leucine, arginine, phenylalanine and valine (Leal et al. 2010).

The use of hydrolyzed protein in diets for aquatic animals involves some particularities that should be taken into account. For example, diets with high content of hydrolyzed protein can lead to a loss of nutrients by lixiviation, because it contains high levels of small peptides that are usually more soluble (Leal et al. 2010). The extent of hydrolysis can result in protein hydrolysates with different contents of amino acids, di- and tripeptides, oligopeptides, polypeptides and proteins, which will ultimately result in differences in the quality of the diet.

The nutritional value of a diet is not based simply on its chemical composition, but also on the ability physiological fish to digest and absorb, according to these animals' eating habits. During digestion, these proteins are broken down into simpler compounds (peptides and free amino acids) and then are absorbed by specific membrane proteins specialized in peptide transport (e.g., PepT1) and utilized by the body (Sangaletti et al. 2009; Terova et al. 2009; Veri et al. 2011). This degradation in the digestive tract is performed with the assistance of enzymes. A number of authors have analyzed digestive enzyme activity in aquatic organisms and dietary composition, reporting divergent results. Nagase (1964) and Kohla et al. (1992) found enhanced trypsin activity, corresponding to an increase in

Fig. 3 Inhibition zymogram of digestive proteases (3 % casein as substrate) of intestine enzyme extracts from juvenile Nile tilapia *O. niloticus* fed the SPH 0 diet. Lanes correspond to treatments: control without inhibitors, TPCK, PMSF, Benzamidine, TLCK and Bestatin. Further details in "Materials and methods" and Fig. 2



feeding rates for the species Mozambique tilapia, *Oreochromis mossambicus* (Peters) and *Colossoma macropomum* (Cuvier), respectively. Papoutsoglou and Lyndon (2006) found an increase in chymotrypsin activity in Spotted wolffish, *Anarhichas minor* (Olafsen), when the protein concentration was reduced, indicating the adaptation of this species to low protein concentrations as a way to better absorb nutrients from the diet. Studies about the effect of protein hydrolysates on the digestive enzymes from aquatic organisms are available in the literature (Córdova-Murueta and García-Carreño 2002; Cahu et al. 2004; Zambonino Infante and Cahu 2007). However, there is little information regarding the specific effects of dietary supplementation with shrimp protein hydrolysate on the digestive enzymes in Nile tilapia.

In the present paper, no statistical differences in proteases activities were observed among the SPH 0 (control) and treatments with higher concentrations of SPH (SPH15, SPH30 and SPH60) (Table 5), or in growth parameters (final weight, survival, weight gain, average daily gain, specific growth rate, feed conversion ratio, protein efficiency ratio, or apparent net protein utilization) (Table 4) (Leal et al. 2010). Only the activity of total alkaline protease was significantly higher ($P < 0.05$) in the fish under SPH15 and SPH 60 treatments, when compared to SPH0 (control) (Table 5). No correlation was observed between different concentrations of hydrolysate in the diets and the activity of any of the enzymes studied.

European seabass, *Dicentrarchus labrax* (L) were fed diets containing enzymatic hydrolyzed replacement for fish meal (native protein) in the proportions of 0, 20 and 40 %. Zambonino Infante et al. (1997) demonstrated that fish to the trypsin activity was enhanced by the native protein, whereas chymotrypsin activity was enhanced by the diets containing di- and tripeptides. On the other hand, Kotzamanis et al. (2007) also evaluated the inclusion effect of protein hydrolysate on the digestive enzyme activities in European seabass, *D. labrax* larvae fed on diets with 10 and 19 % protein hydrolysate. They found no statistical differences in amylase and trypsin activity between treatments. However, aminopeptidase activity was higher when 10 % protein hydrolysate was administered. Thus, the absence of statistical difference found for most enzymes, and growth parameters can be related to the loss of SPH by lysis. In addition, commercial substrates used in this study showed no sensitivity to detect differences in most of the enzymatic activities; in other words, it was observed that the effects of SPH on the digestive enzymes were comparably small and not dose-dependent.

The different results observed in the literature and those obtained here led us to consider other methods. The zymograms afforded to observe the dose-dependent effect of SPH on some proteases (caseinolytic bands). In fact, zymograms appear to be a more sensitive biochemical tool in comparison with other methods for detecting the proteinase composition of crude extracts from tissues and to allow the determination of enzyme activity zones (García-Carreño et al. 1993).

Through analyses of inhibition zymograms, it was possible to determine the following: one caseinolytic band with aminopeptidase activity (inhibited only by Bestatin, P5) and probably another aminopeptidase (P9) inhibited by Bestatin and by PMSF; some other active bands, P3, P4, P6, P7, P8 and P11, seemed to be proteases of low specificity, with trypsin/aminopeptidase activity (P3, P4, P6, P7 and P8 which were inhibited by trypsin inhibitors and by Bestatin) and chymotrypsin/trypsin activity (P11 which was inhibited by PMSF, TPCK and TLCK). It was not possible to identify P1, P2, P10 and P12, as none of the inhibitors had any effect on these proteases. Comparing the zymogram of enzyme activity according to dietary treatment to the inhibition gel, at increased concentration of shrimp protein hydrolysate, there was an overall slowdown of proteases with trypsin

activity (P6, P7 and P8). On the other hand, it was identified that the caseinolytic bands P5 and P9 (identified as aminopeptidase activity, Fig. 3) grew in intensity with increasing SPH contents. The caseinolytic bands P10 and P12 (not identified) and P11 (with chymotrypsin-like/trypsin-like activity) also exhibited an increase in intensity with the inclusion of SPH in the diet. Cahu et al. (2004) found that trypsin secretion was higher in larvae of the *D. labrax* fed on diets with 14 % protein hydrolysate, being reduced with increasing concentrations (46 %).

Zymograms in Figs. 2 and 3 show that the classical protease inhibitors herein employed were not effective upon some caseinolytic bands (P1, P2, P10 and P12), and in some cases, the inhibitory effect was not conclusive (P3, P4, P6, P7, P8, P9 and P11), suggesting a low compatibility of these enzymes with mammalian proteases. In fact, these commercial inhibitors are generally synthesized based on the mammalian enzyme mechanism, and as the fish digestive system may be considerably different from that of mammals, differences in the mechanisms of their enzymes may also occur. In spite of the results of the present work, tilapia digestive enzymes were not considerably affected by classical trypsin inhibitors. Bezerra et al. (2005) showed that purified Nile tilapia trypsin could be strongly inhibited by TLCK and Benzamidine. PMSF was shown able to inhibit about 50 % of trypsin activity.

Although there were differences in digestive enzyme activity (total alkaline proteases) between fish fed different experimental diets, there was no clear relationship between enzyme activity and different concentrations of shrimp protein hydrolysate in these diets. Substrate-SDS-PAGE zymogram proved to be efficient in detecting changes in enzyme activity in fish submitted to different diets. Through this method, different protease profiles could be detected according to experimental diet. These data underscore the suggested ability of tilapia to adapt to different food sources. Actually, as an omnivorous opportunistic fish, tilapia is able to assimilate nutrients from different sources, like benthic algae, phytoplankton, macrophytes, zooplankton, small invertebrates, detritus and cyanobacteria. The feed choice is mainly related to the availability of food items in the environment (Bowen 1982; Fitzsimmons 1997; Stickney 1997; Beveridge and Baird 2000; Lowe-McConnell 2000). To benefit from all these sources, the enzyme arsenal of tilapia should be diverse enough to digest the wide range of ingested food.

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6. CAPÍTULO II

ARTIGO CIENTÍFICO:

Use of condition factor and its derivatives to evaluate the effect of different culture systems on the growth of Nile tilapia (*Oreochromis niloticus* L.)

ARTIGO A SER SUBMETIDO A REVISTA AQUACULTURE ISSN: 0044-8486



Use of condition factor and its derivatives to evaluate the effect of different culture systems on the growth of Nile tilapia (*Oreochromis niloticus* L.)

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4 Juliana Ferreira Santos^{1,2}, Karollina Lopes Siqueira Soares¹, Marcelo Osmar Freire da Silva Sá³,
5 Caio Rodrigo Dias Assis¹, Daniel Lemos⁵, Ranilson Souza Bezerra^{1*}

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¹ Laboratório de Enzimologia (LABENZ), Departamento de Bioquímica and Laboratório de Imunopatologia Keizo Asami (LIKA), Universidade Federal de Pernambuco, Cidade Universitária, 50670-901, Recife-PE, Brazil

² Universidade Federal Rural de Pernambuco, Unidade Acadêmica de Serra Talhada, Fazenda Saco,
S/N - Caixa Postal 063 CEP 56900-000 Serra Talhada - PE, Brazil.

¹² ^{3.} Prefeitura de Floresta, Praça Coronel Fausto Ferraz, 183, Centro, 56400-000, Floresta-PE, Brazil.

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16 Running title: Influence of culture systems on the growth parameters of tilapia

17 *Ranilson S. Bezerra.

18 Laboratório de Enzimologia – LABENZ, Departamento de Bioquímica, Universidade Federal de
19 Pernambuco. CEP 50670-420, Cidade Universitária, Recife-PE, Brazil,
20 Tel.: + 55-81-21268540; Fax: + 55-81-21268576
21 E-mail address: ransoube@uol.com.br

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23 Highlights

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- 25 • Even though no statistical differences in growth parameters have not been observed between
26 fish reared in semi-intensive and intensive systems, protein and feed efficiencies and
27 condition factor were better in semi-intensive system in 31 days of culture.
- 28 • In this work, the relative weight of fish grown in semi-intensive and intensive systems was
29 higher in 9 out of 14 comparisons with natural conditions and other cultures.
- 30 • The equation of condition factor and its derivatives (Kmean and Krel) proved to be useful
31 tools when assessing growth.

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48 **Abstract**

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50 Nile tilapia is a fish species that can be produced in different culture systems and scales, including
51 the semi-intensive systems in ponds and the intensive system in cages. Their handlings are
52 completely different, so it is required management to stabilize the growth parameters and
53 production costs. Therefore, the present work intends to evaluate the influence of semi-intensive
54 system and intensive system on growth parameters of Nile tilapia and compare the relative weight
55 with tilapia in natural environment and farming by the method of Froese. Juvenile Nile tilapia
56 ($87.61 \pm 1.52g$) were cultured in semi-intensive and intensive system during 90 days. Two
57 nutritional plans were adopted: triphasic for semi-intensive systems (diet with 35, 32 and 28% crude
58 protein - CP) and biphasic for intensive system (35 and 32% CP). Biometrics measurements were
59 performed monthly for analysis of growth parameters and adjustments in diet. The water quality
60 indicators were monitored continuously and were in accordance with the requirements for animal
61 welfare. The presence of natural food was evaluated in the two systems, being more abundant in
62 semi-intensive systems. Survival, protein efficiency, feed efficiency and condition factor with 31
63 days of cultivation were better in semi-intensive system. The weight-length relationship of tilapia
64 was not altered by the systems. Comparing the fish relative weight from semi-intensive systems and
65 intensive systems with tilapia from other environments, they were higher in 9 of 14 comparisons.
66 The calculations of Fulton's condition factor (K), K_{mean} , K_{rel} and relative weight (W_{rm}) demonstrated
67 by Froese, proved to be useful tools when assessing growth. Furthermore, the relative weight
68 equation allowed a comparison of tilapia from the natural environment and other cultures,
69 expanding the growth studies comparisons. However, one can realize the importance of continuous
70 monitoring of these parameters, since differences were observed in the relative weight of the
71 studied species of tilapia when compared to other environments or cultures, leading us to reflect on
72 how to improve the cultivation conditions of Nile tilapia.

73 **Keywords:** Nile tilapia, culture systems, semi-intensive and intensive systems, growth parameters.
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98 **1.Introduction**

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100 The increase in demand for fisheries resources for human consumption has generated a
101 growth in aquaculture production worldwide. In the last three decades (1980-2010), world
102 aquaculture production has expanded by almost 12 times, with an average annual growth of 8.8%,
103 with emphasis on production in freshwater, which represents about 60% of the total cultivated
104 (FAO, 2012).

105 Among cultured fishes of the world, tilapia is ranked third in terms of production, only after
106 carps and salmonids (El-Sayed, 2006). According to Wing-Keong and Hanin (2007), *Oreochromis*
107 *niloticus* represents 80% of the species of tilapia cultured in the world and is considered the most
108 important. The demand for tilapia has been increasing in recent years and this could only be met by
109 increasing supply and expansion of aquaculture production (Fasakin et al., 2005). However, this
110 growth must be sustainable and in accordance with Abery et al. (2005) if there is no production
111 planning, negative impacts can be generated not only on the system, but also on aquatic ecosystems.

112 Tilapia can be produced in different culture systems and scales, with higher or lower
113 dependence on inputs, especially feed. According to El- Sayed (2006), semi-intensive culture is the
114 production of fish using natural food, through pond fertilization, or fertilization and supplemental
115 feeding, It is a low-cost means of production, which contributes to hunger alleviation especially in
116 rural areas of developing countries. However this system has some barriers, such as the limitation
117 and shortage of supplies of fresh water besides the competition for this resource with agricultural,
118 industrial and other urban requirements, especially in arid regions.

119 Intensive system is characterized by high densities (25-300 fish.m⁻³), diets with high protein
120 levels and high productivity (Mohammad, 2006), furthermore, their technology is relatively
121 inexpensive when compared to other intensive farming systems. Nevertheless, El-Sayed (2006)
122 cites some limitations in cage culture: risk of theft, disease outbreaks, loss due to cage damage

123 caused by predators or storms; low tolerance of fish to poor water quality; dependence on high-
124 quality feeds; the accumulation of faeces and metabolites underneath the cage has a negative
125 environmental impact, among others.

126 The present survey focuses on the following aspects: influence of culture systems /
127 management on the growth parameters of Nile tilapia; different concentrations of protein along with
128 the presence of natural food can contribute to the growth of tilapia in different systems; condition
129 factor of the fish may make changes in cultivation in cages and ponds; condition factor, length-
130 weight relationship and relative weight can be useful tools in growth compared with other
131 populations.

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133 **2.Materials and methods**

134

135 **2.1 Obtaining experimental specimens / Animals and culture conditions**

136

137 The experiment was conducted at the Aquaculture Station of Unidade Acadêmica de Serra
138 Talhada (UAST) / Instituto Agronômico de Pernambuco (IPA) located in the city of Serra Talhada,
139 semi-arid region of Pernambuco State, Brazil, during the period January-May 2011.

140 Juvenile sex-reversed Nile tilapias (variety QAAT1/Arizona-USA) were obtained from the
141 fish farming at the city of Paulo Afonso - BA - Brazil. The fish were stocked in concrete tanks (40
142 fish. m^{-2}) and then submitted to a 20-day acclimatization period for both diets and experimental
143 conditions. Before acclimation, fish were weighed ($38.4 \pm 11.9g$) and measured ($12.7 \pm 1.1cm$).
144 Individuals were fed a ratio of 8% of biomass per day, adjusted every seven days, offered 2 twice a
145 day. The experiment consisted of two treatments (intensive and semi-intensive cultures) and three
146 replicates. In the intensive treatment fish were cultured in cages, while in the semi-intensive they
147 were grown in ponds.

148 For the intensive culture, the dimensions of the cages were 2m x 2m x 1.5m, capacity of 5 m³
149 and 30 mm mesh nylon net. The animals were stocked at a density of 110 fish.m⁻³ (three replicates)
150 and 30 days after the density was decreased to 70 fish.m⁻³ (four replicates). The cages were
151 mounted in reservoir Saco I - Serra Talhada – PE.

152 For the semi-intensive culture were used three ponds with an area of 338 m² and a maximum
153 depth of 1.5 m, supplied by water from the reservoir Saco I. The animals were stocked at a density
154 of 1 fish.m⁻² (0.67 fish.m⁻³) and remained so until the end of the experiment. Before stocking the
155 ponds were dried in the sun and went through the process of liming (CaCO₃ - 15 kg per 1000 m²)
156 and fertilization (bovine manure - 1 ton.ha⁻¹) according to Woynarovich and Horváth (1983) for the
157 growth of natural food such as phytoplankton and zooplankton.

158 The ponds and cages were stocked with juvenile Nile tilapia with an initial average weight of
159 88.6 ± 1.5g and an average total length of 16.3 ± 0.12cm (defined as zero time). 10% of the
160 population were sampled monthly (defined as time 1, 2 and 3) to monitor growth and make feeding
161 adjustments. For this purpose, the fish were anaesthetized using AQUI-S® solution (0.5mL. L⁻¹) for
162 easy handling and return to the ponds/cages after each sampling. Both cultures lasted 94 days.

163

164 **2.2 Feed management**

165

166 Commercial diets containing 35, 32 and 28% crude protein (CP) were used. Composition
167 described in Table 1. Intensive culture used nutritional biphasic plan (35 and 32% CP) and semi-
168 intensive triphasic plan (35, 32 and 28% CP), as described in Table 2. The fish were fed 2 twice a
169 day (8 a.m. and 3 p.m) with feeding rates of 5, 4 and 3% biomass, adjusted monthly after
170 biometrics.

173 **2.3 Water quality**

174

175 During both experiments, temperature ($^{\circ}\text{C}$), dissolved oxygen ($\text{mg} \cdot \text{L}^{-1}$), pH, salinity (‰) and
176 electric conductivity ($\mu\text{S} \cdot \text{cm}^{-1}$) were monitored daily, twice a day *in situ* ($n = 188$) using a
177 multiparameter analyzer Hanna[®].

178 Water samples were taken monthly from each pond and cage and transported to the laboratory
179 of Limnology in the Universidade Federal Rural de Pernambuco to perform the analyzes ($n = 4$).
180 Ammonia ($\text{mg} \cdot \text{L}^{-1}$) was measured according Koroleff (1976); nitrite and nitrate ($\text{mg} \cdot \text{L}^{-1}$) in
181 accordance with Mackereth et al. (1978); orthophosphate ($\text{mg} \cdot \text{L}^{-1}$) under Strickland and Parsons
182 method (1965); alkalinity and hardness ($\text{mg CaCO}_3 \cdot \text{L}^{-1}$) according to Golterman et al. (1978),
183 wherever total hardness was determined from the concentrations of Ca^{+2} and Mg^{+2} .

184 To estimate plankton (phytoplankton and zooplankton) abundance in experimental tanks, 50 L
185 of water were collected in three points of each pond and cage and filtered through 50 mm plankton
186 net to obtain a concentrated sample and stored in plastic bottles of 500 mL. Samples were collected
187 weekly and preserved immediately with 5% formalin. Samples of 2 mL were placed on the counting
188 chamber of the Sedgewick Rafter cell, and then organisms on 10 randomly selected fields of the
189 chamber were identified and counted (Keshavanath, et al., 2002). Phytoplankton ($\text{cel} \cdot \text{mL}^{-1}$) and
190 zooplankton ($\text{ind} \cdot \text{L}^{-1}$) were identified in large groups or families, according to Bicudo and Menezes
191 (2006), Brusca and Brusca (2007) and Hickman et al. (2001). Phytoplankton biomass in terms of
192 pheophytin and chlorophyll *a* concentration ($\mu\text{g} \cdot \text{L}^{-1}$) of water tanks were determined monthly
193 according to Nusch (1980).

194

195

196

197

198

2.4 Growth, survival and feed utilization

199

200 At the end of the experiment, growth was assessed by determination of weight gain rate
 201 (WG), average daily gain (ADG), specific growth rate (SGR), final biomass (FB), Survival (S),
 202 feed conversion ratio (FCR), protein efficiency ratio (PER) and feed efficiency (FE) as follows:

203 Weight gain rate ($WG_{(g)}$) = $BW_f - BW_i$

204 Average daily gain (ADG) = $WG_{(g)} / \text{time (days)}$

205 Specific growth rate (SGR) = $100 (\ln BW_f - \ln BW_i) / \text{time (days)}$

206 Final biomass ($FB_{(Kg)}$) = Final mean weight_(kg) x Final number of fish

207 Survival ($S_{(%)}$) = (Final number of fish/ fish stocked) x 100

208 Feed conversion ratio (FCR) = dry feed offered_(g)/wet weight gain_(g)

209 Protein efficiency ratio (PER) = wet weight gain_(g)/ protein fed_(g)

210 Feed efficiency (FE) = wet weight gain_(g)/dry feed offered_(g),

211 in which BW_i and BW_f = average initial and final body weight_(g) of fish.

212

2.5 Weight-length relationships (WLR) / Condition factor (K)

214

215 Weight-length relationships were estimated at the end of experiment using the equation:

216 $W = aL^b$,

217 in which W = weight of fish in grams, L = total length of fish in centimeters, a is the intercept and b
 218 is the allometric coefficient (Froese, 2006).

219 Fulton's condition factor (K) was determined in time 0 (zero), 31, 63 and 94 for each culture

220 using the equation:

221 $K = 100 W / L^3$,

222 where K = condition factor, W = total weight and L = total length (Froese, 2006).

223 Clark's mean condition factor (K_{mean}) was determined at times 0 (zero), 31, 63 and 96 days,
224 which relates Fulton's condition factor (K) with a and b and which represents the mean condition
225 factor for a given length derived from the respective WLR:

226
$$K_{mean} = 100 aL^{b-3},$$

227 where K_{mean} = mean condition factor for a given length, where a is the intercept and b is the
228 allometric coefficient, both estimated by method of least squares (Clark, 1928).

229 Le Cren's relative condition factor (K_{rel}) was calculated for both systems at times described
230 above, according to Le Cren (1951), which compensates for changes in form or condition with
231 increase in length, and thus measures the deviation of an individual from the average weight for
232 length in the respective sample (Froese, 2006):

233
$$K_{rel} = (W / aL^b) * 100,$$

234 relative condition factor comparing the observed weight of an individual with the mean weight for
235 that length, the factor 100 is used to bring the value for percentage.

236 This same equation (relative condition factor) was used to compare different populations of
237 Nile tilapia in natural environment (using the parameters a and b from FISHBASE for *Oreochromis*
238 *niloticus* and the geometric mean of them) and cultivation (with weight-length relationships in
239 published papers). However it appears as relative weight (W_{rm}):

240
$$W_{rm} = (W / aL^b) * 100 \text{ or } W_{rm} = (W / a_m L^{b_m}) * 100,$$

241 where a_m and b_m are the geometric mean of WLR.

242

243 **2.6 Statistical analysis**

244

245 All data was tested for normal distribution (Shapiro-Wilk test) and homogeneity of variances
246 (Bartlett's test). Student's t -test was used for normally distributed data, whereas Wilcoxon-Mann-
247 Whitney test was used for non-normally distributed data. To assess whether weight-length

248 relationships were affected by culture systems was used General Linear Model. Differences were
249 reported as statistically significant when $P < 0.05$, using OriginTM version 8.0, Assistat 7.6,
250 GraphPad Prism 6.0, IBM SPSS Statistics (Version 19) and Excel 2007.

251

252 **2.6.1 Descriptive analysis**

253

254 For descriptive analyzes of the K_{rel} , total length and weight of Nile tilapia in the two culture
255 systems were used box-plot graphs. The degree of asymmetry between the samples was evaluated
256 according Bisquerra et al. (2004), considering for symmetric data: Mean = Median = Moda; for
257 asymmetric data positive: Mean \geq Median \geq Moda and for asymmetric data negative Mean \leq
258 Median \leq Moda.

259

260 **3. Results**

261

262 The composition of diet and nutritional plan are shown in Tables 1 and 2, respectively.
263 Table 3 shows the parameters of water quality in the intensive and semi-intensive systems. Most of
264 the analyzes showed a significant difference, except ammonia, nitrite, nitrate and orthophosphate.
265 The values of temperature, hardness, Chlorophyll *a* and pheophytin were higher in semi-intensive
266 system; while pH, dissolved oxygen and conductivity were higher in the intensive system.

267 In Table 4 has shown the analyzes of phytoplankton (Chloroficeae, Cyanophyceae and
268 Bacillariophycea) performed in the two farming systems in time "0" (zero), 31, 63 and 94 days. At
269 time "0" the chlorophytes were higher in ponds and at 31, 63 and 94 days of culture, the
270 chlorophytes and bacillariophytes also had their highest densities in semi-intensive system, but the
271 cyanophytes showed higher density in the intensive system at the end of cultivation.

272 Figure 1 shows the development of phytoplanktonic groups during the cultivation. Can be
273 observed a decrease in the total concentration of algae (cel.mL^{-1}) in the two farming systems with
274 the ponds always presenting higher concentrations. The densities of chlorophytes and cyanophytes
275 in intensive system, demonstrated an inverse relationship in time "0", 31 and 63 days.

276 The concentration of zooplankton (rotifers, copepods and cladocera) in different cultures can
277 be seen in Table 5 (Ind.L^{-1}). As for phytoplankton, statistical differences were evaluated by
278 comparison within the times during breeding systems. The concentration of rotifers was higher in
279 semi-intensive system at the times "0", 63 and 94 days. The Cladocera also showed higher density
280 in ponds in the days 31, 63 and 94 days. While copepods were higher in all periods analyzed in
281 semi-intensive system. The zooplankton also decreased their concentrations during the two cultures,
282 but was always higher in semi-intensive farming (Figure 2).

283 The growth parameters of Nile tilapia are shown in Table 6. Only survival, protein
284 efficiency and feed efficiency were better in the semi-intensive system, the remainder of the data
285 showed no significant difference.

286 The parameters of the mathematical models for the evaluation of weight-length relationships
287 of fish under different cultures are displayed in Figure 3. An analysis of the models revealed that the
288 total weight-length relationships were not affected by culture system ($P<0.05$).

289 The Fulton's condition factors (K) of the experimental specimens are shown in Table 7. The
290 only statistical difference was noted with 31 days of culture when the fish of the semi-intensive
291 system showed better conditions when compared to intensive system.

292 Figure 4 shows a comparison between Fulton's condition factor (K) of the population in the
293 present study with the respective K_{mean} . At the beginning of the culture, condition factor of fish in
294 intensive system was similar to the K_{mean} of the population, however this factor decreased in 31
295 days, then increased and stabilized with K_{mean} . The condition factor of fish in the semi-intensive

296 system also showed similarity to the respective K_{mean} in the times "0", 63 and 94 days, however, this
297 condition factor was greater than its corresponding K_{mean} with 31 days of cultivation.

298 The K_{rel} of cultured fish under intensive and semi-intensive system were analyzed by Box-
299 plot graphics (Figure 5). At time "0" the intensive system showed 25% of the fish with a K_{rel}
300 between 82 and 97% (between the minimum and the first quartile), 50% exhibited K_{rel} between 97
301 and 107% (1st quartile to 3rd quartile) and the remainder of the animals (25%) showed a K_{rel}
302 between 107 and 119% (3rd quartile to the maximum value). The mean and median are presented
303 overlapping (approximately 102%) and most fish exhibited a K_{rel} of 95% for intensive culture. The
304 fish in the semi-intensive the mode show overlapping the mean and median, and minor fluctuations
305 that time.

306 Evaluating fish K_{rel} with 31 days of cultivation were observed changes in condition. The
307 semi-intensive system showed 75% of its population between 96 and 119% of relative condition,
308 mean, mode and median higher than in intensive system. Seventy-five percent of the fish in the
309 intensive system exhibited K_{rel} between 93 and 113%.

310 This distribution pattern remained until at least day 63 of cultivation in the semi-intensive
311 system. However, an improvement was observed in the intensive system with 94 days of
312 cultivation. The same showed 75% of its population with a K_{rel} between 96 and 120% compared
313 with 93 to 114% of semi-intensive system. At the end of the experiment, most fish have had a K_{rel}
314 of 93% for the intensive system and 88% for semi-intensive system presenting mean and median
315 similar in the two farming systems (approximately 100%).

316 The total length and weight of fish in intensive and semi-intensive farming systems can be
317 seen in Figure 6A and 6B, respectively, in Box-plot format.

318 A similar increase was observed in both systems up to 63 days of culture (Figure 6A).
319 However, the minimum and maximum growth in the intensive showed a higher variation when
320 compared to semi-intensive system. Was observed a symmetry (mean = median = mode) at the

321 times "0" and 31 days for both systems. From 63 days of cultivation occurred changes in the
322 consistency of growth of the animals in intensive system. They began to show positive asymmetric
323 growth (mean \geq median \geq mode) whereas animals in semi-intensive remained in symmetric pattern
324 until the end of the experiment. However, both systems exhibited outliers in the course of
325 cultivation.

326 Evaluating increase in weight of fish in different farming systems (Figure 6B) was observed
327 in the time "0" and 31 days similarity in growth. However, a larger weight variation was noticed in
328 intensive system from 63 days culture. Evaluating the symmetry in the two culture systems, up to
329 63 days, the animals exhibited positive asymmetry and the fish of the semi-intensive cultivation
330 exhibited up to 94 days. Fish intensive system exhibited negative asymmetry with 94 days culture.

331 The relative weight of fish with 94 days of cultivation in the intensive and semi-intensive
332 was compared with total weight-length relationships of fish obtained in other environments (natural
333 and cultivated) (Table 8).

334 The relative weight of fish in the present study was higher in 9 of the 14 cases analyzed, and
335 3 cases with similar relative weight. Our relative weight found for tilapia in comparison with the
336 tilapia from natural environment was higher in 4 of the 6 cases analyzed, and one situation similar
337 to the relative weight.

338 Comparing studies with tilapia cultured in cages with the two systems described here, the
339 relative weights obtained were higher in the present work in 1 of the 2 situations analyzed. This
340 comparison was also made with tilapia cultured in ponds. The tilapia cultured in the present study
341 had higher relative weights in 2 of the 4 studied cases and two similar situations.

342 Nile tilapia cultured in aquaria subjected to different diets were also compared with the
343 present study and the relative weight found here was also greater than those of fish from aquaria.

344

345

346 **4. Discussion**

347

348 According to the Brazilian Council for Environment – CONAMA, resolution N° 357 of March
349 17, 2005, which provides on the classification and environmental guidelines of surface waters, the
350 water farming systems analyzed were defined as brackish (waters with salinity higher than 0.5 ‰
351 and lower than 30 ‰), corroborating with the value found, the electrical conductivities of the water
352 in intensive and semi-intensive systems were $1,356 \pm 468.1$ and $1,285 \pm 249.1$ mS. cm⁻¹
353 respectively, being considered a brackish water (Bouvy et al., 1998). Tilapia species and hybrids
354 tolerate a wide range of salinity, they can grow and reproduce normally in brackish water and was
355 selected for this type of cultivation (El-Sayed, 2006). Alfredo and Hector (2002) define a limit of
356 salinity for optimal growth of Nile tilapia of 15 ‰ and can tolerate 36 ‰ with acclimation (Al-
357 Amoudi, 1987).

358 According to El-Sayed (2006), temperature is one of the most important factors affecting the
359 physiology, growth, reproduction and metabolism of tilapia. This fish is thermophilic and known to
360 tolerate a wide range of water temperatures. Balarin and Haller, (1982); Chervinski, (1982);
361 Philippart and Ruwet, (1982) state that the optimum temperature for development of the species is
362 about 25-30°C, however, the temperature range for the normal development, reproduction and
363 growth of tilapia is about 20 to 35°C, values found in the present study.

364 Nile tilapia can survive at a pH range of 4 – 11, however, this variation was not found in this
365 study, where the values of pH were in accordance to CONAMA (357/2005) resolution.

366 Only a minimum value of dissolved oxygen in the semi-intensive system ($3.27 \text{ mg} \cdot \text{L}^{-1}$), was
367 out of the requirements of CONAMA, which advocates a value equal or higher than $5 \text{ mg} \cdot \text{L}^{-1}$.
368 However, according to Coche (1982), Boyd and Tucker (1998), the minimum condition of
369 dissolved oxygen for cultivation of tilapia is $3.0 \text{ mg} \cdot \text{L}^{-1}$. El-Sayed (2006) mentions that tilapia can

370 tolerate levels between 0.1 - 0.5 mg. L⁻¹ dissolved oxygen, however these values can cause decrease
371 in growth.

372 Ammonia levels in the two farming systems were compatible with a good growth performance
373 of Nile tilapia. El-Shafai et al. (2004) evaluated the effect of prolonged exposure to sublethal
374 Ammonia nitrogen on the growth performance of Nile tilapia. They found that the toxic level of
375 NH₃-N and its negative effect on the growth performance ranges from 0.07 to 0.14 mg . L⁻¹. They
376 further recommended that the NH₃-N concentration should be maintained below 0.1 mg . L⁻¹.

377 The amounts of nitrite and nitrate in the present study are at acceptable levels according to
378 CONAMA. Nitrite is highly toxic to fish, because it disturbs the physiological functions of the fish
379 and leads to growth retardation. Nitrate is relatively non-toxic to tilapia, however their exposure to
380 high levels for a long time, can decrease the immunity of fish (El-Sayed, 2005).

381 The orthophosphates are directly available for biological metabolism without conversion to
382 simpler forms. Nevertheless, to characterize a system as eutrophic this must have total P values
383 higher than 0.05 mg.L⁻¹. According to the CONAMA polyphosphate recommendations (presenting
384 structure more complex than the orthophosphate) should be less than 0.02 mg. L⁻¹, value found in
385 this study.

386 The CONAMA do not provide optimal values of alkalinity and hardness, however, Boyd and
387 Tucker (1998) suggest that these parameters should be above 20mg.L⁻¹. In the present study, the
388 intensive system had lower alkalinity (13.95 ± 1.86 mg.L⁻¹ CaCO₃) vs. 161.28 ± 106.97 mg.L⁻¹
389 CaCO₃ in semi-intensive, thus demonstrating that the addition of CaCO₃ in the semi-intensive
390 promoted effect "buffer" necessary to minimize variations in pH (7.15 ± 0.4).

391 Water hardness reflects mainly the concentration of Ca⁺ and Mg⁺ which are combined with
392 carbonate and/or bicarbonate, may also be associated with sulphate and chloride. The hardness
393 values found in intensive system (20.06 ± 1.09 mg. L⁻¹ CaCO₃) were reduced significantly from the
394 values found in the semi-intensive (28.50 ± 4.37 mg. L⁻¹ CaCO₃), furthermore they were at the limit

395 defined by Boyd and Tucker (1998). The hardness values presented in this study define water
396 culture like soft, though these values are constantly changing due to the characteristics of the semi-
397 arid environment (relative humidity around 40%, long droughts and occasional rains concentrated
398 in a few months, with precipitation between 400 and 800 mm annually). This study was conducted
399 between January and May 2010, months where the occasional rains are more concentrated.

400 Levels of chlorophyll *a* indicate the degree of eutrophication of an environment, and provide
401 the phytoplankton biomass. According to CONAMA recommended levels of chlorophyll *a* (for
402 freshwater aquaculture) are up to $30 \mu\text{g. L}^{-1}$, presenting no reference to brackish water. Addition of
403 feed and manure to culture ponds resulted in different biomass of phytoplankton in the ponds. The
404 levels found in chlorophyll *a* for intensive and semi-intensive system (17.75 ± 5.92 and $88.49 \pm$
405 $31.16 \mu\text{g. L}^{-1}$, respectively) show clearly the effect of fertilization performed in ponds, which
406 favored the development of natural food. Uddin et al., (2007) assessed the concentration of plankton
407 in monoculture (tilapia) and polyculture (tilapia and shrimp), and observed values of chlorophyll
408 "*a*" ranging from 50 to $500 \mu\text{g.L}^{-1}$ (minimum and maximum between treatments) but these values
409 were not statistically different, however, he noted a decrease in values over time.

410 According to Wetzel (1981) and Lewis (1995), pheophytin is a result of degradation of
411 chlorophyll *a* after the loss of magnesium atom. Thus, the determination of chlorophyll *a*
412 concentrations provides an estimate of chlorophyll whereas pheophytin determination indicates
413 their physiological status, i.e. high values of pheophytin, indicates a larger degradation of
414 chlorophyll *a*, suggesting death of phytoplankton and high decomposition. Phaeophytin values were
415 higher in semi-intensive system ($274.97 \pm 151.95 \mu\text{g. L}^{-1}$) (which also had higher chlorophyll *a*),
416 suggesting a high renewal of phytoplankton.

417 With 14 weeks of cultivation in ponds Keppler (2005) found average values of phaeophytin
418 ranging between 585 ± 789 and $624 \pm 707 \mu\text{g. L}^{-1}$, in the morning and afternoon respectively,
419 however no statistical difference. Keppler (2005) also found a positive correlation between

420 chlorophyll *a* and pheophytin, demonstrating a renewal of plankton during cultivation. Lima (2010)
421 found phaeophytin values in nursery ponds with *Astyanax lacustris* ranging from 8.585 to 73.238
422 µg.L⁻¹, however time the experiment was lower than in the present work.

423 The microalgae contain high concentration of soluble fiber and fatty acids of the omega-3
424 series and can contribute positively in the feeding of aquatic organisms (Azaza et al., 2007).
425 Generally was observed a greater abundance of the three groups analyzed during cultivation in
426 semi-intensive system. Analyses of chlorophyll *a* confirmed this observation. This shows the
427 importance of fertilization in ponds, as well as maintenance of the main biotic and abiotic
428 parameters that promote equilibrium in the system.

429 The largest groups of algae found were chlorophyceae and cyanophyceae. However, the
430 intensive system showed a particularity. Analyzing Figure 1A, the cyanophyceae and
431 Chlorophyceae showed an inverse relationship development with 31 and 63 days of cultivation.
432 This may be related to changes in the environment that promoted an ecological succession of these
433 algae in the months analyzed.

434 In ponds this characteristic was not observed since the occurrence of Chlorophyceae was
435 predominant. This fact is important, since cyanophyceae can produce biotoxins that are significant
436 hazards to animals (Molica et al., 2002) and in addition, is an indicative of eutrophication of an
437 environment. The CONAMA does not define the maximum amount of this algae in brackish water,
438 but freshwater for aquaculture its maximum allowed is 50,000 cel.mL⁻¹. The greatest value of this
439 alga in the studies was in intensive system with 31 days of cultivation ($46,000 \pm 13,600$ cel.mL⁻¹)
440 (maximum value above recommended). However, Deblois et al. (2008) cite the tilapia as an
441 excellent filtrating of cyanophyceae, demonstrating a preference for this group compared to
442 bacilariofíceas (Uddin et al., 2007). Furthermore, Uddin et al. (2009) reported that tilapia can exert
443 a very effective biological control over plankton in semi-intensive system.

444 The abundance of zooplankton was also observed during the cultivation of Nile tilapia. These
445 organisms play an important role in the trophic web, serving as a link between primary producers
446 and higher consumers. According Roche and Rocha (2005), the availability of these organisms in
447 fish farming becomes in most cases essential for fish feeding.

448 The densities of rotifer, copepod and cladocera were significantly higher in semi-intensive
449 system in the majority of times analyzed and the copepods reached higher values in both systems.
450 Despite the diversity of feed from primary producers, consumers also depend on the primary
451 continuous supply of organic detritus as a source of energy. Thus, the predominance of copepods in
452 the analyzed systems can be linked to the eating habits of cyclopoid - the main order of Copepoda
453 found during this study (data not shown). The cyclopoid are raptorial predators who thrive in
454 eutrophic environments rich in decomposing organic matter (Landa et al., 2007).

455 Actually, as an omnivorous opportunistic fish, tilapia is able to assimilate nutrients from
456 different sources, like benthic algae, phytoplankton, macrophytes, zooplankton, small invertebrates,
457 detritus and cyanophyceae. The feed choice is mainly related to the availability of food items in the
458 environment (Bowen, 1982; Fitzsimmons, 1997; Stickney, 1997; Beveridge and Baird, 2000; Lowe-
459 McConnell, 2000).

460 When *O. niloticus* in semi-intensive ponds are supplemented with low-protein feeds, dietary
461 protein is largely provided by the natural food. Liti et al. (2006) mentions that the protein content of
462 natural food ranges between 550 and 700 g. kg⁻¹ on a dry matter basis. This is far above the range
463 (270 - 350 g.kg⁻¹) recommended for intensive culture of Nile tilapia (El-Sayed, 2006).

464 Complementing this observation there were no statistical differences in final weight of Nile
465 tilapia grown in the intensive and semi-intensive, i.e. the triphasic nutrition plan (35, 32 and 28%
466 CP) applied to fish in the pond (Table 2), together with the natural food available, was sufficient to
467 obtain the same weight gain of intensive cultivation (biphasic plan with 35 and 32% CP). Liti et al.

468 (2006) mentions that natural food contributed between 300 and 500 g.kg⁻¹ of growth when tilapia
469 were supplemented with artificial feeds in fertilized ponds.

470 The effectiveness of natural food was also observed in protein efficiency ratio (PER) and feed
471 efficiency factor (FE). Even the two systems with the same feed conversion rates, evaluating the FE
472 and PER of fish the semi-intensive system showed better utilization of provided food. This means
473 that they had the same weight gain of fish in intensive system with less protein in the provided diet.
474 In addition, both systems exhibited growth rates, feed conversion and survival in agreement with
475 commercial scale production.

476 However, the intensive system showed advantages with respect to productivity, precisely
477 because it is a system that enables higher stocking densities, compensating the greatest amount of
478 protein supplied in the diet. In fact, the constant movement of the cages by water changes, high fish
479 densities and low densities of phytoplankton and zooplankton, hinder consumption of natural food
480 in this system. So diets provided for intensive systems must meet all the nutritional requirements of
481 the produced species.

482 Corroborating data growth, weight-length relationship ($W = aL^b$) of Nile tilapia in the two
483 cultures was not affected by the systems. The weight-length relationship showed the exponent $b >$
484 3.0 for both systems. According to Froese (2006), an exponent less than 3 indicates a "decrease in
485 condition or form" or elongated shape and fish presenting $b > 3$ means an increase in the condition
486 or form, i.e., the specimens analyzed further increased in height/width than length. Evaluating the
487 exponent b of the weight-length relationship of Nile tilapia available in the literature (natural and
488 cultivated), were found values ranging between 2.7 and 3.3.

489 In fact Nile tilapia appears to exhibit allometric growth (growth which does not comply to
490 cube law). For weight-length relationship, there are cases in which the slope was significantly
491 below or above the expected value of three for fish growth in general to be assumed as isometric.

492 Different exponents could also suggest that the fish we are dealing with are from different stocks
493 (Njiru et al., 2006).

494 In zootechnical interest of obtaining the condition of live animals, non-destructive techniques
495 have been applied enabling individuals to be cultivated at lower stress levels. This can be done by
496 the use of structure and weight, which combined as indices provide indirect estimates of energy
497 storage for animals (Camara et al., 2011). Condition factor studies take into account the health and
498 general well-being of a fish in relation to its environment (Olurin and Aderibigbe, 2006).
499 Morphometric indices which assume that heavier fish of a given length are in better condition are
500 simple indicators of energy storage (Lloret et al., 2000). This is a common practice in studies with
501 fish, where the conditions are estimated from its weight and length (Camara et al., 2011).

502 Table 7 shows the Fulton's condition factor found in fish submitted to different cultures,
503 where was observed statistical difference in the factor of fish with 31 days of cultivation. The best
504 conditions found at this time (semi-intensive system) may be associated with the consumption of
505 natural food, as currently feeding rate and CP concentration in the diet were the same for both
506 systems. Diana et al. (1994) evaluated the effects of adding supplemental food in ponds fertilized on
507 the growth of Nile tilapia, and observed that the combination of feeding and fertilizers were more
508 efficient than the growth based on these two items alone.

509 Assessing the condition factor found with population K_{mean} (Figure 4) with 31 days of
510 cultivation, condition factor of fish in semi-intensive system increased according to its
511 corresponding K_{mean} and the opposite occurred for fish in intensive system. This reinforces the
512 importance of natural food in the early stages of cultivation. However, with decreasing
513 concentrations of crude protein in diets for fish of the semi-intensive system along with decreased
514 availability of natural food, the condition factor of the fish in the ponds decreased (not decaying
515 K_{mean} below) whereas in the intensive increased, making the two systems not statistically different

516 in the course of cultivation. The condition factor in both cultures remained close to K_{mean} of the
517 population by the end of cultivation, demonstrating the efficiency of management.

518 Offem et al. (2007) assessed the condition factor in male Nile tilapia in a tropical floodplain
519 River and observed values ranging from (0.554-1.376). They considered these values as efficient for
520 tilapia and ascribed this condition to the abundance of natural food in the area. Huchette and
521 Beveridge (2003) also observed an improvement in the condition of Nile tilapia cultured in cages
522 with presence of natural food and diet. Olurin and Aderibigbe (2006) found a mean condition factor
523 of 1.14 for males of tilapia cultured in ponds, considering the value acceptable.

524 Condition factor can influence the reproduction and survival of populations (Lambert and
525 Dutil, 2000) reflecting interactions between biotic and abiotic factors and the physiological
526 condition of the fish. It shows the well-being of the population during the various stages of the life
527 cycle (Lizama and Ambrósio, 2002) and since it is a simple calculation can be used to manage fish
528 farms.

529 The relative condition factor (K_{rel}) represents the ratio of the observed weight and the weight
530 estimated by the weight-length relationship (Le Cren, 1951; Froese, 2006). The use of Box-plot
531 graphs in an evaluation of K_{rel} provided interesting information about the two populations under
532 study (Figure 5). With these tools together is possible to identify the lowest and highest value of K_{rel}
533 in farming systems, as well as variations of K_{rel} between the 1st and 3rd quartile, identifying in
534 more detail the possible changes in population. For the same distribution, measures of central
535 tendency (mode, mean and median) of K_{rel} should be overlapped, and minimum and maximum
536 values near the 1st and 3rd quartile, respectively.

537 Since the beginning of the cultivation the values of K_{rel} ranged between 80 and 120%,
538 indicating a large variation in the condition of farmed fish. However, the mean and median showed
539 up near 100%. The mode was not viewed in a standardized manner, in three cases it was below
540 average/median (in intensive system, with zero and 94 days of cultivation in both systems).

541 However, these values were close to 95%. The variations found in K_{rel} of populations ($\pm 20\%$) may
542 be related to uneven growth of the Nile tilapia, since after 31 days of culture values outside the box
543 plot (between the 1st and 3rd quartile) began to be more evident.

544 In Figure 6 is more evident the variation in length and weight of Nile tilapia during the
545 cultivation and the measures of central tendency were still closer. However, positive and negative
546 asymmetries were observed in both systems. McGinty (1986) evaluated the effect of density on
547 weight gain of Nile tilapia and noted that the growth of small individuals was being depressed by
548 the larger ones because of competition.

549 A comparative study of the relative weight of Nile tilapia was done with specimens of natural
550 environment and culture. In general, we observed a higher relative weight of experimental
551 specimens (9 of 14 cases). Better growth is expected in the cultivations, as well as a good carcass
552 yield, feed conversion and homogeneity, since it works with improved strains of Nile tilapia.
553 However, a decrease in genetic variability can reverse the situations described above, since a
554 substantial loss of this variability is expected, due to the characteristics of the reproduction of tilapia
555 and the appearance of a small size population when there is no proper genetic management of
556 breeding stock (Moreira et al., 2007). The relative weight of the fish in this experiment was greater
557 than the weight of tilapia from Lake Tana (Ethiopia), Ouémé River Basin (Benin), Hippopotamus
558 Pond (Burkina Faso) and Lake Victoria (Kenya).

559 In Lake Tana, Tadesse (1997) observed a seasonal variation in condition factor, assigning
560 lower values to energy expenditure after spawning. This variation in the condition found by the
561 author may have generated differences in the WLR, decreasing their relative weight. A similar fact
562 may have occurred in Ouémé River Basin (Benin), (Laleye, 2006). The same author also mentions
563 that the WLR can change according to fish, sex, maturity, season and time (because of gastric
564 fullness). The relative condition factor was also lower in fish from Hippopotamus Pond (Béarez,
565 2003). This lake is characterized by having a high fishing effort, with total absence of large fish.

566 The fishing activity during the dry season has affected spawning, as a consequence dwarfism
567 becomes more evident in populations.

568 According to Njiru et al. (2006) Nile tilapia from Lake Victoria showed good adaptation to the
569 site including little seasonal variation in their condition, becoming the most commercially important
570 fish in the lake and the third most captured. In this case, fish farmed presented here also showed
571 improved relative weight (approximately 15% more). The authors report that there was an increase
572 in fertility, suggesting a tactic to maximize reproductive success in Lake Victoria. This tactic may
573 be a response to overfishing of the population whose effects probably led to stunting.

574 However, Ahmed et al. (2003) indicate that in the Kaptai Reservoir (Bangladesh), fishing
575 pressure is below the capability of the stock to be increased by about 16%. They also mention that
576 the introduction of *O. niloticus* was accidental, however it has successfully established itself in the
577 reservoir. In this case the relative weight was found to be similar to the present study.

578 In Kainji Lake (Nigeria), Feu and Abiodun (1998) mentioned that anthropogenic activity was
579 great in that lake and the fish were captured below the ideal size, but with actions such as the
580 eradication of purse seine and adjustments in the size of meshes allowed the growth of the fish.
581 Moreover, according FISHBASE, today there are native species of Nile tilapia that can reach up to 74
582 cm in total length. These factors may have contributed to a poor relative weight of tilapia in the
583 present work compared with the tilapias from Kainji Lake.

584 One can see that action/awareness conducted by the authorities towards the population
585 behavior, contributed to the growth of the fish in Lake Kainji. From the ecological point of view
586 this fact is essential to maintaining the population status of the species. However, this
587 greater/similar relative weight in fish from natural environment must be observed with caution
588 (when compared with strains of aquaculture), since the species of tilapia used in fish farming went
589 through genetic improvements aiming to improve the quality of the breeding.

590 Comparing the tilapia described in the present work with those cultured in cages was observed
591 one relative weight major and other minor. Saraiva (2004) evaluated the effect of stocking density
592 on growth of juvenile Nile tilapia. He noted that there was no significant difference in the growth of
593 his specimens. However, when comparing with the present study the relative weight of tilapia
594 cultured in this work was higher than Saraiva (2004). He mentions that he worked with the strain of
595 tilapia chitalada or Thai and the various squads probably have their genetic variability decreased,
596 causing reduction in growth.

597 However, cultivation performed by Medri et al. (1999) showed better relative weight in
598 comparison with the work described here. Unlike Saraiva (2004), Medri et al. (1999) evaluated the
599 growth of tilapia with the inclusion of an alternative ingredient in feed (distillers yeast -
600 *Saccharomyces cerevisiae*), and noted that the rates used showed no statistical difference. This
601 change in diet may have influenced for a better relative weight.

602 Olurin and Aderibigbe (2006) analyzed the condition factor and WLR of Nile tilapia grown in
603 nurseries in Ijebu-Ode, Nigeria. When compared with the present study it was observed that the
604 tilapia produced here showed better relative weight. Also greater than the tilapia grown by Carmo et
605 al. (2008) in a Station of aquaculture in Brazil. Olurin and Aderibigbe (2006) did not mention if
606 they used tilapia with genetic modification, whereas Carmo et al. (2008) worked with a common
607 strain. This fact may have contributed to better growth observed in this study.

608 Carmo et al. (2008) also evaluated the growth of chitalada and red strain in ponds. The two
609 cultures performed by him had a relative weight similar to that observed in the present study,
610 indicating that there was not much difference in growth patterns of the three strains.

611 Leal et al. (2009) evaluated the growth of juvenile Nile tilapia in aquaria, fed with different
612 concentrations of hydrolyzed protein and a commercial feed. It was observed that the tilapia of this
613 study had a higher relative weight of tilapia fed commercial feed, however, when compared with
614 tilapia fed with 5% hydrolyzed protein diet (better WLR according to the author) observed a similar

615 growth. Leal et al. (2009) reported no strain used, however it can be seen that the feeding influence
616 the results.

617 El-Sayed (2006) cites that variations in body size of tilapia in natural habitat may be related to
618 the different environmental factors rather than genetic differences. Mair et al. (2002) evaluated the
619 performance of tilapia farms in the Philippines (mainly Nile tilapia) and noted that growth has
620 declined after the newly introduced pure strains, probably due to loss of genetic variation through
621 founder effect (*bottleneck effect*) and introgression with Mozambican tilapia (*Oreochromis*
622 *mossambicus*). In fact genetic and environmental variations may cause changes in growth. Dey et
623 al. (2000) analyzed the performance of the GIFT strain (Genetically Improved Farmed Tilapia)
624 compared to non-GIFT on farms in Bangladesh, China, Philippines, Thailand and Vietnam, and
625 observed a better growth of the GIFT strain. Santos et al. (2008) evaluated the growth of Nile tilapia
626 in Brazil (GIFT and Thai strains) and also noted the improved performance of GIFT. Tenório et al.
627 (2012) compared the performance of three strains of Nile tilapia defined as common chitalada and
628 mestizo (crossroads of common with chitalada) and found better growth with chitalada and
629 mestizo.

630

631 **5. Conclusion**

632

633 At the end of cultivation, the two farming systems showed the same trend of growth, i. e., the
634 adopted management promoted the same growth performance for both systems.

635 The calculations of Fulton's condition factor (K), Clark's mean condition factor (K_{mean}), Le
636 Cren's relative condition factor (K_{rel}) and relative weight (W_{rm}) proved to be useful tools when
637 assessing growth. Furthermore, the relative weight equation allowed a comparison of tilapia from
638 the natural environment and other cultures, expanding the growth studies comparisons. However,
639 one can realize the importance of continuous monitoring of these parameters, since differences were

640 observed in the relative weight of the studied species of tilapia when compared to other
641 environments or cultures, leading us to reflect on how to improve the cultivation conditions of Nile
642 tilapia.

643 The intensification of cultivation generates a greater need for skilled workers and therefore
644 implies in an increase in the risks of production, especially with regard to the genetic variability of
645 these animals. The lack of proper selection and genetic information from families can lead to an
646 inbreeding framework reflecting in decreasing productivities.

647

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649

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935 **7. Tables**

936

937 Table 1: Proximate analysis of the commercial diets used in the experiment

Proximate analysis (as-fed basis, g . kg ⁻¹)*	Crude Protein		
	35%	32%	28%
Pellet diameter (mm)	3 - 4	4 - 6	6 - 8
Moisture	130	130	130
Crude protein	350	320	280
Ether extract	65	50	50
Crude fiber	50	50	50
Ash	120	120	120
Calcium	15	15	15
Phosphorus	6	6	6
Vitamin C	0.5	0.5	0.5

938 *Data from private company

939

940 Table 2: Nutritional plan used as feed for Nile tilapia (*O. niloticus*) subjected to different culture
941 systems

Time (days)	Nutrition plan	
	Intensive	Semi-intensive
	culture	culture
0 – 31	35% CP	35% CP
32 – 63	35% CP	32% CP
64 – 94	32% CP	28% CP

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947 Table 3: Quality indicators of water in intensive and semi-intensive cultured Nile tilapia (*O.*
 948 *niloticus*)

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Parameters	Intensive culture	Semi-intensive culture
Salinity (‰)	0.62 ± 0.14 ^a	0.64 ± 0.1 ^a
Electric conductivity (µS . cm ⁻¹)	1,356 ± 468.1 ^b	1,285 ± 249.1 ^a
Temperature (°C)	29.40 ± 1.4 ^b	30.15 ± 2.6 ^a
pH	8.07 ± 0.2 ^b	7.15 ± 0.4 ^a
Dissolved oxygen (DO) (mg . L ⁻¹)	8.19 ± 1.9 ^b	6.67 ± 3.4 ^a
Ammonia nitrogen (mg . L ⁻¹)	0.065 ± 0.032 ^a	0.126 ± 0.061 ^a
Nitrite (mg . L ⁻¹)	0.053 ± 0.050 ^a	0.017 ± 0.013 ^a
Nitrate (mg . L ⁻¹)	0.001 ± 0.000 ^a	0.002 ± 0.001 ^a
Orthophosphate (mg . L ⁻¹)	0.016 ± 0.016 ^a	0.001 ± 0.000 ^a
Alkalinity (mg CaCO ₃ . L ⁻¹)	13.95 ± 1.86 ^b	161.28 ± 106.97 ^a
Hardness (mg CaCO ₃ . L ⁻¹)	20.06 ± 1.09 ^b	28.50 ± 4.37 ^a
Chlorophyll <i>a</i> (µg . L ⁻¹)	17.75 ± 5.92 ^b	88.49 ± 31.16 ^a
Pheophytin (µg . L ⁻¹)	19.53 ± 1.86 ^b	274.97 ± 151.95 ^a

950 Values are mean ± SE of mean. Within a row, means with different superscript letters differ
 951 significantly (P < 0.05) by Wilcoxon-Mann-Whitney test and Student's *t* test.

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959 Table 4: Abundance of phytoplankton (cel.ml^{-1}) in water of intensive and semi-intensive systems Nile tilapia (*O. niloticus*) in time "0" (zero), 31, 63
 960 and 94 days.

Groups	Time (days)							
	0 day		31 days		63 days		94 days	
	Intensive culture	Semi-intensive culture	Intensive culture	Semi-intensive culture	Intensive culture	Semi-intensive culture	Intensive culture	Semi-intensive culture
Chlorophyceae	$32,500 \pm 12,200^b$	$73,300 \pm 22,300^a$	$5,000 \pm 2,300^b$	$55,000 \pm 12800^a$	$36,300 \pm 3,000^b$	$48,000 \pm 53^a$	$8,300 \pm 1,200^b$	$26,000 \pm 5,700^a$
Cyanophyceae	$24,000 \pm 15,700^a$	$12,600 \pm 9,600^a$	$46,000 \pm 13,600^a$	$26,000 \pm 5700^a$	$2,000 \pm 700.5^a$	$2,200 \pm 600^a$	$7,400 \pm 2,200^a$	$1,200 \pm 200^b$
Bacillariophyceae	600 ± 500^a	$2,200 \pm 1,200^a$	300 ± 200^b	$1,300 \pm 173^a$	200 ± 150^b	$1,100 \pm 300^a$	700 ± 600^b	800 ± 200^a

961 Values are mean \pm SE of mean. Within a row, at the same times, means with different superscript letters differ significantly ($P < 0.05$) by Wilcoxon-
 962 Mann-Whitney test and Student's *t* test (n=4).

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964 Table 5: Abundance of zooplankton (Ind.L^{-1}) in water of intensive and semi-intensive systems Nile tilapia (*O. niloticus*) in time "0" (zero), 31, 63 and
 965 94 days.

Groups	Time (days)							
	0 day		31 days		63 days		94 days	
	Intensive culture	Semi-intensive culture	Intensive culture	Semi-intensive culture	Intensive culture	Semi-intensive culture	Intensive culture	Semi-intensive culture
Rotifera	$9,503 \pm 2,370^b$	$33,490 \pm 8,478^a$	$16,020 \pm 3,703^a$	$18,521 \pm 4,718^a$	16 ± 29^b	714 ± 198^a	123 ± 32^b	753 ± 276^a
Cladocera	$12,521 \pm 2,401^a$	$15,512 \pm 3,902^a$	$7,500 \pm 2,302^b$	$20,036 \pm 552^a$	3.5 ± 3.0^b	80 ± 55^a	3.0 ± 2.0^b	220 ± 92^a
Copepoda	$88,029 \pm 12,200^b$	$147,496 \pm 33,201^a$	$86,498 \pm 2,622^b$	$113,527 \pm 2,237^a$	182 ± 25^b	$1,423 \pm 740^a$	172 ± 7.0^b	767 ± 326^a

966 Values are mean \pm SE of mean. Within a row, at the same times, means with different superscript letters differ significantly ($P < 0.05$) by Wilcoxon-
 967 Mann-Whitney test and Student's *t* test (n=4).

968 Table 6: Growth performance and nutrient utilization of Nile tilapia (*O. niloticus*) cultivated in
 969 intensive and semi-intensive system in 94 days.

Parameters	Intensive culture	Semi-intensive culture
Density (m ³)	110	0.67
Initial weight (g)	88.33 ± 1.54 ^a	86.90 ± 1.50 ^a
Final weight (g)	543.58 ± 48.11 ^a	501.74 ± 4.37 ^a
WG (g) ¹	455.24 ± 41.68 ^a	414.84 ± 3.48 ^a
ADG (g day ⁻¹) ²	4.84 ± 0.44 ^a	4.41 ± 0.04 ^a
SGR (% day ⁻¹) ³	1.96 ± 0.08 ^a	1.87 ± 0.01 ^a
Final biomass (kg)	228.46±16.18 ^a	143.99±2.74 ^b
Survival (%)	75.0 ^b	85.0 ^a
FCR ⁴	1.59 ± 0.12 ^a	1.57 ± 0.11 ^a
PER ⁵	4.23 ± 0.37 ^b	6.51 ± 0.30 ^a
FE ⁶	1.6±0.02 ^b	2.30±0.11 ^a

970 Values are mean ± SE of mean. Within a row, means with different superscript letters differ
 971 significantly (P < 0.05) by Wilcoxon-Mann-Whitney test and Student's *t* test

972 ¹ Weight gain, ² Average daily gain, ³ Specific growth rate, ⁴ Feed conversion ratio, ⁵ Protein
 973 efficiency ratio, ⁶ Feed efficiency.

974

975 Table 7: Fulton's condition factor of Nile tilapia cultivated in intensive and semi-intensive systems
 976 at times "0" (zero), 31, 63 and 94 days

Time (days)	Intensive culture	Semi-intensive culture
0	1.97±0.06 ^a	2.00±0.01 ^a
31	1.95±0.02 ^b	2.17±0.07 ^a
63	2.03±0.06 ^a	2.11±0.04 ^a
94	2.10±0.08 ^a	2.08±0.01 ^a

977 Values are mean ± SE of mean. Within a row, means with different superscript letters differ
 978 significantly (P < 0.05) by Student's *t* test

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981 Table 8: Relative weight (W_{rm}) of fish with 94 days of cultivation in the intensive and semi-intensive systems compared with total weight-length
 982 relationships of fish obtained in other environments (natural and cultivated) of the experimental specimens in the analyzed environments.

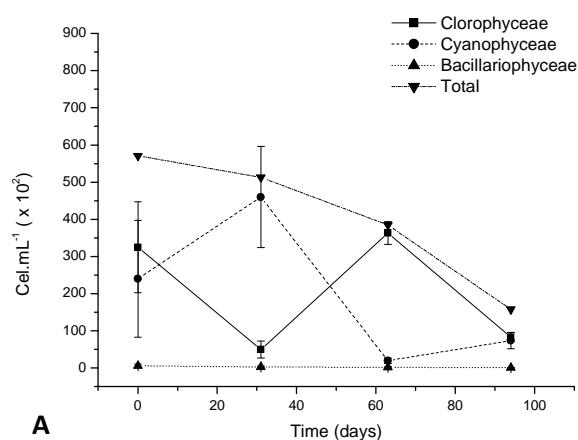
Country	Environment	Locality	Length class (cm)	Sex	a	B	R^2	References	W_{rm} Intensive system	W_{rm} Semi-intensive system
Brazil	Cage culture	Aquaculture Station UAST/IPA	16.0 - 30.0	Male ^a	0.0139	3.117	0.985	Present work	101.6 ± 3.5	101.2 ± 0.8
Brazil	Ponds culture	Aquaculture Station UAST/IPA	16.0 - 30.1	Male ^a	0.0160	3.081	0.987	Present work	99.79 ± 3.5	99.3 ± 0.8
Ethiopia	Natural	Lake Tana	13.0 - 32.0	Unsexd	0.0424	2.740	0.941	Tadesse (1997)	118 ± 5.1	117 ± 0.7
Benin	Natural	Ouémé River Basin	7.5 - 26.5	Unsexd	0.0330	2.790	0.961	Lalèyè (2006)	126 ± 5.2	124 ± 0.8
urkina Faso	Natural	Hippopotamus Pond	9.7 – 22.1	Unsexd	0.0377	2.800	0.986	Béarez (2003)	110 ± 4.5	108 ± 0.7
Kenia	Natural	Lake Victoria	10.0 - 60.0	Male	0.0118 ^d	3.130 ^d	0.987 ^d	Njiru et al. (2006)	115 ± 3.9	114 ± 0.9
Bangladesh	Natural	Kaptai Lake	15.0 - 53.0	Unsexd	0.0366	2.840	0.980	Ahmed et al.(2003)	97 ± 3.9	96 ± 0.7
Nigeria	Natural	Kainji Lake	8.3 - 19.5	Unsexd	0.0169	3.130	0.958	Feu and Abiodun (1998)	79 ± 2.7	79 ± 0.6
Brazil	Cage culture	Fish Culture Association /Xingó	3.0 - 17.3	Male ^b	0.0272 ^d	2.840 ^d	0.096 ^d	Saraiva (2004)	132.7 ± 5.4	131.4 ± 0.9
Brazil	Cage culture	Fish Culture Station / UEL	3.6 - 18.8	Male	0.0195 ^d	3.040 ^d	0.984 ^d	Medri et al. (1999)	94.2 ± 3.4	93.7 ± 0.7
Nigeria	Ponds culture	Ijebu-Ode	5.5 - 11.4	Male	0.0093	3.140	uninformed	Olurin and Aderibigbe (2006)	140 ± 4.7	140 ± 1.1
Brazil	Ponds culture	Aquaculture Station / UFRPE	13.0 - 28.2	Male	0.0342	2.801	0.998	Carmo et al. (2008)	120.3 ± 5.0	119.0 ± 0.8
Brazil	Ponds culture	Aquaculture Station / UFRPE	13.0 - 28.0	Male ^b	0.0230	2.968	0.998	Carmo et al. (2008)	101.4 ± 3.8	100.7 ± 0.8
Brazil	Ponds culture	Aquaculture Station / UFRPE	13.0 - 28.1	Male ^c	0.0171	3.053	0.997	Carmo et al. (2008)	102.4 ± 3.7	101.9 ± 0.8
Brazil	Aquarium	Laboratory of Aquaculture/UFPE	uninformed	Male	0.0163 ^e	2.990 ^e	0.988 ^e	Leal et al. (2009)	129.0 ± 4.8	128.2 ± 1.0
Brazil	Aquarium	Laboratory of Aquaculture/UFPE	uninformed	Male	0.0132 ^f	3.116 ^f	0.995 ^f	Leal et al. (2009)	107.4 ± 3.7	107.0 ± 0.9

983 ^a Male Nile tilapia strain GAAT-1, ^bMale Nile tilapia strain Chitalada, ^c Male Nile tilapia red strain, ^d Geometric mean of a, b and R^2 , ^e Values of a, b
 984 and R^2 from Nile tilapia fed commercial diet, ^f Values of a, b and R^2 from Nile tilapia fed diet containing 5% shrimp protein hydrolysate.
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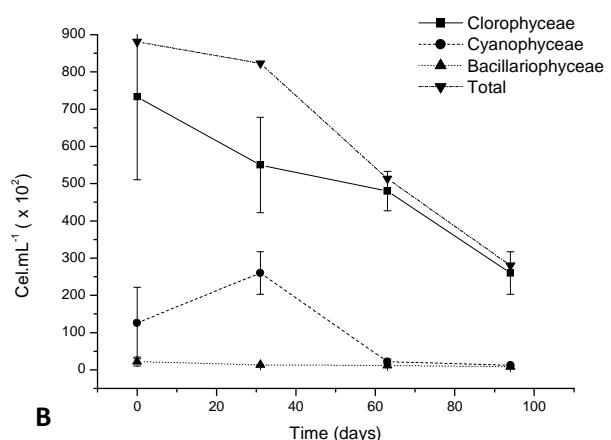
986 **8. Figures**

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994 Figure 1: Mean \pm SE of mean of the evolution of the abundance of phytoplankton ($\text{cel.mL}^{-1} \times 10^2$)
995 intensive (Figure 1A) and semi-intensive cultivation (Figure 1B) of Nile tilapia (*O. niloticus*).

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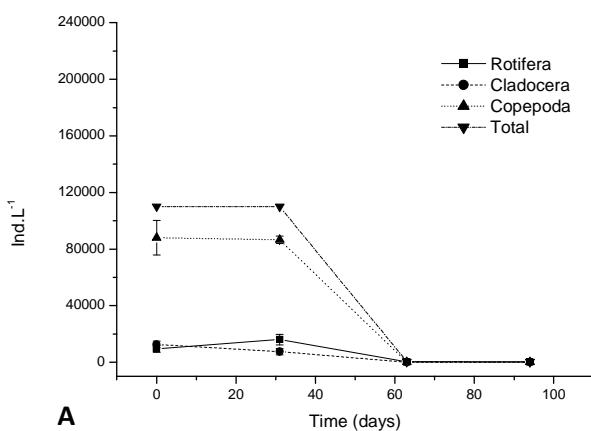
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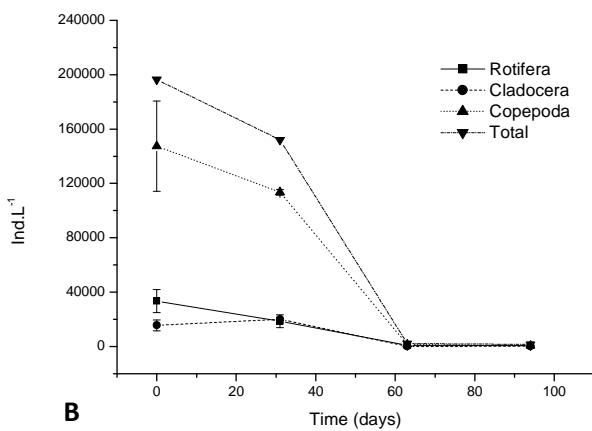
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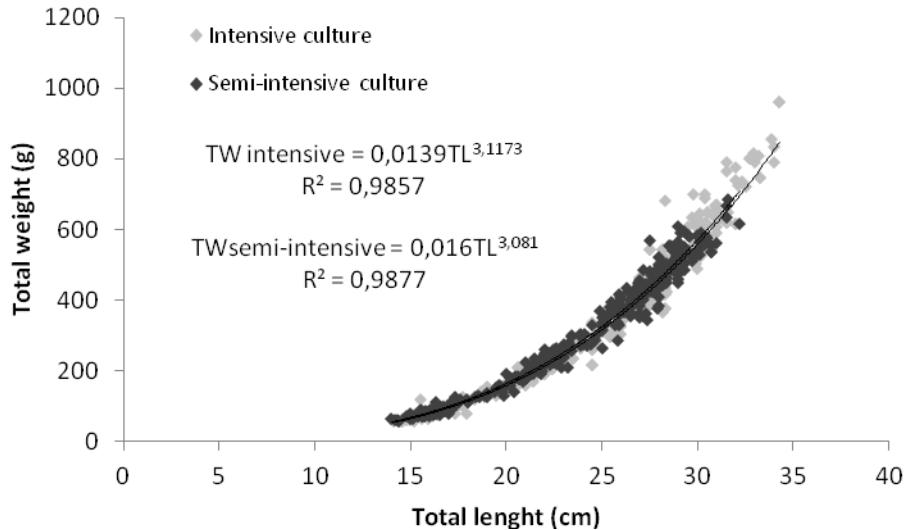
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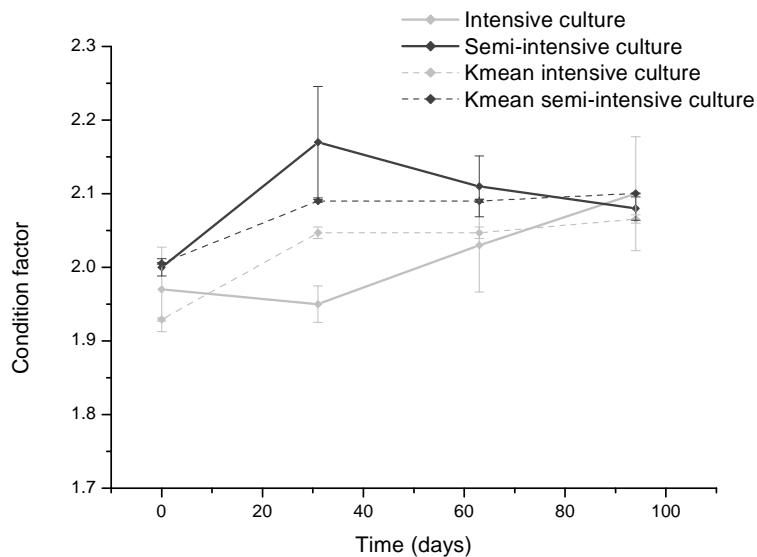
1005 Figure 2: Mean \pm SE of mean of the evolution of the abundance of zootoplankton (Ind.L^{-1}) intensive
1006 (Figure 2A) and semi-intensive cultivation (Figure 2B) of Nile tilapia (*O. niloticus*).



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1010 Figure 3: Parameters of the mathematical models of Nile tilapia ($W_t = a L_t^b$) adjusted to total
1011 weight-length data from fish under different culture systems.

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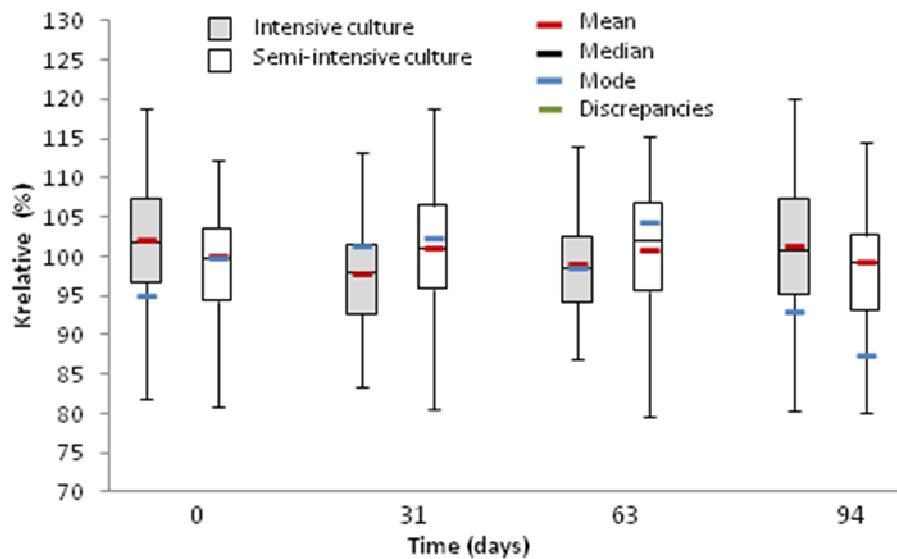
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1014 Figure 4: Mean of the evolution of Fulton's condition factor (K) of fish subjected to different
1015 cultures in comparison with the corresponding population K_{mean} .

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1028 Figure 5: Boxplot of the evolution in the values of $K_{\text{rel}}(\%)$ from fish under different cultures.
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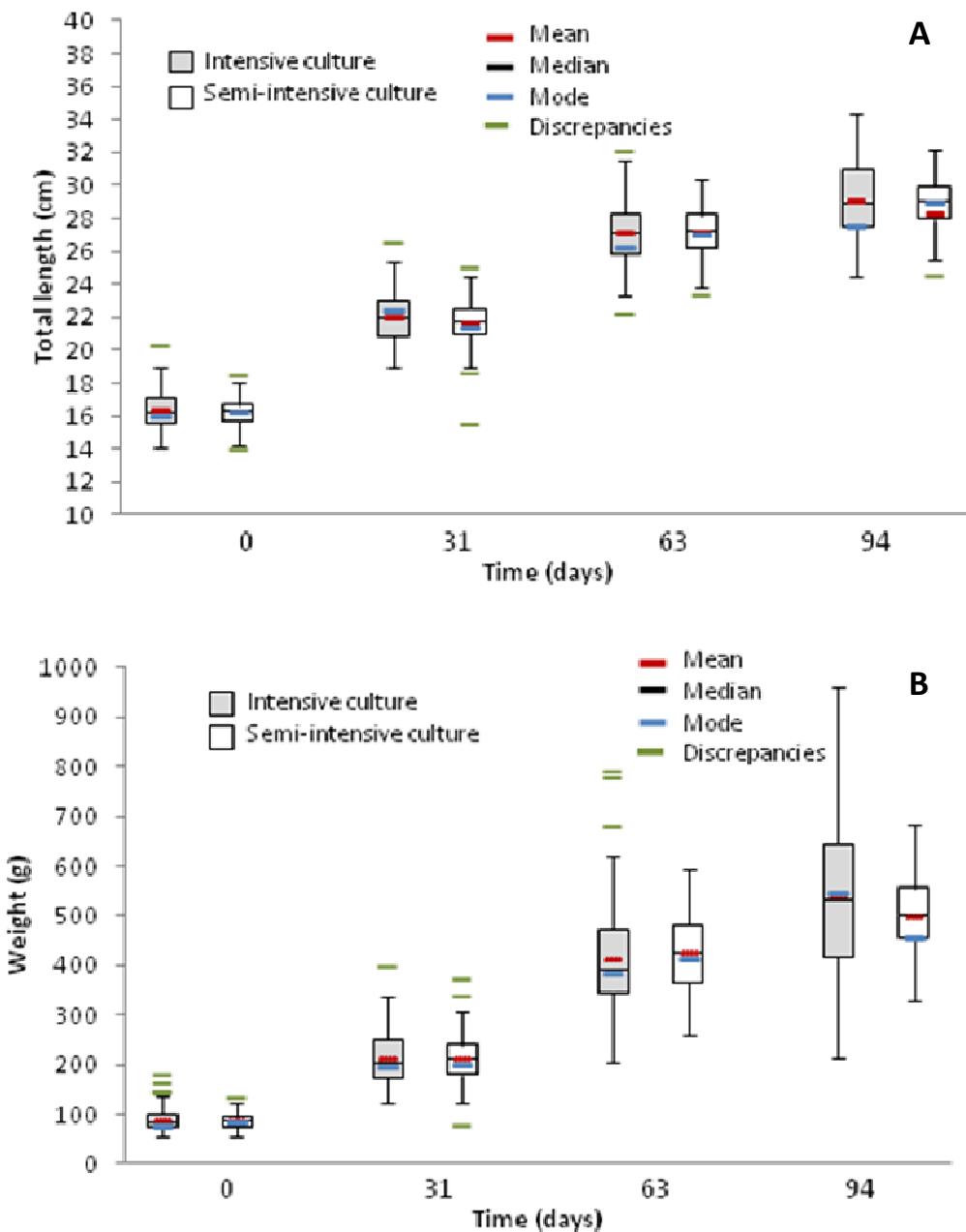


Figure 6: Boxplot of the evolution of Total length (cm) (Figure 6A) and Weight (g) (Figure 6B) of fish under different cultures.

7. CAPÍTULO III

ARTIGO CIENTÍFICO:

Enzymatic changes in the intestine of Nile tilapia (*Oreochromis niloticus* L.) on intensive and semi-intensive culture systems

ARTIGO A SER SUBMETIDO A REVISTA AQUACULTURE ISSN: 0044-8486



Enzymatic changes in the intestine of Nile tilapia (*Oreochromis niloticus* L.) under intensive and semi-intensive culture systems

4 Juliana Ferreira Santos^{1, 2}, Karollina Lopes Siqueira Soares¹, Caio Rodrigo Dias Assis¹, Carlos
5 Augusto Martins Guerra³, Daniel Lemos⁴, Luiz Bezerra Carvalho Jr¹, Ranilson Souza Bezerra^{1*}

² Universidade Federal Rural de Pernambuco, Unidade Acadêmica de Serra Talhada, Fazenda Saco,
S/N - Caixa Postal 063 CEP 56900-000 Serra Talhada - PE, Brazil.

³ Instituto Agronômico de Pernambuco (IPA), Fazenda Saco, CEP 56900-000 Serra Talhada - PE, Brazil.

¹⁴ Laboratório de Aqüicultura Marinha (LAM), Instituto Oceanográfico, Universidade de São Paulo,
¹⁵ Cidade Universitária, 05508-120 São Paulo-SP, Brazil.

16
17 Running title: Influence of culture systems on the digestive enzymes of tilapia

18 *Ranilson S. Bezerra.
19 Laboratório de Enzimologia – LABENZ, Departamento de Bioquímica, Universidade Federal de
20 Pernambuco. CEP 50670-420, Cidade Universitária, Recife-PE, Brazil,
21 Tel.: + 55-81-21268540; Fax: + 55-81-21268576
22 E-mail address: ransoube@uol.com.br

24 **Highlights**

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- 26 • Total proteolytic, amylase and lipase activity did not statistically differ between treatments
27 whereas trypsin and chymotrypsin activity increased with 31 and 63 days of culture.
- 28 • A positive correlation was observed between the recommended concentration of essential
29 amino acids for Nile tilapia and specific aminopeptidases activity.
- 30 • PAGE revealed 12 active proteolytic bands in both systems, however integrated density
31 values were higher in the bands of semi-intensive.
- 32 • Specimens of the intensive and semi-intensive exhibited five bands of amylolytic activity,
33 with higher integrated density at 31 days of cultivation for both systems.
- 34 • The two-dimensional electrophoresis of the intestine of cultured animals in intensive and
35 semi-intensive systems showed a total of 1,233 spots, among them 731 different.

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49 **Abstract**

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51 The effect of different culture systems on digestive enzyme activities of Nile tilapia was evaluated.
52 Juvenile Nile tilapia (87.61 ± 1.52 g) were cultured in semi-intensive and intensive system during 90
53 days. Two nutritional plans were adopted: triphasic for semi-intensive systems (diet with 35, 32 and
54 28% crude protein - CP) and biphasic for intensive system (35 and 32% CP). Biometric
55 measurements were monthly performed for adjustments in diet and removal of intestine tissues to
56 evaluate the performance of enzyme activities. Total proteolytic, amylase and lipase activity were
57 not statistically different between the treatments along of the periods analyzed (31, 63 and 94 days
58 of culture). However, trypsin and chymotrypsin activity were higher with 31 and 63 days of culture
59 in fish from semi-intensive system, suggesting that natural food may have influenced these
60 activities. A positive correlation was observed between the recommended concentration of essential
61 amino acids for Nile tilapia and specific aminopeptidases activity in fish intensive system.
62 Substrate-SDS-PAGE revealed 12 active proteolytic bands in both systems. However, integrated
63 density (*ID*) values were higher in the bands of semi-intensive. Specimens of the intensive and
64 semi-intensive exhibited five bands of amyloytic activity. Fish from intensive and semi-intensive
65 systems showed the highest values of *ID* with 31 days of cultivation. The lowest values of *ID* were
66 observed with fish from semi-intensive system with 63 and 94 days of cultivation. The comparison
67 of the proteomic profile of the intestine of cultured animals in intensive and semi-intensive system
68 showed a total of 1,233 spots, among them 731 different. It can be noticed that the natural food and
69 diets used greatly influenced the culture systems. The expression/activity of these proteins appears
70 to have a form of performance according to environmental and feeding conditions, so that the
71 digestive process can be as efficient as possible in accordance with the conditions of cultured
72 animals.

73 **Keywords:** Nile tilapia, culture systems, semi-intensive and intensive systems, digestive enzymes.

74 **1. Introduction**

75

76 In aquaculture, good rates of growth are essential to ensure the success of the production and
77 for such achievement, an efficient management becomes necessary. Inadequate management in
78 addition to ecologically unsustainable production can destabilize growth parameters directly
79 affecting production costs (El-Sayed, 2006).

80 The growth depends mainly on the food provided. However, the nutritional value of a diet is
81 not based simply on their chemical composition, but also the physiological ability of the fish to
82 digest and absorb, according to its feeding habits and nutritional requirements (Santos et al., *in*
83 *press*). According to Tengjaroenkul et al. (2000), the ability of fish to utilize ingested nutrients
84 depends on the presence of appropriate enzymes along of the digestive tract. The study of the
85 digestive enzymes in fish may clarify some aspects of their physiology and, therefore, can also
86 afford to solve some nutritional problems in fish feeding (Ribeiro et al., 1999). Ferron and Leggett
87 (1994) suggested that digestive enzymes can be interesting as condition indices due to their
88 specificity and sensitivity. Furthermore, according to Cara et al. (2007) its value as an indicator can
89 be justified if its activity clearly shows genetic variations (between different populations) or shows
90 a response to environmental changes.

91 Enzymatic activities appear to be related to age, feeding habits of fish and other factors
92 (Falcón-Hidalgo et al., 2011). Santos et al. (*in press*) suggest that Nile tilapia has the ability to adapt
93 to different food sources. For optimum utilization of all these sources, the enzymatic arsenal of
94 tilapia should be sufficiently diversified to digest the wide range of ingested food.

95 Enzymes have been studied over the last decades for their applications in nutritional
96 physiology and biochemistry for aquaculture (Tengjaroenkul et al. 2000; Muhlia-Almazan et al.,
97 2003; Buarque et al., 2009a; Buarque et al., 2009b; Santos et al., *in press*). However, changes in
98 enzyme activity resulting from management are not well elucidated. A better knowledge of the

99 enzymatic activity during growing is relevant for the understanding of fish nutritional needs
100 (Ribeiro et al., 1999). Since the growth is directly dependent on the absorption of the nutrients from
101 the diet, it is very important to attempt that the digestive enzymes may play a key role on the
102 nutrient incorporation. Therefore, investigations conducted during the cultivation, could provide
103 valuable information for a better understanding of the physiological processes associated with the
104 culture system employed.

105 Nile tilapia is often commercially cultivated in intensive or semi-intensive system, under
106 different stocking densities, management strategies and environmental conditions. Furthermore,
107 many features make them an ideal candidate for aquaculture, especially in developing countries (El-
108 Sayed, 2006). Thus, the aim of this work is to detect enzymatic changes in the intestine of Nile
109 tilapia cultured in the intensive and semi-intensive culture systems, relating these activities to the
110 production system adopted.

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112 **2. Materials and methods**

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114 **2.1 Culture conditions**

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116 The animals were cultivated at the Aquaculture Station of Unidade Acadêmica de Serra
117 Talhada (UAST) / Instituto Agronômico de Pernambuco (IPA) located in Serra Talhada, semi-arid
118 region of Pernambuco State, Brazil.

119 The experiment consisted of two treatments (intensive and semi-intensive cultures) and three
120 replicates. In the intensive treatment fish were cultured in cages, while in the semi-intensive they
121 were grown in ponds.

122 For the intensive culture, the animals were stocked at a density of 110 fish.m⁻³ (three
123 replicates), the cages were located in the reservoir Saco I - Serra Talhada – PE.

124 The semi-intensive culture was carried out using three ponds with an area of 338 m² and a
125 maximum depth of 1.5 m, supplied by water from the reservoir Saco I. The animals were stocked at
126 a density of 1 fish.m⁻² (0.67 fish.m⁻³). Before stocking, the ponds were dried in the sun light and
127 went through the process of liming (CaCO₃ - 15 kg per 1000 m²) and fertilization (bovine manure -
128 1 ton.ha⁻¹) according to Woynarovich and Horváth (1983) for the growth of natural food such as
129 phytoplankton and zooplankton.

130 The ponds and cages were stocked with juvenile sex-reversed Nile tilapia (variety
131 QAAT1/Arizona-USA) with an initial average weight of 88.6 ± 1.5g and an average total length of
132 16.3 ± 0.12cm. An aliquot of 10% of the populations were sampled monthly to monitor growth and
133 make feeding adjustments. Both cultures lasted 94 days.

134 Commercial diets containing 35, 32 and 28% crude protein (CP) were used (composition
135 described in Table 1). Intensive culture used nutritional biphasic plan (35 and 32% CP) and semi-
136 intensive triphasic plan (35, 32 and 28% CP), as described in Table 2. The fish were fed twice a day
137 (8 a.m. and 3 p.m.) with feeding rates of 5, 4 and 3% biomass, adjusted monthly after biometrics.

138 Water temperature, dissolved oxygen, pH, ammonia nitrogen, nitrite, nitrate, orthophosphate,
139 alkalinity and hardness were monitored and the average for intensive and semi-intensive system
140 were 29.40 ± 1.4 and 30.15 ± 2.6 °C (mean ± SD); 8.19 ± 1.9 and 6.67 ± 3.4 mg L⁻¹; 8.07 ± 0.2 and
141 7.15 ± 0.4; 0.065 ± 0.032 and 0.126 ± 0.061 mg L⁻¹; 0.053 ± 0.050 and 0.017 ± 0.013 mg L⁻¹; 0.001
142 ± 0.000 and 0.002 ± 0.001 mg L⁻¹; 0.016 ± 0.016 and 0.001 ± 0.000 mg L⁻¹; 13.95 ± 1.86 and
143 161.28 ± 106.97 mg L⁻¹; 20.06 ± 1.09 and 28.50 ± 4.37 mg L⁻¹, respectively. Data described in
144 Santos (2013).

145 At the end of the experiment, the main growth parameters were calculated from the
146 following equations:

147 Weight gain rate (WG_(g)) = BW_f - BW_i

148 Survival (S_(%)) = (Final number of fish/ fish stocked) x 100

149 Feed conversion ratio (FCR) = dry feed offered_(g)/wet weight gain_(g)
150 Protein efficiency ratio (PER) = wet weight gain_(g)/ protein fed_(g)
151 Feed efficiency (FE) = wet weight gain_(g)/dry feed offered_(g),
152 Fulton's condition factor (K): K = 100 W / L³,
153 in which BW_i and BW_f = average initial and final body weight_(g) of fish; W = total weight in
154 the times 0, 31, 63 and 94 days and L = total length at the same times. Data described in Santos
155 (2013).
156

157 **2.2 Extraction of digestive enzymes**

158
159 In the times 31, 63 and 94 days of cultivation, fish remained under fasting for 24 h and five
160 individuals from each replicate were sacrificed according to Pedrazzani et al. (2007) by sectioning
161 of medulla and bleeding by cutting the gills. Intestines were immediately collected and
162 homogenized (40 mg tissue.mL⁻¹) in 0.15 M NaCl, using a tissue homogenizer. The resulting
163 preparations were centrifuged at 10,000 x g for 10 min at 4 °C to remove cell debris and nuclei. The
164 supernatants (crude enzyme extracts) were frozen at -20 °C and used in further assays (Bezerra et
165 al., 2005).

166
167 **2.3 Protein concentration**
168

169 Protein concentration was determined according to Bradford (1976) using bovine serum
170 albumin (BSA) as the standard.

171
172
173

174

2.4 Enzyme assays

175

176

2.4.1 Total proteolytic activity

177

178

For total proteolytic activity 1% (w/v) azocasein was used as substrate according Bezerra et al. (2005). One unit (U) of enzyme activity was defined as the amount of enzyme capable of hydrolyzing azocasein to produce a 0.001 change in absorbance per minute.

181

182

2.4.2 Trypsin and chymotrypsin activity

183

184

Trypsin and chymotrypsin activity were determined using 8.0 mM BApNA ($\text{N}\alpha$ -benzoyl-DL-arginine-p-nitroanilide) and 8.0 mM SApNA (Suc-Ala-Ala-Pro-Phe p-nitroanilide) as substrate, respectively, according Bezerra et al. (2005). One unit (U) of activity was defined as the amount of enzyme required to produce 1 μmol *p*-nitroaniline. min^{-1} . Specific activity was expressed as $\text{U}.\text{mg}^{-1}$ of protein.

189

190

2.4.3 Aminopeptidase activity

191

192

Aminopeptidase activity was evaluated using 4.2 mM aminoacyl- β -naphthylamide as substrate. The aminoacyl moieties of substrates used were Leu, Arg, Ala, His, Tyr, Gly, Pro, Iso and Ser, as described in Buarque et al. (2009a). One unit of enzyme activity was defined as the amount of enzyme required to hydrolyse 1 μmol β -naphthylamine. min^{-1} . Specific activity was expressed as $\text{mU}.\text{mg}^{-1}$ of protein.

197

198

199

200 **2.4.4 Amylase activity**

201

202 Amylase activity was evaluated according to Bernfeld (1955) using 2% (w/v) starch as
203 substrate. One unit of enzyme activity was defined as the amount of enzyme required to liberate 1
204 μg of maltose per milligram of protein per min (specific activity).

205

206 **2.4.5 Lipase activity**

207

208 Lipase activity was assayed according to Aryee et al. (2007) using 8mM *p*-nitrophenyl
209 palmitate (*p*-NPP), as substrate. One unit of activity (U) was defined as the amount of enzyme that
210 catalyzed the release of 1 μmol of *p*-nitrophenol (*p*-NP) per min under the assay conditions. Specific
211 activity was expressed as U.mg⁻¹ of protein.

212

213 **2.5 Enzyme characterization in substrate-SDS-PAGE (zymograms)**

214

215 Zymogram of protease activity (aliquots of 50 μg of protein, *n* = 3) were carried out based
216 on Garcia-Carreño et al. (1993) after electrophoresis. The gels were immersed in 2.5 % Triton X-
217 100 dissolved in 0.1 M Tris–HCl buffer pH 8.0 to remove the SDS and incubated with 4 % casein
218 (w/v) in 0.1 M Tris–HCl buffer pH 8.0 for 30 min at 4 °C. The temperature was raised to 25 °C and
219 maintained for 90 min to allow the digestion of casein by the active fractions. Finally, the gels were
220 stained overnight for protein in 0.18 % (w/v) Coomassie Brilliant Blue R250 prepared in acetic acid
221 and methanol (10:25 % v/v), and the background of the gel was distained in acetic acid and
222 methanol (10:25 % v/v). Clear bands in blue background denoted protease bands by digestion of
223 casein substrate.

224 Zymogram of amylase activity ($n = 3$) were carried out according to the modified
225 methodology described by Castro et al. (2012). 30 µg of protein was applied to a 7.5% (w/v)
226 separating gel. Electrophoresis was performed at a constant current of 15 mA per gel at 4 °C for 150
227 minutes, for better visualization of the bands. The gels were immersed in 2.5 % Triton X-100
228 dissolved in 0.1 M Tris–HCl buffer pH 8.0 to remove the SDS and incubated with 2 % starch (w/v)
229 in 0.1 M Tris–HCl buffer pH 8.0, containing 1 mM CaCl₂ for 30 min at 37°C, to allow the
230 digestion of starch by the active fractions. Finally, the gels were washed with distilled water and
231 stained with an iodine/KI solution (15 and 12 mM) for 5 minutes. The gels were washed with
232 distilled water to stop the reaction and then with an acetic acid and methanol solution (10:25 % v/v).
233 Clear bands in black background denoted amylase bands by digestion of starch substrate.

234

235 **2.6 Densitometry of the bands**

236

237 Zymograms images ($n = 3$) were obtained and the grayscale intensity of the bands were
238 evaluated by densitometry using ImageJ software (Image Processing and Analysis in Java, USA).
239 Values were analyzed in integrated density (*ID*) which is the sum of the values of the pixels in the
240 image or selection. This is equivalent to the product of a given area and the respective mean gray
241 value (Rasband, 2012; Schneider et al., 2012).

242

243 **2.7 Two-dimensional electrophoresis**

244

245 **2.7.1 Extraction and protein quantification**

246

247 Samples of intestine tissues of Nile tilapia cultivated in the intensive and semi-intensive
248 systems were macerated in liquid nitrogen. 100 mg of tissue was added to 1 mL of Trizol

249 (Invitrogen[®]) and this material was incubated for five minutes. Then, after the addition of 200 µL of
250 100% chloroform, the material was incubated for 3 minutes. The samples were centrifuged at
251 12,000 x g for 20 minutes at 4 °C. 300 µL of cold 100% ethanol was added to the precipitate and
252 then, incubated for 3 minutes. The material was centrifuged again under the same conditions and
253 the supernatant was collected and added to 1 ml of cold 100% isopropanol, waiting 10 minutes at
254 27°C. The samples were centrifuged and the supernatant discarded. The pellet was washed three
255 times, at each wash was added 1 ml of hypochlorite solution of guanidine and 95% ethanol. The
256 pellet was left in the solution under stirring for 20 minutes at 27°C and then centrifuged at 8,000 x g
257 for five minutes at 4 °C. After the three washes 1 ml of ethanol was added to the pellet and a
258 stirring was performed and allowed to incubate for 20 minutes at room temperature. The material
259 was again centrifuged at 8,000 x g for 10 minutes at 4 °C. The supernatant was discarded and the
260 precipitate was dried for 10 minutes at room temperature. Then, the pellet was dissolved with
261 200µL of 1% SDS and 1X PBS (phosphate buffered saline) and incubated for two hours at 50 °C in
262 a water bath (Kirkland et al., 2006).

263 The protein quantification was performed using the 2D Quant Kit (GE Healthcare) according to
264 manufacturer's instructions.

265

266 **2.7.2 Precipitation of proteins**

267

268 The extracted proteins were precipitated using 20% trichloroacetic acid (TCA). An aliquot
269 containing 600 mg of protein (according to the previous quantification) was separated and added to
270 a volume three-fold higher of 20% TCA. This material remained at 4 °C for two hours and
271 subsequently centrifuged at 14,000 x g for 30 minutes at 4 ° C. The supernatant was discarded and
272 the centrifugate was washed twice with 1 ml of 100% cold acetone. Between the washes (after the

273 addition of acetone) the samples were incubated for 10 minutes and then centrifuged at 14,000 x g
274 for five minutes.

275

276 **2.7.3 Two-dimensional electrophoresis in polyacrylamide gels**

277

278 For two-dimensional electrophoresis was used methodology Kirkland et al. (2006), adapted to
279 our conditions.

280

281 **2.7.3.1 First Dimension - Isoelectric focusing**

282

283 After precipitation of proteins, 250 µl of sample buffer Dithiothreitol (DTT) and 1% ampholyte
284 (GE Healthcare) were added. 13 cm Strips (Immobiline DryStrip pH 3-10 - GE Healthcare) were
285 hydrated for 16 hours. The isoelectric focusing was performed in the Ettan IPGphor III (GE
286 Amersham Biosciences) for about 6 hours.

287

288 **2.7.3.2 Second Dimension - SDS-PAGE**

289

290 Before electrophoresis (SDS-PAGE) strips were equilibrated in equilibration buffer according to
291 Kirkland et al. (2006), in two phases: the first with equilibration buffer with addition of 1% DTT
292 (Dithiothreitol) and the second with the addition of 2.5% iodoacetamide, each phase with 20
293 minutes of incubation under stirring at room temperature. The strips were applied over a 12.5%
294 polyacrylamide concentration gel (16 x 18 cm height-width and 1 mm thick). The electrophoresis
295 was performed by applying two voltages: 15 mA for 30 minutes and after 45 mA until the end of
296 the run.

297 For staining, the gels were incubated with staining solution (20% methanol, 5% acetic acid and
298 0.2% Coomassie Blue) for two hours under stirring and at room temperature and destained with
299 solution (20 % methanol and 5% acetic acid) for 16 hours under stirring and also at room
300 temperature.

301

302 **2.8 Statistical analysis**

303

304 All data was tested for normal distribution (Shapiro-Wilk test) and homogeneity of variances
305 (Bartlett's test). Student's *t*-test and One-way analysis of variance (ANOVA) followed Tukey tests
306 were used for normally distributed data, whereas Wilcoxon-Mann-Whitney and Kruskal-Wallis test
307 were used for non-normally distributed data. Correlations were tested using Pearson's Product
308 Moment Correlation following regression analysis. Densitometry to quantify the bands in gels was
309 done with the software ImageJ from imagej.nih.gov/ij/. Differences were reported as statistically
310 significant when $P < 0.05$, using OriginTM version 8.0, Assistat 7.6 and GraphPad Prism 6.0.

311 The two-dimensional gels were subjected to analysis with Platinum 2D ImageMaster software
312 (Amersham Biosciences) for the detection of quantitative and qualitative differences of proteins
313 between treatments ($n = 3$).

314

315 **3. Results**

316

317 The proximate analysis of the commercial diets used in the experiment and nutritional plan
318 is shown in Tables 1 and 2, respectively. The main growth parameters calculated for Nile tilapia can
319 be seen in Table 3. There was no statistical difference between the final weight, weight gain and
320 feed conversion ratio of experimental specimens. However, the best values of survival, protein
321 efficiency ratio and feed efficiency were observed for fish in semi-intensive system. The condition

322 factor found for Nile tilapia systems in semi-intensive and intensive during cultivation can be seen
323 in Table 4. At 31 days of cultivation statistical difference was observed between the systems,
324 reaching the best value for semi-intensive system (Santos, 2013).

325 Figure 1 shows the alkaline proteolytic activity of Nile tilapia cultivated under intensive and
326 semi-intensive systems. Total proteolytic activity (Figure 1A) was not statistically different between
327 the treatments ($P \geq 0.05$) along of the periods analyzed (31, 63 and 94 days of culture). However,
328 trypsin and chymotrypsin activity (Figures 1B and 1C) were higher with 31 and 63 days of culture
329 ($P < 0.05$) in fish from semi-intensive system, but no statistical difference between the treatments
330 was observed for these enzymes at 94 days of cultivation. Specific aminopeptidase activities (Figure
331 2) were measured using the following substrates: Leu-, Arg-, Tyr-, Gly-, Ser-, Ala-, His-, Iso- and
332 Pro- β -naphthylamide (Figure 2A-I, respectively). After 31 days of culture, most activities of
333 aminopeptidases showed no statistical difference except for Pro- and Iso- that showed higher
334 activity in fish of semi-intensive and intensive systems, respectively. After 63 days of cultivation
335 Iso- showed no statistic difference between treatments, the other enzymes showed higher values in
336 intensive system. At 94 days of culture Leu-, Arg-, Tyr-, Gly-, Ser- and Pro- aminopeptidases
337 remained higher in the intensive system, while Ala-, His-, and Iso- showed no statistical difference
338 between the systems. There were no significant differences in amylase (Figure 3) and lipase activity
339 (Figure 4) in the crude extract of the intestine of Nile tilapia cultured in intensive and semi-intensive
340 systems. A correlation was observed between the recommended concentration of essential amino
341 acids for Nile tilapia (Fagbenro, 2000; Celik, 2012) and specific aminopeptidases activity in fish
342 intensive system, using aminoacyl- β -naphthylamide as substrates (Figure 5).

343 Proteolytic activities in fish from the intensive and semi-intensive system were detected in
344 substrate (casein) gel electrophoresis (Figure 6A). Intensive and semi-intensive system specimens
345 exhibited twelve intense activity bands (P1 to P12) with 31, 63 and 94 days of culture. Evaluating
346 the integrated density (*ID*) of the caseinolytic bands of proteases during the cultivations, the semi-

347 intensive system showed constant values for the times analyzed, while in intensive system the value
348 decreased at 63 days, increasing again at 94 days of cultivation (Figure 6B). The Figure 6C can be
349 seen the intensity of pixels values of the bands along the lanes of zymogram of farmed fish in the
350 intensive (subfigure 1, 2 and 3) and semi-intensive (subfigure 4, 5 and 6) system. Images can
351 confirm, with increasing intensity of the pixels values, the twelve bands observed in Figure 6A. In
352 their respective subfigures are integrated density values (*ID*) of the bands in cultivation systems at
353 the time periods analyzed (corresponding lanes of Figure 6A). Statistical differences were observed
354 between the cultures. Fish from the semi-intensive system showed the highest values of *ID* at the
355 three times (31, 63 and 94 days of cultivation) corresponding to 228.9 ± 6.96 , 229.27 ± 7.12 and
356 231.10 ± 4.93 respectively. The lowest value of *ID* was observed for fish from the intensive system
357 with 63 days of cultivation (183.17 ± 1.72).

358 Table 5 shows *ID* related to the caseinolytic bands from intestine of Nile tilapia cultured in
359 the intensive and semi-intensive systems. The bands followed the profile found by Santos et al. (*in*
360 *press*). Therefore, the previously results were employed to identify the caseinolytic bands found in
361 this study. The bands with trypsin / aminopeptidase activity (P3, P4, P6, P7 and P8) showed the
362 following results: P3 showed the highest *ID* values in semi-intensive system at 63 and 94 culture
363 days (19.84 ± 0.57 and 19.37 ± 0.43 , respectively). However, no statistically significant difference
364 was observed between *ID* values in the semi-intensive with 31 days and in the intensive with 31 and
365 94 days of culture. P4 also showed a higher value in the semi-intensive system with 63 days of
366 culture (16.51 ± 0.51). Nevertheless, this value did not statistically differ with 94 days of culture
367 (15.29 ± 0.60). P6 and P7 bands presented a higher *ID* in semi-intensive system in the three times
368 studied. P8 showed a higher value in this system at 94 days of culture (13.80 ± 0.20), but this value
369 did not differ statistically from the times 31 and 63 days on this same system, and was also equal to
370 the intensive system with 31 days of culture. The P5 band (aminopeptidase activity) showed higher
371 values in fish from semi-intensive system at 94 days of culture (28.36 ± 1.40), but this value did not

372 statistically differ from 63 days of culture (26.91 ± 1.12). P9 showed higher aminopeptidase activity
373 in intensive system with 94 days of culture (11.23 ± 0.21), but was not statistically different from the
374 semi-intensive system in the three times studied. The P11 band with trypsin/chymotrypsin activity
375 presented a higher value in fish from the semi-intensive system with 31 days of culture
376 (17.03 ± 0.89). The bands P1 and P10 demonstrated higher values of *ID* with 31 days of culture in
377 semi-intensive system (4.60 ± 0.37 and 18.32 ± 0.78 , respectively). The P2 band showed no
378 statistical difference in most times analyzed. Finally, the band P12 showed a higher value of *ID* at
379 63 days of culture in semi-intensive system (12.80 ± 0.74), but showed no significant difference
380 compared to days 31 and 94 and neither differed from intensive system at 94 days.

381 Amylase activities in fish from the intensive and semi-intensive systems were detected in
382 substrate (starch) gel electrophoresis (Figure 7A). Intensive and semi-intensive system specimens
383 exhibited five activity bands (A1 to A5) related to 31, 63 and 94 days of culture. The integrated
384 density (*ID*) of the amylase bands during the cultivations (Figure 7B), showed a similar trend at the
385 times analyzed. At 31 days *ID* values were higher in both systems and decreased at 63 and 96 days
386 of cultivation. In the Figure 7C can be seen the intensity of pixels of the bands along the lanes of
387 zymogram of farmed fish in the intensive (subfigure 1, 2 and 3) and semi-intensive (subfigure 4, 5
388 and 6) system. The images also confirm the five bands observed in Figure 7A. Integrated density
389 values (*ID*) in their respective subfigures showed a statistical difference between the systems. Fish
390 from intensive and semi-intensive systems showed the highest values of *ID* with 31 days of
391 cultivation (95.95 ± 8.63 and 88.24 ± 12.65 respectively). The lowest values of *ID* were observed with
392 fish from semi-intensive system with 63 and 94 days of cultivation (60.71 ± 6.57 and 58.49 ± 5.99
393 respectively).

394 Table 6 shows *ID* values related to the bands of amylase zymogram from intestine of Nile
395 tilapia cultured in the intensive and semi-intensive systems. Amylase 1 (A1) showed the highest
396 values for intensive and semi-intensive system with 31 days of culture (20.15 ± 1.40 and 18.92 ± 2.76 ,

397 respectively). The highest value for amylase 2 (A2) was observed for the intensive system with 31
398 days (22.05 ± 0.62), but did not differ statistically from the value of 94 days of cultivation for this
399 system and neither from semi-intensive with 31 days. Amylase 3 (A3) showed the highest values
400 for intensive system with 31 days (20.40 ± 1.86). However, no statistical difference was observed
401 with 63 and 94 days of cultivation and also for the semi-intensive system with 31 days. The highest
402 integrated density for Amylase 4 (A4) was also observed for fish in intensive system (6.12 ± 0.42)
403 with 31 days, being statistically equal to the values obtained with 63 and 94 days of culture. In the
404 evaluation of Amylase 5 (A5), the intensive system showed the highest values for 31 and 63 days of
405 cultivation (3.70 ± 0.52 and 3.71 ± 0.45 , respectively).

406 The two-dimensional electrophoresis (Figures 8 and 9) showed well-defined spots without
407 trawling (horizontal and vertical) that could prevent detection. The proteins are distributed
408 throughout the pH range of 3-10 and the molecular weights between 10 and 100 kDa.

409 The comparison of the proteomic profile of the intestine of animals cultured in intensive
410 (Figure 8) and semi-intensive system (Figure 9) showed a total of 1,233 spots, 731 being different.

411

412 **4. Discussion**

413

414 Liti et al. (2005) showed that an inefficient management can promote reduction in weight gain
415 in tilapia. However, animals can demonstrate some physiological compensation, thus the quality of
416 food and adopted management may contribute for this kind of adjustment. Despite of the similarity
417 observed between the final weight of fish cultured in both systems, the food and protein efficiency
418 were better for animals cultured on semi-intensive system. Although it was observed a decrease in
419 condition of fish from intensive system, experimental specimens reached similar final weight, as a
420 way of adapting to the culture system employed (Santos, 2013). The digestive physiology of Nile
421 tilapia may have contributed to this fact. Muhlia-Almazan et al. (2003) describe that changes in diet

422 of aquatic organisms can cause alterations in the expression and activity of digestive enzymes.
423 Consequently, changes in management may induce animals physiological adaptations to maintain or
424 achieve satisfactory levels of specimens nutritional status.

425 The Nile tilapia is considered an opportunistic omnivore species, able to assimilate nutrients
426 from different sources such as benthic algae, phytoplankton, macrophytes, zooplankton, small
427 invertebrates, detritus and cyanobacteria. The choice of feeding is primarily related to food
428 availability in the environment, indicating that this species has high trophic plasticity (Bowen,
429 1982; Philippart and Ruwet, 1982; Fitzsimmons, 1997; Beveridge and Baird, 2000; Lowe-
430 McConnell, 2000; Sampaio and Goulart, 2011). Therefore, the enzymatic arsenal of tilapia should
431 be sufficiently diversified to digest the wide range of food sources.

432 This fact was taken into account to evaluate the digestive enzymes of Nile tilapia under
433 different cultures. This study demonstrated a great diversity in the gut enzymes of Nile tilapia with
434 the presence of trypsin, chymotrypsin, aminopeptidase, amylase and lipase. All substrates generated
435 typical products of their respective reactions, however differences were detected in enzyme activity
436 in the cultures analyzed. Total proteolytic activity remained constant as seen in Figure 1A. The
437 different managements adopted were insufficient to cause detectable changes using azocasein as
438 substrate. Buarque et al. (2009a), comparing total proteolytic activity between two stages of
439 subadult shrimp (*Farfantepenaeus subtilis*), also found no statistical difference using the same
440 substrate. Nevertheless, changes could be seen with the use of specific substrates.

441 Trypsin and chymotrypsin showed higher values in fish from semi-intensive system when
442 compared to intensive system at the initial time (31 and 63 days of culture) even with a decrease in
443 concentration of crude protein in the feed of semi-intensive system. In fact, this system is
444 characterized by high natural food, especially at the beginning of cultivation (El Sayed, 2006).
445 According to Liti et al. (2006) when tilapia feeding is supplemented with low protein concentrations
446 in semi-intensive farming, the deficit is supplied by natural food. Luz and Zaniboni-Filho (2001)

447 emphasize that natural food has high nutritional value, and according to Kuz'mina and Golovanova
448 (2004) one of the major advantages of this diet is the presence of a high concentration of digestive
449 enzymes. França et al. (2010) also cite the importance of enzymes in the digestive processes of fish,
450 suggesting its presence and functional performance in the live food. Lauff and Hofer (1984) found
451 higher trypsin activity in *Coregonus sp.* larvae fed with *Monia sp.* Abolfathi et al. (2012) also
452 mention that the large proteolytic activity in omnivorous fish may be related to the need to digest
453 plant foods.

454 At 94 days of culture there was no statistical difference between the systems (activity of
455 trypsin and chymotrypsin), it is worth highlighting that the fish were in the intensive system with a
456 higher percentage of CP in the diet compared to semi-intensive and this fact may have contributed
457 to the increased enzyme activity in this system concomitantly with a possible decrease in the
458 concentrations of natural food in the semi-intensive system. Protein levels above 25% in the diet
459 induced proteolytic activity in *Labeo rohita* fingerlings (Debnath et al., 2007). Perez-Jimenez et al.
460 (2009) indicate that, independent of eating habits, the response of the digestive system of the fish
461 closely correlates with diet.

462 However, it should be noted that the activity of chymotrypsin showed an increase at 94 days of
463 cultivation for both systems, despite lower concentrations of CP in the diet and natural food
464 decrease. Some authors indicate trypsin, trypsin/chymotrypsin or chymotrypsin as nutritional
465 indicators in fish, commenting that its increase may occur to compensate for dietary deficiencies
466 (Cara et al. 2007). Papoutsoglou and Lyndon (2006) also observed an increase in the activities of
467 trypsin and chymotrypsin when there was a decrease in the concentration of protein in the diet for
468 *Anarhichas minor*. Corroborating this fact, Kumar et al. (2007) suggest that herbivorous fish can
469 compensate a low protein diet increasing its enzymatic activity. The authors mentioned that some
470 herbivorous fish have similar or higher trypsin activities when compared to carnivorous species,
471 possibly to maximize the efficiency of protein digestion. This can be extended for omnivorous fish

472 like tilapia, mainly because this species is characterized as opportunistic omnivorous (Bowen, 1982;
473 Ruwet and Philippart, 1982; Fitzsimmons, 1997; Beveridge and Baird, 2000; Lowe-McConnell,
474 2000; Goulart and Sampaio, 2011).

475 According to Wilson (2002) the fish have no nutritional requirement of protein *per se*, but
476 need minimum quantities and proportions of essential and non-essential amino acids in the diet for
477 the deposition of muscle protein and other body proteins. After hydrolysis of proteins by proteases
478 into long chain peptides, aminopeptidases further degrade into smaller peptides and free amino
479 acids, increasing the absorption of these nutrients (Buarque et al. 2009b). According to Zambonino-
480 Infante and Cahu (2007), activity of peptidases facilitates assimilation of amino acids in marine fish
481 larvae. Thus, understanding the regulation of aminopeptidases becomes quite important from the
482 nutritional and physiological point of view. Intestinal aminopeptidases of Nile Tilapia were capable
483 of hydrolyzing substrates leucine, arginine, tyrosine, glycine, serine, alanine, histidine, isoleucine
484 and proline β -naphthylamide. However, the activities of these enzymes differed between farming
485 systems, not showing a logical relationship, except for His-, Iso-, Leu-, Arg- aminopeptidase that
486 showed a significant correlation with the requirement of amino acids, based on the ideal protein
487 concept for Nile tilapia (Fagbenro, 2000; Celik, 2012), suggesting that the activity of these enzymes
488 has a relationship with essential amino acid requirement for this species. From physiological and
489 productive perspectives this fact is very important for aquaculture, as it may provide subsidies for
490 works regarding digestibility and nutritional requirements for other species. Buarque et al. (2009a;
491 2009b) also found a correlation between the aminopeptidases of *Farfantepenaeus subtilis* and *F.*
492 *paulensis* and their respective amino acid requirements.

493 Most aminopeptidases analyzed showed no difference between systems in beginning of the
494 cultivation, except Pro- and Iso- aminopeptidase. However, with 63 and 94 days of culture the
495 majority of these enzymes had higher activity in fish intensive system. In intensive system was not
496 present abundant natural food (less exogenous enzymes), this fact may have contributed to different

497 regulation of these enzymes in this system and the nutrients ingested by the fish need to be better
498 utilized since the contribution of natural food is minimal. In order to assess the function of digestive
499 enzymes in *Salmo salar*, Refstie et al. (2006) fed the animals with different concentrations of
500 protein made from yellow or narrow leafed lupins and verified no statistical difference in Leu-
501 aminopeptidase activity. Moreover, Ezquerra et al. (1999) observed an increase in Met-, Val-, Pro-,
502 Lys- and Leu- aminopeptidase activity in white shrimp fed with menhaden fish meal although this
503 was not in accordance with the nutritional standards required for this species. The authors also
504 mention that when the shrimps were fed with soybean meal, the activities of these enzymes
505 decreased mainly Gly- and Met- aminopeptidase. Methionine is usually the first limiting amino acid
506 in diets based on soybean protein to fish (Furuya et al. 2001).

507 Several animal and vegetable foods can be used in feed for aquatic organisms, partially or
508 totally replacing fishmeal traditionally used in aquaculture feed. Thus, several studies have been
509 conducted to evaluate new protein sources that could be used in the manufacture of feed (Jackson et
510 al., 1982; Gaber, 1996; Olvera-Novoa et al., 1997; Olvera-Novoa et al., 2002 ; Jatomea-Plascencia
511 et al., 2002, El-Saidy and Gaber, 2003; Gaber, 2006, Leal et al., 2009). Thus, in diets for Nile
512 tilapia, several sources can be used in the manufacture of feed, due to its ability to digest proteins
513 from plant and animal origin, therefore contributing to changes in enzymatic profile.

514 In order to observe the gut proteases of Nile tilapia under different culture conditions, more
515 sensitive tools were used to detect possible changes in the activities. According to Santos et al. (*in*
516 *press*) and Lemos et al. (2000) zymogram can be considered a more sensitive technique than
517 conventional quantitative analyzes. In fact, this technique allowed a good view of the behavior of
518 proteases from Nile tilapia, especially when the bands were analyzed using the intensity of pixel
519 values (integrated density - *ID*). Corroborating the data observed in quantitative analysis, the values
520 of *ID* observed for Nile tilapia were higher at all times in semi-intensive system. Making a
521 comparison of these results with the condition factor, both remained constant for these specimens.

522 The fish of intensive system showed the lowest values of *ID* and also had a decrease in the
523 condition with 31 days of cultivation. This suggests that the adopted management can interfere with
524 the condition/enzymatic activity of the experimental specimens. However, Nile tilapia through his
525 great capacity of adaptation, overcome the changes in management as evidenced by results of
526 growth. But it is noteworthy the importance of natural food in semi-intensive cultures, since
527 zooplankton can assist in the digestion process through their digestive enzymes, either by autolysis
528 or as zymogens that activate endogenous digestive enzymes (Kolkovsk, 2001). Furthermore, in
529 terms of cost/benefit, the use of natural foods in farm ponds is one of the most efficient ways to
530 reduce costs and environmental impacts of aquaculture (Beyruth et al., 2004). Its continuous
531 availability allows the fish to adjust the intake and nutritional supply to their metabolic needs. The
532 increase in feed frequency implies in more regular nutrient supply, increasing the efficiency of
533 digestion and metabolism of nutrients (Beyruth et al., 2004).

534 Santos et al. (*in press*) identified eight out of twelve proteases found in the intestine of Nile
535 tilapia. Based on this information, was assessed the integrated density of bands in intensive and
536 semi-intensive farming systems. Several bands showed similarity between the systems,
537 demonstrating the sensitivity of the method. Most enzymes from fish of semi-intensive system
538 presented higher *ID*, including enzymes with aminopeptidase activity (not shown in the quantitative
539 analyzes, excepting Pro- aminopeptidase activity with 31 days of cultivation). However, some
540 identifications cited in Santos et al. (*in press*) were not conclusive, therefore the authors suggest that
541 the Nile tilapia enzymes may have low compatibility with mammalian protease inhibitors.

542 In quantitative terms, amylase and lipase showed no statistical difference between the systems
543 (Figures 3 and 4, respectively). Bowyer et al. (2012) report that the lipase activity of *Seriola lalandi*
544 was statistically lower in fish fed with canola oil.

545 Through the amylase zymograms some differences in enzyme activity could be detected.
546 Figure 7B shows a similar trend of activity for the two culture systems, where the values of *ID*

547 decreased with the course of time. The intensive system showed higher values of integrated density,
548 though these values showed no statistical difference when compared to the mean of semi-intensive
549 system with 31 days (Figure 7C). Evaluating the bands separately, most showed higher values of *ID*
550 in intensive system (Table 6), excepting amylases 1, 2 and 3 of the semi-intensive system with 31
551 days of cultivation.

552 According to Le Moullac and VanWormhoudt (1994) amylase activity increases with the
553 increase of carbohydrates in the diet. Kuz'mina et al. (1996) also found a correlation between
554 amylase activity and concentration of carbohydrates in the diet for freshwater fish. This leads to a
555 very important point: the carbohydrates can be digested as they may have a positive relationship
556 with amylases. This may reflect the satiation of animals harming their growth. In fact the condition
557 of the animals in intensive system was minor than that of the semi-intensive with 31 days of culture,
558 suggesting a better energy: protein ratio for the semi-intensive system at this time. However,
559 according to Ceccaldi (1997) when optimal levels of dietary carbohydrates are overcome, the
560 enzyme activity tends to decrease. Nakamura (1970) and Reed et al. (2010) also reported the
561 inhibition of amylases by their own substrates. Sands and Lukes (1974) also mention that in some
562 microorganisms, the carbohydrate may even prevent expression of the enzyme. Thus, increases in
563 the carbohydrate diet may have caused a regulation/inhibition of enzyme activity. In addition, the
564 fact of amyloytic activities were slightly higher in the intensive system can be related to results
565 obtained by Ribeiro et al. (2002). These authors evaluated the enzymatic activities of *Solea*
566 *senegalensis* fed with live food and diet observed that amylase secretion was significantly higher in
567 the treatment with diet. In fact, the use of natural food for fish farmed in cages is more difficult than
568 in cultured fish in ponds due to density, fish movement, etc. Corroborating the results, Fountoulaki
569 et al. (2005) showed that differences in enzyme profiles are related to the nutrients present in feed.

570 These differences could also be confirmed by the analysis of proteomic profile of culture
571 systems, demonstrating how the environment/diet can promote changes in protein profile.

572 Moreover, in semi-intensive system there is a greater occurrence of microorganisms in the water.
573 Such microorganisms besides serving as food for fish, can also form symbiotic relationships with
574 them. Therefore, it is possible the existence of proteins belonging to these microorganisms in the
575 digestive tract, which could not be present in fish intensive system. Lucitt (2008) compared
576 proteomic profiles of two different embryonic developmental stages of zebrafish using two-
577 dimensional electrophoresis and found more than 660 proteins, being 477 different. The level of
578 changes that may occur in the proteomic profile of a species is very variable. Gómez-Requeni
579 (2011) subjected the zebrafish feeding unbalanced in only one amino acid (lysine) and the
580 proteomic profile of the body showed 207 proteins in total, being 45 significantly different.

581 Following the advances of technology in the areas of genomics and proteomics, the study of
582 the fish physiology has increased, making possible to analyze the effects of stress on these animals
583 (Parrington et al., 2002). Thus, not only the food, but also the environment in which animals are
584 exposed may cause quantitative and qualitative changes in global protein expression of a species.

585

586 **5. Conclusion**

587

588 The complexity of digestive functions allows the animal to modify the expression/activity of
589 the enzymes to achieve more efficiently the reaction products necessary for their metabolism. This
590 fact could be observed in both production systems through adaptations made for Nile tilapia. Thus,
591 the expression of these proteins appears to have a performance according to environmental and
592 feeding conditions, so that the digestive process can be as efficient as possible in accordance with
593 the conditions of cultured animals.

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598

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867 **7. Tables**

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869 Table 1: Proximate analysis of the commercial diets used in the experiment.

Proximate analysis (as-fed basis, g . kg ⁻¹)*	Crude Protein		
	35%	32%	28%
Pellet diameter	3 - 4	4 - 6	6 - 8
Moisture	130	130	130
Crude protein	350	320	280
Ether extract	65	50	50
Crude fiber	50	50	50
Ash	120	120	120
Calcium	15	15	15
Phosphorus	6	6	6
Vitamin C	0.5	0.5	0.5

870 *Data from manufacturer

871

872 Table 2: Nutritional plan used as feed for Nile tilapia (*O. niloticus*) subjected to different culture
873 systems.

Time (days)	Nutrition plan	
	Intensive	Semi-intensive
	culture	culture
0 – 31	35% CP	35% CP
32 – 63	35% CP	32% CP
64 – 94	32% CP	28% CP

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879 Table 3: Growth performance and nutrient utilization of Nile tilapia (*O. niloticus*) cultivated in
 880 intensive and semi-intensive system.

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Parameters	Intensive culture	Semi-intensive culture
Initial weight (g)	88.33 ± 1.54 ^a	86.90 ± 1.50 ^a
Final weight (g)	543.58 ± 48.11 ^a	501.74 ± 4.37 ^a
WG (g) ¹	455.24 ± 41.68 ^a	414.84 ± 3.48 ^a
Survival (%)	75.0 ^b	85.0 ^a
FCR ²	1.59 ± 0.12 ^a	1.57 ± 0.11 ^a
PER ³	4.23 ± 0.37 ^b	6.51 ± 0.30 ^a
FE ⁴	1.6±0.02 ^b	2.30±0.11 ^a

882 Values are mean ± SE. Within a row, means with different superscript letters differ significantly (P
 883 < 0.05) by Wilcoxon-Mann-Whitney test and Student's *t* test

884 ¹ Weight gain, ² Feed conversion ratio, ³ Protein efficiency ratio, ⁴ Feed efficiency.
 885 Data described in Santos (2013).

886

887 Table 4: Fulton's condition factor of Nile tilapia cultivated in intensive and semi-intensive systems
 888 at times "0" (zero), 31, 63 and 94 days.

889

Time (days)	Intensive culture	Semi-intensive culture
0	1.97±0.06 ^a	2.00±0.01 ^a
31	1.95±0.02 ^b	2.17±0.07 ^a
63	2.03±0.06 ^a	2.11±0.04 ^a
94	2.10±0.08 ^a	2.08±0.01 ^a

890 Values are mean ± SE. Within a row, means with different superscript letters differ significantly (P
 891 < 0.05) by Student's *t* test.

892 Data described in Santos (2013).

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895 Table 5: Integrated density (*ID*) of the bands of protease zymograms from the intestine of Nile
 896 tilapia (*O. niloticus*) cultured in intensive and semi-intensive systems.

Bands	Intensive culture			Semi-intensive culture			Enzyme*
	31 days	63 days	94 days	31 days	63 days	94 days	
P1	2.98±0.24 ^b	3.48±0.56 ^c	4.29±0.48 ^b	4.60±0.37 ^a	4.39±0.25 ^b	4.67±0.39 ^b	Not identified ₁ **
P2	15.71±0.39 ^{ab}	15.88±0.71 ^{ab}	15.15±0.19 ^b	15.70±0.36 ^{ab}	16.24±0.35 ^{ab}	16.34±0.38 ^a	Not identified ₂ **
P3	18.57±0.15 ^{ab}	17.21±0.36 ^b	18.71±1.24 ^{ab}	18.89±0.73 ^{ab}	19.84±0.57 ^a	19.37±0.43 ^a	Trypsin/Aminopeptidase
P4	13.77±0.58 ^{bc}	12.27±0.60 ^c	14.87±0.32 ^b	13.26±0.76 ^c	16.51±0.51 ^a	15.29±0.60 ^{ab}	Trypsin/Aminopeptidase
P5	23.43±0.63 ^c	22.68±0.44 ^c	24.19±0.89 ^c	24.91±1.06 ^{bc}	26.91±1.12 ^{ab}	28.36±1.40 ^a	Aminopeptidase
P6	7.90±0.09 ^b	6.87±0.20 ^c	8.34±0.19 ^b	9.26±0.50 ^a	9.29±0.11 ^a	9.81±0.51 ^a	Trypsin/Aminopeptidase
P7	18.81±0.35 ^c	17.96±0.44 ^d	21.04±1.21 ^b	21.70±0.73 ^a	21.72±0.74 ^a	21.77±0.51 ^a	Trypsin/Aminopeptidase
P8	13.11±0.08 ^{ab}	11.04±0.22 ^c	12.17±0.42 ^{bc}	12.95±0.81 ^{ab}	13.01±0.96 ^{ab}	13.80±0.20 ^a	Trypsin/Aminopeptidase
P9	9.82±0.40 ^{bc}	9.42±0.22 ^c	11.23±0.21 ^a	10.88±0.67 ^{ab}	10.79±0.59 ^{ab}	10.68±0.31 ^{ab}	Aminopeptidase
P10	15.93±0.31 ^c	9.66±0.74 ^e	14.42±0.56 ^d	18.32±0.78 ^a	17.34±1.03 ^b	17.10±0.84 ^b	Not identified ₁₀ **
P11	16.12±0.37 ^b	11.49±0.24 ^f	14.62±0.48 ^e	17.03±0.89 ^a	15.58±0.63 ^d	15.92±0.33 ^c	Trypsin/Chymotrypsin
P12	9.51±1.20 ^{bc}	8.53±0.22 ^c	10.40±0.50 ^{abc}	11.19±0.52 ^{ab}	12.80±0.74 ^a	11.48±1.73 ^{ab}	Not identified ₁₂ **

897 Values are mean ± SE of Integrated density mean. Within a row, means with different superscript
 898 letters differ significantly ($P < 0.05$) by Kruskal-Wallis test or One-way analysis of variance
 899 (ANOVA) followed Tukey test.

900 *Santos et al. (*in press*).

901 **Numbering defined by the authors.

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903 Table 6: Integrated density (*ID*) of the bands of amylase zymograms from the intestine of Nile
 904 tilapia (*O. niloticus*) cultured in intensive and semi-intensive systems.

Bands	Intensive system			Semi-intensive system			Enzyme*
	31 days	63 days	94 days	31 days	63 days	94 days	
A1	20.15±1.40 ^a	7.88±2.02 ^b	12.98±1.7 ^b	18.92±2.76 ^a	11.59±1.73 ^b	9.61±2.14 ^b	Amylase 1
A2	22.05±0.62 ^a	16.65±1.67 ^{bc}	17.77±1.64 ^{abc}	21.19±2.21 ^{ab}	15.35±1.90 ^c	15.29±1.55 ^c	Amylase 2
A3	20.40±1.86 ^a	18.45±2.13 ^{ab}	16.58±1.02 ^{abc}	18.37±3.20 ^{ab}	13.01±0.46 ^c	14.52±1.66 ^{bc}	Amylase 3
A4	6.12±0.42 ^a	5.55±1.02 ^{ab}	4.86±0.62 ^{ab}	4.17±0.79 ^{bc}	2.40±0.27 ^d	3.08±0.32 ^{cd}	Amylase 4
A5	3.70±0.52 ^a	3.71±0.45 ^a	2.47±0.46 ^b	2.66±0.35 ^b	2.59±0.36 ^b	1.93±0.09 ^b	Amylase 5

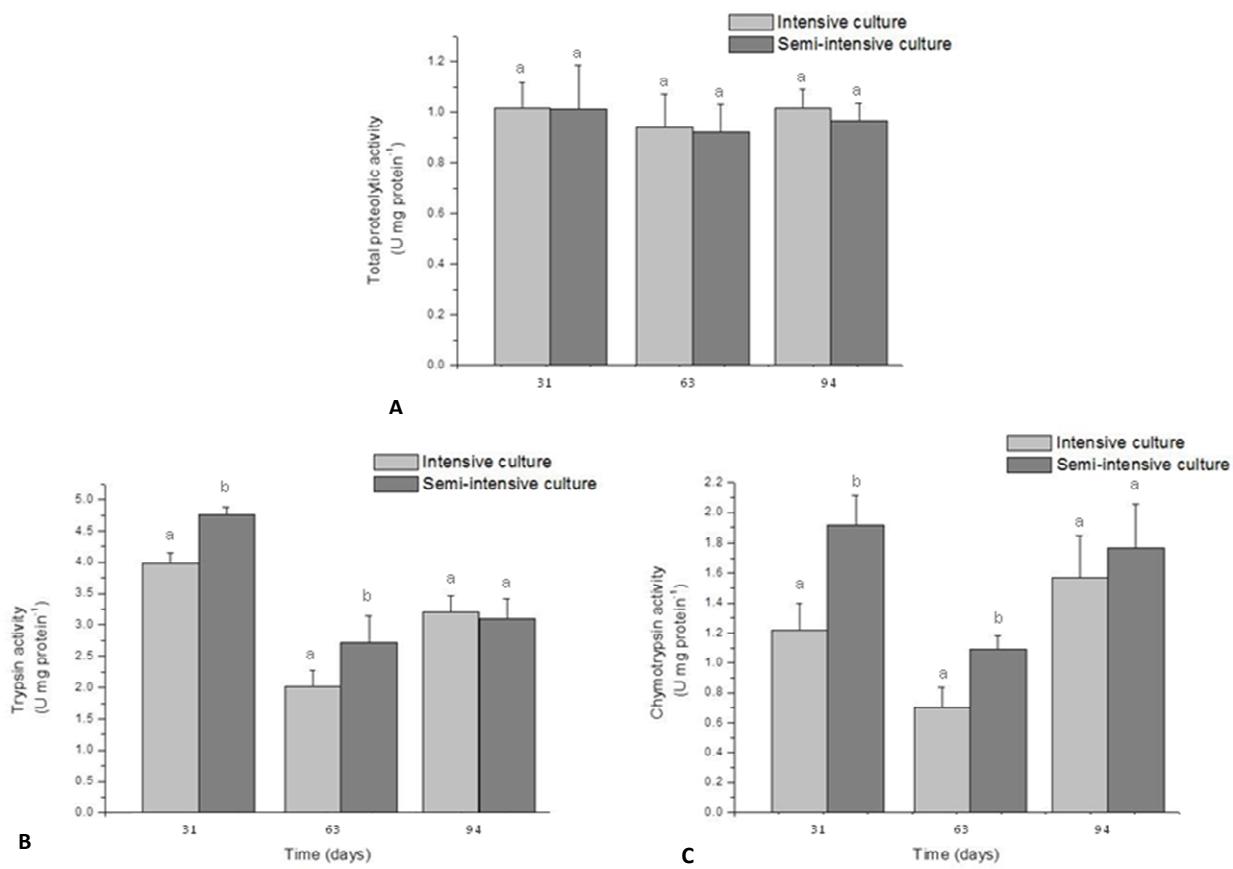
905 Values are mean ± SE of Integrated density mean. Within a row, means with different superscript
 906 letters differ significantly ($P < 0.05$) by One-way analysis of variance (ANOVA) followed Tukey
 907 test.

908 *Nomenclature defined by the authors.

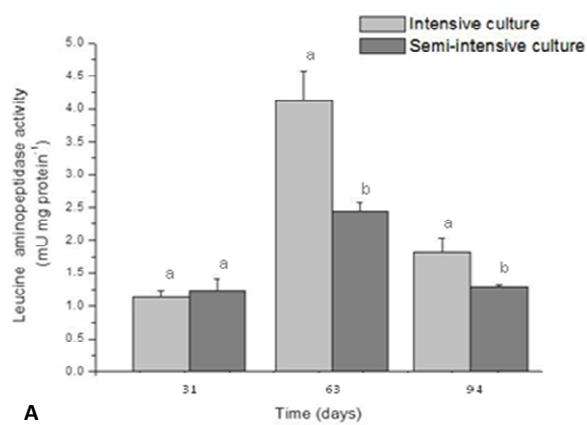
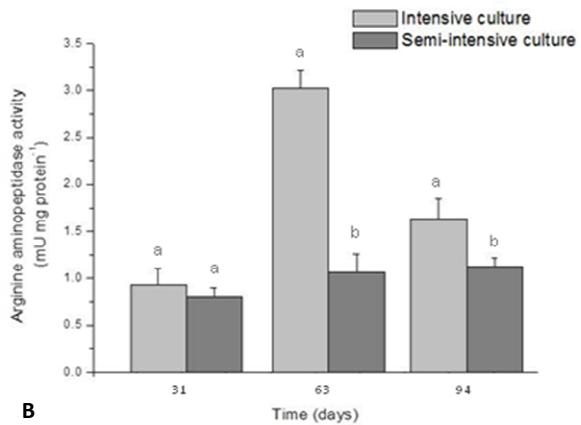
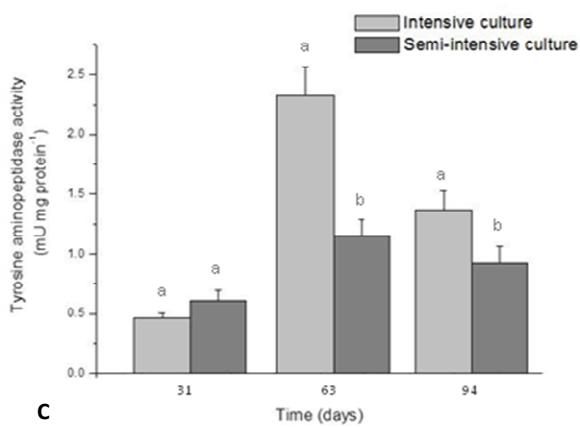
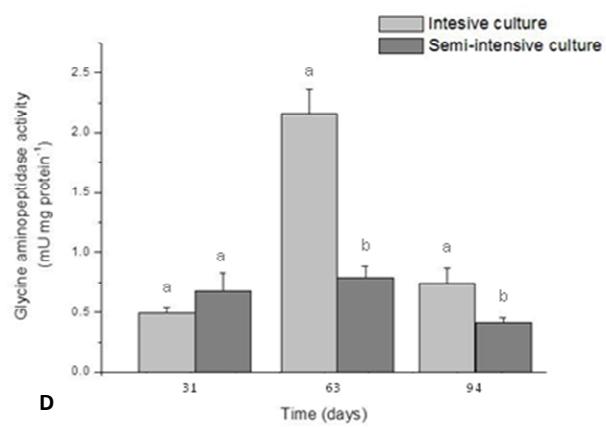
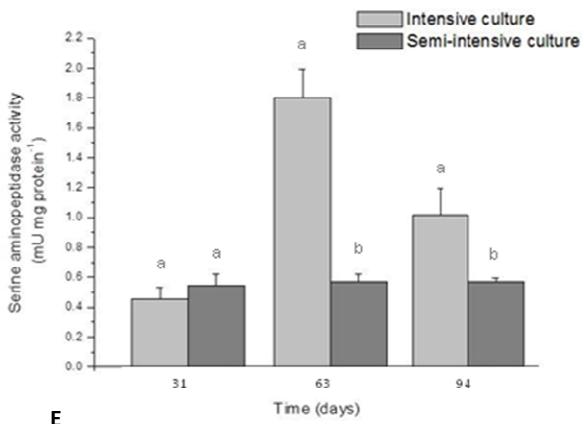
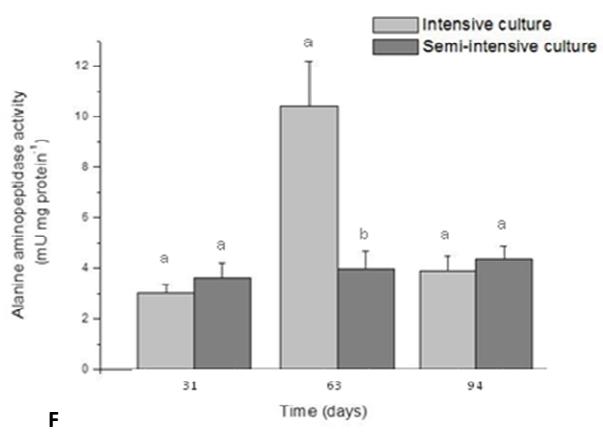
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911 **8. Figures**



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 926 Figure 1 : Alkaline proteolytic activity of the crude extract from the intestine of Nile tilapia
 927 (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD,
 928 using azocasein (A), BApNA (*Na*-benzoyl-DL-arginine-*p*-nitroanilide) (B), and SApNA (Suc-Ala-
 929 Ala-Pro-Phe-*p*-nitroanilide) (C) as substrate, expressed as U mgProtein⁻¹. For the same times,
 930 means with different superscript letters differ significantly ($P < 0.05$) by Student's t test ($n = 3$).

**A****B****C****D****E****F**

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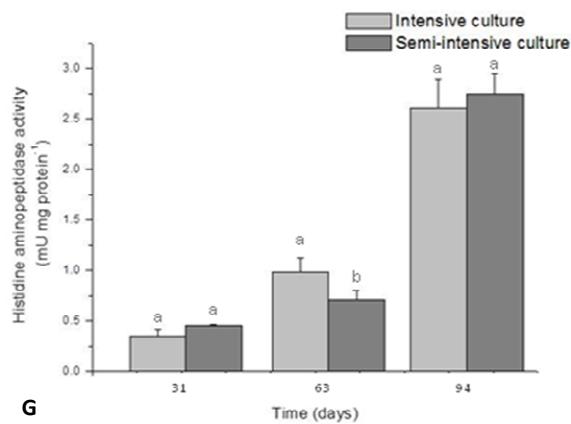
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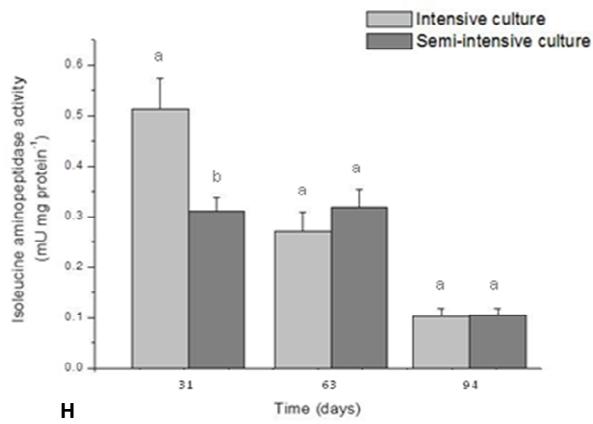
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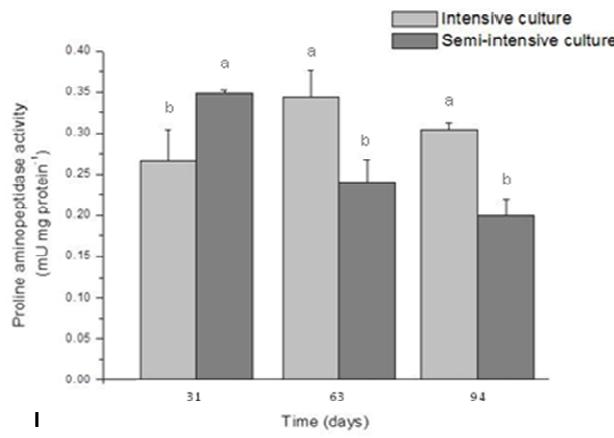
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979 Figure 2: Aminopeptidase activity of the crude extract from the intestine of Nile tilapia
980 (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD,
981 using aminoacyl- β -naphthylamide as substrates with Leu (A), Arg (B), Tyr (C) Gly (D), Ser (E),
982 Ala (F) His (G) Iso (H) and Pro (I) as specific substituents and expressed as mU mgProtein⁻¹. For
983 the same times, means with different superscript letters differ significantly ($P < 0.05$) by Wilcoxon-
984 Mann-Whitney test and Student's t test ($n = 3$).

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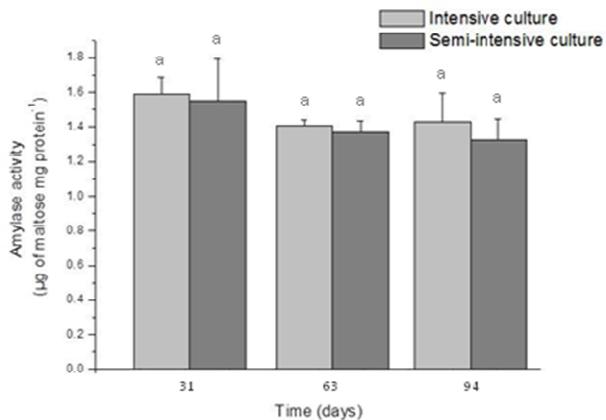


Figure 3: Amylase activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD, using starch as substrate, expressed as $\mu\text{g of maltose} \cdot \text{min}^{-1} \cdot \text{mg Protein}^{-1}$. For the same times, means with different superscript letters differ significantly ($P < 0.05$) by Student's t test ($n = 3$).

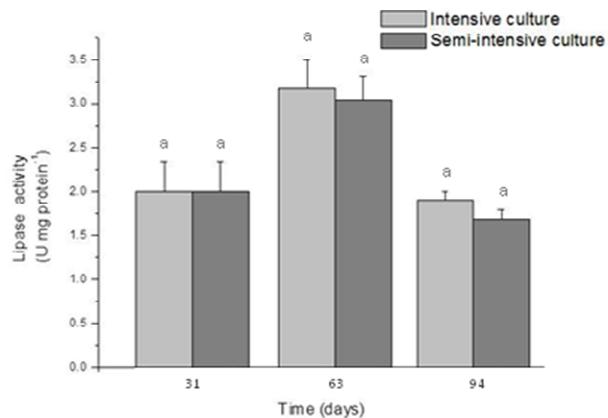


Figure 4: Lipase activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system. Values are mean \pm SD, using *p*-nitrophenyl palmitate (*p*-NPP), as substrate, expressed as U mgProtein^{-1} . For the same times, means with different superscript letters differ significantly ($P < 0.05$) by Wilcoxon-Mann-Whitney test ($n = 3$).

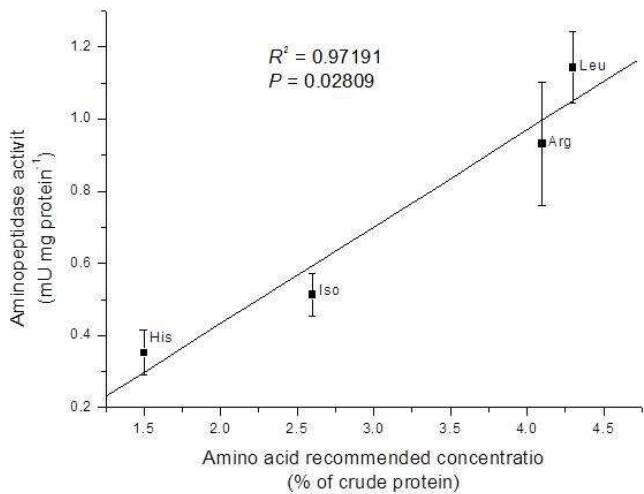
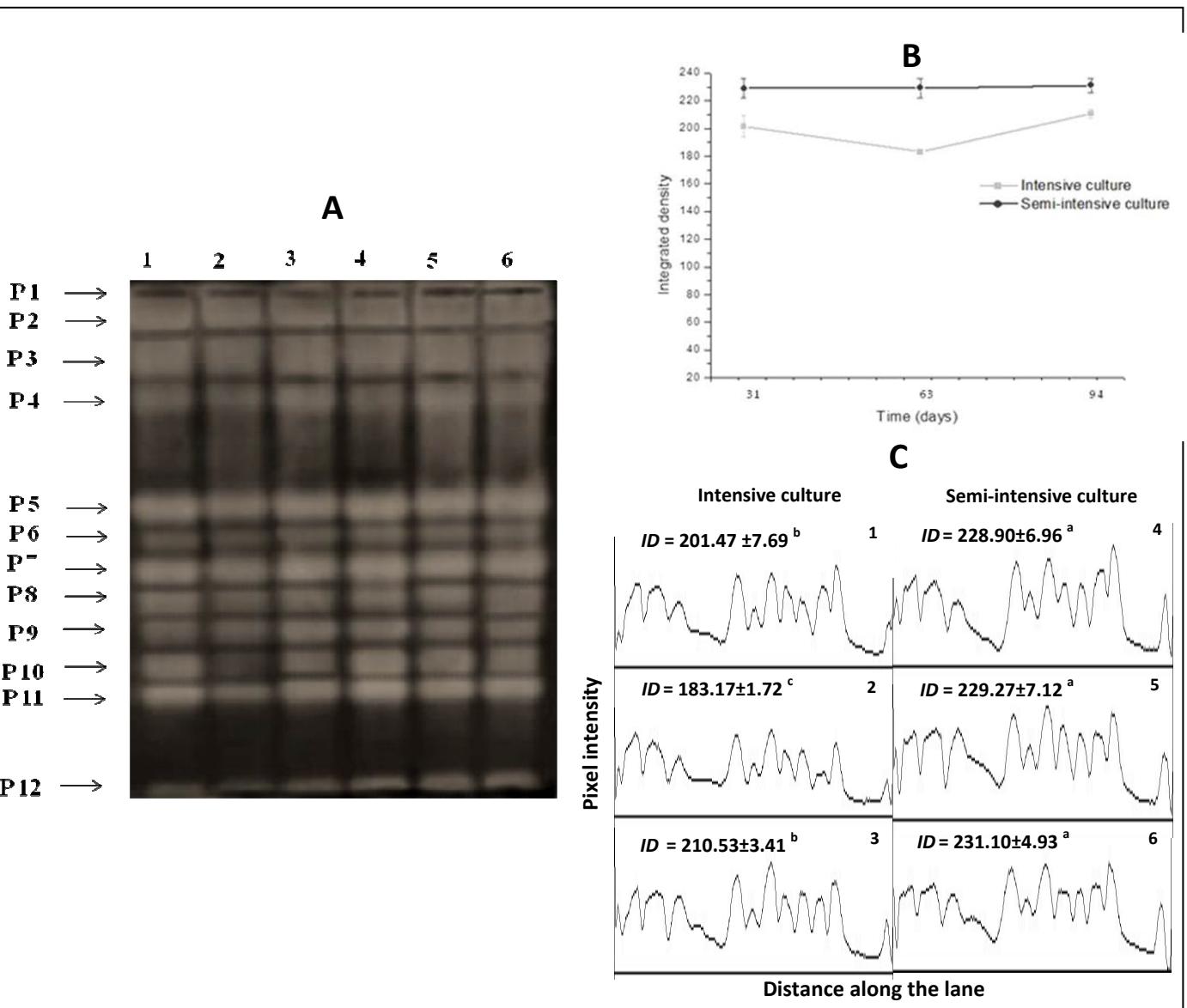


Figure 5: Correlation between aminopeptidase activity of Nile tilapia (*Oreochromis niloticus*) using aminoacyl- β -naphthylamide as substrates and recommended concentration of essential amino acids for Nile tilapia assessed by the Ideal Protein Concept. Values are shown as mean \pm SD of three crude extracts obtained from intensive culture. Concentration of amino acids expressed in % of dietary protein: arginine (4.1), leucine (4.3), isoleucine (2.6) and histidine (1.5) *Celik (2012), based on Fagbenro (2000).



1084 Figure 6: (A) Zymogram of digestive proteases of the crude extract from the intestine of Nile tilapia
1085 (*Oreochromis niloticus*) cultured in intensive and semi-intensive system, using casein as substrate.
1086 Lanes 1, 2 and 3 correspond to intensive culture in the times 31, 63 and 94 days, respectively; Lanes
1087 4, 5 and 6 correspond to semi-intensive culture in the times 31, 63 and 94 days, respectively. (B)
1088 Evolution of the integrated density of the bands of proteases zymograms from the intestines of Nile
1089 tilapia cultivated in the intensive and semi-intensive system. (C) Pixel intensity and respective
1090 means values of integrated density (*ID*) for each lane of protease bands (lanes 1-6), subfigures 1, 2
1091 and 3 correspond to the intensive system and 4, 5 and 6 semi-intensive system, means with different
1092 superscript letters differ significantly ($P < 0.05$) using One-way analysis of variance (ANOVA)
1093 followed by Tukey test ($n = 3$).

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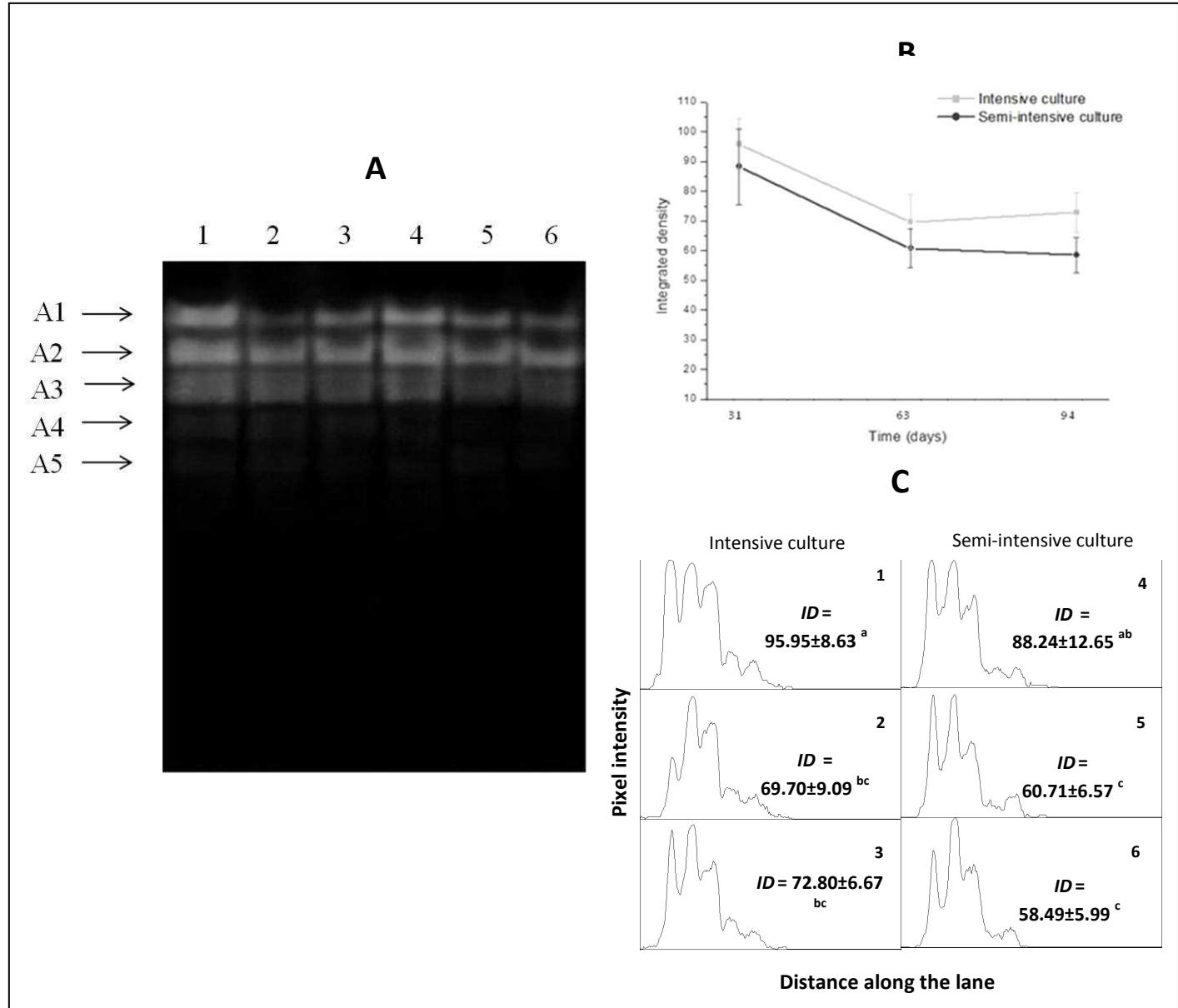


Figure 7: (A) Zymogram of amylase activity of the crude extract from the intestine of Nile tilapia (*Oreochromis niloticus*) cultured in intensive and semi-intensive system, using starch as substrate. Lanes 1, 2 and 3 correspond to intensive culture in the times 31, 63 and 94 days , respectively; Lanes 4, 5 and 6 correspond to semi-intensive culture in the times 31, 63 and 94 days, respectively. (B) Evolution of the integrated density of the bands of amylase zymograms from the intestines of Nile tilapia cultivated in the intensive and semi-intensive system. (C) Pixel intensity and respective means values of integrated density (ID) for each lane of amylase bands (lanes 1-6), subfigures 1, 2 and 3 correspond to the intensive system and 4, 5 and 6 semi-intensive system, means with different superscript letters differ significantly ($P < 0.05$) using One-way analysis of variance (ANOVA) followed by Tukey test (n=3).

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1149 Figure 8: Two-dimensional gel electrophoresis profile of the crude extract from the intestine of Nile
1150 tilapia (*Oreochromis niloticus*) cultured in intensive system. First proteins were separated by charge
1151 in the first dimension (immobiline Drystrip pH 3–10) and then by molecular weight in the second
1152 dimension (12% SDS-PAGE).

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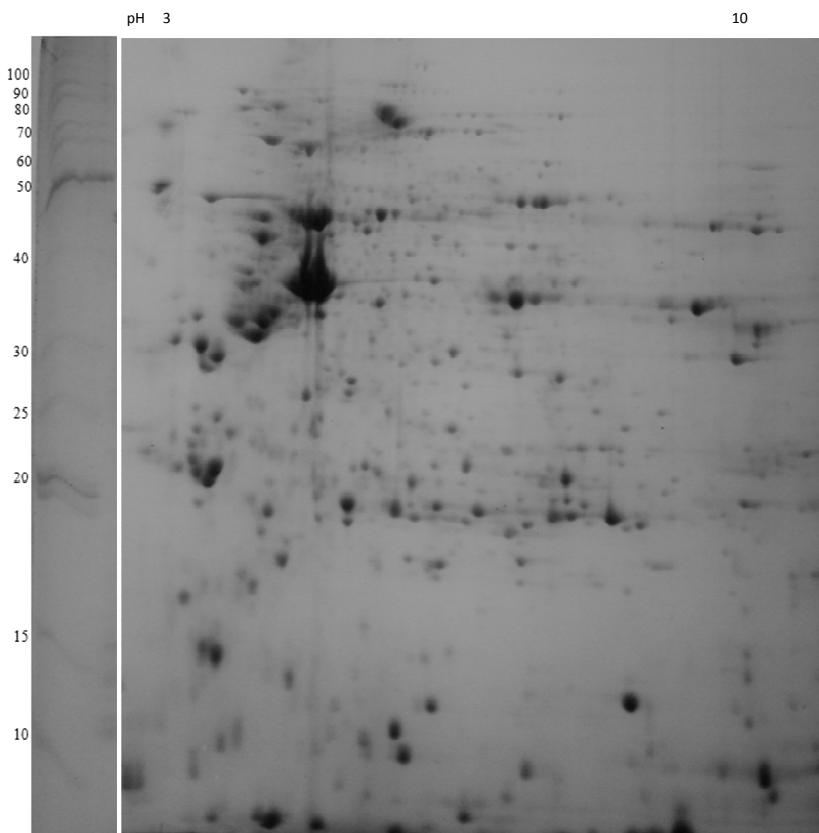
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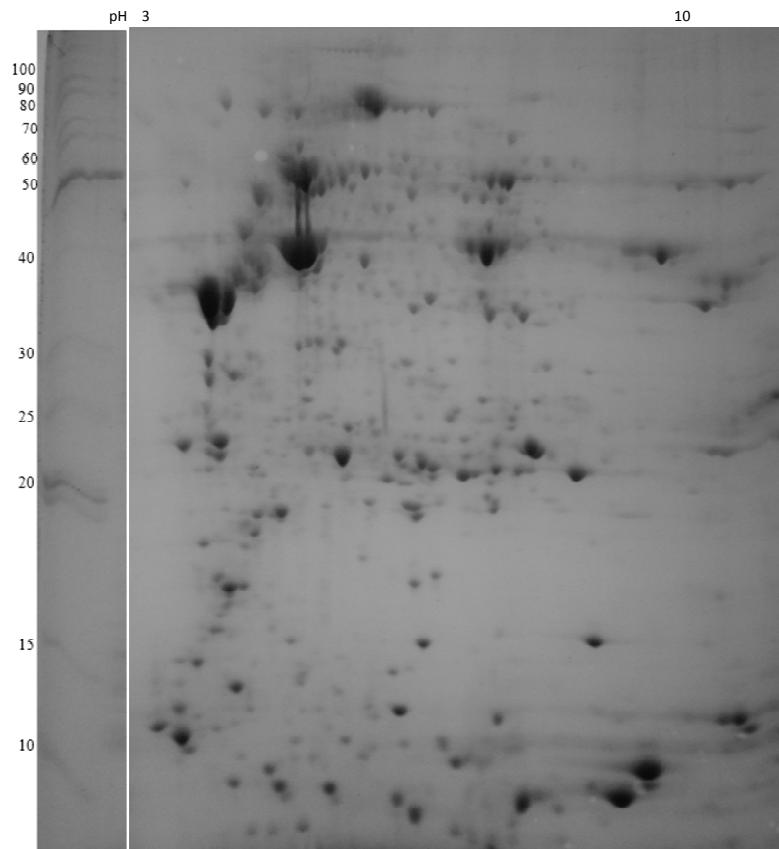
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1182 Figure 9: Two-dimensional gel electrophoresis profile of the crude extract from the intestine of Nile
1183 tilapia (*Oreochromis niloticus*) cultured in semi-intensive system. First proteins were separated by
1184 charge in the first dimension (immobiline Drystrip pH 3–10) and then by molecular weight in the
1185 second dimension (12% SDS-PAGE).

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8. CONCLUSÃO

Avaliando o desempenho das tilápias do Nilo alimentadas com diferentes concentrações de hidrolisado proteico de camarão e cultivadas em sistemas semi-intensivo e intensivo, foram observadas as mesmas tendências de crescimento.

Contudo, avaliando os sistemas intensivo e semi-intensivo a partir de outros parâmetros (fator de condição e seus derivados) algumas mudanças puderam ser observadas. Os peixes do sistema semi-intensivo apresentaram melhores resultados no início do cultivo, contudo os espécimes experimentais demonstraram compensações que levaram a uma similaridade no crescimento na avaliação final dos sistemas. O monitoramento contínuo destes parâmetros aliados ao estudo do peso relativo nos leva a refletir sobre a qualidade dos espécimes de tilápia do Nilo cultivados atualmente, já que este peixe mostrou, em algumas situações, peso relativo inferior a espécimes do ambiente natural.

As funções digestivas permitiram que a tilápia pudesse modificar suas atividades enzimáticas de forma a obter mais eficientemente os produtos das reações necessárias para o seu metabolismo, demonstrando que este peixe pode se beneficiar de várias fontes alimentares, já que seu arsenal enzimático se apresentou diversificado/adaptado suficientemente para digerir a grande variedade de comida ingerida.

9. ANEXOS

9.1 Normas da Revista Aquaculture

INTRODUCTION

Types of paper

Original Research Papers should report the results of original research. The material should not have been previously published elsewhere. Articles are expected to contribute new information (e.g. novel methods of analysis with added new insights and impacts) to the knowledge base in the field, not just to confirm previously published work.

Review Articles can cover either narrow disciplinary subjects or broad issues requiring interdisciplinary discussion. They should provide objective critical evaluation of a defined subject. Reviews should not consist solely of a summary of published data. Evaluation of the quality of existing data, the status of knowledge, and the research required to advance knowledge of the subject are essential.

Short Communications are used to communicate results which represent a major breakthrough or startling new discovery and which should therefore be published quickly. They should not be used for preliminary results. Papers must contain sufficient data to establish that the research has achieved reliable and significant results.

Technical Papers should present new methods and procedures for either research methodology or culture-related techniques.

The *Letters to the Editor* section is intended to provide a forum for discussion of aquacultural science emanating from material published in the journal.

Contact details for submission

Papers for consideration should be submitted via the electronic submission system mentioned below to the appropriate Section Editor:

Nutrition:

D.M. Gatlin

The Nutrition Section welcomes high quality research papers presenting novel data as well as original reviews on various aspects of aquatic animal nutrition relevant to aquaculture. Manuscripts addressing the following areas of investigation are encouraged:

- 1) determination of dietary and metabolic requirements for various nutrients by representative aquatic species. Studies may include environmental/stress effects on animal's physiological responses and requirements at different developmental stages;
- 2) evaluation of novel or established feedstuffs as well as feed processing and manufacturing procedures with digestibility and growth trials. Such studies should provide comprehensive specifications of the process or evaluated ingredients including nutrients, potential anti-nutrients, and contaminants;
- 3) comparison of nutrient bioavailability from various ingredients or product forms as well as metabolic kinetics of nutrients, food borne anti-nutrients or toxins;
- 4) identification of key components in natural diets that influence attractability, palatability, metabolism, growth reproduction and/or immunity of cultured organisms;
- 5) optimization of diet formulations and feeding practices;
- 6) characterization of the actions of hormones, cytokines and/or components in intracellular signaling pathway(s) that influence nutrient and/or energy utilization.
- 7) evaluation of diet supplementation strategies to influence animal performance, metabolism, health and/or flesh quality.

Manuscripts concerning other areas of nutrition using novel or advanced methods are also welcome. Please note that in regard to various diet additives such as probiotics, prebiotics, herbal extracts, etc., a very large number of papers have already been published. Therefore, Aquaculture will not continue to accept manuscripts that present initial and preliminary investigations of such additives. Manuscripts addressing these and other feed additives will be accepted for review only if they are of the highest scientific quality and they represent a significant advance in our knowledge of the mechanisms involved in their metabolism. Manuscripts may also be considered if they present clinical efficacy data generated in large-scale trials and economic cost-benefit analysis of these applications.

Aquaculture Production Science:

B.Costa-Pierce

AQUACULTURE PRODUCTION SCIENCE (PS) is one of 5 sections of the international journal AQUACULTURE dedicated to research on improvements and innovations in aquatic food production. Worldwide dissemination of the results of innovative, globally important, scientific research on production methods for aquatic foods from fish, crustaceans, mollusks, amphibians, and all types of aquatic plants. Improvement of production systems that results in greater efficiencies of resource usage in aquaculture. Effective applications of technologies and methods of aquaculture

production for improved stocking regimes, the use of new species and species assemblages, and research on the efficient and sustainable usage of system space with the objective of minimizing resource usage in aquaculture. Investigations to minimize aquaculture wastes and improve water quality, technologies for nutrient recycling in aquaculture ecosystems, and the synergy of aquaculture and other food production systems using methods such as polyculture and integrated aquaculture.

Physiology and Endocrinology:

E.M. Donaldson

The Physiology and Endocrinology Section welcomes high quality research papers presenting novel data as well as original reviews, on various aspect of the physiology and endocrinology of cultured aquatic animals and plants, providing that their content is relevant to solving aquaculture problems. Manuscripts submitted to the journal must have a valid hypothesis or objective, clearly state the relevance to aquaculture, have proper experimental design with appropriate controls and utilize appropriate statistical analysis. When a study involves the administration of a pharmaceutical or other commercial product, please use generic or scientific names rather than trade names especially in the Title. The trade name can be mentioned in the Materials and Methods together with an exact description of its composition.

Topics covered by this section include, but are not limited to, physiological and endocrine aspects of:

- Reproductive development including both endocrine and environmental control
- Induced ovulation and spermiation
- Gamete quality, storage and cryopreservation
- Control of sex differentiation
- Physiology and endocrinology of gynogenetic and triploid aquatic organisms
- Physiology and endocrinology of transgenic organisms
- Molecular genetic assessment of physiological processes
- Larval physiology and the ontogeny of endocrine systems
- Melanism
- Metamorphosis, smolting (salmonids) and molting (crustacea)
- Nutritional physiology
- Osmoregulation
- Stress
- Anesthesia, transport, handling

- Physiology of harvest techniques, product quality, flesh quality and pigmentation
- Endocrine and environmental regulation of growth
- Rearing density
- Temperature tolerance
- Disease physiology
- Pollutant physiology and water quality (when directly relevant to aquaculture)
- Respiratory, muscle and exercise physiology of cultured organisms
- Immunology (manuscripts on the physiological effects of probiotics must be of high scientific quality with statistically valid conclusions)

Diseases:

B. Austin

The Diseases Section welcomes high quality research papers presenting novel data as well as original reviews, on various aspect of the diseases of aquatic animals and plants, so long as their content is relevant to solving aquaculture problems.

Please note, however, with respect to the probiotic potential of various bacteria and the antibacterial or immunostimulatory effects of herbal extracts a very large number of papers have already been published. As a result, Aquaculture will not continue to accept manuscripts that present further initial and preliminary investigations of these phenomena. Manuscripts addressing these topics will be accepted for review only if they are of the highest scientific quality and they represent a significant advance in our knowledge of the mechanisms involved. Manuscripts may also be considered if they present clinical efficacy data generated in large-scale trials and economic cost-benefit analysis of these applications.

Genetics:

G. Hulata

The Genetics Section welcomes high-quality research papers presenting novel data, as well as critical reviews, on various aspects of selective breeding, genetics and genomics, so long as the content is relevant to solving aquaculture problems. Please note, however, that Aquaculture will not accept manuscripts dealing with the application of well-described techniques to yet another species, unless the application solves a biological problem important to aquaculture production. Aquaculture will not accept manuscripts dealing with gene cloning, characterizing of microsatellites, species identification using molecular markers, EST papers with small collections, or mapping papers with a small number of markers, unless the papers also deal with solving a biological problem that is relevant to aquaculture production. Where appropriate, linkage maps should include co-dominant

markers, such as microsatellite DNA and SNP markers, to enable application to other populations and facilitate comparative mapping. Aquaculture will not accept manuscripts focusing mainly on population genetics studies that are based on RAPD and AFLP markers, since the dominance and multilocus nature of the fingerprints are not suitable for making inferences about population genetic diversity and structure. There may be other journals that are more suitable for manuscripts not meeting these requirements.

Sustainability and Society:

D.C. Little

The Sustainability and Society section of the journal Aquaculture invites articles at the interface of natural and social sciences that address the broader roles of aquaculture in global food security and trade. Aims and scope of the Sustainability and Society section are the: global dissemination of interdisciplinary knowledge regarding the management of aquatic resources and resulting impacts on people. Interconnections with other sectors of food production; resource management and implications for societal impact. Going beyond a narrow techno-centric focus, towards more holistic analyses of aquaculture within well-defined contexts. Enquiry based on understanding trajectories of change amid the global challenges of climate change and food security. Mixed methods and approaches that incorporate and integrate both social and natural sciences. Relevance for the diverse range of policy makers, practitioners and other stakeholders involved. Articles that take a value chain approach, rather than being wholly production orientated, are encouraged.

Page charges

This journal has no page charges.

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This must be stated at an appropriate point in the article.

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