1	Title
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3	The effects of combined phytogenics on growth and nutritional physiology of Nile tilapia
4	Oreochromis niloticus
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Abstract

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24 This study investigated whether dietary supplementation of phytogenic compounds 25 limonene and thymol had synergistic or additive effects on growth and selected nutritional 26 physiology pathways in Nile tilapia. A 63-day feeding experiment was conducted using 27 fish of 1.5 ± 0.0 g (\pm standard error) fed on a commercial diet coated with either 0 ppm 28 limonene and thymol (control), 400 ppm limonene (L), 500 ppm thymol, (T) or a 29 combination of 400 ppm limonene and 500 ppm thymol (LT). Final fish weight (FW) was 30 significantly improved to similar extents by diet LT (16.7 \pm 0.3 g) and L (16.6 \pm 0.4 g). Dietary thymol alone and the control did not enhance FW (15.0 \pm 0.4 g and 13.7 \pm 0.4 g 31 respectively). Dietary thymol had shown a strong tendency to improve somatic growth 32 33 (P=0.052). The analysed candidate genes involved in the pathways of nutrient digestion, 34 absorption and transport (muc), lipid metabolism (lpl), antioxidant enzymes (cat) and 35 somatotropic axis growth (igf-I) were also up-regulated to similar extents in Nile tilapia by 36 diet L and LT (P<0.05), above the regulation observed with the diet supplemented 37 exclusively with thymol. This suggests lack of synergistic or additive effects on growth and nutritional physiology pathways when limonene and thymol are supplied in the diet. 38

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Keywords

- 41 Growth promoters, limonene, Nile tilapia, nutritional physiology, phytogenic compounds,
- 42 thymol

1. Introduction

Phytogenic compounds are natural bioactive compounds derived from herbs, shrubs and spices with essential oils extracted from the plant parts being the major source of phytogenic compounds (Yang et al., 2015; Sutili et al., 2017; Upadhaya and Kim, 2017). Each phytogenic compound contains several bioactive components or molecules in different proportions with bioactive components present in higher proportions largely determining the biological properties of the essential oils (Santos et al., 2011; Chakraborty et al., 2014; Yitbarek, 2015). Bioactive components in the plants are mainly hydrocarbons (e.g. terpenes, sesquiterpenes), oxygenated compounds (e.g. alcohol, aldehydes, ketones) and a small percentage of non-volatile residues (e.g. paraffin, wax) (Losa, 2000). Some active compounds such as thymol, carvacrol, limonene, cinnamaldehyde and eugenol from the plants thyme, oregano, citrus, cinnamon and clove respectively have been noted to exert positive effects on nutrition, performance or health of monogastric animals (Wallace et al., 2010; Chakraborty et al., 2014; Sutili et al., 2017; Upadhaya and Kim, 2017). When mixtures of phytogenic compounds are used in animal feed, they can either have synergistic, additive, indifferent or antagonistic effects on growth and other response

synergistic, additive, indifferent or antagonistic effects on growth and other response indicators in monogastric animals (Bassole and Juliani, 2012; Costa et al., 2013; Abd Elhack et al., 2016; Valenzuela-Grijalva et al., 2017; Amer et al., 2018; Youssefi et al., 2019). Synergistic or additive effects of phytogenic compounds on growth performance lead to

enhanced growth of animals above the levels attained when the compounds are supplied individually (Windisch et al., 2008; Yang et al., 2015). In fish, the combination of thymol and carvacrol is arguably the most investigated blend of phytogenic compounds for its beneficial effects on growth (Zheng et al., 2009; Ahmadifar et al., 2011; Hyldgaard et al., 2012; Chakraborty et al., 2013; Ahmadifar et al., 2014; Peterson et al., 2014; Perez-Sanchez et al., 2015). For instance, Peterson et al. (2014) reported that channel catfish (*Ictalurus punctatus*) fed on a diet supplemented with a combination of limonene, thymol, carvacrol and anethol gained 44% more weight than the control attributed to synergistic or additive effects of the phytogenic compounds.

Often though indifferent effects can be observed when combination of phytogenic compounds show no differences compared to treatments consisting of their individually supplied compounds or the controls (Bassole and Juliani, 2012). Indifferent effects have also been noted when combination of phytogenic compounds exerts significantly higher effects to similar extents with only some treatments composed of their individually supplied phytogenic compounds. Zheng et al. (2009) supplemented diets with 500 ppm of either carvacrol, thymol or a mixture of carvacrol and thymol, and found a significantly

higher weight gain of channel catfish with the dietary mixture of carvacrol and thymol compared to the control and diet with thymol alone, but not with the diet supplemented only with carvacrol. In addition, antagonistic effects occur when individual phytogenic compounds might have positive effects but their combination results in negative effects compared to the controls (Bassole and Juliani, 2012). Such antagonistic effects derived from phytogenic blends have been often attributed to high concentrations of these compounds that potentially result in unpleasant taste and smell and thereby retarding feed intake and consequently growth (Windisch et al., 2008; Steiner, 2009; Costa et al., 2013; Colombo et al., 2014).

The different responses to phytogenics mentioned above highlights the importance of identifying combinations and doses of phytogenic compounds resulting in additive and synergistic effects on fish growth. In our previous study (Aanyu et al., 2018), two phytogenic compounds, namely limonene and thymol, classified as monoterpene and diterpene, respectively, were found to have growth-promoting tendencies in Nile tilapia (*Oreochromis niloticus*). We hypothesised that combinations of limonene and thymol can potentially have additive or synergistic effects on the growth of Nile tilapia. Consequently, this study aimed to investigate the effects of a blend of limonene and thymol, compared with each of the compounds individually, on the growth, feed efficiency and nutritional physiology of Nile tilapia. The study followed a candidate gene approach to investigate physiological pathways underpinning the response of fish to the phytogenic compounds. A selection of marker genes within the pathways of somatotropic axis-mediated growth, nutrient absorption and transport, lipid metabolism and antioxidant enzyme status that showed potential to be regulated by limonene and/or thymol were analysed (Aanyu et al., 2018).

2. Materials and Methods

- 106 2.1 Ethical statement
- All experiments were subjected to ethical review and approved by the University of
- Stirling through the Animal and Welfare Ethical Review Body. The project was conducted
- under the UK Home Office in accordance with the amended Animals Scientific Procedures
- 110 Act implementing EU Directive 2010/63.

112 2.2 Experimental design

113 The feeding trial was carried out at the Aquaculture Research and Development Center (ARDC), Uganda between March and May 2015. Nile tilapia juveniles from the same cohort were obtained from the ARDC fish farm, acclimatized and size graded to 1.54 ± 0.0 g (mean \pm standard error). Thirty eight (38) fish were stocked in each of the 16 experimental tanks. Each tank had a water holding capacity of 60 L, in a flow through system with a flow rate of 1-2 L min⁻¹. The water in each tank was aerated using air stones and heated using aquaria water heaters to 25.0 - 26.6 °C. A photoperiod of 12h light-12h

dark was maintained.

Water quality was monitored routinely to ensure that it was within the requirements for Nile tilapia growth (Lim and Webster, 2006). A multi-parameter meter (HQ40D model, Hach Ltd Germany) was used to measure dissolved oxygen, pH and water temperature. The level of ammonia-nitrogen was assessed using a fresh water test kit from API Company Ltd UK following the user guide from the manufacturer. Water flowing into the fish rearing tanks had 6.6 ± 0.6 mg L⁻¹ of dissolved oxygen, pH 6.8 ± 0.3 , undetectable levels (< 0.05 mg L⁻¹) of ammonia-nitrogen and water temperature ranging from 23.3 - 24.3 °C before heating with aquaria water heaters.

2.3 Experimental diets

A standard commercial feed (CP35%, Kampala, Uganda) for juvenile Nile tilapia produced at the ARDC was supplemented with limonene (97 % purity) and/or thymol (95 % purity) from Sigma Aldrich, Kampala, Uganda using concentrations found to have growth-promoting potential in Nile tilapia (Aanyu et al., 2018). The diets included: 0 ppm limonene and thymol (Control); 400 ppm limonene (L); 500 ppm thymol (T); and a combination of 400 ppm limonene and 500 ppm thymol (LT). In order to supply the above concentrations of phytogenic compounds to the feed, each concentration of phytogenic compounds was prepared in 100 mL of absolute ethanol and sprayed onto 1 kg of feed. The control was also coated with a similar amount of ethanol but no phytogenic compound was added. All diets were air-dried for one day, packed in airtight polythene bags and stored at room temperature until use.

Each experimental diet was tested in quadruplicate tanks and the treatments were distributed randomly. The fish were fed the experimental diets twice a day to apparent satiation and the feed intake was recorded.

145 The proximate nutritional composition (dry matter, moisture, protein, lipid, fibre, ash 146 and gross energy) of the standard diet was determined according to the methods of the 147 Association of Official Analytical Chemists (AOAC, 1990) and the joint technical 148 committee of the International Organisation for Standardisation and International 149 Electrotechnical Commission (ISO/IEC 17025). Briefly, dry matter content was estimated by drying a sample of feed in an oven at 105-110°C to a constant weight and the 150 151 percentage retained weight from the original sample was the amount of dry matter whereas 152 the percentage loss in weight of the sample was the moisture content. Crude protein 153 content was determined using the Kjeldahl method and crude lipid by petroleum ether 154 extraction using the Soxhlet method. Crude fibre content was analysed by acid / alkaline 155 hydrolysis of a sample and the amount of insoluble residues resistant to hydrolysis was the 156 fibre content. Crude ash was determined by combustion of a sample in a furnace at 600 °C 157 for 24 h. The gross energy using determined using the bomb calorimetry method. 158 Proximate composition of the standard diet used in this study is shown in Table 1.

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2.3 Fish measurements and sample collection

161 Growth of the fish was estimated by measuring the weight (accuracy of 0.1 g) and total

length (0.1 cm) of all fish in each tank every three weeks and at the end of the experiment

163 (63 days). This procedure was carried out while fish were anesthetised using 0.02 g L⁻¹ of

164 clove oil for 3-5 min, after which the fish were placed in aerated water and taken back to

the experimental tanks. At the end of the experiment, the number of live fish in each tank

was recorded for survival estimation, and sections of liver and fore intestine were dissected

167 from 16 fish per treatment (n=4 per tank) and placed in 1.5 mL tubes containing

168 RNAlater® (Sigma Aldrich, Kampala, Uganda). The liver and fore intestine samples were

kept at 4 $^{\circ}$ C overnight, shipped to the UK and transferred to a -20 $^{\circ}$ C freezer until RNA

was extracted.

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172 *2.4 Fish performance calculations*

- 173 The effects of the experimental diets on fish performance were assessed by calculating the
- performance indicators below using the following formulae:

- Final average fish weight (FW, g) = total fish biomass at end of the trial (g) / number of
- 177 fish at end of the experiment;
- Percentage (%) weight gain (WG, %) = ((final average fish weight (g) initial average fish
- weight (g)) / initial average fish weight (g)) \times 100;

- Condition factor (CF) = (final average fish weight / final average total length³) \times 100;
- Fish survival rate (SR, %) = (number of alive fish at end of trial / initial number of fish
- 182 stocked) \times 100 %;
- Feed intake (FI, % body weight per day) = $(100 \times (average feed intake fish^{-1} / ((initial)^{-1} + (average feed intake fish^{-1} + (average feed intake feed intake feed intake fish^{-1} + (average feed intake feed inta$
- average body weight \pm final average body weight)/2))) / duration of trial (d);
- Feed conversion ratio (FCR) = average feed intake fish⁻¹ / weight gain;
- Protein efficiency ratio (PER) = weight gain / protein intake.

- 188 *2.5 Molecular analyses*
- 2.5.1 RNA extraction and complementary DNA (cDNA) synthesis
- 190 Tissue samples from the liver and fore intestine were homogenised in TRI Reagent (Sigma
- 191 Aldrich, Dorset, UK) with a mini bead-beater 16 (Biospec Bartlesville, OK, USA), and
- 192 total RNA was extracted from the samples (n=16 per tissue and treatment). The
- 193 concentration and purity of the RNA was measured by spectrophotometry with an ND-
- 194 1000 Nanodrop (Nanodrop 1000, Thermo Scientific, Glasgow, UK). The integrity of the
- 195 RNA (aliquots fo 200 ng total RNA) from each sample was further assessed by agarose gel
- 196 (1 %, v/v) electrophoresis.
- From a total of 16 RNA samples extracted per tissue and treatment, eight RNA samples
- were derived by pooling together two samples from the same tank and treatment, taking
- 199 equal quantity (2.5 µg µl⁻¹) of RNA from each of the two samples being pooled (adapted
- from Glencross et al., 2015). Thus the final mixture (vortex mixed and centrifuged) had a
- 201 50 % contribution of each of the two samples that were pooled.
- A high capacity reverse transcription kit without RNase inhibitor from AB Applied
- Biosystems (Warrington, UK) was used to reverse transcribe 2 µg µl⁻¹ RNA from each
- 204 pool sample (n = 8 per treatment) to cDNA following the protocol provided by the
- 205 manufacturer.

- 207 2.5.2 Quantitative real-time Polymerase Chain Reaction (qPCR)
- The mRNA expression levels of selected genes of interest in the pathways of somatotropic
- axis-mediated growth, nutrient absorption and transport, lipid metabolism and antioxidant
- 210 enzyme status were analysed by quantitative real-time polymerase chain reaction (qPCR)
- 211 in tissue samples (liver or fore intestine) in which they perform their major biological
- 212 functions. The selected target genes included mucin-like protein (muc), oligo-peptide
- 213 transporter 1 (pept1), lipoprotein lipase (lpl), sterol regulatory element binding
- 214 transcription factor 1 (*srebf1*), alkaline phosphatise (*alp*), phospholipase A2 (*pla2*),

215 catalase (cat), growth hormone (gh), and insulin growth factor I (igf-I). Efficiency of the 216 primers was first tested by generating standard dilution curves, assessing the melting 217 curves and cycle threshold (Ct) values (Larionov et al., 2005). Efficient primers were 218 considered to have values between 0.80 - 1.10, a single melting peak, Ct value below 30 219 and one clear band under 1 % agarose gel electrophoresis. The details of the primers used 220 for qPCR analyses are provided in Table 2. Each qPCR contained duplicate samples (total 221 volume 20 µl each) containing 5 µl of 20-fold diluted cDNA, 3 µl nuclease-free water, 1 µl 222 (10 pmol) each for the forward and reverse primers, and 10 µl of Luminaris color higreen 223 gPCR Mix (Thermo Scientific, Hemel Hempstead, UK). All the reactions were run in 96 224 well-plates using a Biometra TOptical Thermocycler (Analytik Jena, Goettingen, 225 Germany). A calibrator sample (20-fold dilution of all samples pooled cDNA) and a 226 negative control with no cDNA (non-template control-NTC) were included in each plate. 227 The qPCR thermocycling program involved pre-heating samples at 50 °C for 2 min 228 followed by 35 cycles, initial denaturing at 95 °C for 10 min, denaturation at 95 °C for 15 s, 229 annealing at 60 °C for 30 s, and an extension at 72 °C for 30 s.

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- 2.5.3 Gene expression computations
- 232 Ct values for the duplicate runs of each sample were averaged. The data were normalised
- using the geometric mean expression of the references genes (β -actin and ef-1 α) and the
- relative expression of each gene was calculated according to the equation of Pfaffl (2001).
- Heat maps enabling cluster analysis and visualisation of the expression patterns of the
- analysed genes were generated but not based on statistical differences. Java Tree View
- 237 Software was used to plot the data and perform cluster analysis based on Euclidean
- 238 distance. Expression level of each gene was natural log transformed and normalised
- 239 against two reference genes (β -actin and ef-1 α).

- 241 2.6 Statistical analysis
- The data were analysed using the Statistical Package for the Social Sciences (SPSS)
- version 19 (Chicago, USA) (Landau and Everitt, 2004). The fish performance and qPCR
- results are expressed as means \pm standard error. Normality of distribution of the data was
- assessed using Kolmogorov-Smirnov's tests. Data not normally distributed were subjected
- 246 to natural logarithm ln (qPCR data) and arcsin square-root (WG, CF, SR, FI, FCR, and
- 247 PER) transformation. Differences among dietary treatments were analysed by one-way
- ANOVA followed by Tukey's test. When heterogeneity of variances occurred, Welch's test
- 249 was performed with Game-Howell's test to determine differences between treatments.

Significant differences were considered at P value < 0.05. In addition, Pearson's correlation analysis was performed to indicate the relationship and degree of correlation between FI and FCR, FW and SR, FW and PER. The significance level of correlation was set to P < 0.05.

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3. Results

256 3.1 Fish performance

257 Table 3 shows the performance of Nile tilapia fed on diets supplemented with 400 ppm 258 limonene (L), 500 ppm thymol (T) and the combination of 400 ppm limonene with 500 259 ppm thymol (LT) for 63 days (9 weeks). There was a significant increase (P < 0.01) in the 260 final weight of fish fed on the diets supplemented with limonene, that is, diets L and LT, 261 compared to the control. No significant increases in fish weight were observed with diet L 262 and LT during day 1, 21, and 42 of the feeding experiment (data not shown). The diet 263 supplemented exclusively with thymol (T) did not significantly improve the final weight of 264 the fish (P = 0.052). There was a significantly higher percentage weight gain (% WG; P = 0.01) of fish fed on diets L and LT compared to the control, whilst fish fed diet T did not 265 266 show significant differences compared with the control. Despite there were no significant differences in the final survival of the fish among treatments, there was a strong significant 267 268 positive correlation (r = 0.967, P = 0.033) between the survival rate and final weight of the 269 fish (Table 3). Condition factor (CF) was not significantly different among treatments. 270 The protein efficiency ratio (PER) was significantly higher in fish fed the diets L and LT 271 compared to the control. While no significant differences in PER were observed among 272 fish fed on diets L, T and LT, PER had a strong significant positive correlation (r = 0.974, 273 P = 0.026) with final fish weight and thus higher PER corresponded with higher final fish 274 weight. This study found no significant differences in feed conversion ratio (FCR) among 275 the treatments supplemented with L, T and LT, but fish fed diets L and LT had

intake (% FI) not being significantly different among fish fed diets L, T and LT, significantly lower (P = 0.019) FI was obtained with the fish fed on diets L and LT compared with the control. In addition, there was a strong positive significant correlation (r = 0.996, P = 0.004) between FI and FCR, and lower FI corresponded with low FCR and

significantly lower FCR (P = 0.006) than the control-fed fish (Table 3). Despite % feed

therefore better feed utilisation efficiency.

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- 283 3.2 Relative mRNA gene expression
- 284 The heat map in Figure 1 represents the relative expression patterns (not based on
- statistical differences) of genes analysed in the liver (a) and fore intestine (b) of Nile tilapia
- 286 fed on the experimental feeds. There were more genes with patterns of higher relative
- expression levels (red) among the fish fed on diets L and LT compared with diet T, when
- all dietary treatments were compared to the control.

- 3.2.1 Expression of genes involved in somatotropic axis in liver
- Insulin growth factor I (igf-I) was significantly (P = 0.025) up-regulated in the liver of fish
- 292 fed on diets L and LT compared with control fish (Figure 2). However, the expression of
- 293 igf-I did not differ significantly between the fish fed on diets L, T and LT (Figure 2). In
- addition, the relative expression levels of gh was not significantly different in the livers of
- fish fed on diets L, T, LT and the control.

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- 297 3.2.2 Expression of genes involved in lipid metabolism in liver
- The expression of *lpl*, *alp* and *srebf1* in the liver of Nile tilapia fed on the experimental
- diets is shown in Figure 3. Levels of lpl mRNA were significantly (P = 0.003) higher in
- 300 fish fed on diet LT compared with the control. The expression of *lpl* in the fish fed on diets
- 301 L and T was not significantly different from the control. Similarly, no significant
- differences in the relative expression of *alp* and *srebf1* were found among the experimental
- 303 treatments (Figure 3).

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- 3.2.3 Expression of genes regulating nutrient digestion, absorption and transport in the fore
- 306 intestine
- The mRNA levels of muc were significantly higher (P = 0.025) in the fore intestine of fish
- fed on diet LT compared with the control (Figure 4). Besides, the expression of *muc* in the
- fish fed on diets L and T did not differ significantly from the control (P = 0.097). The
- expression of *pla2* also did not statistically differ among the dietary treatments (P = 0.086).
- Oligo-peptide transporter 1 (pept1) expression was significantly (P = 0.047) up-regulated
- in the fish fed on diet LT compared with the control, although expression levels in fish fed
- diets L and T did not differ statistically compared with the control (Figure 4).

- 3.2.4 Expression of antioxidant enzymes in liver
- The expression of cat was significantly higher (P = 0.006) in the liver of fish fed on diets L
- and LT compared with the control (Figure 5).

4. Discussion

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The present study investigated the effects of diets containing limonene and thymol, supplemented both individually and in combination, on growth and nutritional physiology of Nile tilapia. The goal was to establish whether blends of limonene and thymol had synergistic and/or additive effects on the growth of Nile tilapia. A selection of gene markers regulating nutrient digestion, absorption and transport, lipid metabolism, antioxidant enzymes and somatotropic axis growth were investigated.

Fish fed diets supplemented with limonene alone (L) and the blend of limonene and thymol (LT) had significantly higher FW, % WG, PER and lower FI and FCR than the control. On the contrary, fish fed on the diet supplemented with only thymol (T) did not show any statistical difference in such parameters compared with the control. Similarly, Zheng et al. (2009) found no synergistic or additive effects of a combination of carvacrol and thymol on the weight gain of channel catfish. The fish fed on the diet supplemented with only carvacrol, and the blend of carvacrol and thymol attained statistically higher weight gain compared with the diet with only thymol and the control, but the dietary mixture of carvacrol and thymol did statistically increase weight gain to the same extent as the diet with only carvacrol.

Among the feed utilisation parameters, the present study found enhanced feed efficiency (i.e., lower FCR) in fish fed diets L and LT, and a strong significant positive correlation between PER and FW. The correlation between PER and FW showed that, as the utilisation of protein from the feed was enhanced (high PER), FW of the fish was increased. This could have contributed to the significantly improved somatic growth of the fish fed on diets L and LT, both treatments with increased PER compared to the control. The increased WG and lowered FI levels observed with diet L and LT fed fish are in agreement with Hashemipour et al. (2013) who found lower FI corresponding with the highest WG and feed efficiency in broiler chicken fed on a diet with a mixture of 200 ppm of thymol and carvacrol compared to the control. It is known that efficient growth in fish does not necessarily coincide with maximum or higher FI because fish adjust their FI according to their energy requirements (Ali and Jauncey, 2004), with better feed efficiency occurring below maximum FI (Rad et al., 2003; Sawhney, 2014). Conversely, some studies with phytogenic compounds (thymol and carvacrol) in pig diets found low FI corresponding with low WG (Lee et al., 2003a; Lee et al., 2003b; Zhai et al., 2018). While it is difficult to identify the exact causes of such an apparent discrepancy with the present results, one possible reason might stem from the pungent odour of thymol and carvacrol

that can affect palatability and ultimately feed intake since, compared to fish, pigs are more sensitive to smell (Michiels et al., 2012; Muthusamy and Sankar, 2015).

The actions of genes regulating growth in the pathways within nutritional physiology are complementary to each other (Hashemi and Davoodi, 2010; Steiner and Syed, 2015). In this study, insulin growth factor I (*igf-I*), which plays a core role in regulating growth in the somatotropic axis, was up-regulated to a similar extent in the liver of fish fed diets L and LT, corresponding also to higher final FW and feed utilisation efficiency (FCR) than the control. This observation implies that *igf-I* was largely activated by limonene suggesting that there was no synergistic or additive effect of limonene and thymol in influencing somatotropic axis-mediated growth.

Key mechanisms underlying feed utilisation efficiency include nutrient digestion, absorption and transport, in which mucin-like protein (*muc*) and oligo-peptide transporter 1 (pept1) are important components (Verri et al., 2011; Fascina et al., 2012). The present study found a significantly higher expression of *muc* in the fore intestine of fish fed on diet LT compared with the control, with diets L and T showing no differences in expression of muc with the control and diet LT. The high expression of muc found with diet LT can be associated with an increase in the secretion/quantity of mucus, which then serves as a lubricant aiding absorption of nutrients into the bloodstream through which they are transferred to tissues for various functions including growth (Kamali et al., 2014). Moreover, high expression of *muc* corresponded with enhanced somatic growth of the fish in the LT treatment. Despite that Tsirtsikos et al. (2012) did not specifically investigate muc expression, their study on broilers fed on diets containing a blend of limonene, carvacrol and anethol also reported an increase in mucus volume in the fore intestine. Additionally, Jamroz et al. (2006) found higher mucus secretion in the fore intestine of broilers fed diets supplemented with a combination of phytogenics including carvacrol, cinnamaldehyde and capsicum oleoresin. The present results for Nile tilapia are consistent with these terrestrial animal studies, suggesting that the mechanism of action is somewhat conserved across vertebrates.

The movement of nutrients from the lumen of the intestine, aided by mucus, into epithelial cells takes place through diffusion and/or active transport regulated by nutrient transporters (Rust, 2003). The nutrient transporter *pept1* that aids the transport of protein in the form of di/tri peptides through the above process (Verri et al., 2011), was significantly regulated by diet LT compared with the control. Moreover, the higher expression of *pept1* in fish fed diet LT corresponded with significantly improved feed efficiency (lower FCR) and PER compared with the control, with diet L also having enhancing feed efficiency

(lower FCR) and PER compared with the control. This suggested that limonene drove the improved protein absorption, which could have contributed to increased growth. Similarly, dietary peppermint and Digestarom P.E.P (Biomin GmbH, Herzogenburg, Austria), a commercial matrix-encapsulated phytogenic mixture, improved protein utilisation in broilers (Upadhaya and Kim, 2017) and gilthead seabream *Sparus aurata* (Goncalves and Santos, 2015).

In order to maximise the use of dietary protein for somatic growth, energy for supporting metabolic processes can be derived from non-protein sources, particularly lipids (Nankervis et al., 2000). Lipid metabolism including, among others, processes such as lipid catabolism or fatty acid and triglyceride synthesis occurs along with lipid transport and deposition with the liver as the main active site (He et al., 2015). In the present study, diet LT activated lipid metabolism as reflected by significantly increased expression of lipoprotein lipase (lpl) in comparison to the control. Since the expression of lpl in the fish fed diet T did not differ from that of the control, it is reasonable to deduce that limonene, not thymol, is the compound that triggers such metabolic response in fish fed diet LT. Given that lpl plays a pivotal role in breaking down plasma lipids into free fatty acids and transporting them for use in energy production (Tian et al., 2015), the high gene expression of lpl found in this study suggests that dietary limonene increased the energy level of the fish, thereby providing sufficient energy for running metabolic processes and sparing protein, which significantly improved fish growth in the dietary treatments L and LT. Such effect of limonene to regulate lpl and a corresponding somatic growth enhancement further confirmed the results obtained by Aanyu et al. (2018) in the same teleost species.

Metabolic processes in the body result into production of reactive oxygen intermediates (ROIs), which can induce damage to cells and tissues if their levels are not maintained low (Covarrubias et al., 2008; Costa et al., 2013) This can ultimately impair adequate physiological function and subsequently negatively affect growth. In this study, the expression of catalase (*cat*), a key antioxidant enzyme that breaks down the ROI hydrogen peroxide, was significantly increased by dietary treatment with limonene (i.e., treatments L and LT) to similar extents. These results suggest that the enhanced antioxidant status by *cat* could reduce the hydrogen peroxide levels and thus result in improved somatic growth of the fish fed on diet L and LT. Recent research has shown that, when ROIs are at low concentrations, they are vital molecules mediating physiological processes including somatic growth (Covarrubias et al., 2008; Barbieri and Sestili, 2012). The herein reported action of dietary limonene on *cat* up regulation (catalase enzyme activity) has not been observed for other phytogenic compounds. For instance, Zheng et al. (2009) did not find

enhanced activity of catalase enzyme in channel catfish fed on diets containing thymol, carvacrol or their mixture, although the fish attained higher weight with the diet containing both compounds and carvacrol alone. Thymol did not appear to have an obvious role in regulation of antioxidant enzymes such as *cat*, and thus it can be assumed that, as noted above, limonene exerts a major action in up-regulating *cat*.

5. Conclusions

This study confirmed that dietary limonene and the blend of limonene and thymol improved somatic growth and feed utilisation efficiency of Nile tilapia to similar extents, although thymol individually showed no effects on enhancing growth performance. This indicated that dietary limonene was the major contributor towards the enhanced fish growth observed, suggesting lack of synergistic or additive effects of the combined compounds. The gene expression of biomarkers for nutrient digestion, absorption and transport, lipid metabolism, antioxidant enzymes and somatotropic axis growth also largely showed lack of synergistic or additive effects of the dietary combination of limonene and thymol in Nile tilapia.

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- Government of Uganda through the Agricultural Technology and Agribusiness Services
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Data Availability Statement

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

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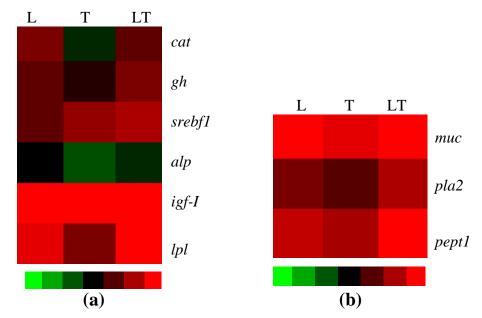
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FIGURES



data from Nile tilapia fed on diets supplemented with limonene (L), thymol (T) and their combination (LT). Data were plotted using Java Tree View and rows were clustered according to Euclidean distance. The columns represent mean data values of three dietary treatments L (400 ppm limonene), T (500 ppm thymol) and LT (400 ppm limonene and 500 ppm thymol). The rows indicate each of the analysed genes in the liver (a) and fore intestine (b) of Nile tilapia. Expression level of each gene was natural log transformed and normalised against two reference genes. The colour bars at the bottom represent the mean relative expression levels as low (green), neutral (black) or high (red). The black colour represents the control to which the relative expression of the other treatments was determined. *cat*, catalase; *gh*, growth hormone; *srebf1*, sterol regulatory element binding transcription factor 1; *alp*, alkaline phosphatase; *igf-1*, insulin growth factor 1; *pla2*, phospholipase A2; *lp1*, lipoprotein lipase; *muc*, mucin-like protein; *pept1*, oligo-peptide transporter 1.

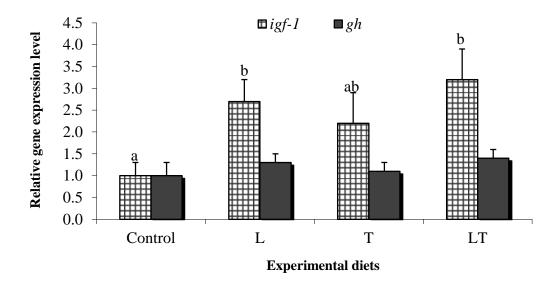
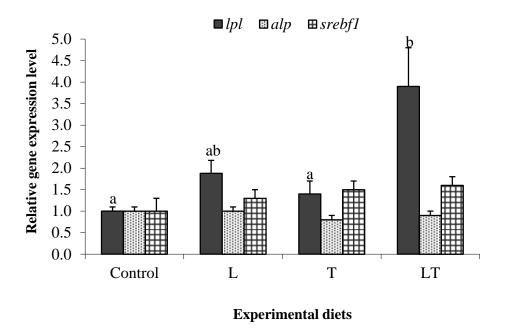


FIGURE 2 Expression of insulin growth factor I (igf-I) and growth hormone (gh) in the liver of Nile tilapia fed on diets supplemented with 400 ppm limonene (L), 500 ppm thymol (T) and the combination of 400 ppm limonene and 500 ppm thymol (LT). All values are means of treatments \pm standard error (n=8). Different superscript letters denote significant differences in the expression of igf-I between the treatments.



expression of *lpl* between the treatments.

FIGURE 3 Expression of lipoprotein lipase (lpl), alkaline phosphatase (alp), and sterol regulatory element binding transcription factor 1 (srebf1) in the liver of Nile tilapia fed on diets supplemented with 400 ppm limonene (L), 500 ppm thymol (T) and a combination of 400 ppm limonene and 500 ppm thymol (LT). All values are means of treatments \pm standard error (n=8). Different superscript letters denote significant differences in the

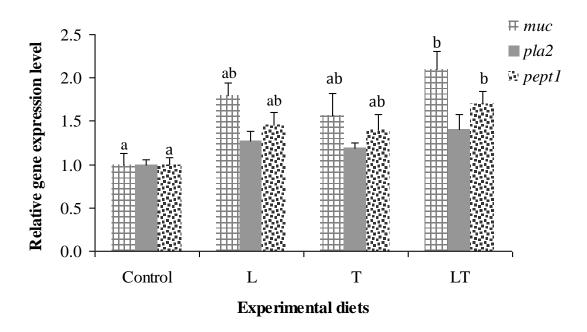


FIGURE 4 Expression of mucin-like protein (*muc*), phospholipase A2 (*pla2*), and oligopeptide transporter 1 (*pept1*) genes in the fore intestine of Nile tilapia fed on diets supplemented with 400 ppm limonene (L), 500 ppm thymol (T) and a combination of 400 ppm limonene and 500 ppm thymol (LT). All values are means of treatments ± standard error (n=8). For each gene (*muc* or *pept1*), different superscript letters denote significant differences in the expression of each gene between the treatments.

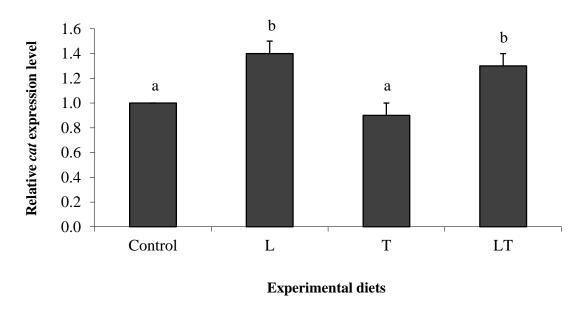


FIGURE 5 Expression of the antioxidant enzyme catalase (cat) in the liver of Nile tilapia fed on diets supplemented with 400 ppm limonene (L), 500 ppm thymol (T) and a combination of 400 ppm limonene and 500 ppm thymol (LT). All values are means of treatments \pm standard error (n=8). Different superscript letters denote significant differences in the expression of cat between the treatments.

TABLES

TABLE 1 Proximate analysis of the nutritional composition of the diet (CP35, ARDC - Uganda) used in the present trial to feed Nile tilapia (*Oreochromis niloticus*) for 63 days.

7sis	Quantity	
Dry matter (%)	89.1	
Moisture (%)	10.9	
Crude protein (%)	33.1	
Crude fat (%)	3.3	
Crude ash (%)	10.9	
Crude fibre (%)	9.9	
Gross energy (Kj g ⁻¹)	16.9	

TABLE 2 Details of the primers used for quantitative real-time PCR analyses.

Functional group	Gene symbol	Primer / oligonucleotide sequences (5'-3')	Size (base pairs)	Accession number*
Nutrient digestion,	тис	F: TGCCCAGGAGGTAGATATGC	101	XM_005466350
absorption and		R: TACAGCATGAGCAGGAATGC		
transport	pept1	F: CAAAGCACTGGTGAAGGTCC	196	XM_013271589
		R: CACTGCGTCAAACATGGTGA		
Lipid metabolism	lpl	F: TGCTAATGTGATTGTGGTGGAC	217	NM_001279753
		R: GCTGATTTTGTGGTTGGTAAGG		
	srebf1	F: TGCAGCAGAGAGACTGTATCCGA	102	XM_005457771
		R: ACTGCCCTGAATGTGTTCAGACA		
	alp	F: CTTGGAGATGGGATGGGTGT	200	XM_005469634
		R: TTGGCCTTAACCCCGCATAG		
	pla2	F: CTCCAAACTCAAAGTGGGCC	177	XM_005451846
		R: CCGAGCATCACCTTTTCTCG		
Antioxidant activity	cat	F: TCCTGGAGCCTCAGCCAT	79	JF801726
		R:		
		ACAGTTATCACACAGGTGCATCTTT		
Somatotropic axis-	gh	F: TCGGTTGTGTGTTTTGGGCGTCTC	90	XM_003442542
aided growth		R: GTGCAGGTGCGTGACTCTGTTGA		
	igf-I	F: GTCTGTGGAGAGCGAGGCTTT	70	NM_001279503
		R: CACGTGACCGCCTTGCA		
Reference genes	ef-1α	F: GCACGCTCTGCTGGCCTTT	250	NM_001279647
		R: GCGCTCAATCTTCCATCCC		
	β -actin	F: TGGTGGGTATGGGTCAGAAAG	217	XM_003443127
		R: CTGTTGGCTTTGGGGTTCA		

muc, mucin-like protein; pept1, oligo-peptide transporter 1; lpl, lipoprotein lipase; srebf1, sterol regulatory element binding transcription factor 1; alp, alkaline phosphatase, pla2 phospholipase A2; cat, catalase; gh, growth hormone; igf-I, insulin growth factor I; $ef-1\alpha$, elongation factor 1α ; β -actin beta-actin.

*GenBank (http://www.ncbi.nlm.nih.gov/); bp, base pairs

TABLE 3 Growth, feed utilisation efficiency and survival rate of Nile tilapia fed on diets with 400 ppm limonene (L), 500 ppm thymol (T) and a combination of 400 ppm limonene and 500 ppm thymol (LT) for 63 days.

Parameter	Experimental diet					
rarameter	Control	L	T	LT	P value	
Initial mean weight (g)	1.5 ± 0.0	1.6 ± 0.0	1.5 ± 0.0	1.6 ± 0.0	NS	
Final mean weight (g)	13.7 ± 0.4^a	16.6 ± 0.4^b	15.0 ± 0.4^a	16.7 ± 0.3^b	0.001	
% WG	793.2 ± 29.1^{a}	957.3 ± 51.9^{b}	887.0 ± 16.1^{ab}	980.0 ± 41.3^{b}	0.011	
CF	1.8 ± 0.0	1.8 ± 0.0	1.9 ± 0.0	1.9 ± 0.0	NS	
% Survival	94.1 ± 3.5	97.4 ± 1.5	94.8 ± 3.0	98.1 ± 0.7	NS	
% FI (% body weight d ⁻¹)	4.5 ± 0.1^{b}	3.9 ± 0.1^a	4.3 ± 0.2^{ab}	4.0 ± 0.1^a	0.019	
FCR	$1.8 \pm 0.1^{\rm b}$	1.5 ± 0.0^a	1.7 ± 0.1^{ab}	1.5 ± 0.0^a	0.027	
PER	1.7 ± 0.1^{a}	2.0 ± 0.1^b	1.9 ± 0.1^{ab}	2.0 ± 0.1^b	0.009	

All values are means of treatments \pm standard error. Mean values with different superscript in the same row are significantly different from each other at P < 0.05. NS, refers to not significantly different values. For each treatment, n =152 for initial fish weight, for final fish weight, n = number of alive fish at the end of the trial, and n = 4 for percentage of weight gain (% WG), condition factor (CF), survival rate (% survival), food intake (% FI), feed conversion ratio (FCR) and protein efficiency ratio (PER).