Feed utilization and rumen microbial ecology of lambs consuming *Daniellia Oliveri* seed based diet

Okunade S.A. ^{*1}, Isah O.A.², Odedara O.O. ³, Adebayo K.O ², and Olafadehan O.A. ⁴

^{*}Corresponding author: saokunade2013@gmail.com

¹Department of Animal Production Technology, Federal College of Wildlife Management, New Bussa, affiliated to Forestry Research Institute of Nigeria

²Department of Animal Nutrition, Federal University of Agriculture, Abeokuta, Nigeria

³Department of Microbiology, Federal University of Agriculture, Abeokuta, Nigeria

⁴Department of Animal Science, University of Abuja, Abuja, Nigeria.

ABSTRACT

This study investigated the effects of replacing processed *Daniellia oliveri* seed meal (PDS) for groundnut cake (GNC) in the concentrate diets of Yankasa male lambs (6 months old, body weight of 13.70 ± 0.1 kg (body weight)) on feed intake, rumen fermentation parameters and microbial population. Four diets were formulated to contain 0, 50, 100 and 150 g/kg PDS to replace 0, 25, 50 and 75% GNC (dry matter (DM) basis) in a complete randomized design with five animals per diet. Total DM intake (g/kg BW^{0.75}), ruminal volatile fatty acids, valerate and fungal population linearly increased but ruminal NH₃-N, acetate, acetate: propionate, methane production and *Entodinium* (as % of the total protozoa count) linearly decreased with increasing PDS level ($P < 0.01$). Whereas, intake of concentrate, total DM (g/d and g/kg BW), average daily weight gain, ruminal pH, butyrate, iso-valerate, total bacterial count, proteolytic and amylolytic bacterial counts increased linearly and quadratically, basal roughage intake, protozoal count, cellulolytic bacterial count, methanogen count, *Diplodinium* and *Holotrichs* (as % of the total protozoa count) decreased linearly and quadratically ($P < 0.05$) with increasing PDS level. Feed conversion was lowest (L and Q:P < 0.01) for PDS15. Ruminal *Ophryoscolex* (as % of the total protozoa count) was lower (Q:P < 0.0001) for PDS15 than for other diets. Dietary inclusion of 150 g/kg *Daniellia oliveri* seed meal to replace 75% DM of groundnut cake in the diet of Yankasa lambs improved intake, rumen fermentation and microbial ecology and weight gain.

Keywords: condensed tannins, weight gain, rumen fermentation, rumen microbiota, methane, *Daniellia*

INTRODUCTION

The rumen is endowed with a composite community of microbes (anaerobic bacteria, protozoa, fungi, methanogenic archaea and phages) whose major function is fermentation of plant materials ingested by the animals. This symbiotic relationship furnishes the host animal with metabolic energy and concomitant methane production in the rumen which is energy loss to the animal. The understanding of dynamic rumen ecosystem can be harnessed in ruminant nutrition for an increased production to satisfy global protein demand, whilst ensuring minimal environmental impact. Many studies report several attempts to exploit plant secondary metabolites (PSM) as natural feed additives to modulate rumen ecology for improved rumen fermentation efficiency, protein and energy metabolism, and improved animal health and productivity and methane mitigation (Benchaar et al., 2007). The anti-microbial and anti-nutritional properties of PSM made them to be regarded as anti-nutritional factors for decades (Patra et al., 2017). Different processing methods including milling, soaking, toasting, decortications, cooking etc have been reported to reduce anti-nutritional factors. However, polyphenols and tannins retain their inhibitory properties even after heat treatment. Since the PSM mode of anti-microbial action is highly specific, they can be exploited in modulating rumen microbial ecosystem for better feed utilization and performance by the host animal.

Daniellia oliveri, a tanniniferous evergreen leguminous tree belonging to the family Caesalpiniaceae, is a copiously available tree in the savannah area of Nigeria (Okunade et al., 2014b, 2019; Olafadehan et al., 2020). Previous studies (Okunade and Olafadehan, 2019) on the nutritive value of *Daniellia seed meal* indicated its potential as an alternative protein source to the expensive and occasionally scarce conventional protein sources such as groundnut cake, soybean cake, etc. in livestock production. However, like other tropical legume seeds, it contains high levels of PSM (tannins, saponins and other anti-nutritive factors). Plant secondary metabolites, particularly condensed tannins (CT), could induce intoxication if consumed in excess (Olafadehan et al., 2014a; Okunade et al., 2014a), altering ruminal microbiota population and activity and affecting performance of animals. Due to its content of PSM, processed *Daniellia oliveri* seed meal (PDS) can be used as a protein source in the diet of ruminants to modulate rumen microbiota and thus improve feed utilisation, ruminal fermentation efficiency and growth performance, and mitigate methane production. Few studies have considered the effect of replacing secondary metabolites free conventional protein source, such as groundnut cake (GNC), with PSM containing alternative protein ingredient. Information on the effect of feeding tannin-containing PDS, as a

protein source, to modulate ruminal microflora with the aim of altering ruminal fermentation and improve performance of sheep is very scarce. The current study evaluated the effect of replacing GNC with PDS in the supplementary diet of growing lambs on feed utilisation, ruminal microbiota and fermentation and growth performance.

MATERIALS AND METHODS

Experimental site, animals and management

The experiment was carried out at the small ruminant unit of the Teaching and Research Farm, Federal College of Wildlife Management, New Bussa, Niger State Nigeria. The study area location (latitude: 9° 53' and 10° 00" N and longitudes 4° 31' and 40° 33" E) is in the guinea savannah agro-ecological zone of North central part of Nigeria. It has a mean annual temperature of about 37.2° C, with relative humidity of about 60% and the mean annual rainfall value is about 1109mm (Okunade et al, 2019).

Twenty healthy growing Yankasa intact male lambs (6 months old, body weight, 13.7 ± 0.1 kg) were stratified into four groups of five animals of similar body weight (BW) per group in a completely randomized design. They were housed individually in a clean, well-ventilated, half-walled and rammed floored bedded with softwood shaving pens. All male lambs were quarantined and fed with experimental diets for four weeks; they were intramuscularly injected and dewormed with 1 mL/10 kg BW of oxytetracycline and levamisole respectively and sprayed against external parasites with diazuntol.

Seed collection and processing

Seeds of *Daniellia oliveri* were collected from the established plots of *Daniellia oliveri* plants within the premises of the Federal College of Wildlife Management, New Bussa Nigeria. The forest conservation was established many years ago for the purpose of wildlife and soil conservation to prevent desertification and act as wind break. The dehiscent seeds of *Daniellia oliveri* were gathered from the ground, cleaned from extraneous materials and sun dried. The seeds were toasted by placing big round iron cooking pot containing sand upon a tripod stand under which fire was made with wood. Dried clean *D. oliveri* seeds were added to sand in ratio 2:1 (w/w). The sand and seeds mixture were heated under controlled temperature (not exceeding 80°C) and frequent stirring was done to ensure even toasting for 30–35 minutes until the seeds cracked and their white endosperm turned brown with appealing aroma. The seeds were carefully separated from the sand, allowed to cool and cracked exoderm was carefully removed (decorticated) manually. The processed seeds were ground in a hammer mill (Model W-6-H, Buffalo NY 14206) to pass through a 1 mm sieve particle size, and tagged

processed *D. oliveri* seed meal (PDS). The seed meal was stored in air -tight containers until ready for use.

Experimental diets

Four iso-nitrogenous concentrate diets; (160 g/kg CP on DM basis) were formulated such that (GNC, a conventional protein source) in the control diet was replaced with the PDS meals at 0, 25, 50 and 75%, and the dietary treatments were designated PDS0, PDS25, PDS50 and PDS75, respectively. Each of the lambs was offered 1 kg of concentrate once daily at 09:00 h, while the basal diet, a low-quality threshed sorghum tops (TST), was offered ad libitum throughout the twelve weeks experimental period. Clean water was provided ad libitum daily.

Rumen fermentation parameters

About 100 mL of representative rumen liquor was collected before morning feeding on the last day of the experiment from each lamb via stomach tube. Temperature and pH of rumen liquor were determined immediately after collection using a portable digital pH and temperature metre model CT 6020. The rumen liquor was strained through four layers of muslin cloth, preserved by adding a few drops of saturated mercury (II) chloride solution, placed, taken to the laboratory and frozen at -30 °C pending analysis. Samples were analysed for ammonia nitrogen using micro Kjeldahl method (AOAC, 2000) immediately on getting to the laboratory. Total volatile fatty acids in SRL were determined by steam distillation process as described by Keyol and Borilek, (1995). Individual volatile fatty acids (VFA; acetic, propionic and butyric acids) were determined using gas chromatography as described by (Mebrahtu, 1992).

Rumen microbial enumeration and isolation

Samples of strained rumen fluid were analysed in triplicate. Total viable bacteria and fungi were enumerated by the method of Joblin (1981), respectively, using anaerobic roll tubes. Total protozoa population in the rumen fluid was determined using a Sedgewick-Rafter counting chamber (Dehority et al., 1989) with the modifications proposed by D'Agosto and Carneiro (1999), and the protozoa present in each sample were isolated based on the criteria described by Ogimoto and Imai (1981). The bacteria isolation was conducted using conventional techniques of serial dilutions and repeated tubing of the selectively enriched microbial cultures by using the specific media for rumen bacteria. Strict anaerobic conditions were maintained during the isolation procedure by continuous flushing of the rumen fluid with CO₂ gas. The growth media contained 15 mL mineral solution I (KH₂PO₄ 3.0 g; (NH₄)₂SO₄ 6.0 g; NaCl 6.0 g; MgSO₄ 0.6 g; CaCl₂·2H₂O 0.795 g per Litre), 15 ml mineral solution II (K₂HPO₄ 3g/litre), 0.25 g yeast extract, 1 g tryptone, 0.1 ml

resazurine (0.1%), 0.2 ml hemin (0.05%), 0.5 g microcrystalline cellulose, 0.1 g cellobiose, 0.4 g sodium carbonate, 20 ml clear rumen Fluid, 50 ml distilled water and 50 mg cysteine hydrochloride.

Chemical analysis and calculations

Processed *Daniellia oliveri* seed meals (PDS), basal (TST) and the *Daniellia oliveri* seed-based concentrate diets were analysed for their proximate constituents according to the method of AOAC (2000). Neutral detergent fibre (NDF), acid detergent fibre (ADF) and acid detergent lignin (ADL) were determined by methods of Van Soest et al. (1991). Cellulose and hemicellulose were estimated as differences between ADF and lignin and NDF and ADF, respectively. Total condensed tannins were determined as reported by Polshettiwar et al. (2007). Methane production was estimated by the equation of Moss et al. (2000).

Estimated CH₄ production = 0.45 (acetate) – 0.275 (propionate) + 0.5 (butyrate)

Statistical analysis

Data were subjected to one way of ANOVA in completely randomized design using version 6 of SAS software (SAS Institute, 2012). Significance difference between individual means was separated by Duncan's procedure of the same software. Treatments means were considered significant at P<0.05, while polynomial (linear and quadratic) contrasts were used to examine the variable responses to increasing level of PDS in the diets.

RESULTS

Nutrient composition of experimental concentrate diets

Dry matter content ranged from 842.8 g/kg DM in PDS25 to 872.9 g/kg DM in PDS50. Whereas OM and NFC increased, ether extract (EE), NDF and ADF decreased with increasing replacement of GNC with PDS (P<0.05). Diets were isonitrogenous. Condensed tannins were higher (P<0.05) for PDS15 than PDS5 and PDS10. PDS is a good quality feedstuff with a high crude protein (CP), low neutral detergent fibre and moderate CT. The basal TST had a low CP and high CT and NDF (Table 1).

Table 1. Ingredient and chemical compositions of experimental concentrate diets, processed seed and basal diet (g/kg DM)

Nutrient	Level of PDS in the diet (g/kg)				SEM	PDS	TST
	PDS0	PDS5	PDS10	PDS15			
Maize offal	450	450	450	450			
Cowpea husk	255	250	250	250			
Brewers dried grain	50.0	50.0	50.0	50.0			
Groundnut cake	200	150	100	50.0			
<i>D. oliveri</i> seed meal	-	50.0	100	150			
Urea	5.00	10.0	10.0	10.0			
Dicalciumphosphate	10.0	10.0	10.0	10.0			
*Vit-Miniral Premix	10.0	10.0	10.0	10.0			
Sulphur powder	10.0	10.0	10.0	10.0			
Table salt	10.0	10.0	10.0	10.0			
<i>Analyzed compositions (g/kg DM)</i>							
Ash	75.2 ^a	51.9 ^b	44.0 ^c	39.0 ^d	2.14	72.41	61.0
Organic matter	924.8 ^d	948.1 ^c	956.0 ^b	961.0 ^a	0.83	927.6	959.0
Crude protein	181.3	180.5	181.2	181.1	0.28	390.0	68.1
Ether extract	60.9 ^a	53.7 ^b	50.3 ^c	37.5 ^d	0.27	152.8	146.0
Non fibre carbohydrate	238.0 ^d	283.6 ^c	298.2 ^b	329.4 ^a	7.48	122.0	96.4
Neutral detergent fibre	443.6 ^a	431.4 ^b	426.2 ^c	413.1 ^d	1.73	263.3	695.7
Acid detergent fibre	255.0 ^a	223.2 ^b	200.4 ^c	182.3 ^d	2.14	172.2	580.0
Condensed tannins	ND	25.0 ^b	26.0 ^b	30.2 ^a	1.20	29.50	65.5

a,b,c,d, means within a row not bearing a common superscript letter significantly differ, (P<0.05)
PDS, processed Daniellia oliveri seed meal; ND, not detectable

Feed utilisation

Intake of TST and feed conversion decreased (L: $P < 0.0001$; Q: $P < 0.01$) as the inclusion of PDS in the concentrate increased (Table 2). Intakes of concentrate and total DM (g/d) increased linearly (L: $P < 0.0001$) and quadratically ($P < 0.05$) with increasing PDS level. Total DM intake (g/kg LW) and average daily gain (ADG) increased (L: $P < 0.0001$; Q: $P < 0.05$) with increasing proportion of PDS in the diets. Total DM intake (g/kg W^{0.75}) and CP intake were greater (L: $P < 0.0001$) for PDS15 and PDS10 compared with PDS0 and PDS5. Intake of CT was higher ($P < 0.0001$ for L and Q) for PDS15 than for PDS5 and PDS10.

Table 2. Feed utilization of ram-lambs fed processed *Daniellia oliveri* seed meal based diet

Parameters	Diet				SEM	P-value	
	PDS0	PDS5	PDS10	PDS15		L	Q
TST (g/d)	285.7 ^a	279.0 ^a	267.7 ^b	243.0 ^c	3.04	<0.0001	0.003
Concentrate (g/d)	667.0 ^d	683.2 ^c	724.3 ^b	769.1 ^a	2.19	<0.0001	<0.001
Total DM (g/d)	953 ^d	962 ^c	992 ^b	1012 ^a	3.93	<0.0001	0.044
Total DM (g/kg LW)	51.98 ^c	52.01 ^c	53.84 ^b	56.90 ^a	0.62	<0.0001	0.019
Total DM (g/kg W ^{0.75})	108 ^b	108 ^b	111 ^a	112 ^a	0.38	<0.0001	0.287
Crude protein (g/kg W ^{0.75})	18.96 ^d	19.51 ^{cd}	20.23 ^b	20.50 ^a	0.22	0.001	0.501
CT (g/kg W ^{0.75})	0.18 ^c	0.27 ^b	0.29 ^b	0.35 ^a	0.02	<0.0001	<0.0001
ADG (g/day)	104 ^c	105 ^c	113 ^b	120 ^a	1.37	<0.0001	0.035
Feed conversion ratio	9.14 ^a	9.18 ^a	8.77 ^b	8.43 ^c	0.05	<0.0001	0.001

a,b,c, means within a row not bearing a common superscript letter significantly differ, ($P < 0.05$) TST, threshed sorghum tops; PDS, Processed *Daniellia oliveri* seed. CT, Condensed tannins. ADG, Average daily weight gain

Rumen fermentation parameters and enteric methane production

Diet effect on ruminal temperature was marginal. Ruminal pH was greater (L: $P < 0.0001$; Q: $P = 0.003$) in PDS15 than in other diets (Table 3). Ruminal NH₃-N was more pronounced in PDS0 and PDS15, intermediate in PDS10 and lowest in PDS15, resulting in a linear response (L: $P < 0.0001$). Ruminal TVFA

was greater for PDS15 relative to PDS0 and PDS, producing linear trend ($P < 0.007$). Whereas ruminal acetate, acetate to propionate ratio (C2:C3) and methane gas production decreased linearly ($P < 0.0001$), propionate (L: $P < 0.0001$), valerate (L: $P < 0.0001$) and isovalerate (L: $P < 0.0001$; Q: $P = 0.003$) increased with increasing PDS level. Diet effect on ruminal butyrate was higher in PDS0 than other PDS diets (linear and quadratic, $P < 0.01$).

Table 3. Rumen fermentation parameters and enteric methane production of ram-lambs fed processed *Daniellia oliveri* seed meal-based diet

Parameter	Diet				SEM	P-value	
	PDS0	PDS25	PDS50	PDS75		Linear	Quadratic
Tempt. ($^{\circ}\text{C}$)	38.98	38.90	39.00	39.02	2.73	0.560	0.410
pH	6.63 ^a	6.60 ^a	6.58 ^a	6.49 ^b	0.05	<0.0001	0.003
NH ₃ -N (mg/L)	173.0 ^a	170.0 ^a	164.0 ^b	160.0 ^c	1.97	<0.0001	0.645
TVFA (mM)	117.3 ^c	118.1 ^{bc}	120.3 ^{ab}	122.0 ^a	1.02	0.007	0.614
VFA (mol/100 mol)							
Acetate (C ₂)	78.0 ^a	76.0 ^b	73.0 ^c	71.0 ^d	0.71	<0.0001	0.757
Propionate (C ₃)	26.0 ^d	30.0 ^c	34.0 ^b	37.0 ^a	1.07	<0.0001	0.530
Butyrate (C ₄)	9.8 ^a	8.5 ^b	8.2 ^{bc}	8.0 ^c	0.17	<0.0001	0.001
Iso-valerate (C ₅)	1.2 ^d	1.9 ^c	2.3 ^b	2.8 ^a	0.05	<0.0001	0.036
Valerate (C ₆)	1.4 ^d	2.2 ^c	2.5 ^b	3.0 ^a	0.11	<0.0001	0.111
C ₂ :C ₃	3.0 ^a	2.5 ^b	2.2 ^c	1.9 ^d	0.12	<0.0001	0.212
CH ₄ (mol/100 mol)	24.0 ^a	22.6 ^b	20.0 ^c	18.6 ^d	0.16	<0.0001	0.787

a,b,c, means within a row not bearing a common superscript letter significantly differ, ($P < 0.05$)

Rumen microbial population and classification

Total ruminal bacterial population was highest for PDS15 and lowest for PDS0 and PDS5, resulting in linear and quadratic responses (L: $P < 0.0001$; Q: $P = 0.008$) (Table 4). Total protozoal population (linear and quadratic, $P < 0.01$) and *Ophryoscolex* (as % of the total protozoal count) (Q: $P < 0.0001$) were lower for PDS15 than for other treatments. Fungal population was highest in PDS10 and PDS15 and lowest in PDS0 (linear effect, $P < 0.0001$). Proteolytic bacterial numbers were lowest and highest in PDS0 and PDS15, producing linear and quadratic trends (L and Q: $P < 0.0001$) respectively. Diet effect on cellulolytic bacteria and Holotrichs (as % of the total protozoal count) were linear and quadratic (L: $P < 0.0001$; Q: $P < 0.05$), with PDS0 and PDS5 having the highest populations and PDS15 the lowest. Amylolytic bacterial population was higher in PDS15 compared to other diets (L and Q: $P < 0.0001$). The populations of ruminal archaea methanogens and *Entodinium* (as % of the total protozoal count) decreased (linear, $P < 0.0001$; quadratic, P

= 0.0016), as the PDS inclusion level increased. *Diplodinium* (as % of the total protozoal count) was highest in PDS0, intermediate in PDS5 and lowest in PDS10 and PDS15, resulting in linear and quadratic trends (L and Q: $P < 0.01$).

Table 4. Rumen microbial population and classification of ram-lambs fed processed *Daniellia oliveri* seed meal-based diet

Rumen Microbial Biomass	Diet				SEM	P-value	
	PDS0	PDS5	PDS1 0	PDS15		Linear	Quadrat ic
Total viable bacteria, × 10 ⁸ cfu/mL	9.07 ^c	9.32 ^c	10.0 ^b	10.8 ^a	0.24	<0.0001	0.008
Total protozoa, × 10 ⁴ cell/mL	6.86 ^a	6.68 ^a	6.48 ^a	5.07 ^b	0.20	<0.0001	0.002
Total fungi, × 10 ⁶ cfu/mL	6.67 ^c	8.02 ^b	8.70 ^a	9.27 ^a	0.26	<0.0001	0.078
Bacteria class according to substrate fermented							
Proteolytic, × 10 ⁶ /mL	2.30 ^c	2.52 ^b	2.56 ^b	3.08 ^a	0.02	<0.0001	<0.001
Cellulolytic, × 10 ⁷ /mL	6.80 ^a	6.74 ^a	5.87 ^b	4.75 ^c	0.10	<0.0001	0.002
Amylolytic, ×10 ⁶ /mL	0.78 ^c	0.85 ^{bc}	0.89 ^b	1.28 ^a	0.03	<0.0001	<0.001
Methanogenic archaea, × 10 ⁸ /mL	1.02 ^a	0.73 ^b	0.67 ^c	0.43 ^d	0.02	<0.0001	0.016
<i>Protozoa genera (as % of total protozoa count)</i>							
<i>Entodinium</i>	90.0 ^a	87.0 ^b	82.0 ^c	79.5 ^d	0.52	<0.0001	0.264
<i>Diplodinium</i>	23.5 ^a	23.0 ^{ab}	22.0 ^b	21.0 ^b	0.58	<0.001	0.006
<i>Ophryoscolex</i>	6.10 ^a	6.32 ^a	6.42 ^a	6.00 ^b	0.07	0.358	<0.001
<i>Holotrich</i>	48.0 ^a	47.9 ^a	45.8 ^b	42.0 ^c	0.82	<0.0001	0.033

a,b,c,d means within a row not bearing a common superscript letter significantly differ, ($P < 0.05$)

DISCUSSION

The nutrient composition of TST is consistent with previous studies (Isah et al., 2015; Okunade et al., 2016; Okunade et al., 2018). The PDS is tannin-containing feedstuffs, but with much lower contents than the threshold level at which tannins impair nutrient utilization and performance in ruminants. This might probably be due to the effect of heat treatment and decortications of the *Daniellia oliveri* seed during processing. The low CP and high fibre fractions contents of TST confirm it as a typical low-quality roughage diet, with low feeding value that cannot meet the requirements of ruminants in various physiological states without supplementation. The CP content of the

supplementary concentrates met requirement of 180 g/kg DM recommended for growing sheep (MLA, 2007), while NDF content was far below the 650 g/kg DM threshold level at which cell wall inhibits feed intake, digestibility and animal performance (Meissner et al., 1991).

Higher concentrate intake of PDS15 indicates increased palatability and hence, acceptability and consumption (Kholif et al., 2018a; Olafadehan et al., 2018). Higher voluntary intake of PDS15 resulted in the higher CP intake for the lambs. It has been shown that CP intake is a major determinant of ruminant performance due to increased availability of fermentable CP and other nutrients required by rumen bacteria as well as a greater opportunity for dietary protein to escape rumen fermentation (Conte et al., 2010). Although CT concentration and intake increased with increasing PDS in the diet, highest concentrate intake and ADG of sheep on the PDS15 confirms the moderate and beneficial level of its CT because CT at a high concentration is harmful and impairs palatability, feed consumption, protein metabolism, nutrient digestibility and weight gain (Isah et al., 2015; Olafadehan et al., 2014a, 2020). Therefore, the CT concentration of 3.02 g/kg DM of the PDS15 did not compromise palatability as to adversely affect consumption and ADG. The linear increase in the concentrate diets intake suggests that PDS is more palatable than GNC. Toasted legume seeds are characterized by unique flavours which increase their sensory appeal to animals (Raji et al., 2014). Furthermore, the highest total DM intake in PDS15 diet could be attributed to the fibre level, which decreased progressively with increasing PDS replacement for GNC. This must have reduced gut fill and improved the rate and extent of ruminal degradation, resulting in a faster passage of the digesta through the gastrointestinal tract and a shorter retention time (Kang et al. 2012; Okunade et al., 2014b; Olafadehan et al., 2014a). In spite of the lowest basal TST intake of the PDS15 diet supplemented sheep, overall DM intake was highest for the animals. Since both the basal TST and supplementary diet contained CT, the result indicates that the dietary CT was well tolerated by animals and did not induce tannin toxicosis (Olafadehan et al., 2014a) which affects rumen microbiota and fermentation, feed utilization and growth performance. The generally high acceptability of all the diets by the lambs is further confirmed by the relatively high DM consumption (expressed as g/kg LW), which increased from 51.98 to 56.9 g/kg LW as the PDS level increased. Feed intake (g/kg LW), in the current study, is higher than the 3-4 g/kg LW generally recommended for ruminants. Intakes expressed as the metabolic were within the range of 3.9 to 6.3 g/kg LW obtained for the Yankasa lambs (Olafadehan et al., 2014b; Isah et al., 2015; Okunade and Olafadehan, 2019). Dry matter intake is a key determinant of growth of animals; therefore, in response to feed intake, ADG also increased as the PDS increased. The improved FCR of the PDS diets confirm the previous submission that the CT of

the diets was moderate, well tolerated and did not interfere with nutrient metabolism, absorption and utilization, in agreement with earlier reports (Olafadehan et al., 2020).

Tannin-containing diets modified rumen fermentation parameters in this study, in accordance with the reports of Belanche et al. (2019). Ruminant temperature measurement is a suitable and feasible means to detect potential adverse alteration in the health of animals in real time (Wahrmund et al., 2012). The unaffected rumen temperature was maintained within the recommended range of 39 to 39.5°C (Wahrmund et al., 2012) which suggests that lambs had no health challenge. No external factor can alter intraruminal temperature, although it is approximately 0.5°C higher than core body temperature due to the activity of heat-producing rumen microorganisms (Bodas et al., 2014).

Ruminal pH was within optimal range (6.2 - 7.2) for normal rumen ecology and fermentation (Gunun et al., 2013; Isah et al., 2013). The PDS15 diet with the highest NFC and lowest fibre fractions had the lowest ruminal pH. High VFA and lactic acid production, as a result of rapid ruminal fermentation of dietary readily fermentable carbohydrates and reduced chewing time and salivation rate due to low dietary fibre level, have been reported to lower ruminal pH (Olafadehan et al., 2016; Olafadehan and Adebayo, 2016).

Concentration of NH₃-N in the rumen fluid is the net result of NH₃-N production from feed, protein fermentation, absorption rate through the rumen wall, passage out of the rumen and utilization by microbes. Ruminant NH₃-N concentrations in all the treatments were within the optimal ruminal range of 150 - 200 mg/L for optimal ruminal ecology, fermentation and microbial growth (Perdok and Leng, 1990). Decreased ruminal NH₃-N concentration of PDS diets indicates lower ruminal CP degradability of PDS, due to its CT, relative to GNC. The result indicates higher bypass protein value of PDS than GNC.

Increased ruminal VFA concentration of PDS15 diet may be a direct result of its highest readily fermentable NFC content ((Wanapat et al., 2014; Olafadehan et al., 2016; Okunade and Olafadehan, 2019)) and enhanced ruminal fermentation (Kholif et al., 2018b). The ruminal VFA concentrations of all the animals were within the normal range for ruminants. The linear increase in ruminal propionate may be due to progressive increase in feed intake as the PDS proportion increased in the diet. Therefore, low ruminal propionate in animals fed PDS0 diet might be as a result of low DM intake (Isah et al., 2014) and reduced ruminal fermentation due to its relatively low NFC content. The higher rumen propionate concentration of the lambs fed the PDS, particularly PDS15, is beneficial as it must have enhanced energy availability for improved nutrient utilisation and ADG. The linear decrease in

molar proportion of ruminal acetate may be related to the decrease in fibre level of the diets as PDS inclusion level increased. Opsi et al. (2012) affirmed that ruminal acetate production is due majorly to structural carbohydrate fermentation by cellulolytic bacteria, while propionate production is due to NFC degradation by amylolytic bacteria. In the current study, structural carbohydrates were higher in control diet (PDS0) while NFC was higher in the PDS diets. Highest cellulolytic bacteria and lowest amylolytic bacteria numbers in the control diet than in the PDS diets further explained the result.

Propionate formation can be considered as a competitive pathway for CH₄ production (Moss et al., 2000). Therefore, the increased ruminal propionate concentration of the PDS diets led to the reduction of CH₄ production. Condensed tannins suppressed CH₄ production by redirecting hydrogen from the CH₄ pathway to produce propionate (Wanapat et al., 2014). Similarly, Animut et al. (2008) and Olafadehan and Okunade (2018) reported increased propionate and reduced CH₄ production with CT-containing forage. Plant secondary metabolites, particularly CT and saponins, have been shown to reduce protozoa and archaea methanogens, resulting in methanogenesis suppression (Wanapat et al., 2014; Gunun et al., 2016). It is plausible to aver that PDS can be used as an environmentally friendly feedstuff to abate enteric methane emission for sustainable environment and ruminant production.

The microbial population in this study revealed an unequal number among the typical rumen microbes enumerated. Bacterial species had the largest population compared to other microbial populations. It is, therefore, likely that bacteria are responsible for the major part of ruminal feed degradation and are the main drivers behind the observed differences (Kamra, 2005). The highest bacteria population with PDS15 diet could be as a result of its decreased protozoa population. Decreased ruminal protozoa population has been reported to result in an increase in bacterial population (Carberry et al., 2012; Isah et al., 2013) because protozoa engulf bacteria. This implies that CT concentration in the PDS15 diet made a significant reduction in the protozoa population. Since both protozoa and methanogens are closely related, archaea methanogens population was also lowest in PDS15. Earlier studies (Makkar et al., 1995; Guo et al., 2008) reported reduction in protozoa and methanogens populations due to the effect of tannins.

The increased ruminal fungi population with PDS15 could be due to the interaction between protozoa and fungal zoospores. Newbold and Hillman (1990) also noted an increase in the numbers of fungal zoospores when protozoa were removed from the rumen. The increased bacterial and fungal populations and reduced protozoal and methanogens populations with the PDS15 diet are nutritionally and environmental beneficial due to improved

nutrient availability and utilisation for weight gain and methane production abatement.

Proteolytic bacterial population increased progressively with the increasing PDS level, with the highest population in PDS15 diet. This increase might be as a result of reduction in the number of protozoa as well as increase in CP intakes of the PDS diets, as opined by Nguyen et al. (2017). Also, the inhibitory effect of CT may lower the activities of cellulolytic bacteria and consequently allows the proteolytic bacteria to compete more effectively for nutrients, thus increasing their relative abundance (Wolin and Miller, 1988). On the contrary, cellulolytic bacterial population decreased with increasing PDS level, which could be as a result of many factors including interactions between the cellulolytic and proteolytic bacterial populations and changes in digesta composition and rate of outflow from the rumen (Mcsweeney et al., 2001).

Present study showed that *Entodinium* is the most dominant protozoa genera followed by *Holotrichs* then *Diplodinium*, while *Ophryoscolex* was the least in population in the rumen of the lambs. This result corroborates the presence of protozoal population found in rumen liquor of sheep fed *Dicantium annulatum* grass supplemented with *Leucaena* and *Haedwickia* tree leaves (Singh and Kundu, 2011). Both *Diplodinium* and *Ophryoscolex* general had the lowest population, in agreement with the observations of earlier workers (Mughetti et al., 2007) who noted *Ophryoscolex* in the range of 5.51 to 7.17% in rumen of sheep. Lowest populations of the protozoa genera in the PDS diets relative to the control diet could be explained by the reduction in the total protozoal count in the treatment diets, particularly in the PDS15 diet with highest CT concentration. Since protozoa are generally associated with fibre digestion, the highest populations of the protozoa genera in the control diet could be attributed to its pronounced dietary fibre level and consumption of the basal TST roughage diet. Generally, high forage diets favour cellulolytic microbes and make them preponderance while low-forage diets result in lower number of cellulolytic microbes. This may account for the higher population of *Entodinium* and *Diplodinium* in the PDS0 (control diet) relative to the PDS diets. About 16 to 30% of total rumen \microbial fibre digestion was done by protozoa (Jabari et al., 2014) who indicated that ability of *Entodinium*, *Epidinium* and *Diplodinium* to degrade cellulose, hemicellulose and NDF was high. *Ophryoscolex* and *Holotrichs* are ciliate protozoa speculated to help in fibre digestion, although their specific role in rumen ecosystem remains controversial (Newbold, 2015).

CONCLUSION

Processed (toasted and seed coat decorticated) *Daniellia oliveri* seed meal, particularly 150 g/kg inclusion level, improved feed utilisation, ruminal

fermentation efficiency and growth performance, altered ruminal microbiota and reduced methane production. The seed meal can be used as an alternative protein source to replace up to 75% of groundnut cake in a supplementary concentrate for growing lambs consuming low-quality basal diet. Further research on complete replacement of processed *Daniellia oliveri* seed meal for groundnut cake is warranted.

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