

## **An *in-vitro* evaluation of the potentials of turmeric as phytogetic feed additive for rumen modification**

Aderinboye, R. Y. and Olanipekun, A. O.

Department of Animal Nutrition, College of Animal Science and Livestock Production,  
Federal University of Agriculture, Abeokuta



**Corresponding author:** [aderinboyer@funaab.edu.ng](mailto:aderinboyer@funaab.edu.ng); +2349033795787

### **Abstract**

*The potential risk to animal and human health in the use of antibiotic feed additives for modifying rumen fermentation has necessitated the search for natural alternatives which are generally regarded as safe. The aim of this study was to evaluate the potentials of turmeric powder in rumen manipulation using the in vitro method. Substrate of Panicum maximum and concentrate in ratio 6: 4 with turmeric inclusion at four levels of 0, 5, 10 and 15 mg/g dry matter (DM) were used for this study. The experiment was arranged in a completely randomized design. Approximately 200 mg of substrate in each treatment was weighed separately into 100 mL glass syringes into which 30 mL of rumen fluid and buffer solution (1:2 v/v) were added. The quantities of total gas, methane, ammonia, total volatile fatty acids production and substrate degraded were determined 48-h post incubation. Rumen bacteria, protozoa, fungi population were determined and microbial biomass was estimated. Some phytochemical constituents of turmeric were also determined using standard methods. Turmeric had a higher percentage of curcumin relative to other phytochemical contents determined. Turmeric effectively and consistently ( $p < 0.05$ ) reduced gas production at levels above 5 mg/g of substrate inclusion throughout the 48-h incubation period. Similarly, turmeric reduced ( $p < 0.05$ ) methane, carbon-dioxide, ammonia and total volatile fatty acids production, and substrate degradation at 10 – 15 mg/g inclusion. Rumen bacteria and protozoa reduced when turmeric was included at 10 – 15 mg/g while fungi reduction was observed at 15 mg/g of inclusion. Reduction in microbial biomass was observed at 15 mg/g of turmeric inclusion. It can be concluded from this study that turmeric inclusion above 5 mg/g DM of substrate, can modify the rumen by causing a reduction in fermentation end-products. The reduction of ammonia production at 15 mg/g which significantly reduced microbial biomass has implication for lowering microbial protein synthesis.*

**Keywords:** Rumen manipulation, turmeric, incubation, microbial protein

### **Une évaluation in vitro des potentiels du curcuma comme additif phytogénique pour la modification du rumen**



#### **Résumé**

*Le risque potentiel pour la santé animale et humaine dans l'utilisation d'additifs alimentaires antibiotiques pour modifier la fermentation du rumen a nécessité la recherche d'alternatives naturelles qui sont généralement considérées comme sûres. Le but de cette étude était d'évaluer les potentiels de la poudre de curcuma dans la manipulation du rumen en utilisant la méthode in vitro. Substrat de Panicum maximum et concentré dans le rapport 6: 4 avec l'inclusion de curcuma à quatre niveaux de 0, 5, 10 et 15 mg/g de matière sèche (DM) ont été utilisés pour cette étude. L'expérience a été organisée dans une conception complètement randomisée. Environ 200 mg de substrat dans chaque traitement ont été pesés séparément dans des seringues en verre de 100 mL dans lesquelles 30 mL de liquide rumen et de solution*

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*tampon (1:2 v/v) ont été ajoutés. Les quantités totales de gaz, de méthane, d'ammoniac, de production totale d'acides gras volatils et de substrat dégradé ont été déterminées 48 h après incubation. La bactérie Rumen, le protozoaire, la population de champignons ont été déterminés et la biomasse microbienne a été estimée. Certains constituants phytochimiques du curcuma ont également été déterminés à l'aide de méthodes standard. Le curcuma avait un pourcentage plus élevé de curcumine par rapport à d'autres contenus phytochimiques déterminés. Le curcuma a réduit efficacement et systématiquement ( $p < 0,05$ ) la production de gaz à des niveaux supérieurs à 5 mg/g d'inclusion du substrat tout au long de la période d'incubation de 48 h. De même, le curcuma a réduit ( $p < 0,05$ ) le méthane, le dioxyde de carbone, l'ammoniac et la production totale d'acides gras volatils, et la dégradation du substrat à 10 à 15 mg/g d'inclusion. Les bactéries rumen et le protozoaire ont diminué lorsque le curcuma a été inclus à 10–15 mg/g tandis que la réduction des champignons a été observée à 15 mg/g d'inclusion. La réduction de la biomasse microbienne a été observée à 15 mg/g d'inclusion de curcuma. On peut conclure de cette étude que l'inclusion de curcuma au-dessus de 5 mg/g de DM de substrat, peut modifier le rumen en causant une réduction des produits finaux de fermentation. La réduction de la production d'ammoniac à 15 mg/g, ce qui a considérablement réduit la biomasse microbienne, a des répercussions sur l'abaissement de la synthèse des protéines microbiennes.*

**Mots-clés:** Manipulation rumen, curcuma, incubation, protéine microbienne

#### **Introduction**

Modifying the rumen fermentation process to reduce feed nutrient losses is of great significance both to the ruminant animal and to the environment (Tadesse, 2014; Subepang *et al.*, 2019). The process of feed fermentation by rumen microbes generates end-products which include volatile fatty acids, ammonia, hydrogen, carbon-dioxide and methane. The volatile fatty acids are the major source of energy to the ruminant while the ammonia produced from the deamination of dietary nitrogen, are utilized for microbial protein synthesis necessary for microbial proliferation. However, dietary proteins and energy losses occur in the rumen in form of ammonia and methane, respectively (Ulfina *et al.*, 2019). The deamination of dietary nitrogen in the rumen leading to excessive production of ammonia which eventually are absorbed via the rumen wall is a wasteful use of dietary nitrogen, as it results to inefficient nitrogen retention (Eschenlauer *et al.*, 2002). Methanogenic archaea present in the rumen, utilize hydrogen and carbon-dioxide produced during microbial feed

fermentation in the rumen to form methane (Chuntrakort *et al.*, 2014; Patra *et al.*, 2017). A loss of 2 – 15% of gross energy intake associated with enteric methane emission has been reported (Tadesse, 2014; Wanapat *et al.*, 2015). Therefore, in addition to excessive ammonia production, emission of enteric methane is another significant waste of dietary nutrients, which makes the fermentation process not completely effective (Castillo-Gonzalez *et al.*, 2014). Shifting rumen fermentation away from methane and excessive ammonia production, is a beneficial step towards achieving efficient rumen fermentation. Several rumen modification strategies have been employed in this regard, one of which includes the use of phytogetic plant extracts and their secondary metabolites (Tadesse, 2014). Several *in vitro* and *in vivo* studies on different plants, plant extract or essential oils have been carried out in a bid to examine the influence of plant secondary metabolites in ruminants (Hassan *et al.*, 2020). It is well established that plant secondary metabolites have antibacterial

properties (Compeer and Ynalvez, 2005; Reddy *et al.*, 2020) which is the basis for exploring them as rumen modifying agents. Turmeric (*Curcuma longa*) is the rhizome of an herbaceous perennial plant which belongs to the family Zingiberaceae (Thangavel and Dhivya, 2019). It has long been used for medicinal purposes and in various nutritional applications (Chatzinasiou *et al.*, 2019). Turmeric is rich in curcuminoids, a polyphenolic compound which contains 70-77% curcumin, 18-20% demethoxycurcumin and 7-10% bisdemethoxycurcumin (Pradeep *et al.*, 2016). These active constituents of turmeric are known to exhibit a wide range of biological activities which includes antibacterial and antifungal activities (Kebede *et al.*, 2021). In a bid to search for natural and readily available alternatives to antibiotic-based rumen modifiers, this study aimed at exploring the potentials of turmeric as a rumen modifying feed additive with a view of determining its level of inclusion in the diet for achieving rumen fermentation changes.

## **Materials and methods**

### ***Experimental location***

The experiment was carried out at the Animal Nutrition Laboratory of the College of Animal Science and Livestock production, Federal University of Agriculture, Abeokuta, Ogun State, Nigeria. The region is in the derived savannah ecological zone of South-Western Nigeria and it falls within Latitude 7°13'37" N and Longitude 3°27'02" E (Google Earth, 2021). The climatic condition is humid with a mean annual rainfall of 1037mm. The mean annual temperature and humidity are

34.7°C and 82%, respectively.

### ***Collection and processing of test ingredient***

Fresh turmeric rhizomes were sourced from a reputable market within Abeokuta, Ogun State, Nigeria. The rhizomes were washed to remove debris and were air-dried until stable dry weight. Air-dried samples were milled to pass through a 1 mm sieve. Dried and milled samples were then stored at room temperature in air-tight glass container and kept away from sunlight for subsequent use. Aliquot of the air-dried samples were used for dry matter content determination and chemical analysis.

### ***Experimental substrate***

The experimental substrate used for *in vitro* studies consisted of *Panicum maximum* grass and a formulated concentrate supplement in the ratio 6:4 on dry matter basis. Air-dried and milled turmeric was incorporated in the substrate at different levels of 0, 10, 20, 30 mg/g to make four different treatments. *P. maximum* was harvested from an existing pasture at 15cm above ground level after six weeks of re-growth and chopped into sizes of about 2–4 cm. The concentrate feed mixture as formulated as complete feed using agro-industrial by-products. Known weights of the *Panicum maximum* also known as Guinea grass and concentrate samples (n=3) were collected and oven-dried at 65°C to constant weight to obtain dry matter content. Oven dried and milled samples were analyzed for their chemical composition. The composition of *P. maximum* and the concentrate are presented in Table 1.

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**Table 1: Composition (%) of experimental *Panicum maximum* and concentrate**

<b>Ingredient</b>	<b>Percentage composition</b>	<b><i>Panicum maximum</i></b>
Wheat offal	48.00	-
Maize	17.00	-
Brewers' dried grain	32.00	-
Bonemeal	2.00	-
Salt	1.00	-
<b>Total</b>	<b>100</b>	-
<b>Nutrient composition</b>		
Dry matter <sup>1</sup>	91.64	31.43
Crude protein	18.13	10.68
Ether extract	6.86	6.18
Organic matter	93.88	89.80
Neutral detergent fibre	53.95	68.38
Acid detergent fibre	22.51	30.63
Acid detergent lignin	10.68	18.11
GE (MJ/kg DM)	19.05	17.75
ME (MJ/kg DM)	12.55	12.24

<sup>1</sup>Dry matter on as-received basis

***Measurement of in vitro gas production***

The *in vitro* gas procedure of Menke and Steingass (1988) was followed, with slight modification. About 2 L of rumen fluid required for incubation was collected in the morning before feeding from three West African dwarf goats using suction method. The goats were previously fed on 60% grass and 40% concentrate diet. The concentrate contained 40% corn, 10% wheat offal, 10% palm kernel cake, 20% groundnut cake, 5% soya bean meal, 10% dried brewers' grain, 1% common salt, 3.75 oyster shell and 0.25% fish meal. The rumen fluid collected was strained through four-layered cheese cloth into warm (39°C) thermo-flask and taken to the laboratory. A medium containing micro and macro nutrients, buffer solution, reduction solution and resazurin were mixed with the rumen liquor at ratio 2: 1 (v/v) to obtain the inoculum used for digestion. The mixture was handled under continuous anaerobic condition. There were four treatment samples according to the different levels of turmeric inclusion. Approximately 200 mg of the samples (n= 10 per treatment) were weighed into dacron bags (2mm by 5mm

size) of known weights and inserted into 100 ml glass syringes. The syringes were then filled with 30mL inoculum. The piston of the syringes was pushed upwards to eliminate air completely in the inoculum. A silicon tube inserted at the tip of the syringe was tightened by a metal clip to prevent escape of gas during the incubation process. Incubation was carried out at 39°C and the volume of gas produced was measured at three hours (h) interval from 0 to 48-h. Three blanks of syringes containing 30 mL of medium only was included in the run to correct for gas production not arising from substrate degradation. The volume of gas produced from the blanks was deducted from the volume of gas produced per sample.

***Determination of methane production***

At the end of 48-h incubation, the volume of methane gas produced from the fermentation of substrates was determined by dispensing 4 mL of 10N sodium hydroxide into each of the syringes (n=5 for each treatment group) through the silicon tube. Sodium hydroxide was added to absorb carbon-dioxide produced during the process of fermentation and the remaining

volume of gas was recorded as methane according to the method of Fievez *et al.* (2005).

#### **Determination of ammonia nitrogen and total volatile fatty acids**

After 48-h incubation, 10 mL sample (n=20; 4 treatment × 5 replicates) and 5mL sample (n=20; 4 treatment × 5 replicates) of digestion liquor was collected for ammonia nitrogen and total volatile fatty acids, respectively. The concentration of rumen ammonia nitrogen in the digestion liquor was determined using the micro-Kjeldahl method according to A.O.A.C (2000). Total volatile fatty acids were determined following the procedure of Barnett and Reid (1957).

#### **Microbial analysis**

The conventional roll-tube technique (Hungate, 1969) was used for culturing and isolation of bacteria, protozoa and fungi from rumen fluid. Each organism was identified according to Cowan and Steel (1993) method of microbial identification. Microbial count was done according to Galyean (1989) method of total direct count.

#### **Chemical analyses**

Dried samples of *P. maximum*, concentrate and turmeric powder were analyzed for proximate composition (AOAC, 2000) and fibre fractions (Van Soest *et al.*, 1991). Metabolizable energy content was estimated according to De Boever *et al.* (1997). Gross energy estimated according to Weiss and Tebbe (2019) and value obtained in Mcal/kg was multiplied by 4.184 for conversion to MJ/kg. Curcumin content in turmeric was determined following the procedure of Singh *et al.* (2012). Tannin (Makkar, 2003), total phenol (Do *et al.*, 2014), flavonoids (Nasseri *et al.*, 2019). Oxalate (Mishra *et al.*, 2017) and saponin (Mir *et al.*, 2016) contents were also determined in turmeric.

#### **Experimental design and statistical analysis**

The experiment was arranged as a completely randomized design. Data collected under this study were analyzed using the one-way analysis of variance (SAS, 2002). The experimental model is as shown below:

$$Y_{ij} = \mu + T_i + \Sigma_{ij}$$

Where:  $Y_{ijk}$  is the observation,  $\mu$  is the population mean,  $T_i$  is the effect of turmeric levels in substrate ( $i = 1-4$ ), and  $\Sigma_{ij}$  is the residual error. Significant differences among means were compared using Duncan multiple range test (SAS, 2002).

#### **Results and discussion**

The phytochemical composition of turmeric as determined in this study (Table 2) revealed the presence of curcumin, total phenol, tannin, flavonoid, oxalate and saponin. Of the phytochemicals determined, curcumin had the highest percentage composition of 3.52%. This value was comparable to the value of 3.14% reported by Tayyem *et al.* (2006) and within the range of 2.85 and 4.32% reported by Pawar *et al.* (2014) for curcumin content in turmeric samples. Curcumin has been reported as the active ingredient in turmeric known to be responsible for various biological activities (Tayyem *et al.*, 2006; Kebede *et al.*, 2021).

The gas production from *in vitro* degradation of substrate containing varying levels of turmeric powder is shown in Table 3. At the end of 27-h incubation period, turmeric reduced ( $p < 0.05$ ) gas volumes at each level of inclusion relative to the control. However, at the end of 48-h, gas volumes were significantly ( $p < 0.05$ ) reduced only at levels above 5 mg/g of inclusion. Gas production in the rumen is a direct result of microbial fermentation of carbohydrate (Yang, 2017), therefore, the observed reduction effect of turmeric on gas production suggests its potential to inhibit carbohydrate degradation in the rumen. The loss of the reduction effect of turmeric at 5

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mg/g after 27-h of incubation suggests that at lower levels of inclusion, microbes can adapt to turmeric inclusion in substrate during fermentation, over a period. The fermentation end-products from the degradation of substrates incubated with varying levels of turmeric is shown in Table 4. Turmeric altered ( $p < 0.05$ ) fermentation at levels above 5 mg/g of substrate. Turmeric decreased ( $p < 0.05$ ) the concentration of methane, carbon-dioxide, ammonia nitrogen and total volatile fatty

acids at all levels of inclusion. The observed reduction in total volatile fatty acids might be reflection of reduced acetic acid and butyrate since gas production is assumed to occur when substrate carbohydrate is fermented to acetate and butyrate (Yang, 2017). The reduction in ammonia was attributed to inhibition of substrate protein degradation by microorganisms. This reduction effect of turmeric on ammonia could be beneficial in optimizing the utilization of dietary proteins in the rumen.

**Table 2: Phytochemical composition of experimental turmeric**

Constituents	Composition <sup>1</sup>
Curcumin (%)	3.52
Total phenols (mg/kg)	3107.83
Tannin (mg/kg)	1455.75
Flavonoid (mg/kg)	465.26
Oxalate (mg/kg)	370.5
Saponin (%)	0.82

<sup>1</sup>To convert from mg/kg to %, divide value by 10, 000

**Table 3: Total gas production (m L/200 mgDM) from degradation of substrate in response to turmeric inclusion**

Incubation hour	Turmeric inclusion levels in substrate degraded, mg/g				SEM±
	0	5	10	15	
3	0.47 <sup>a</sup>	0.40 <sup>ab</sup>	0.00 <sup>c</sup>	0.13 <sup>bc</sup>	0.056
6	2.47 <sup>a</sup>	2.00 <sup>b</sup>	1.40 <sup>c</sup>	0.60 <sup>bc</sup>	0.090
9	3.00 <sup>a</sup>	2.87 <sup>a</sup>	2.20 <sup>b</sup>	2.00 <sup>b</sup>	0.099
12	5.13 <sup>a</sup>	4.53 <sup>a</sup>	3.87 <sup>b</sup>	3.73 <sup>b</sup>	0.127
15	8.40 <sup>a</sup>	7.53 <sup>b</sup>	6.47 <sup>c</sup>	6.27 <sup>c</sup>	0.173
18	11.20 <sup>a</sup>	9.87 <sup>b</sup>	8.40 <sup>c</sup>	8.00 <sup>c</sup>	0.265
21	17.67 <sup>a</sup>	12.20 <sup>b</sup>	10.60 <sup>c</sup>	9.80 <sup>c</sup>	0.294
24	20.07 <sup>a</sup>	16.33 <sup>b</sup>	14.80 <sup>bc</sup>	12.80 <sup>c</sup>	0.391
27	22.25 <sup>a</sup>	18.50 <sup>b</sup>	17.75 <sup>bc</sup>	16.00 <sup>c</sup>	0.511
30	25.13 <sup>a</sup>	22.43 <sup>ab</sup>	21.80 <sup>b</sup>	18.75 <sup>c</sup>	0.535
33	28.88 <sup>a</sup>	26.38 <sup>ab</sup>	25.13 <sup>b</sup>	20.88 <sup>c</sup>	0.550
36	30.13 <sup>a</sup>	28.75 <sup>ab</sup>	28.50 <sup>b</sup>	24.00 <sup>c</sup>	0.643
39	34.38 <sup>a</sup>	31.88 <sup>ab</sup>	30.63 <sup>b</sup>	25.38 <sup>c</sup>	0.629
42	36.00 <sup>a</sup>	34.13 <sup>ab</sup>	32.00 <sup>b</sup>	26.88 <sup>c</sup>	0.557
45	36.50 <sup>a</sup>	36.25 <sup>ab</sup>	34.80 <sup>b</sup>	30.88 <sup>c</sup>	0.653
48	38.78 <sup>a</sup>	36.88 <sup>ab</sup>	35.38 <sup>b</sup>	32.71 <sup>c</sup>	0.742

<sup>a,b,c</sup>Means along the same row with different superscript are significantly different ( $p < 0.05$ ), SEM: standard error of mean

According to Bach *et al.* (2005), dietary rumen degradable proteins results in ammonia production and if more than what rumen microbes can utilize, leads to the ammonia absorption of across the rumen wall, conversion to urea in the liver and excretion in the urine, which is a wasteful process. Manipulation of rumen protein degradation is considered the most effective strategy for reducing nitrogen N losses (Bach *et al.*, 2005; Casalmiglia *et al.*, 2010). The potentials of turmeric to reduce ammonia in the rumen was evident in this study and this reduction did not alter microbial biomass up to 10 mg/g of inclusion in the substrate. However, levels up to 15 mg/g consequently reduced microbial biomass. The reduced ammonia production with concomitant decrease in microbial biomass at 15 mg/g of turmeric inclusion implied that the amount of ammonia released was insufficient for efficient microbial protein synthesis in the rumen. Microbial protein synthesis is dependent on degradable nitrogen or proteins to supply ammonia for microbial utilization (Andrade-Montemayor *et al.*, 2009). Although, fungi population was

reduced at 15 mg/g of turmeric inclusion, the role of fungi in substrate degradation is not clearly understood. It is however, associated with the degradation of lignified plant tissues which are not degraded by other rumen microbes (Krause *et al.*, 2003). The reduced effect of turmeric on microbial population was linked with the phytochemicals in turmeric which have been associated antimicrobial activities. Studies of Tyagi *et al.* (2015) reported strong antibacterial activity of curcumin against gram-positive and gram-negative bacteria. The antimicrobial effect of tannin, flavonoids, saponin and oxalate are well documented in literature (Anantasook *et al.*, 2014; Kim *et al.*, 2015; Wang *et al.*, 2009; Benbati *et al.*, 2013). Hence, the observed fermentation and microbial population reductions were assumed to be directly related to the effect of curcumin and other plant secondary metabolites present in turmeric. Plant secondary metabolites have been reported to possess diverse antimicrobial mechanisms which include the disruption of cell membrane, enzyme inhibition, substrate deprivation, and the prevention of bacterial colonization (Reddy *et al.*, 2020).

**Table 4: Rumen fermentation end -products (mL/200 mgDM) from degradation of substrate in response to turmeric inclusion**

Incubation hour	Turmeric inclusion levels in substrate degraded, mg/g				SEM±
	0	5	10	15	
Methane production, %	32.27 <sup>a</sup>	30.05 <sup>ab</sup>	23.75 <sup>bc</sup>	20.20 <sup>c</sup>	1.74
Carbon-dioxide, %	61.78 <sup>a</sup>	57.64 <sup>ab</sup>	53.98 <sup>ab</sup>	49.74 <sup>b</sup>	1.80
Ammonia nitrogen, mg%	30.94 <sup>a</sup>	29.18 <sup>a</sup>	22.74 <sup>b</sup>	21.66 <sup>b</sup>	1.29
Total volatile fatty acids, mM/100 mL	20.61 <sup>a</sup>	18.50 <sup>ab</sup>	16.31 <sup>b</sup>	15.11 <sup>b</sup>	0.79
Substrate degraded, mg	139.11 <sup>a</sup>	130.50 <sup>ab</sup>	123.18 <sup>bc</sup>	116.36 <sup>c</sup>	2.80
Bacteria, ×10 <sup>6</sup> CFU/mL	4.2 <sup>a</sup>	3.9 <sup>a</sup>	3.0 <sup>c</sup>	2.6 <sup>c</sup>	0.20
Protozoa, ×10 <sup>6</sup> CFU/mL	1.7 <sup>a</sup>	1.6 <sup>a</sup>	0.9 <sup>b</sup>	0.8 <sup>b</sup>	0.14
Fungi, ×10 <sup>6</sup> CFU/mL	0.9 <sup>a</sup>	0.8 <sup>ab</sup>	0.7 <sup>ab</sup>	0.7 <sup>b</sup>	0.03
Estimated microbial biomass, mg/g DM	47.44 <sup>a</sup>	45.44 <sup>ab</sup>	41.78 <sup>ab</sup>	37.89 <sup>b</sup>	1.60

<sup>a,b,c</sup>Means along the same row with different superscript are significantly different (p<0.05), SEM: standard error of mean

### Conclusion

This study revealed that turmeric inclusion above 5 mg/g DM in substrate degraded *in vitro* altered rumen fermentation by reducing methane, carbon-dioxide, ammonia nitrogen and total volatile fatty acid production. Rumen bacteria and protozoa were also reduced at these levels of inclusion. Fungi population and microbial biomass were reduced. This effect of turmeric suggests it as a potential natural feed additive to modify rumen fermentation in ruminants. However, levels up to 15 mg/g DM inclusion can have a detrimental effect on microbial protein synthesis in the rumen.

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