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Abstract: Daidzein is one of common metabolites in plants and has chemotactic effect on soil bacteria that colonize the plants. There are several tests to assess bacterial chemotaxis, but none focused on rumen bacteria. Therefore, the aim of this study was to test the chemotactic response of the rumen microflora towards daidzein using a standardized bacterial chemotaxis assay. It consisted in a modifying capillary technique and employing technology for measuring in vitro gas production. Ruminal fluids and cellulose were used as controls. The response of bacteria to daidzein was greater than the response to cellulose, supporting the hypothesis that when fodder is chewed by the ruminant it releases daidzein which can attract rumen bacteria towards feed particles (chemotaxis) for attachment and subsequent degradation.

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1 Chemotactic response of the flavonoid daidzein and its effect on the composition of the

2 rumen bacterial community

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18 Abstract

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Daidzein is one of common metabolites in plants and has chemotactic effect on soil bacteria 19 that colonize the plants. There are several tests to assess bacterial chemotaxis, but none 20 focused on rumen bacteria. Therefore, the aim of this study was to test the chemotactic 21 response of the rumen microflora towards daidzein using a standardized bacterial 22 23 chemotaxis assay. It consisted in a modifying capillary technique and employing 24 technology for measuring in vitro gas production. Ruminal fluids and cellulose were used 25 as controls. The response of bacteria to daidzein was greater than the response to cellulose, 26 supporting the hypothesis that when fodder is chewed by the ruminant it releases daidzein 27 which can attract rumen bacteria towards feed particles (chemotaxis) for attachment and 28 subsequent degradation. 29 Keywords: Daidzein, cellulose, chemotaxis, ruminal bacteria, rumen

30 1. Introduction

31 The rumen is a highly dynamic environment, and none of the changes are permanent within 32 the influence due to microbial species (bacteria, archaea, fungi and protozoa) which are found in the rumen, since each species has an affinity for a substrate and / or fermentation 33 byproducts (Church, 1974; Janssen and Kirs, 2008). The potentially bioactive compounds 34 35 found in plants which constitute ruminant diet have become an area of interest in animal nutrition (Hammes and Hertel, 2002). Recent studies have shown that extracts of plants 36 37 containing secondary metabolites (i.e. saponins, tannins and essential oils) can modify rumen microbial population. However, it is not clear how these metabolites affect rumen 38 bacteria populations and therefore ruminant production efficiency. Daidzein is one of the 39 40 most common isoflavonoid in plants, and it has chemoattractant effects on soil bacteria which colonize plants (Gough et al., 1997; Peck et al., 2006). Chemotaxis is a mechanism 41

by which the bacteria respond quickly and efficiently to an attractant concentration 42 43 gradient, either towards (positive chemotaxis) or away (negative chemotaxis) of such compound (Murialdo et al., 2009). Daidzein can stimulate the growth of some rumen 44 bacteria and lactobacilli in the intestine of piglets (Yu et al., 2004), and may have 45 46 stimulating effects on rumen microbial fermentation, suggesting an interaction between 47 rumen microorganisms and this isoflavonoid (Mao et al., 2007). However, many aspects 48 remain to be elucidated to understand how bacteria sense and respond in the rumen 49 environment. Therefore the aim of this work was to elucidate the chemotactic effect of daidzein over the rumen microflora. To achieve that, we standardized a chemotaxis assay 50 51 for rumen bacteria by modifying a capillary method (Adler, 1973) developed for aerobic 52 bacteria by combining it with technology commonly used for measuring in vitro gas 53 production (Theodorou et al., 1994). We included parallel assays for studying the 54 isoflavone daidzein as a possible attractant and looked through the profile fingerprints of Denaturing Gradient Gel Electrophoresis (DGGE) the effect of different chemoattractants 55 (sterile rumen fluid, cellulose and daidzein) on ruminal bacterial consortium. 56

57 2. Material and methods

58 2.1. Culture medium (artificial saliva)

The medium was prepared according to Menke and Steingass (1988). It consisted of
various components: trace minerals solution, buffer solution, a solution of macrominerals,
cysteine as reducing solution and resazurin as an indicator of anaerobiosis, prepared under
CO₂ (100%).

63 2.2. Obtaining rumen fluid inoculums

| 64 | The rumen fluid content (solid and liquid) was collected from 3 fistulated cows with 12 |
|----|--|
| 65 | hours of fasting, placed in sealed plastic bags (to maintain anaerobiosis) and transported at |
| 66 | 39 °C to the Nutrition Laboratory (100m away), Faculty of Veterinary Medicine, |
| 67 | Universidad Autónoma de Yucatán, México. Ruminal fluid of 3 cows was leaked (together) |
| 68 | using sterile cheese cloth and collected in a beaker under constant flow of CO ₂ . The |
| 69 | remaining solids were blended under CO_2 (100%) for 20 seconds with artificial saliva in a |
| 70 | volume equal to that extracted as rumen liquour (ratio 1:1) (Menke and Steingass 1988). |
| 71 | Rumen solids were filtered again and liquor added to the initial rumen liquid obtained. |
| 72 | Then, rumen liquor was placed in 45 ml Falcon tubes and centrifuged at 10,000 rpm |
| 73 | (16,770 x g) min at 4 °C. The pellet containing the bacteria was recovered and resuspended |
| 74 | in 80 ml of artificial saliva medium and incubated at 39 °C overnight. All procedures were |
| 75 | carried out under constant flow of 100% CO ₂ . |
| 76 | 2.3. Chemotaxis assay |
| 77 | The capillary method of Adler (1973) was adapted to material for measuring in vitro gas |
| 78 | production (Theodorou et al., 1994). There is evidence of chemotaxis in the rumen bacteria |
| 79 | towards cellulose (Miron et al., 2001, Morris, 1998) and compounds in the rumen fluid |
| 80 | (Orpin and Bountiff, 1978). Thus, these compounds were chosen as positive controls. Then, |
| 81 | 50 mg/L daidzein (attractant tested), 50 mg/L of cellulose (positive control), artificial saliva |
| 82 | (negative control), and rumen fluid (positive control) were prepared. These solutions |
| 83 | (daidzein, cellulose, rumen fluid and artificial saliva) were sterilized by filtration (0.22 μm |
| 84 | pore size). Then, capillaries (75 mm lenght, 1.1-1.2 mm and 1.5-1.6 mm inside and outside |
| 85 | Ø respectively, Marienfeld, Germany) were filled with with 60 μ l of the solutions and one |

end placed inside a serum bottle (100 ml nominal capacity) and one end was kept outside. 86 Capillaries were sealed with clay and inserted trough the septum until the open end was in 87 88 contact with the culture media (80 ml) containing rumen bacteria (Figure 1). The septa were sealed with parafilm "M" and incubated at 39 °C for 1 h. Subsequently, bacteria that 89 had been attracted by the flavonoid and entered the capillary tubes were transferred into 90 91 individual sterile eppendorf tubes and transported in a cooler at 4 °C to the Laboratory of Biotechnology, Faculty of Chemical Engineering, University of Yucatan. Samples were 92 93 centrifuged for 30 min at 13000 rpm (28,341 x g), and then the cell pellet was resuspended in 30 µl sterile distilled water. For each attractant (cellulose, daidzein, rumen liquour) Each 94 95 treatment had 30 replicates, 15 were stored at -20 °C for subsequent DNA extraction and 15 repetitions were kept at 4 °C for subsequent cell count by direct counting using the 96

97 Nuebauer camera.

98 2.4 Statistical analysis

99 The chemotactic effect of attractants: ruminal fluid and cellulose (controls), and daidzein (unknown

100 response substance) on rumen bacteria, was performed by analysis of variance (ANOVA).

101 Differences in means were determined by Fisher's Least Significant Difference (LSD) test, using

102 a significance level of $\alpha = 0.05$ and the Minitab statistical program (2007).

103 2.5. PCR-denaturing gradient gel electrophoresis (DGGE) analysis

104 For each attractant replicate, the cells pellets were lysed by five consecutive cycles of

105 freezing (-20 °C) and heating (65 °C) for 3 min. Two microliters of cell lysate were used

- 106 for amplification. The primer set, 338f (5'ACT CCT ACG GGA GGCAGC AG-3') and
- 107 518r (5'ATT ACC GCG GCT GCTGG-3'), spanned the V3 region of the 16S rDNA. The

| 108 | 338f GC primer has a GC clamp (5'CGC CCG CCG CGC GCG GCG GGC GGG GCG |
|-----|--|
| 109 | GGG GCA CGA GGG G3') attached to the 5'end of primer 338f (Cocolin et al., 2001). |
| 110 | The PCR (25 μ L reaction mixtures with appropriate template) amplification program |
| 111 | consisted of preheating at 94 °C for 5 min and 10 cycles of denaturing (94 °C, 1 min.), |
| 112 | annealing (65 °C, 1 min. decresing 1°C per cycle) and extension (72 °C, 1 min), continued, |
| 113 | with three steps 20 cycles of denaturing (94 °C, 1 min.), annealing (55 °C, 1 min. decresing |
| 114 | 1°C per cycle) and extension (72 °C, 1 min), followed by final extension at 72 °C for 10 |
| 115 | min. The DGGE analysis of PCR amplicons was performed (DCode Universal Detection |
| 116 | System, Bio-Rad). The amplicons were separated in 8% polyacrylamide gel containing a |
| 117 | 100 to 40% gradient of 8 M urea and formamide increasing in the direction of |
| 118 | electrophoresis. The electrophoresis was conducted in 1× TAE buffer under 70 V at 60 $^\circ \text{C}$ |
| 119 | for 18 h. The DNA bands in gels were visualized by SYBR Gold. The similarities of PCR- |
| 120 | DGGE profiles were analyzed using the program Quantity One (BioRad Imaging Systems) |
| 121 | analysis was performed on the image. The presence (1) or absence (0) of the band through |
| 122 | patterns jolting generated a binary matrix to obtain a similarity dendrogram using the |
| 123 | method Euclidean nearest neighbor and the degree of similarity was represented by a |
| 124 | similarity coefficient was determined, using the program Paleontological Statistics |
| 125 | Software Package for Education and Data Analysis (PAST) (Hammer et al., 2001). |
| 126 | 3. Results |

127 3.1 Chemotaxis assay

The technique proposed by Alder (1973) is based on a capillary tube containing a solutionas an attractant at one end. It is placed in a chamber containing a suspension of motile

130 bacteria in buffer without a carbon source. The positive response is observed as an

accumulation of bacteria near the attractant in the tip of the capillary. However, this 131 132 chemotaxis assay was not designed for the study of anaerobic bacterial consortia, thus, some modification have been developed for different purposes (Bharat et al., 2004). In the 133 present report, the technique was integrated with material used for measuring in vitro gas 134 135 production (Theodorou et al., 1994). Four capillaries per bottle were used simultaneously 136 (although a higher number could be used) to test if daidzein, cellulose and rumen liquour had a chemotactic effect (attractant or repellent). When each capillary content was observed 137 138 under a light microscope, using bright field and differential Gram stain, microscopy 139 revealed Gram negative rod-shaped bacteria, forming palisades groups, diplobacillus and 140 estreptobacillus, while in artificial saliva no bacteria was not found. The higher chemotactic 141 response (P =0.0001) was found with ruminal fluid followed by daidzein and finally 142 cellulose (Table 1).

143 3.2.PCR-DGGE analysis of bacterial profiles

144 Partial sequences of 16s rDNA genes (DGGE) from the capillaries showed the V3 region (Ampe et al., 1999). Profiles of starved bacteria and those found in capillaries with different 145 the chemoattractants tested showed similar patterns (Figure 2). We considered 4 different 146 phylotypes involved in chemotaxis. The phylotype 1 (presente solo en daidzeina) was 147 positioned at the same place in the denaturing gradient as DNA from R. albus 7 (not 148 149 showed information). No similarity was found with F. succinogenes, and R. flavefaciens. This agrees with the results of Galicia Jiménez et al., (2011) where a search of sequences of 150 F. succinogenesis, R. albus and R. flavefaciens in the NCBI database and comparative 151 152 analysis of chemotactic genes encoding a protein involved in chemotaxis was found only in *R. albus* 8 sequences. 153

154 **4. Discussion**

In the present studio, the higher response of ruminal fluid in the test of quimiotaxis was 155 probably due to its rich and complex composition (Orpin and Bountiff, 1978). Several 156 solutes (amino acids, peptides, ammonia, soluble sugars, starch and VFA's) can be found in 157 158 the rumen liqour considering its heterougenous origin, feeds, saliva, microorganisms and 159 digestion byproducts, etc. (Araujo, 2007), some of them might also have chemotactic effect and as a result the response observed with rumen fluid could be an additive effect. In 160 161 analysis of bacterial profiles, to sorrow that the fluid ruminal presented major number of 162 bacterial cells, only he presented two phylotype, whereas in cellulose and daidzeina 3 163 presented phylotype, only the phylotype 1 was present in the daizeina. Provided that the 164 phenomenon quimiotáctico has been little studied in bacteria ruminales, this investigation 165 and the detection of genes responsible for the quimiotaxis in these microorganisms, as the 166 brought in *Ruminococcus albus* (Galicia Jiménez et al., 2011), they provide a tool in the 167 molecular dissection of this phenomenon in ruminants.

168 **5.** Conclusion

169 In summary, the anaerobic chemotaxis assay developed allowed to: 1) work directly with

rumen samples without the need to separate bacteria 2) study chemotaxis testing

simultaneously several rumen bacteria sources or, 3) using multiple chemoattractant with

the same bacterial consortium, 4) quantify microorganisms accumulated inside the capillary

tube and 5) Provide quick and relative clean samples to characterize bacterial consortium

174 chemotactically attracted (to compounds of choice) through the profile of DGGE-

175 fingerprints. 6) Obtain evidence of the chemotactic effect of daidzein upon rumen bacteria

176 consortia.

| 177 | This work tries to contribute in knowledge of the beginning that govern the communication | | |
|-----|---|--|--|
| 178 | of the microbial populations, his principal interactions and products of the microbial | | |
| 179 | metabolism might raise the manipulation of the fermentation ruminal, creating this way the | | |
| 180 | cultures probióticos for the cattle, acting charitably in the intestinal flora of the individual. | | |
| 181 | Conflict of interest | | |
| 182 | All the authors have no conflict interest. | | |
| 183 | Acknowledgment | | |
| 184 | Fibrobacter succinogenes S85, Ruminococcus flavefaciens FD-1, and Ruminococcus albus | | |
| 185 | 7 DNA sequences provided by D.M. Stevenson and Paul J. Weimer (Research | | |
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Livestock Science

I am contacting you to submit to evaluation and publication the following article Chemotactic response of the flavonoid daidzein and its effect on the composition of the rumen bacterial community

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Waiting for your comments. Best Regards

Sincerely, "mare nostrum veritable faciendum"

Dra. Mónica Marcela Jiménez Galicia Research Professor C



Figure 1. Material and setup of a modified capillary chemotaxis assay.



Figure 2 A. DGGE profile of 16S rDNA fragments amplified with primers 338f and 518r. (1) Starved rumen bacteria (2), artificial saliva (negative control), bacterial community accumulated in the capillaries with the attractant (3) sterile rumen fluid (positive control), (4) daidzein flavonoid (5) cellulose (positive control). Negative image of the gel stained with SYBR. **B.** Euclidean distance generated dendrogram with the nearest neighbor method using a binary presence or absence of bands in the banding profiles obtained rDNA DGGE bacterial chemotaxis assay. The numbers indicate the profiles identified phylotype (1 to 4).

| Substrate | Mean | S.D. | Ν |
|----------------------|-----------------------|-------------|----|
| Ruminal Fluid | 17.32251 ^a | 0.1134 | 15 |
| Daidzein | 5.6514 ^b | 0.1215 | 15 |
| Celullose | 5.2869 ^c | 0.1619 | 15 |
| Control ¹ | 0.0284 ^d | 0.0393 | 15 |

Table 1. Ruminal bacteria found in capillary chemotaxis assay (bacterias /microliter).

Different literals (a, b, c, d) in the same column, differs statistically (P < 0.001)

¹Artificial saliva (negative control). Menke and Steingass (1988).