

1 ***In vitro* production of methane with increasing levels of corn or wheat based dried**
2 **distillers' grains with solubles in a barley silage based diet**

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4 **M. HÜNERBERG^{1,2}, K. A. BEAUCHEMIN¹, E. K. OKINE², L. HOLTSHAUSEN¹,**
5 **S. M. MCGINN¹, O. M. HARSTAD³ AND T. A. MCALLISTER^{1*}**

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7 *¹Agriculture and Agri-Food Canada Research Centre, Lethbridge, Alberta, Canada*

8 *²Department of Agricultural, Food & Nutritional Science, University of Alberta,*

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Edmonton, Alberta, Canada

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³Norwegian University of Life Sciences, Ås, Norway

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21 Correspondence: Tim A. McAllister, Agriculture and Agri-Food Canada, Lethbridge

22 Research Centre, Lethbridge, Alberta, Tel: +1-403-317-2240, E-mail:

23 tim.mcallister@agr.gc.ca

24 **ABSTRACT:** Methane production from wheat or corn based dried distillers' grains with
25 solubles (CDDGS, WDDGS) was compared *in vitro*. Wheat DDGS (49 g fat/kg DM) or
26 CDDGS (115 g fat/kg DM) partially or completely replaced whole crop barley silage at
27 200, 400, 600, 800 or 1000 g/kg DM. Production of CH₄ increased linearly and
28 quadratically ($p < 0.01$) with increasing levels of CDDGS. Cumulative CH₄ production at 24
29 h was higher ($p < 0.05$) for WDDGS (12.0 ± 0.5 mg/g DM) than CDDGS up to 800 g/kg
30 DM. Molar proportions of propionate in incubation fluid were higher ($p < 0.05$) for CDDG
31 than for WDDGS at 200, 400 and 600 g/kg DM, respectively. *In vitro* CH₄ production
32 (mg CH₄/g DM; mg CH₄/g DMD) was lower for CDDGS than WDDGS up to 800 g/kg
33 substrate DM. The higher residual oil content in CDDGS compared to WDDGS likely
34 elicited this response.

35 **Keywords:** *in vitro*, dried distillers' grains with solubles, methane

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47 **Introduction**

48 Dried distillers' grains with solubles (DDGS) is a major by-product from the biofuel
49 industry wherein cereal grains are fermented to produce ethanol. As ethanol production
50 has increased considerably in the last decade, large amounts of DDGS are available and
51 predominantly used as feed for ruminant livestock (Klopfenstein et al., 2008). Corn based
52 DDGS (CDDGS) is the most abundant DDGS in the USA whereas in Canada wheat
53 based DDGS (WDDGS) accounts for almost one third of total DDGS production (USDA
54 Foreign Agricultural Service, 2010). As a result of the fermentation process, DDGS is
55 largely starch free, but concentrated three fold in protein, fibre and fat (Spiels et al.,
56 2002). The fat content is higher in CDDGS (~100 g/kg dry matter [DM]; Spiels et al.,
57 2002) than in WDDGS (~50 g/kg DM; Gibb et al., 2008) owing to the higher level of fat
58 in corn. Supplementation of ruminant diets with dietary fat reduces ruminal CH₄ through
59 a number of mechanisms including reduction in ruminal DM digestibility, direct effects
60 of fatty acids on ruminal methanogens and protozoa, and by biohydrogenation of
61 unsaturated fatty acids (Czerkawski et al., 1966; Johnson and Johnson, 1995).
62 Additionally, dietary fats often replace fermentable carbohydrates that otherwise would
63 contribute to an increase in the reducing equivalents available to reduce CO₂ to CH₄
64 (Beauchemin et al., 2008).

65 Replacing a mixture of 350 g/kg barley grain and 50 g/kg canola meal (DM basis)
66 by CDDGS (100 g fat/kg DM) in a growing high-forage diet reduced enteric CH₄
67 emissions of beef cattle from 25.3 to 21.5 g CH₄/kg DM intake, while including 400 g/kg
68 DM WDDGS (41 g fat/kg DM) had no effect on CH₄ emissions (23.9 g/kg DM intake;
69 Hünenberg et al., 2012a). In a second study by Hünenberg et al. (2012b), replacing 400

70 g/kg DM of barley grain with CDDGS (97 g fat/kg DM) in a high-grain finishing diet
71 reduced CH₄ emissions from 16.6 to 13.6 g/kg DM intake; while WDDGS (34 g fat/kg
72 DM) had no effect on enteric CH₄ production (18.4 g CH₄/kg DM intake). Results from
73 both *in vivo* trials indicate that high-fat CDDGS can effectively reduce CH₄ emissions at
74 dietary inclusion level of 400 g/kg DM. However, it is unknown how CDDGS and
75 WDDGS at inclusion level different from Hünerberg et al. (2012a; 2012b) affect CH₄
76 production. Measuring *in vivo* CH₄ production is expensive, labour intensive and time
77 consuming; while *in vitro* batch culture fermentation is an effective technique to screen
78 CH₄ production of several substrates simultaneously under standardized laboratory
79 conditions (Soliva and Hess, 2007).

80 The objective of this study was to compare *in vitro* CH₄ production from CDDGS
81 and WDDGS as these by-products over a range of substitution for whole crop barley
82 silage, and to describe responses of CH₄ and other fermentation parameters to increasing
83 levels of both DDGS types as a substrate.

84

85 **Materials and methods**

86 *Substrates, inoculum and incubation*

87 The substrates used were mixtures of whole crop barley silage and CDDGS or WDDGS
88 in the ratios of 800:200, 600:400, 400:600, 200:800 and 0:1000 (g/kg DM). It has to be
89 acknowledged that DDGS concentrations above 400 g/kg DM to 600 g/kg DM are
90 typically not fed *in vivo* because of adverse effects on feed intake and animal
91 performance. The levels of DDGS used for this study were chosen to characterize *in vitro*

92 CH₄ production and fermentation parameters for a theoretically range of DDGS inclusion
93 level of up to 1000 g/kg DM.

94 All substrate components were dried separately at 55°C for 24 h and ground through
95 a 1 mm screen (Wiley mill standard model 3, Arthur H. Thomas, Philadelphia, PA, USA)
96 before being combined. The incubation included 5 replications for each DDGS type at
97 each inclusion level. The substrates 0.3 ± 0.005 g were weighed into ANKOM bags
98 (model F57, ANKOM Technology, Macedon, NY, USA) and heat sealed. Bags were
99 placed in 125 ml serum vials 1 day prior to incubation.

100 Rumen fluid was obtained from two ruminally cannulated non-lactating Holstein
101 cows 2 h after feeding. Cows were fed a high forage diet (650 g/kg whole crop barley
102 silage, 200 g/kg barley grain, 100 g/kg canola meal and 50 g/kg vitamin/mineral
103 supplement; DM basis) *ad libitum*. Rumen contents were collected from three sites within
104 the rumen (*i.e.*, reticulum and dorsal and ventral sac), thoroughly mixed and squeezed
105 through two layers of PeCAP® polyester 355 µm pore size screen into a preheated and
106 insulated transport bucket. Donor cows were cared for in accordance with the guidelines
107 of the Canadian Council on Animal Care (1993).

108 Rumen fluid was immediately transferred to the laboratory and re-strained through 4
109 layers of cheesecloth. Filtrate was maintained at 39°C in a water bath and the headspace
110 continuously flushed with CO₂. Strained rumen fluid (10 ml) was dispensed into pre-
111 warmed 39°C culture flasks, which were preloaded with a substrate filled ANKOM bag,
112 40 ml of buffer solution and 0.5 ml of cysteine sulfide solution as a reducing agent
113 (Menke et al., 1979). The incubation flasks were sealed with aluminium crimp-sealed
114 rubber stoppers and placed on two rotary shaker platforms (Lab-Line Instruments Inc.,

115 Melrose Park, IL, USA) oscillating at 90 rpm in an incubator (model 1915, Sheldon
116 Manufacturing, Cornelius, OR, USA) at 39°C. Triplicate flasks containing only rumen
117 fluid and buffer solution were used as blank controls. All flasks were incubated for 24 h.

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119 *Gas measurement and sample collection*

120 A pressure transducer (model PX4200-015GI, Omega Engineering, Inc., Laval, QC,
121 Canada) attached to a 22 gauge (0.6 mm) needle was used to measure gas pressure [P_t
122 (kPa)] inside the flasks by inserting the needle into the flasks after 3, 6, 12 and 24 h of
123 incubation. Gas pressures were used to calculate gas production [G_p (ml)] using the
124 equation of Mauricio et al. (1999) as:

125
$$G_p = 0.18 + (3.697 \times P_t) + (0.0824 \times P_t^2)$$

126 Gas production was corrected for the amount of substrate incubated and gas produced
127 from blank controls. After each P_t measurement, a 15 ml gas sample was collected from
128 each flask using a syringe. The gas sample was then injected into a 5.9 ml evacuated
129 Exetainer (Labco Ltd., High Wycombe, Buckinghamshire, United Kingdom) and
130 analyzed for CH_4 . The remaining gas was released from the flask after the gas sample
131 was collected. Gas production (mL/g DM) and CH_4 production per g incubated DM
132 (mg/g DM) or digested DM (mg/g DMD) were summarized and reported for the duration
133 of incubation.

134 After 24 h of incubation, flasks were opened and the pH of the incubation fluid
135 measured using a pH meter (model Accumet 25, Denver Instrument Company, Arvada,
136 CO, USA). Subsequently, flasks were placed on ice and a 1.6 ml subsample of fluid was
137 removed from the bottle, acidified with 400 μ l of metaphosphoric acid (0.25; wt/vol) and

138 stored at -20°C for analysis of VFA. Bags containing the residual substrate were removed
139 from the flasks, washed under cold tap water until the water became clear, dried at 55°C
140 for 48 h and weighed to estimate *in vitro* DM disappearance (IVDMD).

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142 *Laboratory analyses*

143 Methane concentrations were analyzed using a gas chromatograph [GC (model 6890,
144 Agilent Technologies, Wilmington, DE, USA)] coupled to a thermal conductivity
145 detector. The correlation coefficients for all standard curves exceeded 99.9%. The VFA
146 concentrations were determined by GC as described by Holtshausen et al. (2009).
147 Analytical DM was determined by drying at 135 °C for 2 h (AOAC, 2005; method
148 930.15), followed by hot weighing. Organic matter (OM) was calculated as the weight
149 lost upon ignition at 550°C for 5 h (AOAC, 2005; method 942.05). Crude fat was
150 determined by ether extraction (method 920.39; AOAC, 1995) using a hot extraction unit
151 (model E-816 HE, Buchi Labortechnik AG, Flawil, Switzerland). Total N was
152 determined by combustion analysis (model NA 1500, Carlo Erba Instruments, Milan,
153 Italy). Neutral detergent fibre (NDF) and acid detergent fiber (ADF) were quantified as
154 described by Van Soest et al. (1991), using conventional filtration through fritted glass
155 crucibles, and expressed inclusive of residual ash. Neutral detergent fibre was determined
156 with inclusion of a heat stable amylase and sodium sulphite. Starch was determined as
157 described by Rode et al. (1999). Chemical analyses were completed on each sample in
158 duplicate (Table I).

159

160 *Statistical analysis*

161 Data were analyzed using the mixed model procedure of SAS (2001). The incubation
162 flask was the experimental unit for all variables. The statistical model was:

$$163 \quad y_{ij} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \varepsilon_{ij}$$

164 where: y_{ij} was the dependant fermentation variable; μ the overall mean; α_i the fixed effect
165 of type of DDGS i (CDDGS or WDDGS); β_j the fixed effect of DDGS inclusion level j
166 (200, 400, 600, 800 or 1000 g/kg DM); $(\alpha\beta)_{ij}$ the interaction of DDGS type i by inclusion
167 level j ; and ε_{ij} the residual error term. Denominator degrees of freedom were estimated
168 using the Kenward-Roger option in the model statement. Pre-planned comparisons
169 between CDDGS and WDDGS at the same inclusion level were completed using the
170 contrast statement. Polynomial contrasts were used to determine linear and quadratic
171 responses of dependent variables to increasing level of CDDGS or WDDGS. Data are
172 presented as least squares means \pm standard error of means. Differences were declared
173 significant if $p < 0.05$.

174

175 **Results and Discussion**

176 The IVDMD (Table II) decreased linearly ($p < 0.01$) with increasing levels of CDDGS or
177 WDDGS in the diet, likely attributable to the higher concentrations of EE in CDDGS
178 (115 g EE/kg DM) and WDDGS (49 g EE/kg DM) compared to barley silage (25 g
179 EE/kg DM). Elevated dietary fat levels can depress *in vitro* fibre and OM digestion by
180 exerting toxic effects on protozoa and cellulolytic bacteria (Henderson, 1973), and by
181 limiting microbial attachment to feed particles (McAllister et al., 1994). The depression
182 in IVDMD was higher ($p < 0.05$) for CDDGS than for WDDGS at inclusion levels above

183 400 g/kg DM, which corresponds with the lower ($p<0.05$) gas production (as ml/kg DM)
184 for CDDGS compared to WDDGS at all inclusion levels.

185 Production of CH₄ (mg/g DM) increased ($p<0.05$) from 5.7 to 10.0 mg CH₄/g DM as
186 the concentration of CDDGS increased from 200 to 800 g/kg DM. However, this
187 response is not typical of that observed *in vivo* as increased levels of concentrate in the
188 diet are usually associated with lower CH₄ emissions per unit feed intake (Johnson and
189 Johnson, 1995). However, it is important to consider that substitution of DDGS for barley
190 silage also results in a substantial change in both the protein content and the nature of the
191 fibre within the mixed substrate. Our results suggest that substitution of DDGS for barley
192 silage results in an increase in the amount CH₄ produced/g DM fermented.

193 Methane production (mg) per g/DM and g/DMD from CDDGS was lower ($p<0.05$)
194 than from WDDGS when DDGS was included at levels of 200 to 800 g/kg, with the
195 difference being more pronounced at lower DDGS inclusion levels. In contrast, CH₄
196 production (mg/g DM; mg/g DMD) was similar when WDDGS or CDDGS were the sole
197 substrate incubated. Decreased CH₄ emissions (mg/g DM; mg/g DMD) from samples
198 containing 200 to 800 g/kg CDDGS as compared to WDDGS likely reflect the higher fat
199 content in CDDGS, which could have lowered OM fermentation and exerted toxic effects
200 on methanogens and protozoa (Czerkawski et al., 1966). Additionally, biohydrogenation
201 of fatty acids in CDDGS may have directed reducing equivalents away from reduction of
202 CO₂ to CH₄ formation, as previously described *in vitro* (Jenkins 1987; Getachew et al.,
203 2001).

204 Total VFA production and proportions of acetate were consistently higher ($p<0.05$) in
205 samples containing WDDGS compared to CDDGS. Addition of CDDGS increased

206 ($p<0.05$) propionate proportions at levels of 200, 400 and 600 g/kg DM compared to
207 WDDGS. This resulted in higher ($p<0.05$) acetate to propionate ratios for WDDGS
208 compared to CDDGS at levels up to 600 g DDGS/kg DM and likely reflects reduced
209 fibrolytic activity (Getachew et al., 2004) with CDDGS. Higher concentrations of
210 propionate and lower acetate to propionate ratios, in batch culture *in vitro* incubation of
211 200 g/kg DM CDDGS compared to WDDGS have been reported by others (Au et al.,
212 2010; McKeown et al., 2010). Production of CH₄ and propionate are closely linked since
213 both pathways utilize reducing equivalents. Therefore, increased propionate production in
214 diets containing CDDGS compared to WDDGS may have been responsible for the lower
215 CH₄ concentration at DDGS inclusion rates up to 600 g/kg DM. Culture pH remained
216 above 6.4 in all incubations and was only lower ($p<0.05$) in WDDGS *versus* CDDGS at
217 an inclusion level of 200 g/kg DM.

218 Results of this *in vitro* study suggest that compared with WDDGS, adding CDDGS
219 to whole crop barley silage at dietary inclusion levels of up to 800 g/kg DM could reduce
220 CH₄ production *in vivo*. The lower CH₄ production was due to greater reduction in
221 IVDMD/unit CDDGS compared to WDDGS, as well as higher concentrations of
222 propionate when up to 600 g/kg DM CDDGS was included in the diet. These predictions
223 were subsequently confirmed *in vivo* when WDDGS and CDDGS were included in
224 barley silage-based diets at 400 g/kg DM (Hünerberg et al., 2012a; 2012b).

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304 **Table I.** Chemical composition (g/kg DM) of barley silage, corn and wheat dried distillers' grains [CDDGS, WDDGS (means \pm SD; n=2)].

	Barley silage	CDDGS	WDDGS
Dry matter, g/kg	433 \pm 4.3	917 \pm 3.2	917 \pm 2.8
Organic matter	921 \pm 0.1	965 \pm 0.1	937 \pm 0.5
Crude protein	121 \pm 1.4	315 \pm 2.4	457 \pm 1.8
ADF ¹	345 \pm 4.2	143 \pm 5.0	144 \pm 2.7
NDF ²	522 \pm 10.5	474 \pm 14.1	352 \pm 8.5
Crude fat	25 \pm 1.2	115 \pm 4.2	49 \pm 0.6
Starch	247 \pm 7.9	43 \pm 0.4	10 \pm 0.2

305 ¹ADF, acid detergent fibre inclusive residual ash.

306 ²NDF, neutral detergent fibre assayed with heat stable amylase and expressed inclusive residual ash.

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309 **Table II.** Effect of inclusion level of corn or wheat dried distillers' grains with solubles on *in vitro* dry matter disappearance (IVDMD), gas and CH₄ production, pH and volatile fatty acids (VFA) after 24 h *in*
 310 *vitro* incubation.

	Dried distillers' grains with solubles, g/kg DM										Pooled SEM	p-values ⁶						
	200		400		600		800		1000			CDDGS		WDDGS				
	CDDGS	WDDGS	CDDGS	WDDGS	CDDGS	WDDGS	CDDGS	WDDGS	CDDGS	WDDGS		Type ¹	Level ²	Type × Level ³	L ⁴	Q ⁵	L	Q
IVDMD, g/kg DM	492.6	508.5	482.9	496.2	446.9	494.5*	429.9	457.6*	392.2	445.7*	9.74	<0.01	<0.01	0.16	<0.01	0.23	<0.01	0.42
Gas, mL/g DM	122.8	177.9*	130.2	183.2*	143.1	180.9*	147.1	174.1*	146.0	162.0*	4.89	<0.01	<0.01	<0.01	<0.01	0.02	<0.01	0.02
CH ₄ , mg/g DM	5.7	12.5*	7.4	12.4*	8.8	12.2*	10.0	11.5*	9.9	9.5	0.30	<0.01	<0.01	<0.01	<0.01	<0.01	<0.01	<0.01
CH ₄ , mg/g DMD	10.9	23.3*	14.5	23.5*	18.7	23.4*	22.0	23.8*	21.6	20.2	0.74	<0.01	<0.01	<0.01	<0.01	<0.01	0.06	0.02
pH	6.45*	6.41	6.42	6.41	6.43	6.43	6.44	6.45	6.45	6.45	0.005	0.03	<0.01	<0.01	0.11	<0.01	<0.01	0.55
Total VFA, mM	68.3	81.5*	72.1	79.6*	73.4	80.0*	73.5	77.0*	73.7	75.8*	0.85	<0.01	0.16	<0.01	<0.01	<0.01	<0.01	0.56
VFA, mol/100																		
Acetate (A)	49.3	51.4*	50.3	51.6*	50.6	52.0*	51.2	51.9*	51.4	52.2*	0.13	<0.01	<0.01	<0.01	<0.01	<0.01	<0.01	0.55
Propionate (P)	22.3*	19.4	21.3*	19.4	20.3*	19.5	19.7	19.5	19.4	19.7	0.14	<0.01	<0.01	<0.01	<0.01	<0.01	0.04	0.63
Butyrate	18.1	17.9	17.8	17.7	17.8*	17.1	17.6*	17.0	17.5*	16.4	0.09	<0.01	<0.01	<0.01	<0.01	0.41	<0.01	0.52
A:P ratio	2.21	2.65*	2.36	2.66*	2.50	2.66*	2.60	2.66 [†]	2.64	2.65	0.022	<0.01	<0.01	<0.01	<0.01	<0.01	0.95	0.62

311 ¹Type = CDDGS or WDDGS.

312 ²Level = 200, 400, 600, 800 and 1000 g/kg DM of DDGS.

313 ³Type × Level = interaction of DDGS type × inclusion level.

314 ⁴L = linear and

315 ⁵Q = quadratic effects of different types of DDGS.

316 ⁶Means within an inclusion level differ at (*; *p*<0.05).