#### Hulless Barley (Hordeum vulgare L.) with Altered Starch Traits:

#### **Rumen Degradation Kinetics**

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**Key words:** amylose and amylopectin ratio, beta-glucan, hulless barley, rumen degradation and intestinal digestion

#### Abstract:

The objective of this study was to determine the effect of new hulless barley lines (zero-amylose waxy, CDC Fibar; 5%-amylose waxy, CDC Rattan; normal-amylose, CDC McGwire and high-amylose, HB08302) on rumen degradation kinetics and in vitro intestinal nutrient digestion with CDC Copeland included as a hulled control. Three dry Holstein cows fitted with rumen cannula were used for measuring rumen degradation of barley varieties. The estimation of intestinal digestion was determined by a modification of the three-step in vitro procedure described by Calsamiglia and Stern with duplicate of each in situ residue. Among the hulless barley lines, CDC Rattan was greater (P<0.05) in effectively degradable starch (EDST: 461 g/kg DM) but reduced (P<0.05) in rumen undegradable protein (RUP: 55g/kg DM) while CDC Fibar was greater (P<0.05) in effectively degradable crude protein (EDCP: 90 g/kg DM) and total digestible protein (TDP: 147 g/kg DM) than other hulless barley lines. CDC McGwire showed greater (P<0.05) in total digestible bypass starch (TDST: 590 g/kg DM), effectively degradable CHO (EDCHO: 581 g/kg DM) and total digestible CHO (TDCHO: 600 g/kg DM) than other hulless barley lines but no significant difference compared to CDC Copeland (P>0.05). HB08302 was greater (P<0.05) in rumen bypass starch (BST: 218 g/kg DM vs. 146 g/kg DM and 155 g/kg DM) and effective degradable NDF (EDNDF: 74 g/kg DM vs. 49 g/kg DM and 52 g/kg DM) than CDC Fibar and CDC Rattan. HB08302 was also greater in total digestible NDF (TDNDF: 93 g/kg DM vs. 62 g/kg DM and 67 g/kg DM) and intestinal digestible rumen bypass starch (IDBST: 180 g/kg DM vs. 122 g/kg DM and 130 g/kg DM) compared to CDC Fibar and CDC McGwire. HB08302 also showed greater (P<0.05) in bypass CHO (BCHO: 179 g/kg DM) and intestinal digestible bypass CHO (IDBCHO: 31 g/kg DM) than other hulless barley lines. Amylose and amylopectin were significantly correlated (P<0.001) to EDCP (r=-0.71, r=0.64) and TDP (r=-0.85; r=0.77), while amylose, ratio of amylose and amylopectin were positively correlated (P<0.01) to EDNDF, TDNDF, BST, IDBST and TDCHO (P<0.05). BCP, IDP, TDP, BNDF were positively correlated to beta-glucan levels (P<0.001) while EDST, TDST, EDCHO and TDCHO were negatively correlated to beta-glucan levels in hulless barley lines (P<0.01). in conclusion hulless barley lines with altered carbohydrate traits have the potential to increase rumen and intestinal nutrient availability to ruminants. Altered beta-glucan levels had a greater effect on rumen carbohydrates and crude protein degradation than altered starch traits.

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# Introduction

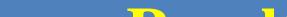
- Hulless barley contained higher energy value and better nutrient availability with its reduced fiber and increased starch content compared to hull cultivars.
- Four hulless barley breeding lines varied in amylose and beta-glucan content were used to determine the difference on carbohydrate utilization and availability in ruminants with hulled barley as reference check.

### **Objectives**

Compare hulled barley (CDC Copeland) with three newly developed hullless barley cultivars (zero-amylose waxy, CDC Fibar; waxy, CDC Rattan; and high-amylose, HB08302) and normal starch hulless barley cultivar (CDC McGwire) in terms of nutrient availability for dairy cattle.

# Conclusion

Hulless barley lines with altered carbohydrate traits have the potential to increase rumen and intestinal nutrient availability to ruminants. Altered beta-glucan levels had a greater effect on rumen carbohydrates and crude protein degradation than altered starch traits.



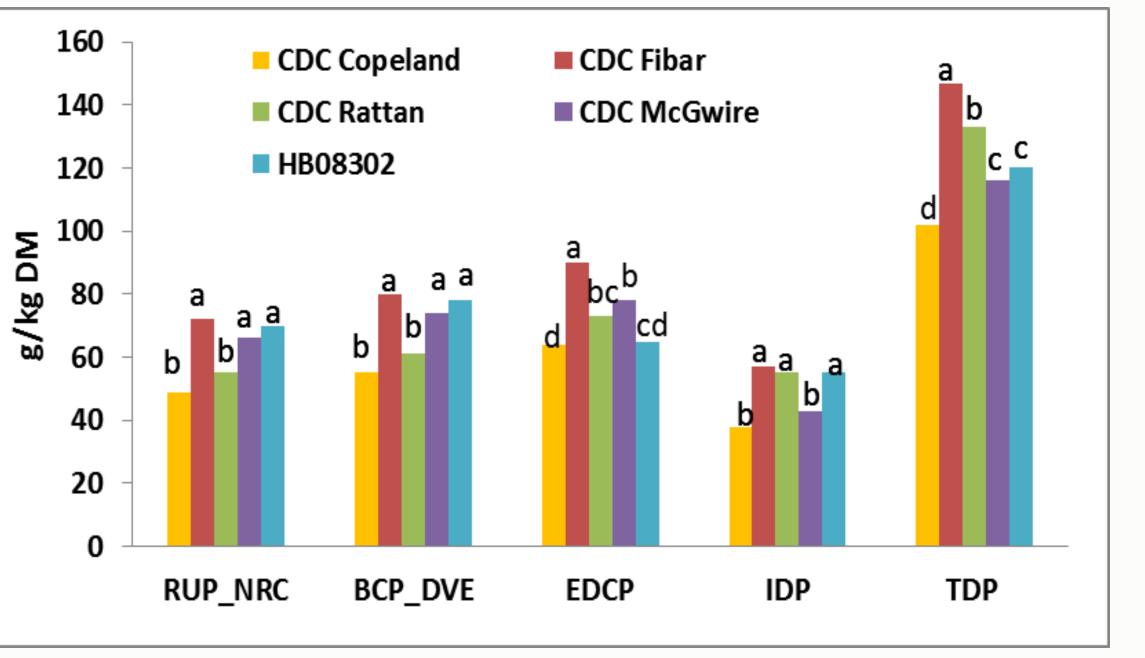
# **Materials and Methods**

Table 1: Breeding targets of four newly developed CDC hulless barleys and CDC hulled barley (control)

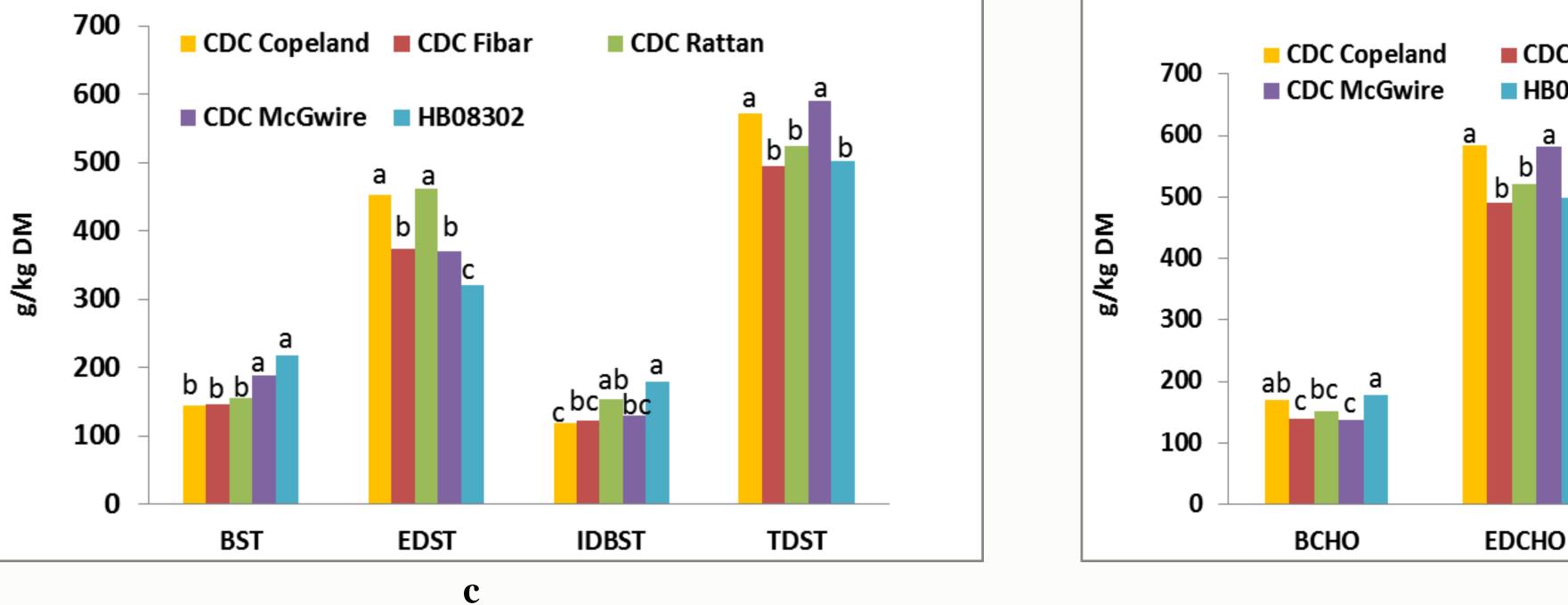
	Hulled	Hulless Barley				
Items	<b>CDC</b> <b>Copeland</b>	CDC Fibar	CDC Rattan	CDC McGwire	HB 08302	
Amylose (% in starch)	25-29	0	5	25	40	
Amylopectin (% in starch)	71-75	100	95	75	60	
Beta-glucan (% in DM)	3.5-4	Very High	High	High	High	
Sample year	2008,2009,2010 09-10					

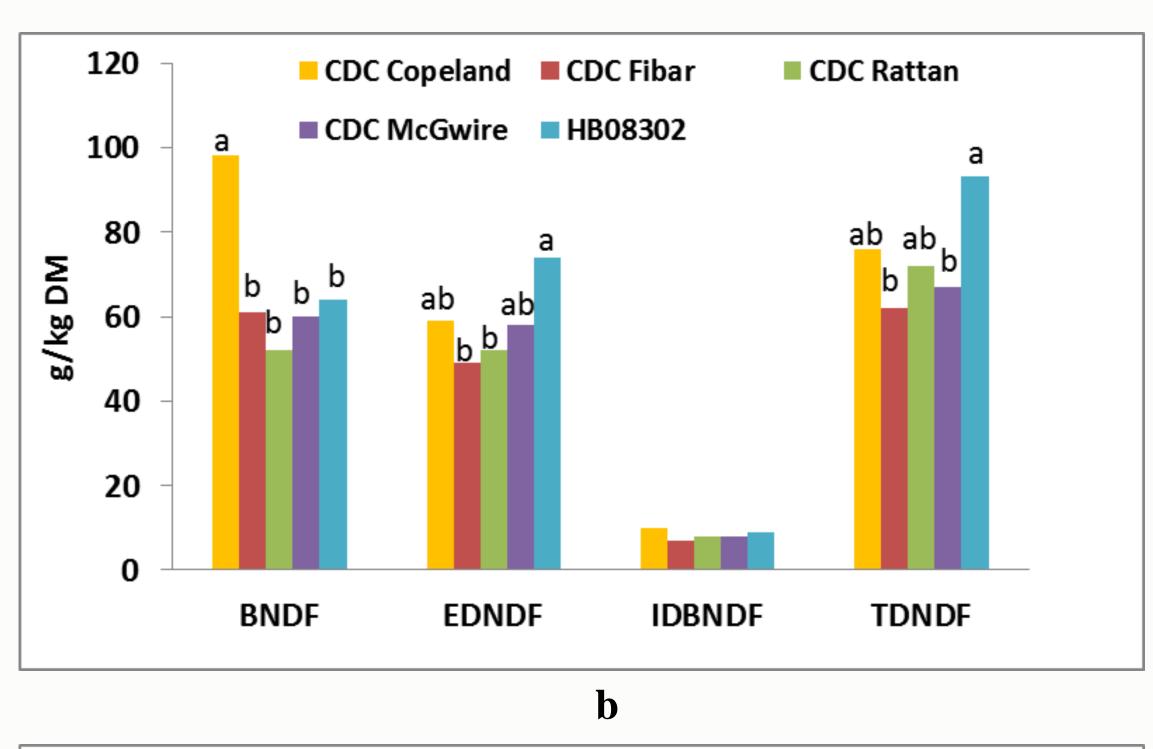


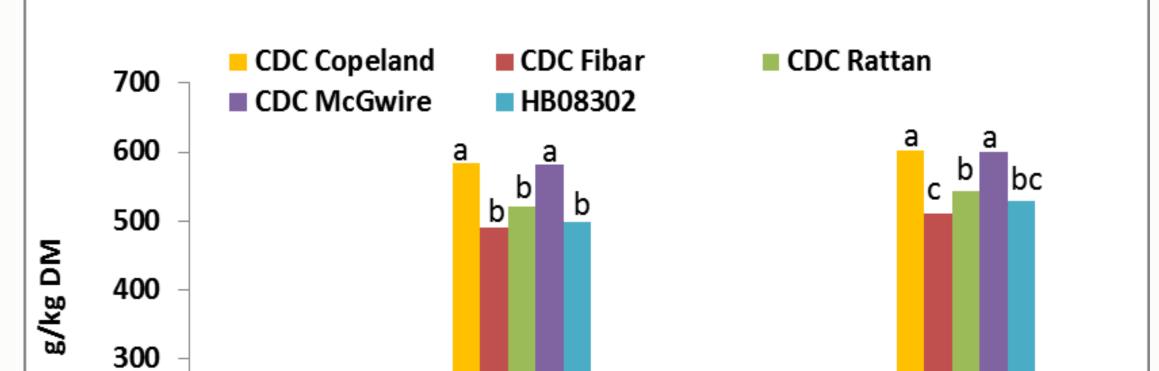










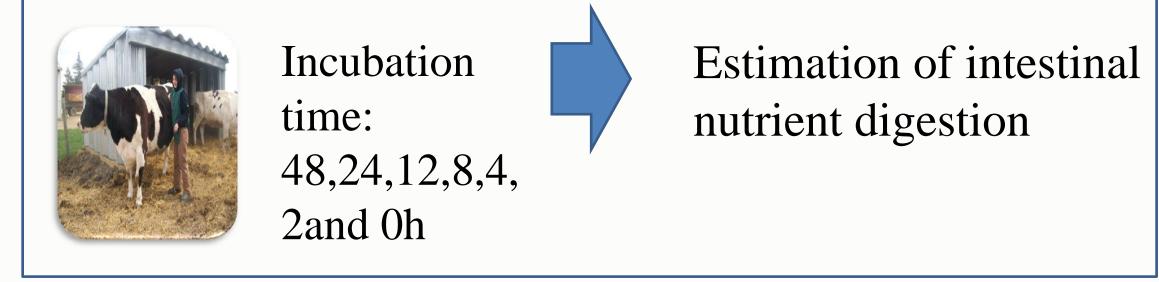


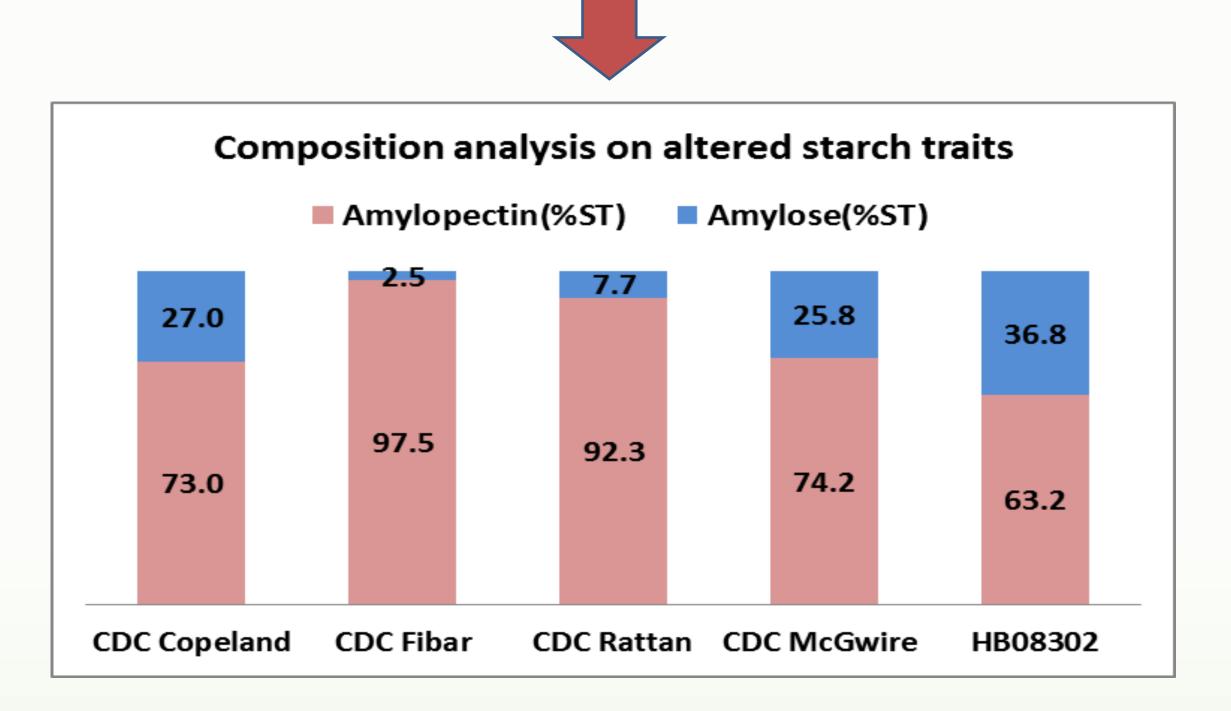
d

b bb b a

IDBCHO

TDCHO





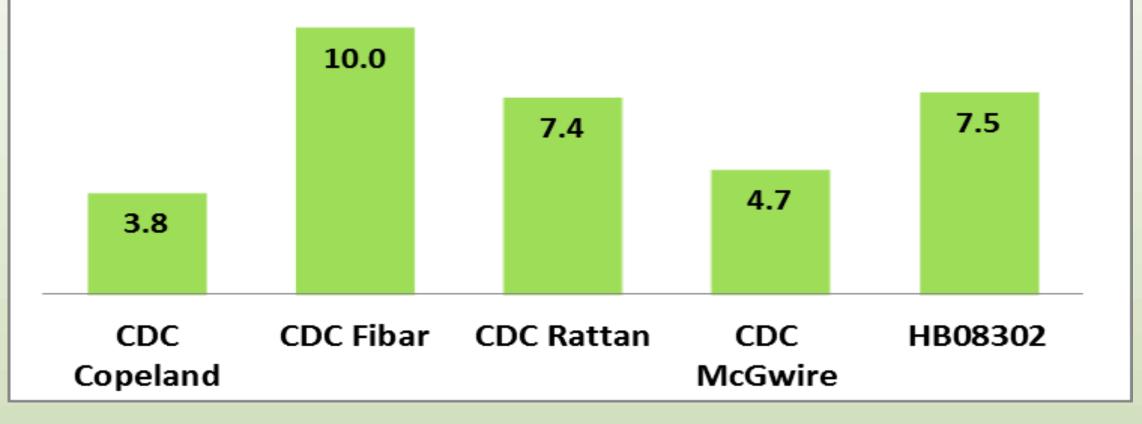
Composition analysis on altered beta-glucan levels

Beta-glucan(%DM)

Figure 1: Rumen bypass(B), effective degradable(ED), intestinal digestible(ID) and total digestible(TD) crude protein(a), **neutral detergent fiber(b), starch( c) and carbohydrate(d)** 

Table 2: Correlation analysis between altered carbohydrate traits [amylose level, amylopectin level, ratio of amylose to amylopectin (Ay:Ap) and beta-glucan level] and rumen degradation, intestinal digestion of CP, NDF, Starch and CHO

Items						
	Amylose (%DM)	Amylopectin(%DM)	Ay:Ap	Beta-glucan (%DM		
	Spearman Correlation R value					
In situ rumen CP degradation a	nd intestinal CP digestion (g	/kg DM)				
BCP_DVE	-0.25	0.08	-0.28	0.63***		
RUP_NRC	-0.25	0.08	-0.28	0.63***		
EDCP	-0.71***	0.64***	-0.70	0.37+		
IDP	-0.39+	0.27	-0.41+	0.76***		
TDP	-0.85***	0.77***	-0.84***	0.74***		
In situ rumen NDF degradation	and intestinal NDF digestion	n (g/kg DM)				
BNDF	-0.25	0.19	-0.25	0.44*		
EDNDF	0.54**	-0.28	0.54**	-0.20		
IDBNDF	0.16	-0.10	0.15	-0.05		
TDNDF	0.58**	-0.45*	0.58**	-0.17		
In situ rumen Starch degradation	on and intestinal starch diges	stion (g/kg DM)				
BST	0.53*	-0.41+	0.51*	0.11		
EDST	-0.07	0.18	-0.04	-0.60**		
IDBST	0.53*	-0.37+	0.51*	0.08		
TDST	0.38+	-0.13	$0.40^{+}$	-0.79***		
In situ rumen CHO degradation	n and intestinal CHO digestic	on (g/kg DM)				
BCHO	0.37+	-0.41+	0.35	0.13		
EDCHO	0.32	-0.07	0.34	-0.80***		
IDBCHO	0.39+	-0.45*	0.36+	0.11		
TDCHO	0.54**	-0.2 9	0.56**	-0.89***		



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Note: + for P<0.10; \* for P<0.05, \*\* for P<0.01, \*\*\* for P<0.001