



Fermentation of Dried Distillers' Grains and Solubles (DDGS) hydrolysates to solvents and value-added products by solventogenic clostridia

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Why DDGS Utilization?

- Number of dry-grind ethanol plants is growing rapidly in the U.S.
- Currently, dry-grind ethanol plants produce the majority of fuel ethanol (ca 60%) in the U.S. By-products from dry-grind ethanol include wet and dry distiller's grains with solubles.
- Approximately 9 and over 1 million metric tons of DDGS were produced in the North America and EU, respectively in 2005
- Some industrial experts are predicting that DDGS production in the U.S. will reach up to 15 million metric tons in few years
- By finding alternative uses for DDGS, ethanol plants can improve their profitability and position themselves to withstand stronger competition (from petrochemical industry)



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Why Bio-Butanol?

Butanol offers significant advantages as a second generation liquid fuel or industrial feedstock chemical:

- Higher energy content than ethanol
- Better solubility characteristics - can be stored under humid conditions
- Can be used in internal combustion and diesel engines
- Niche market in the food and flavor industry as flavor extractant
- Higher overall value as a commodity chemical
- Very large market demand (4.5×10^9 lb/year) which is petrochemical based



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Liquid Fuel Characteristics

Characteristic	Gasoline	Diesel	Methanol	Ethanol	Butanol
Formula	C4-C12 hydrocarbons	C14-C20 hydrocarbons	CH ₃ OH	CH ₃ CH ₂ OH	CH ₃ (CH ₂) ₃ OH
Boiling Point					
°C	32-210	204-343	65	78	118
°F	90-410	400-650	149	173	244
Lower heating value*					
MJ/kg	44.5	43.0	19.6	26.9	33.1
Btu/gal	114,800	140,000	55,610	76,100	96,100



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Objectives:

- Screen various solventogenic clostridia for their ability to ferment mixed hexose and pentose sugar streams for acetone butanol ethanol (ABE) production
- Carry out studies on the utilization pattern of mixed sugar streams i.e. determine whether sugar utilization occurs sequentially, simultaneously and if the microbes have a preference for one sugar over another
- Test for the effect of potential inhibitory compounds [(A). products of hemicellulose hydrolysis, and (B). degradation products] on growth and ABE production by solventogenic clostridia



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Objectives (cont.)

- Test for the ability of DDGS which has been subjected to different pre-treatment regimes to be fermented into ABE



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Composition of DDGS

		Average (95% confidence intervals)
Gross Matter Composition	Moisture Content (% total)	11.2 ± 0.6%
	Dry Matter Content (% total)	88.8 ± 0.6%
Dry Matter Composition	Glucan (total) (% dry matter)	20.9 ± 7.1%
	Cellulose (% dry matter)	16.0 ± 6.6%
	Starch (% dry matter)	5.2 ± 1.0%
	Xylan (% dry matter)	8.2 ± 3.3%
	Arabinan (% dry matter)	5.3 ± 0.7%
	Protein (% dry matter)	26.4%
	Total Dry Matter Mass Closure	92.7%

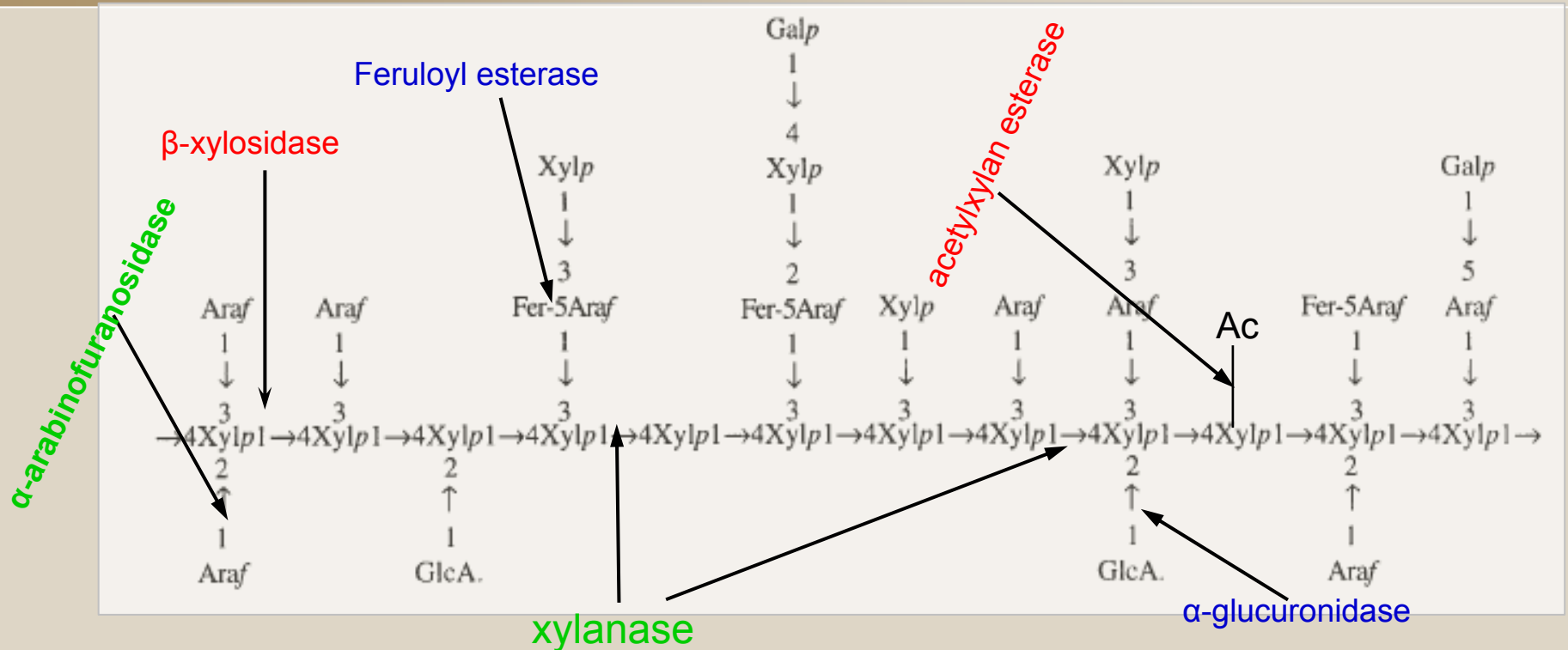
Based on analysis at U. Illinois, USDA NCAUR, Purdue



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Schematic structure of maize bran heteroxylan



Araf: Arabinofuranose, Xylp: Xylopyranose, Galp: Galactopyranose, GlcA: Glucuronic acid, Fer: Ferulic acid, AC: acetylxylan

Source: Saulnier et al, 1999



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Pretreatment and Hydrolysis of DDGS

- Dilute acid
- Alkali
- Hot water/and SO₂ impregnation
- Steam explosion/and SO₂ catalyzed steam explosion
- Ammonia fiber explosion (AFEX)
- Enzymatic



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Composition of DDGS hydrolysates








- Hexose sugars: glucose, mannose, galactose, fructose
- Pentose sugars: xylose and arabinose
- Ferulic and glucuronic acids
- Acetate
- Proteins, Fat and oils



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Composition of DDGS hydrolysates: potential degradation products

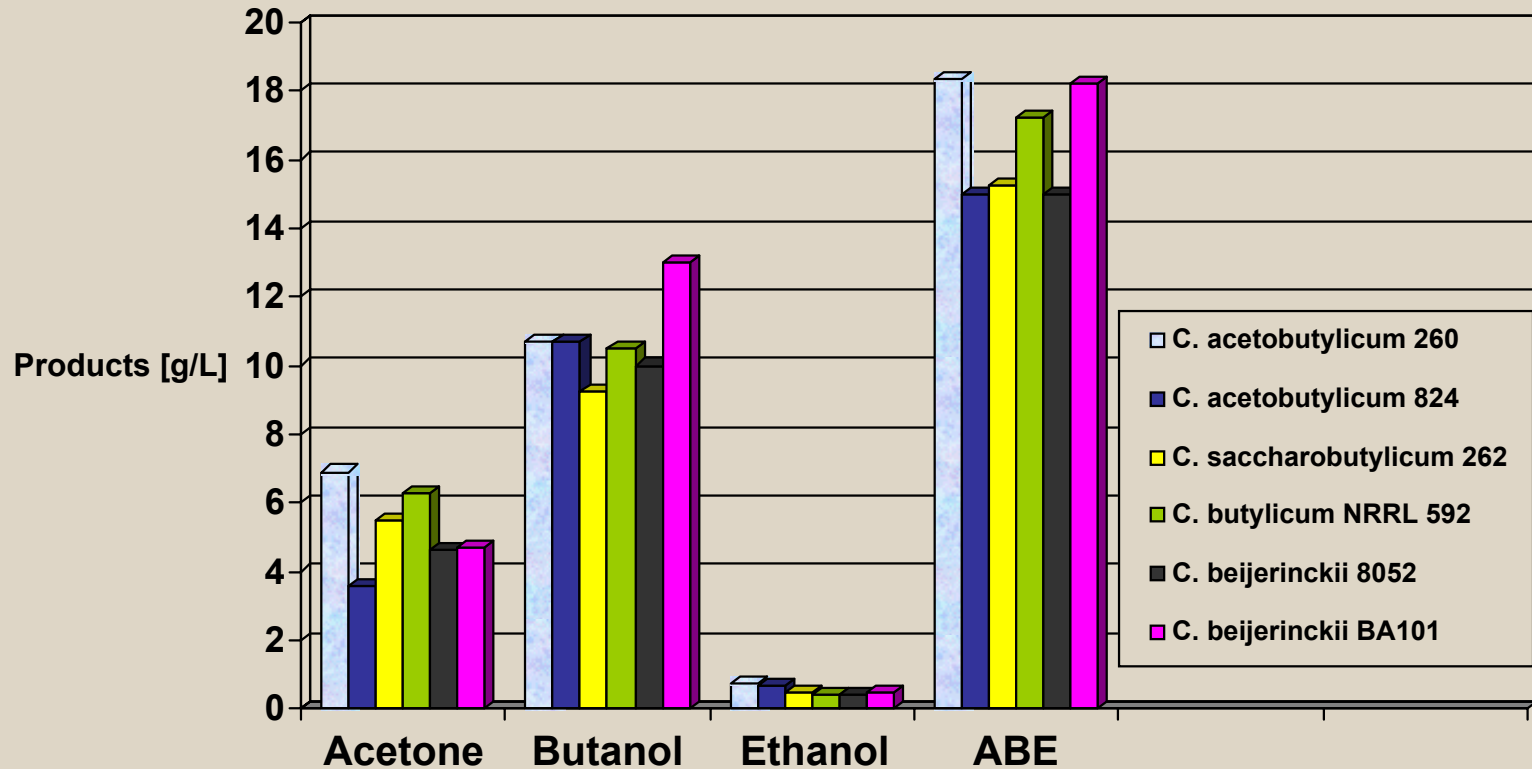
- **Glucose**  Hydroxymethylfurfural  Formic acid  Levulinic acid
- **Xylose**  Furfural  Tars  Other degradation products
- **Lignin** 
 - Syringaldehyde
 - p-hydroxybenzaldehyde
 - Cinnamaldehyde
 - P-coumaric acid
 - Other products



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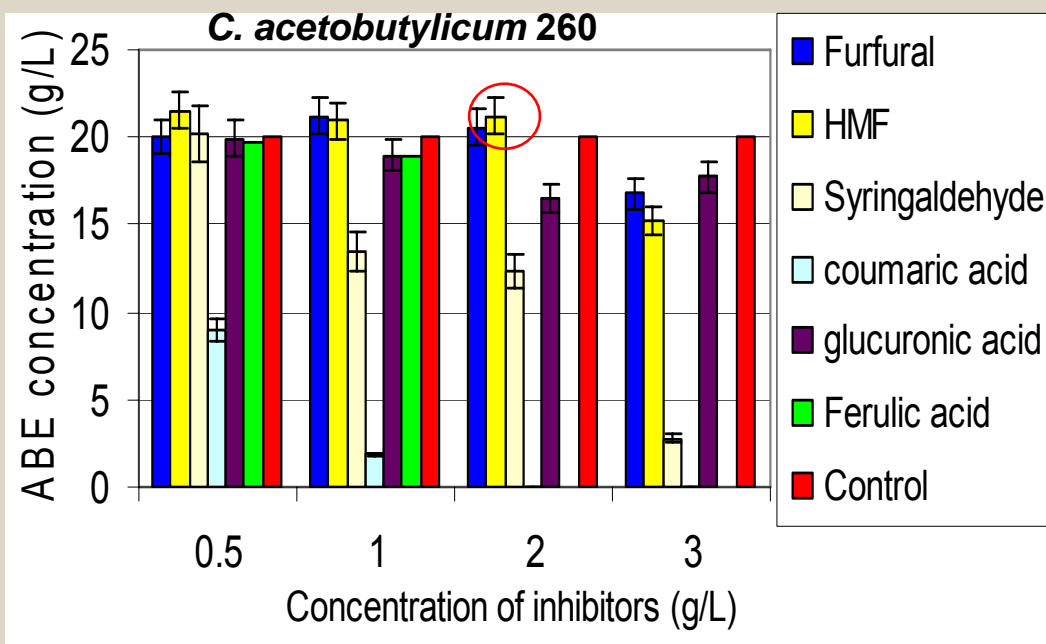
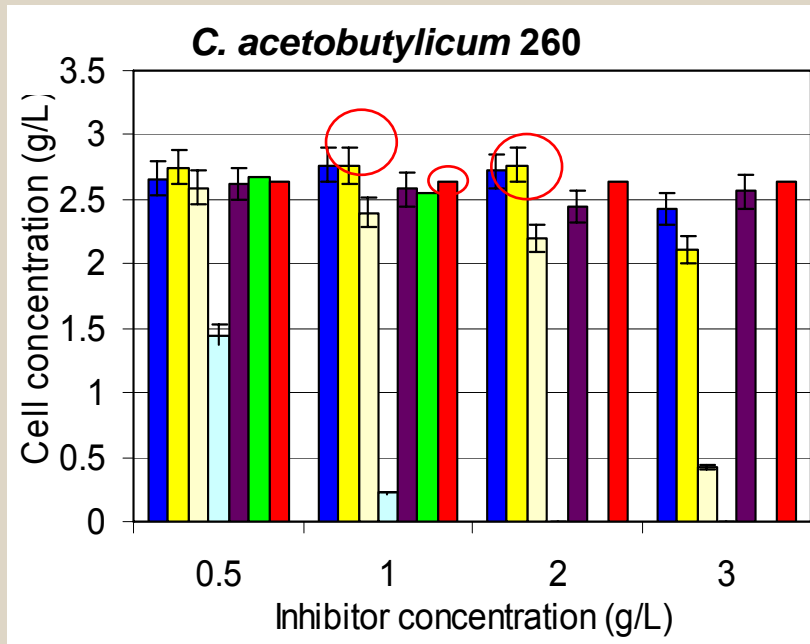
Fermentation of glucose to ABE by solventogenic clostridia



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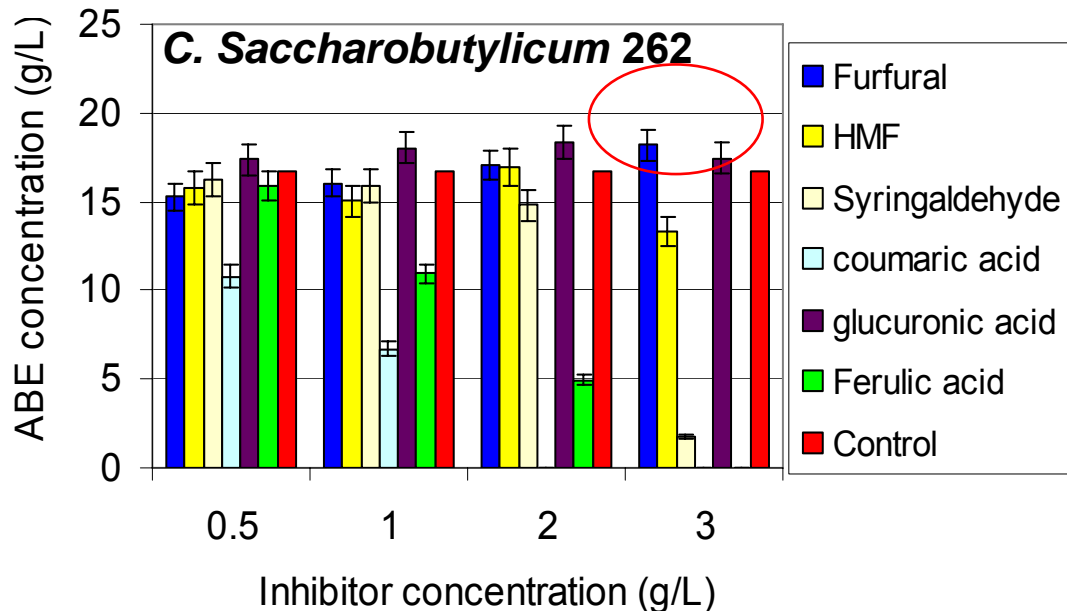
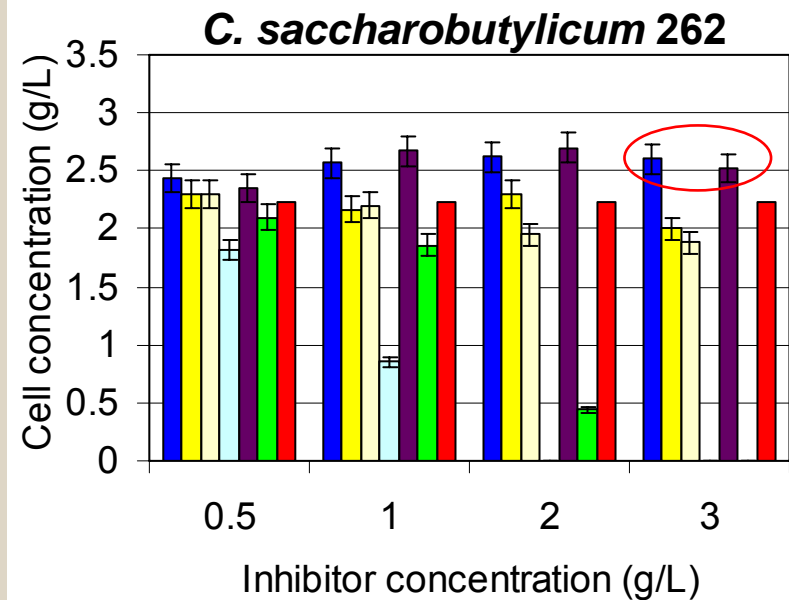
Effect of representative degradation and hydrolysis products on growth and ABE production by Clostridia



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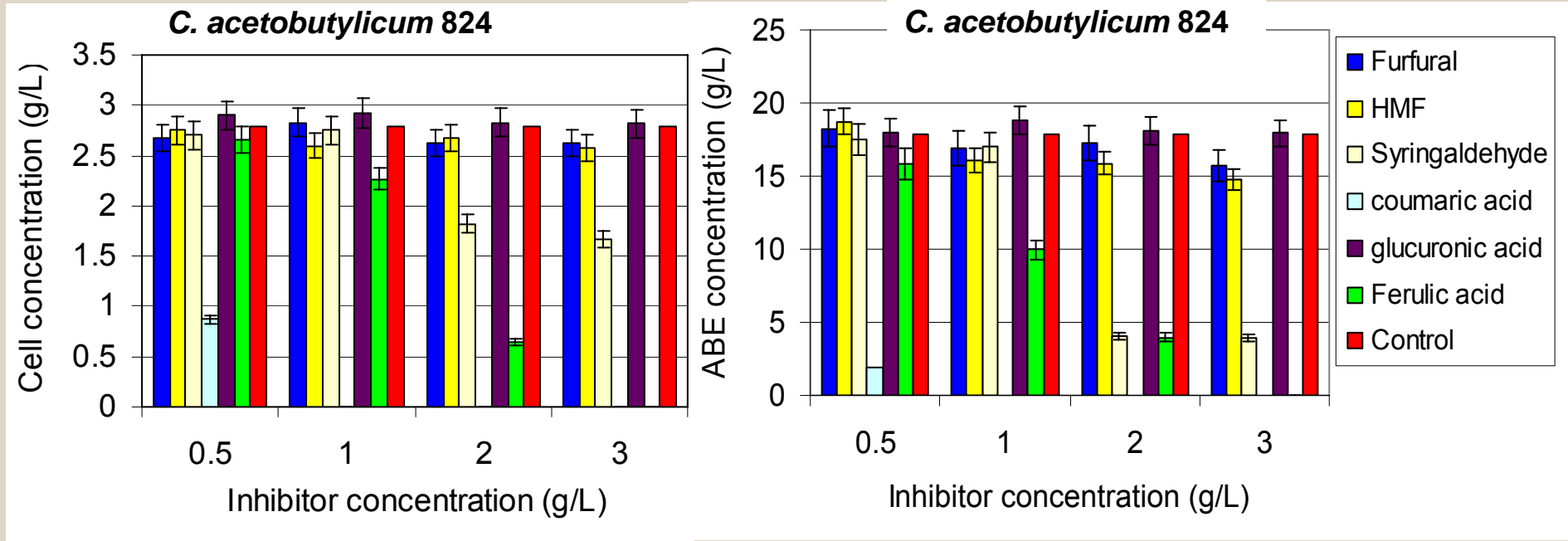
Effect of representative degradation and hydrolysis products on growth and ABE production by Clostridia Contd.



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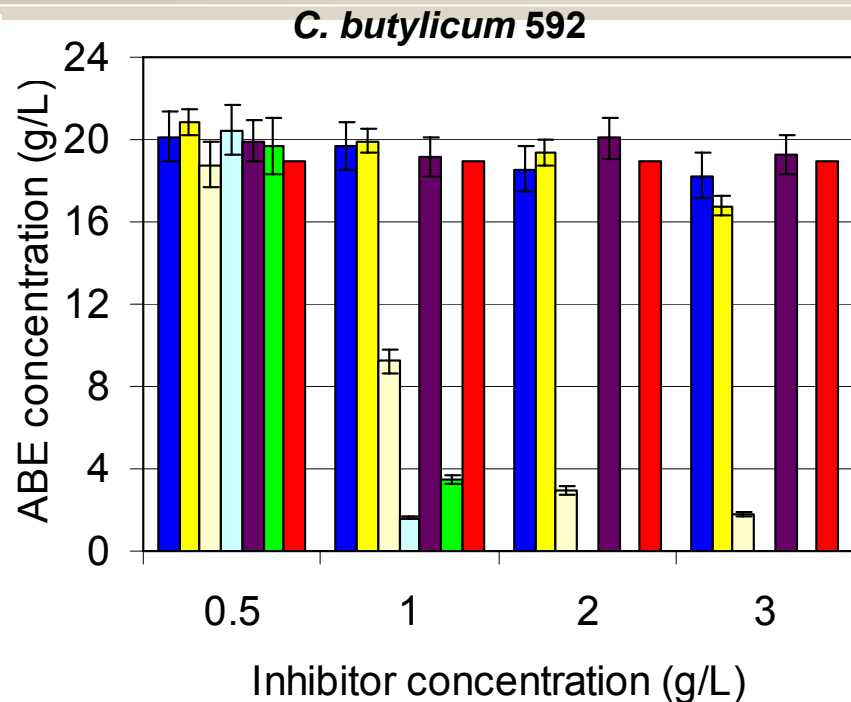
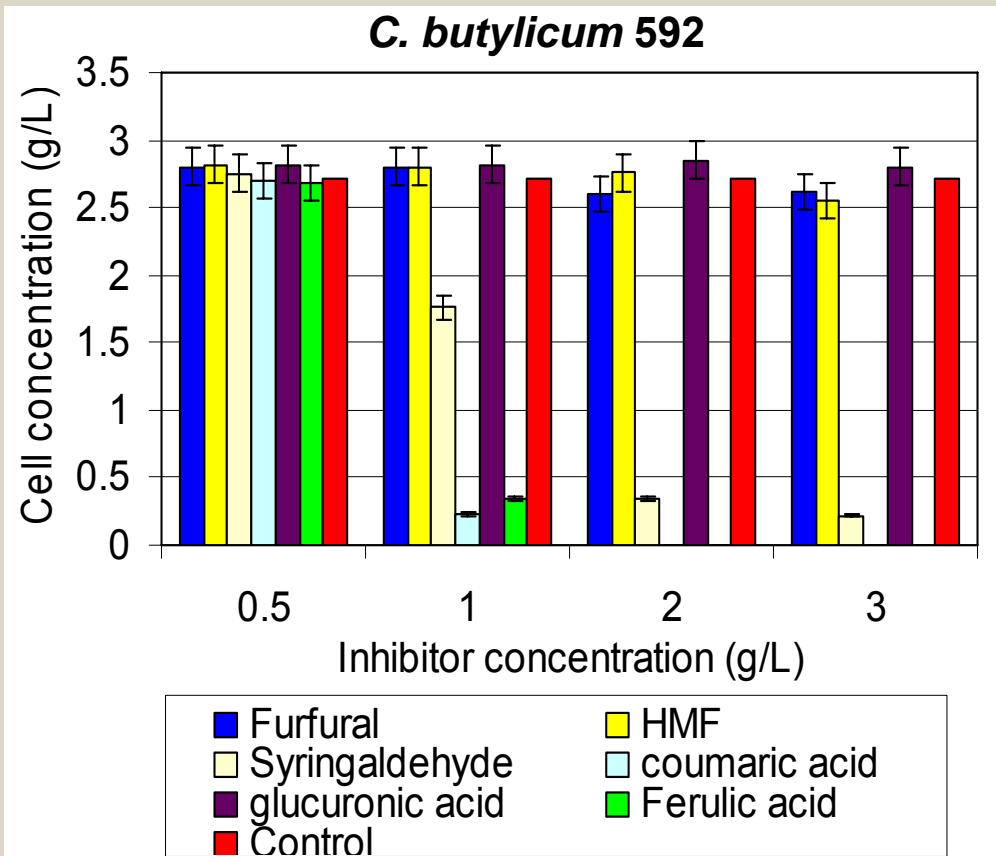
Effect of representative degradation and hydrolysis products on growth and ABE production by *Clostridia* Contd.



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Effect of representative degradation and hydrolysis products on growth and ABE production by *Clostridia* Contd.

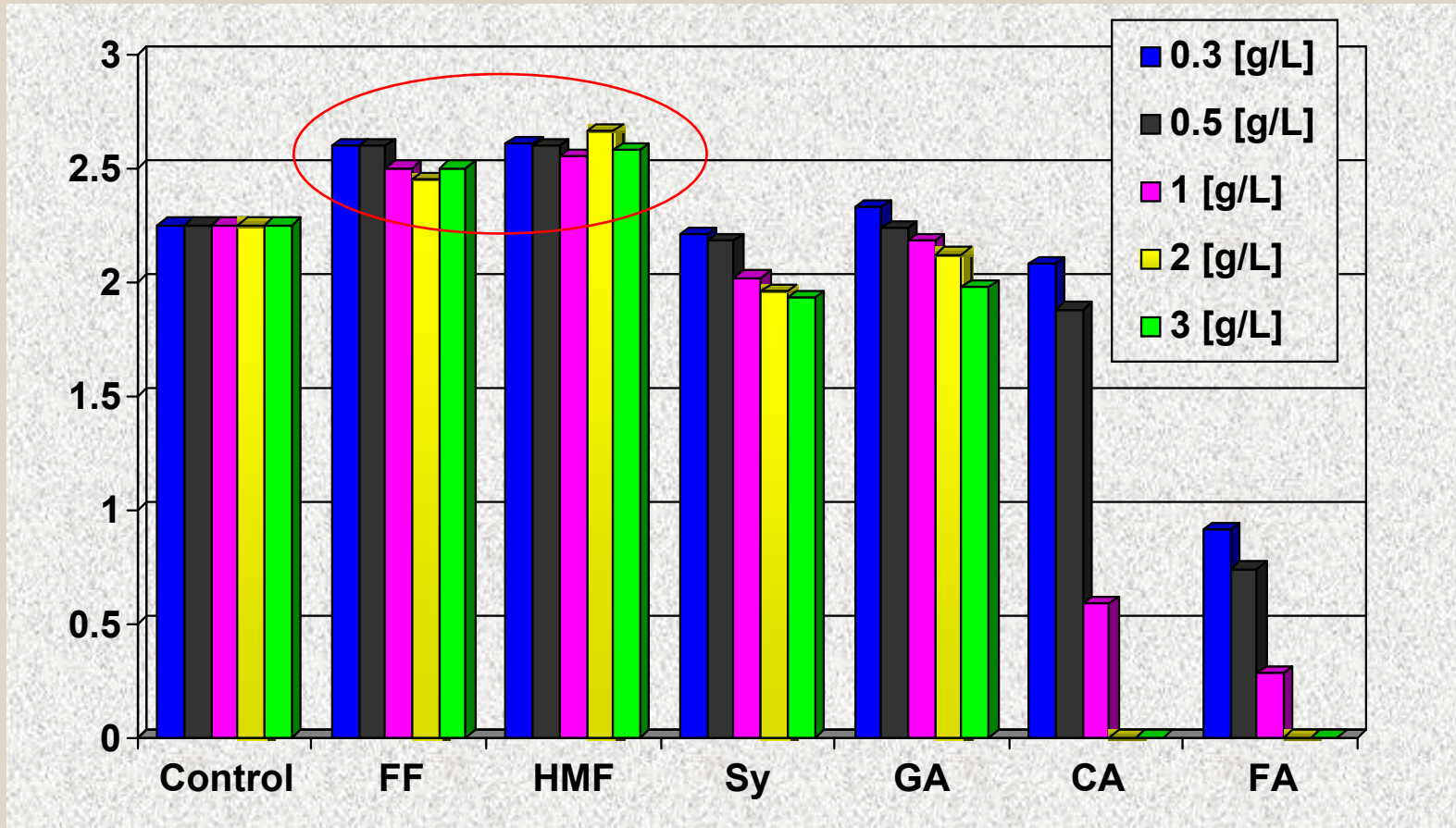


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Effect of representative degradation and hydrolysis products on the growth of *Clostridium beijerinckii* BA101

Cell
Conc.
g/l



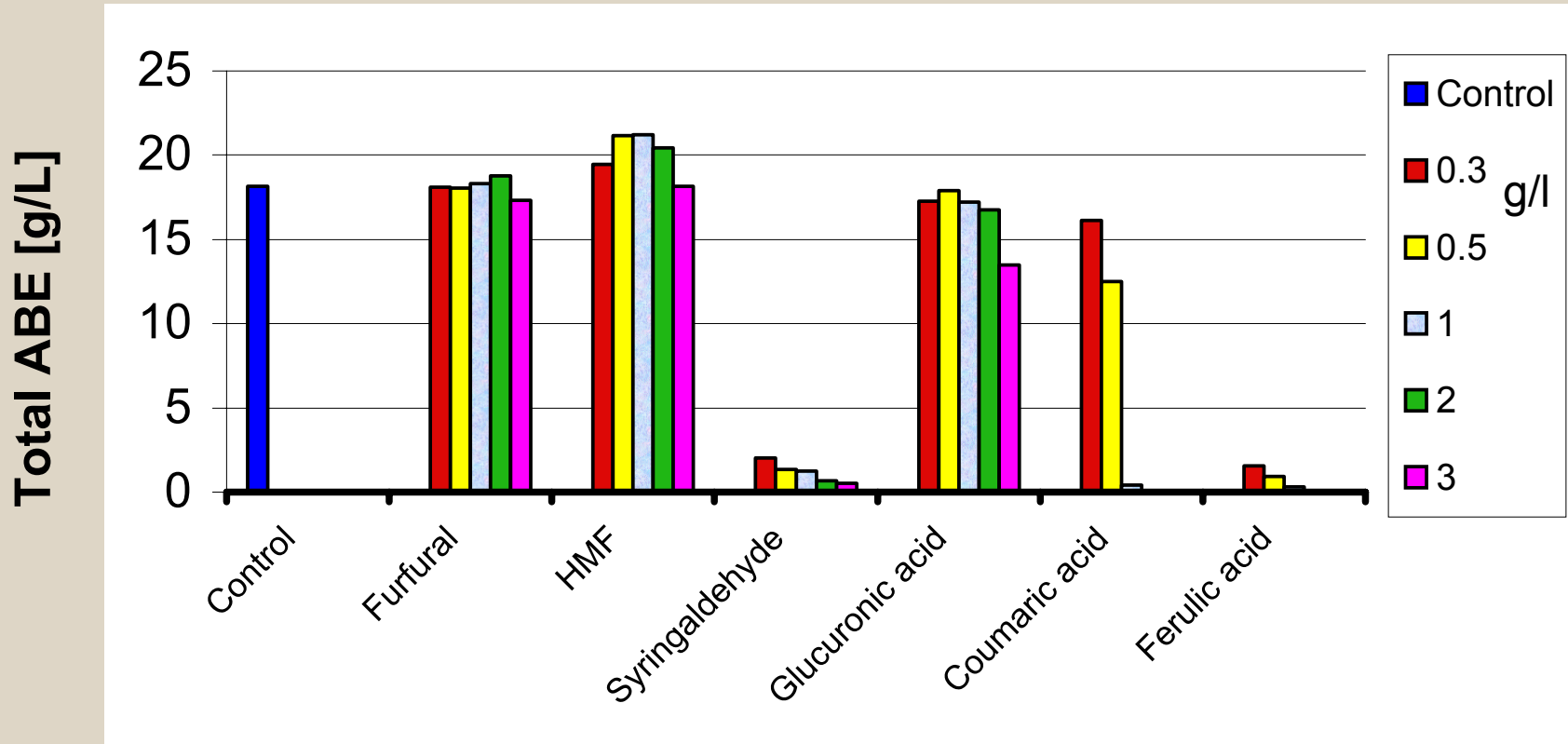
FF: Furfural; HMF: Hydroxymethylfurfural; SY: Syringaldehyde; GA: Glucuronic acid; CA: Coumaric acid; FA: Ferulic acid



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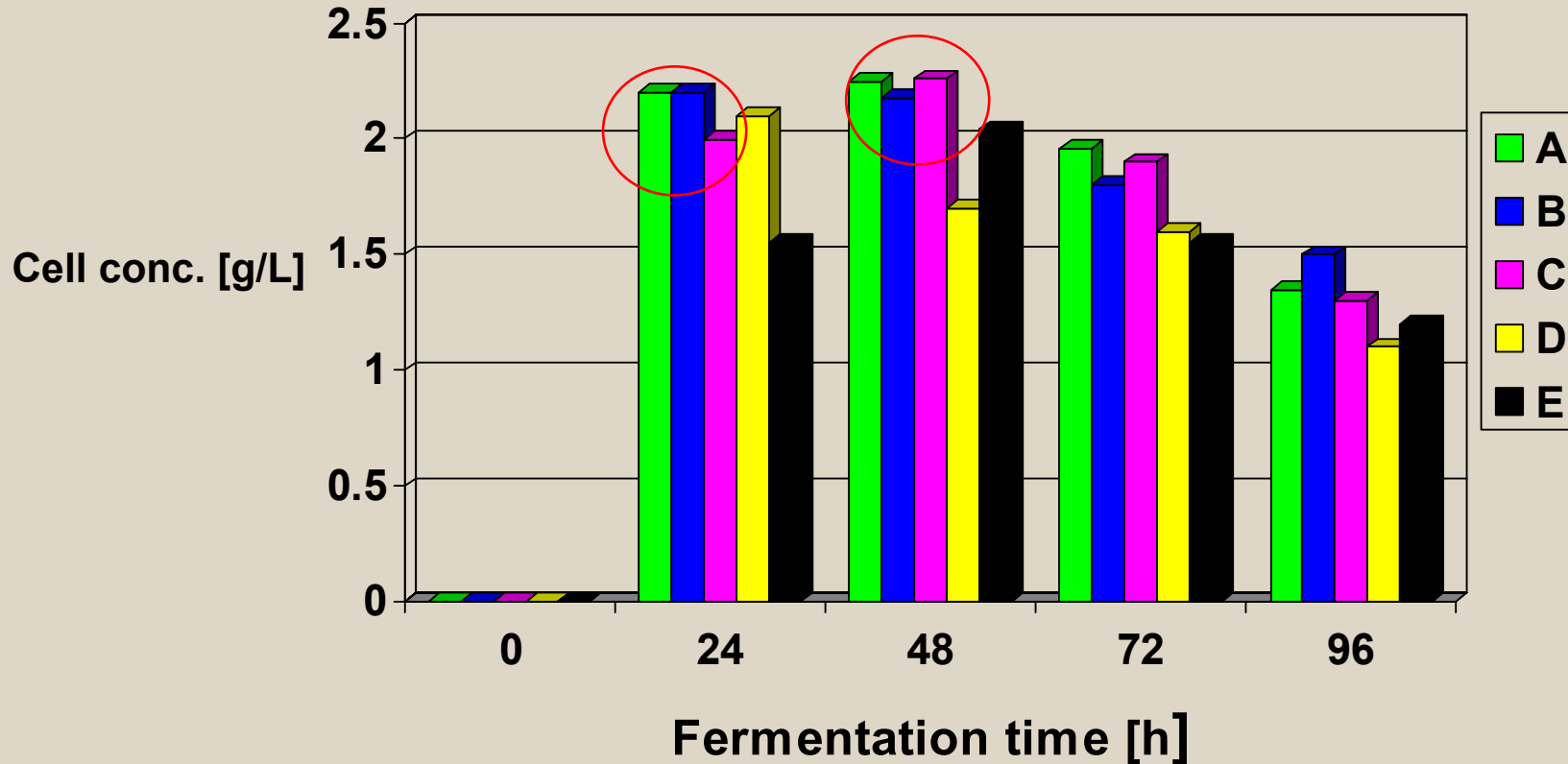
Effect of potential degradation and hydrolysis products on the ABE fermentation by *Clostridium beijerinckii* BA101



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Effect of degradation and hydrolysis products on cell concentration and ABE production by *C. beijerinckii* BA101



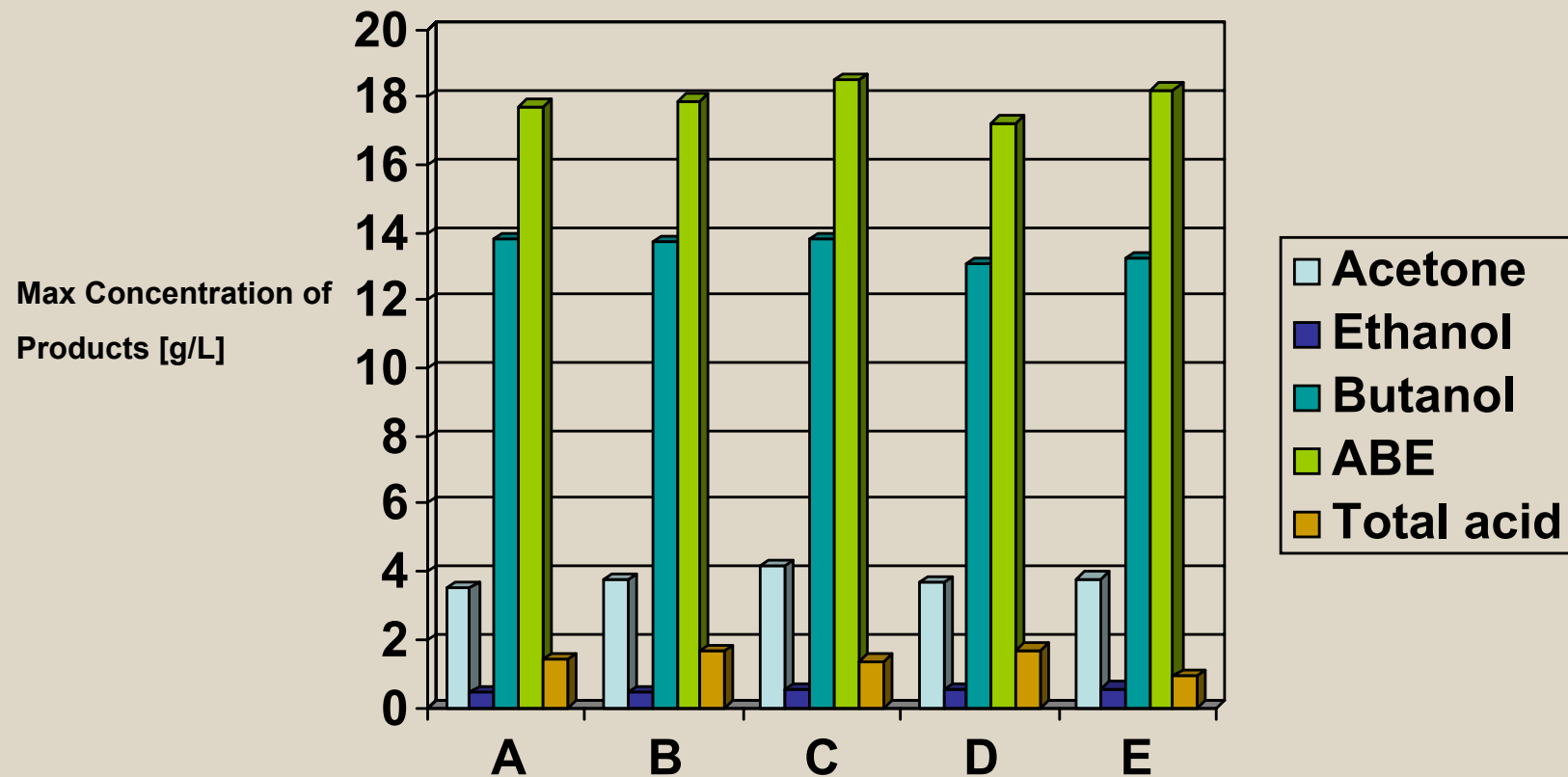
A = Control; B = Furfural + HMF (1.0 g each); C = Furfural + HMF (1.5 g each);
D = Furfural + HMF+ Glucuronic acid (0.67 g each); E = Furfural + HMF + Glucuronic acid (1.0 g each) .



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Effect of degradation and hydrolysis products on cell concentration and ABE production by *C. beijerinckii* BA101



A = Control; B = Furfural + HMF (1.0 g each); C = Furfural + HMF (1.5 g each);
D = Furfural + HMF+ Glucuronic acid (0.67 g each); E = Furfural + HMF + Glucuronic acid
(1.0 g each). Solvent concentration results reported are the maximum produced



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Utilization of hexose and pentose sugars by solventogenic clostridia

Organisms	Sugar preference*
<i>C. acetobutylicum</i> 260	Glucose > cellobiose > mannose > arabinose > galactose > xylose
<i>C. acetobutylicum</i> 824	Glucose > arabinose xylose > cellobiose > galactose > mannose
<i>C. saccharobutylicum</i> 262	Glucose > arabinose > galactose > cellobiose > xylose > mannose
<i>C. butylicum</i> NRRL 592	Glucose > cellobiose > mannose > arabinose > galactose > xylose
<i>C. beijerinckii</i> BA101	Cellobiose > glucose > xylose > arabinose > mannose > galactose

*Preference rated according to maximum ABE produced

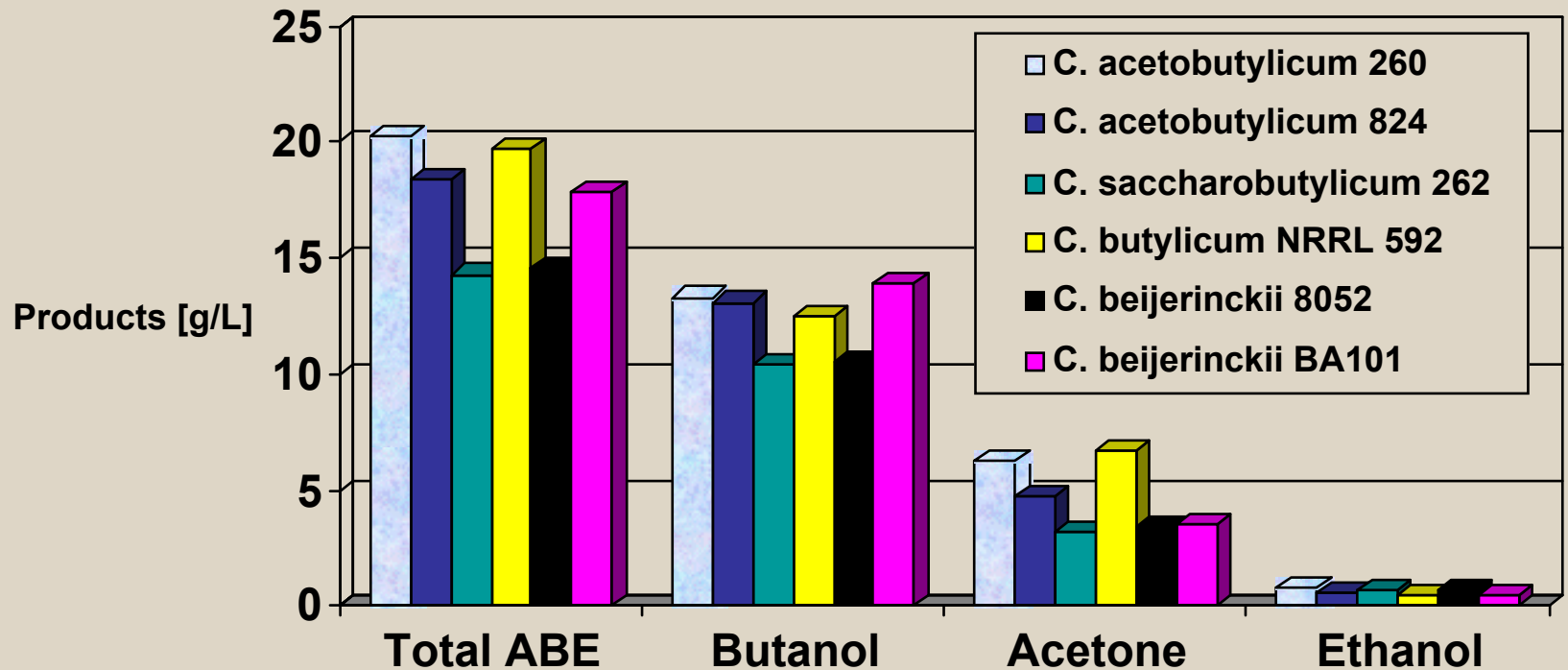


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Fermentation of Hexose and Pentose sugars for ABE production by solventogenic clostridia

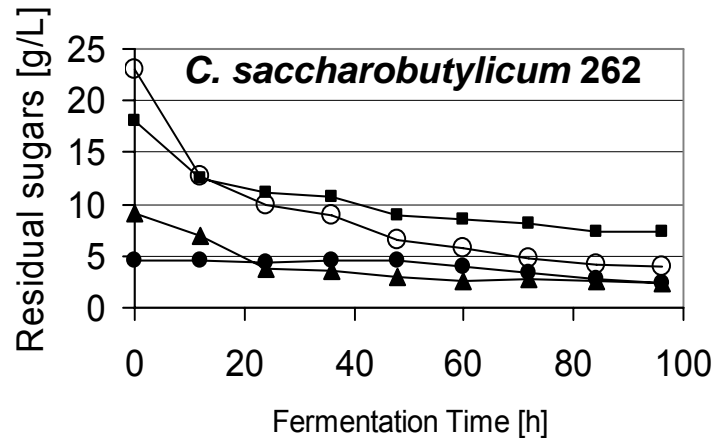
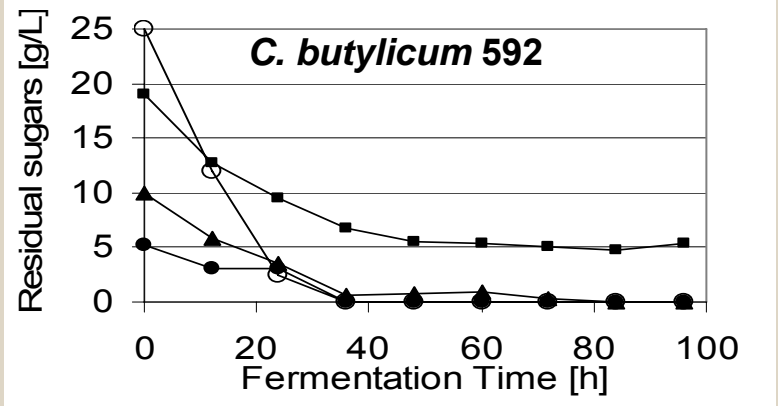
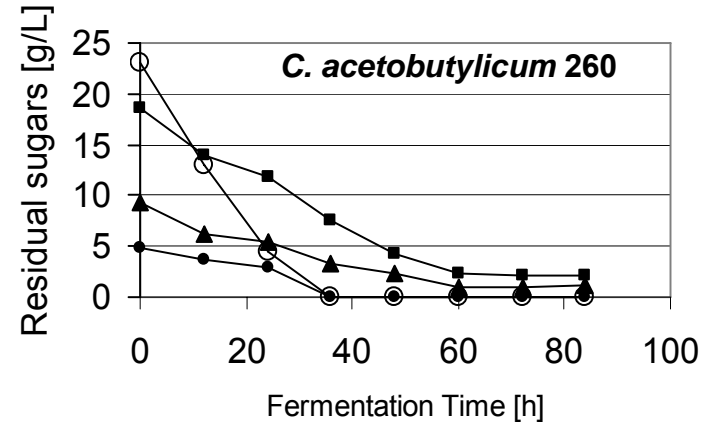
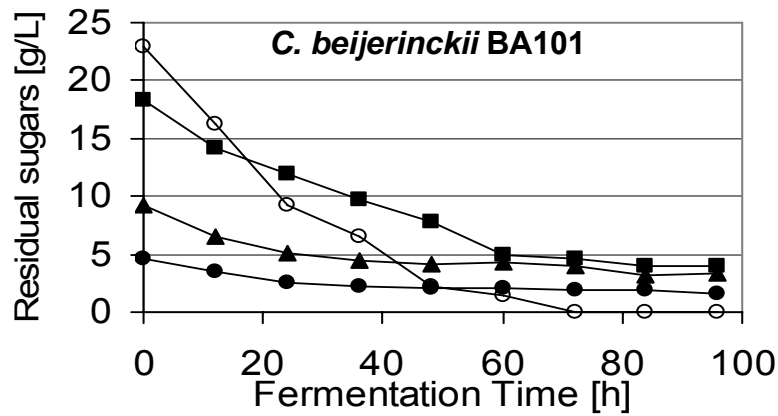
Glucose-mannose-arabinose-xylose composite (g/L: 25:5:10:20)



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Utilization of hexose and pentose sugars by various solventogenic Clostridia



Symbols: ○ , glucose; ■ , xylose; ▲ , arabinose; ● mannose.



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Sugar preference during batch fermentation of mixed sugar streams (grams: glucose 25, mannose 5, arabinose 10, xylose 20) by solventogenic clostridia

Organisms	Fermentation Time [h]*	Sugar preference
<i>C. acetobutylicum</i> 260	60	Glucose> mannose > arabinose>xylose
<i>C. saccharobutylicum</i> 262	72	Glucose > arabinose > xylose> mannose
<i>C. butylicum</i> NRRL 592	60	Glucose > mannose > arabinose > xylose
<i>C. beijerinckii</i> BA101	72	Glucose > xylose > arabinose > mannose

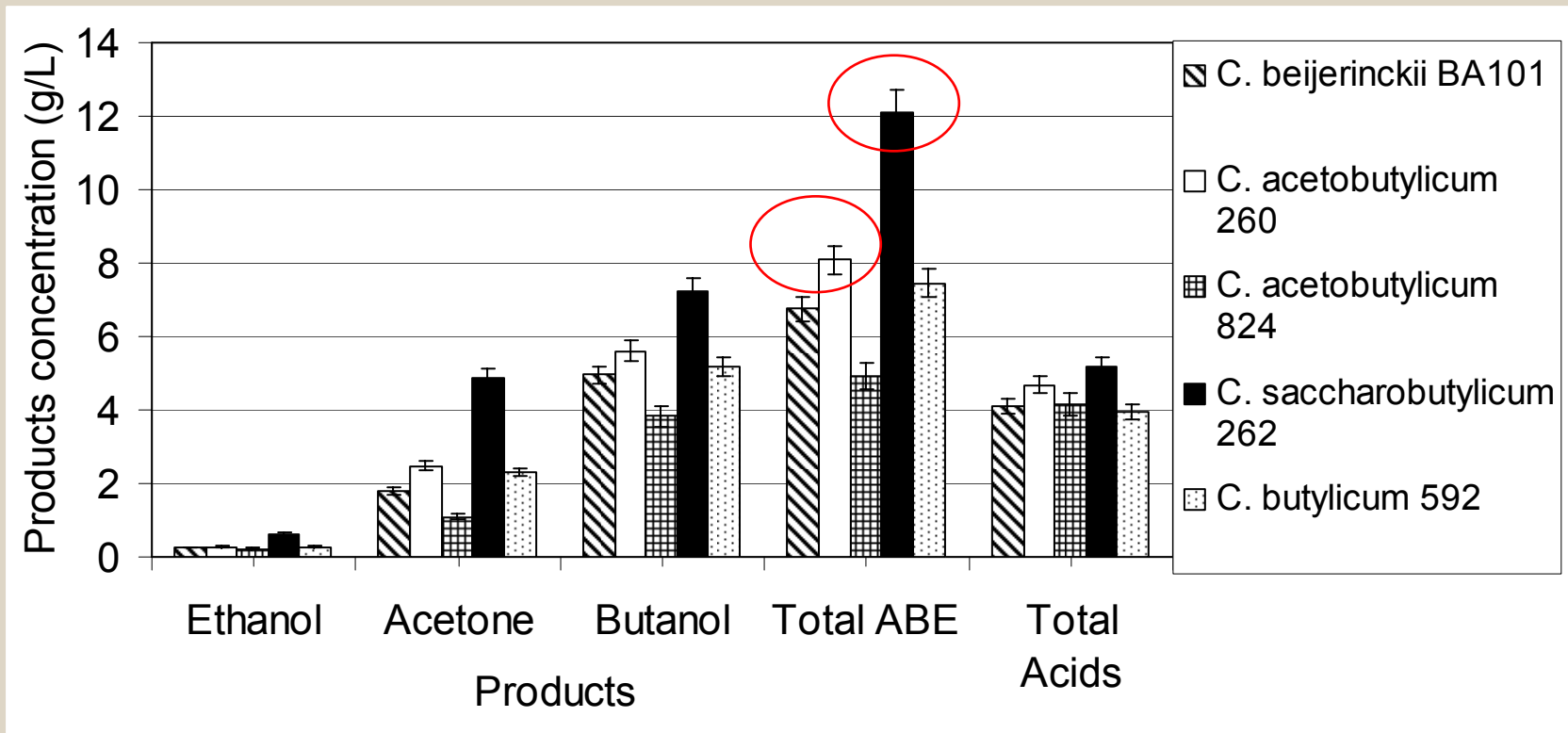
*time required to achieve maximal ABE concentration.



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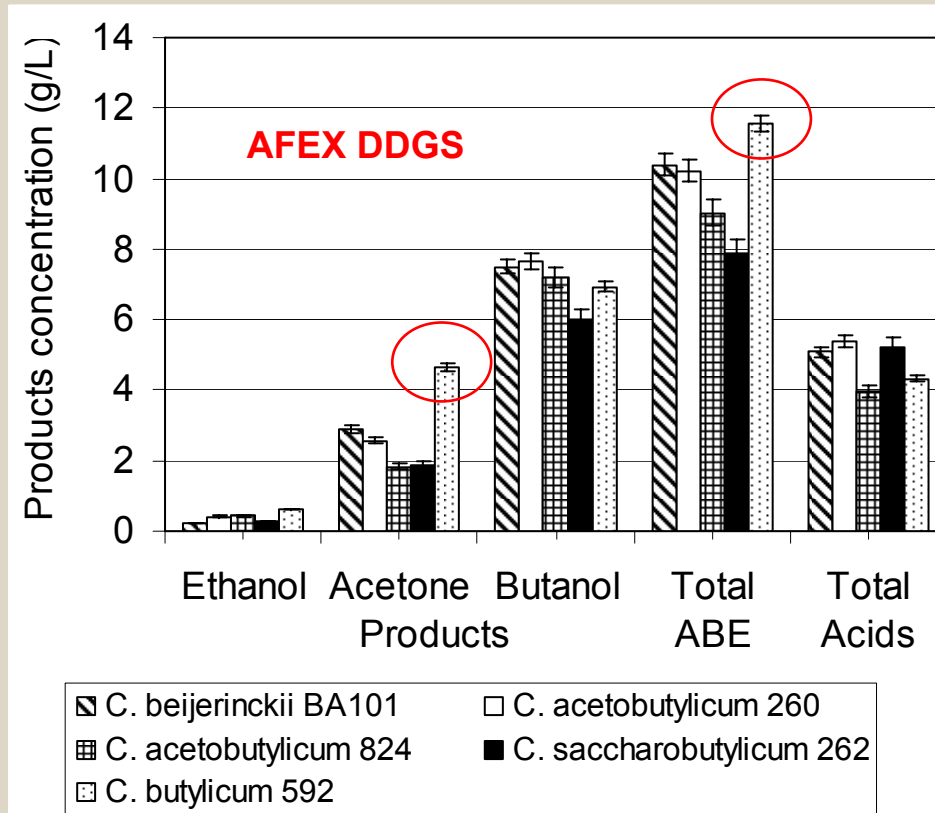
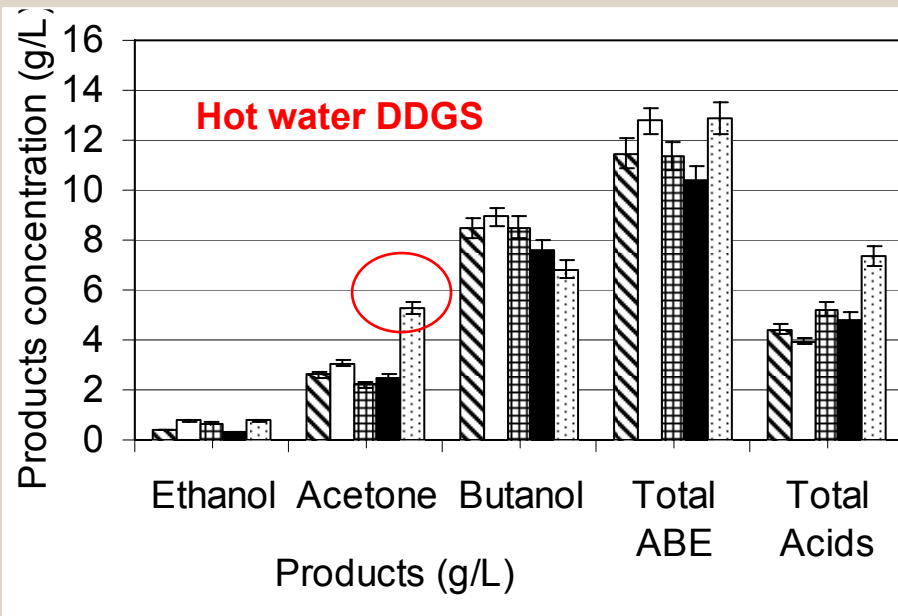
ABE production by solventogenic clostridia using dilute acid pretreated DDGS



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ABE production by solventogenic clostridia using dilute Hot water and AFEX pretreated DDGS



- | | |
|--------------------------------|-----------------------------------|
| ▨ <i>C. beijerinckii</i> BA101 | □ <i>C. acetobutylicum</i> 260 |
| ▩ <i>C. acetobutylicum</i> 824 | ■ <i>C. saccharobutylicum</i> 262 |
| ▤ <i>C. butylicum</i> 592 | |

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Summary

- All the solventogenic clostridia utilized a mixed sugar stream for ABE production. Although glucose is the preferred carbon source, these microbes have the ability to concurrently utilize different (hexose and pentose) sugars for ABE production.
- Furfural and HMF (3g/L) are not inhibitory to *C. beijerinckii* BA101, rather they are stimulatory and the mixture of the two affects the culture negatively.
- Presence of furfural and HMF (2.0 g/L) in the fermentation broth was not inhibitory to other solventogenic clostridia; rather they are stimulatory to some of the clostridia.
- Syringaldehyde, ferulic and p-coumaric acids were potent inhibitors of ABE production by solventogenic clostridia.
- Ferulic acid being the most abundant phenolic acid in maize bran may be the limiting factor in efficient dilute acid treated DDGS and corn fiber fermentation.
- Fermentation of hot water and AFEX pretreated DDGS was successful.



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