Université de Liège

Development of a thermophilic and cellulolytic consortium to improve anaerobic digestion of lignocellulosic biomass. R. Kinet^{1*#}, F. Delvigne¹, J. Destain¹, S. Hiligsmann², P. Thonart¹

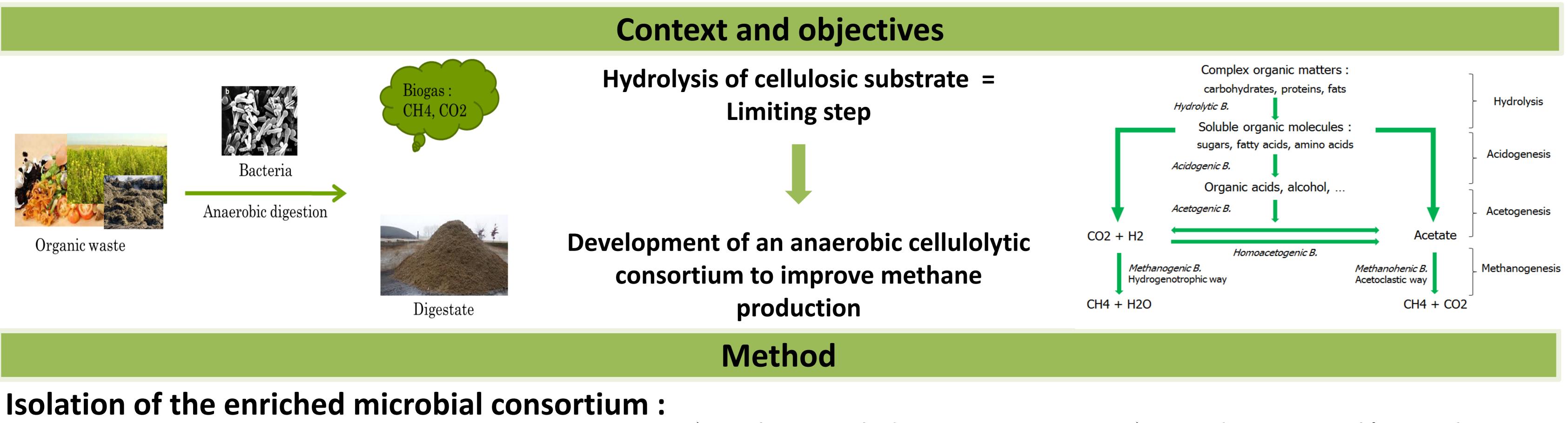


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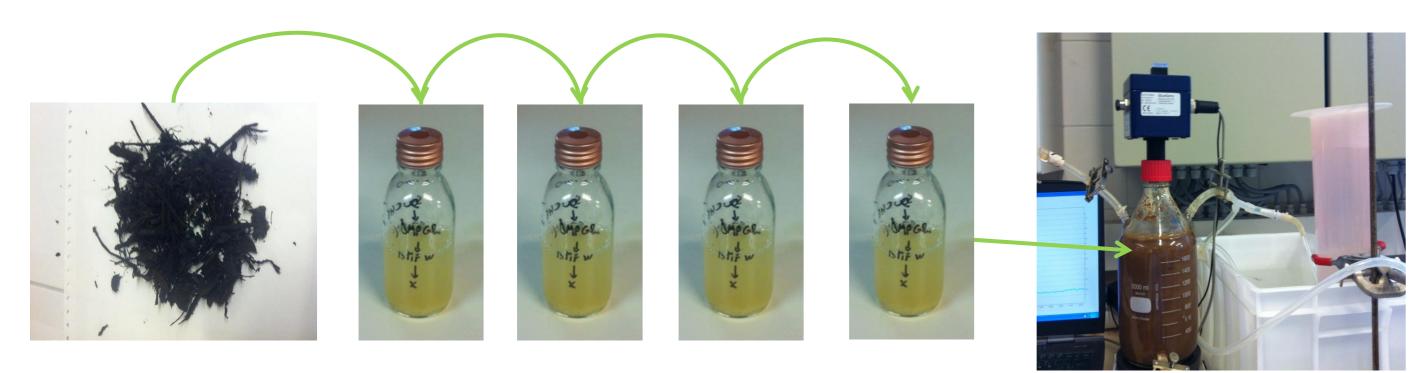


Figure 1: Enrichment method to obtain a cellulolytic, anaerobic and thermophilic consortium from compost. Transfer after 5 days growth at 55°C and in anaerobic static conditions. Scale-up in the fourth transfer.

Enrichment method:

-Compost as microbial inoculum (10%)

-Anaerobia, 55°C, static

-Filter paper 1% (w/v)

-Transfer after 5 days growth

Degradation potential (one week, 10) g/l of substrate) :

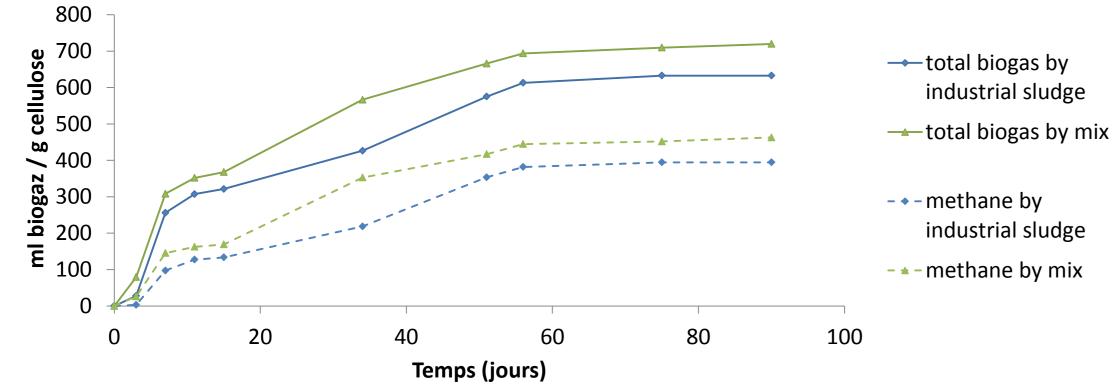
-Filter paper \rightarrow 99%

-Microcrystalline cellulose \rightarrow 98%

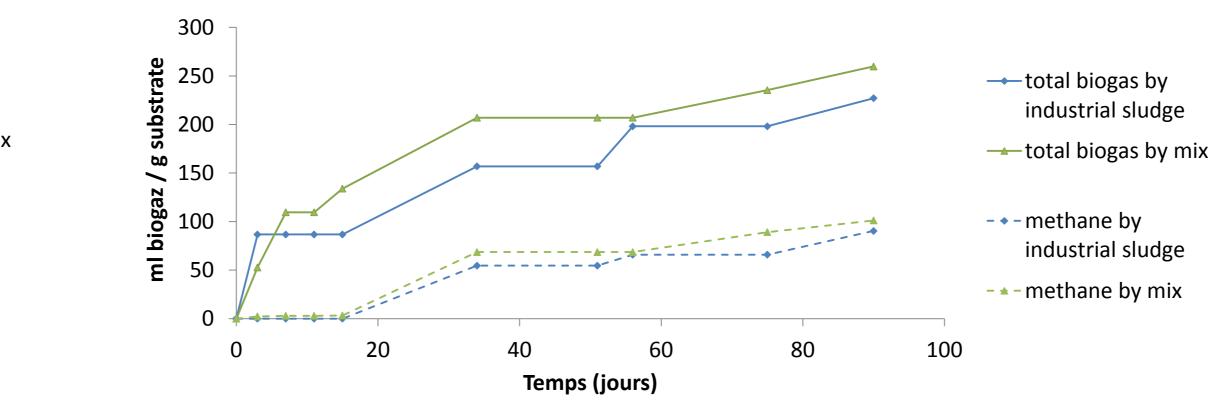
Results

Impact of cellulolytic microbial consortium on biogas production

Cellulosic substrate



🗕 total biogas by industrial sludge Lignocellulosic substrate



 \rightarrow positive effect of the isolated consortium on biogas and methane production by industrial

Figure 2: Evolution of maximal total biogas and methane production (ml/g cellulose) during anaerobic and thermophilic (55°C) digestion of a cellulosic substrate (10 g/l filter paper) by (1) industrial digester sludge (10% v/v) and (2) mix 1:1 of digester sludge and isolated consortium (10% v/v).

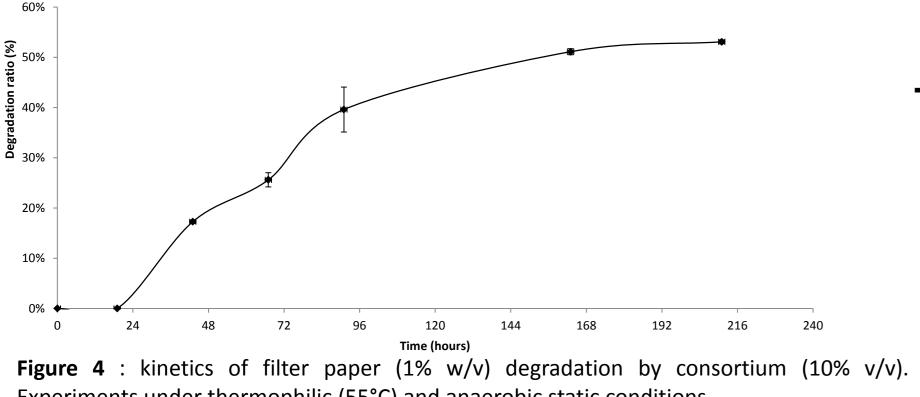
Figure 3 : Evolution of maximal total biogas and methane production (ml/g cellulose) during anaerobic and thermophilic (55°C) digestion of a lignocellulosic substrate (10 g/l mechanical pulp paper) by (1) industrial digester sludge (10% v/v) and (2) mix 1:1 of digester sludge and isolated consortium (10% v/v).

sludge

 \rightarrow gain of 14% biogas in both cases for production and respectively 20% and 12% for methane

Characterization of isolated consortium

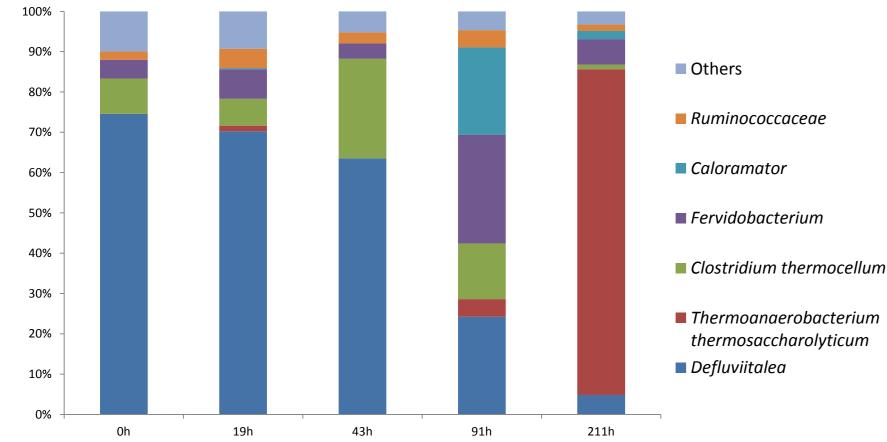
> Cellulose degradation kinetics



 \rightarrow Kinetics similar to sigmoidal microbial growth curve

Experiments under thermophilic (55°C) and anaerobic static conditions.

> Microbial community structure



 \rightarrow pH evolution induces population evolution

> Metabolites of cellulose anaerobic degradation

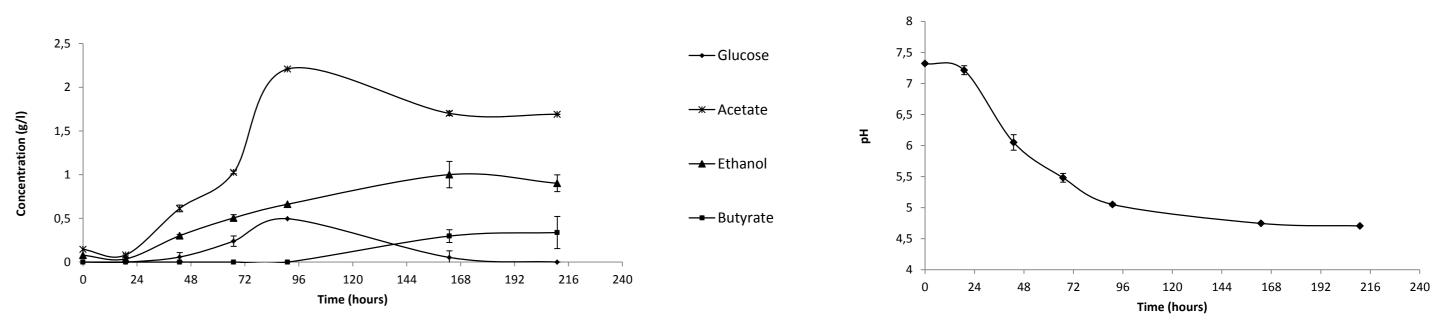


Figure 6 pH evolution during anaerobic thermophilic (55°C) Figure 5 Metabolites' concentration (g/l) evolution during anaerobic thermophilic cellulose (filter paper 10 g/l) degradation by isolated consortium (55°C) cellulose (filter paper 10 g/l) degradation by isolated consortium (10% v/v). (10% v/v).

\rightarrow Glucose accumulation only during the first 91 hours

\rightarrow Acetate and ethanol are predominant metabolites and are produced since the start

Figure 7 Population structure dynamic during anaerobic thermophilic (55°C) degradation of cellulose (filter paper 10 g/l) by isolated consortium (10% v/v).

\rightarrow Cellulose degrading species :

- Clostridium thermocellum
- Thermoanaerobacterium
- thermosaccharolyticum

 \rightarrow Butyrate is measured from 163th hour \rightarrow pH decrease due to VFA accumulation

Conclusions

>Enrichment method, with high temperature compost as microbial source, permitted to obtain an efficient cellulolytic anaerobic and thermophilic consortium

- \blacktriangleright Positive impact of isolated consortium on biogas production.
- \blacktriangleright Acetate and ethanol, are the prevailing metabolites produced during anaerobic cellulose degradation.
- \geq Medium parameters evolution induced important modification of the microbial community structure.

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