



Butanol production by fermentation of *Clostridium acetobutylicum*: solventogenic kinetics

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Acetone-Butanol-Etanol is typically produced during the second stage of batch fermentations of some *Clostridium* strains under selected operating conditions: acids are consumed along with the carbon source and pH increase. The objective of this study was to establish a continuous butanol production system by means of *Clostridium acetobutylicum*, lactose as the sole carbon source, and cell recycling (CSTR equipped with a microfiltration unit). Continuous cultures were carried out under a wide interval of operating conditions (dilution rate and recycle flux) in order to characterize the fermentation process under solventogenesis phase.

Introduction

Depletion of oil resources and concerns regarding both economic and environmental issues have renewed interests for the search of sustainable biofuel produced from renewable resources known as biomass. The spectrum of biofuels includes the butanol, a simple four carbon alcohol characterized by interesting features: low vapour pressure, blending with either gasoline or diesel at any fraction, energy content close to that of the gasoline. Butanol may be produced from renewable resources by the acetone butanol ethanol (ABE) fermentation process [1].

ABE is typically produced during the later stage of batch

fermentations of some *Clostridium* strains under appropriate operating conditions. These strains are able to utilize a wide range of substrates, pentoses, hexoses, mono-, di- and polysaccharides. Under batch

fermentation, the fermentation process proceeds with the production of cells, hydrogen, carbon dioxide and ethanol in the first phase (acidogenesis). As the acid concentrations increases (pH drops) the fermentation shifts to the solventogenesis state with a morphological change.

In the second phase, the cells and endospores are unable to reproduce themselves [2].

The design and the optimisation of continuous bioreactors require to know the kinetics of both phases: acidogenesis and solventogenesis phases [3].

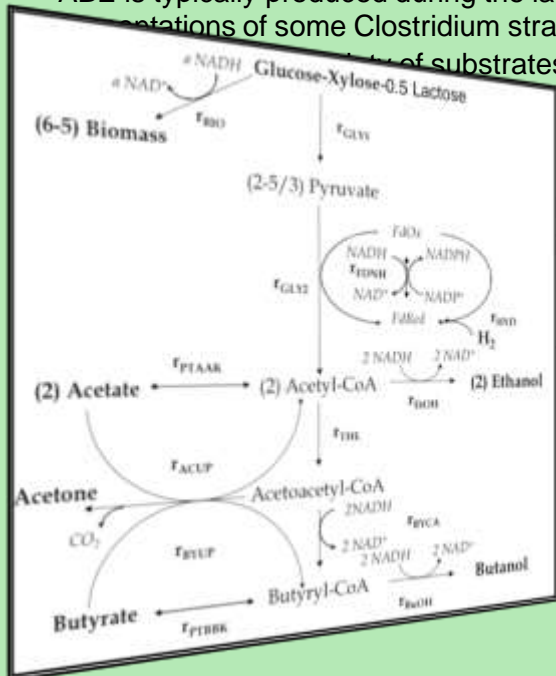


Fig. 1 – *C. acetobutylicum* metabolic pathway and relevant fluxes

Aim

This study is about the continuous butanol production by means of *Clostridium acetobutylicum* adopting lactose as carbon source. A cell recycling reactor system was adopted: CSTR equipped with a microfiltration unit. Continuous cultures were carried out under a wide interval of operating conditions in order to characterize the fermentation process under solventogenesis phase.

Materials & Methods

Continuous fermentations were carried out in the apparatus sketched in Fig. 1 consisting of: a fermenter, a pH controller and a microfiltration unit.

The system was described by the equation set:

$$\begin{cases} D = d_{OUT} + d_p \\ d_{OUT} = \frac{Q_{OUT}}{V} \\ d_p = \frac{Q_p}{V} \\ W_X = D_{OUT} \cdot X_T \end{cases}$$

where W_X is the productivity of biomass and X_T the total biomass.

The cell recycle was characterized in terms of recycle ratio R [4]:

$$R = \frac{Q_p}{Q_0}$$

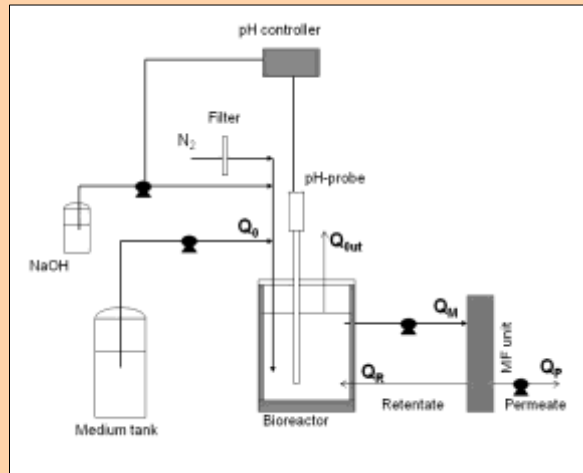
Tests were carried out at:

BIOMASS BALANCE

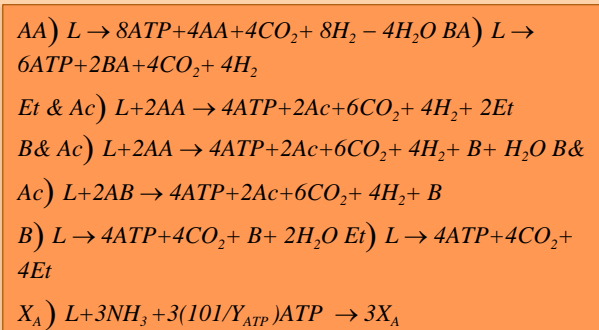
Under continuous solventogenic cultures, the cells may be classified in three groups: acidogenic cells (X_A), solventogenic cells (X_S), and spore (X_D).

$$\begin{cases} W_X = W_X^A + W_X^S + W_X^D \\ W_X^A = \mu_A X_A - \mu_{ly} X_A \\ W_X^S = \mu_S X_S - \mu_{ly} X_S \\ W_X^D = \mu_D X_D - \mu_{ly} X_D \end{cases}$$

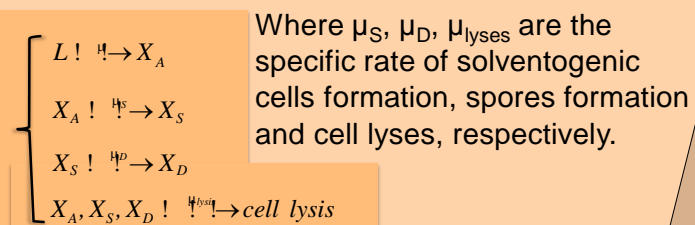
Fig. 1. Outline of the apparatus adopted for continuous tests.



REACTIONS SET



CELLS TRANSFORMATION PATHS



Results & Discussion

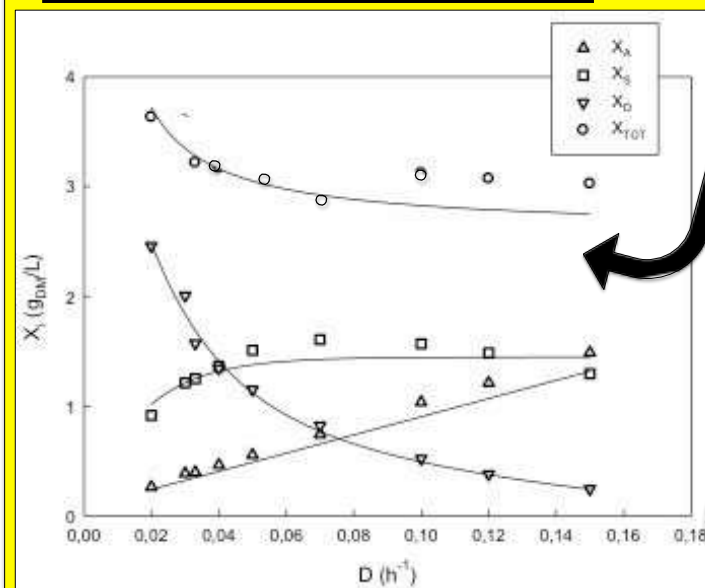


Fig. 3 – Steady state continuous fermentation. Acidogenic cells, solventogenic cells, spore and total biomass vs. dilution rate. $R = 0.88$.

$$\mu = \mu_{max} \cdot \frac{L}{L + K_L} \cdot \frac{AA}{AA + K_{AA}} \cdot \frac{BA}{BA + K_{BA}} \cdot \frac{Ac}{Ac + K_{Ac}} \cdot \frac{Et}{Et + K_{Et}} \cdot \frac{B}{B + K_B}$$

Specific growth rate (μ) of acidogenic cells assessed according to Napoli et al. 2011 [5]

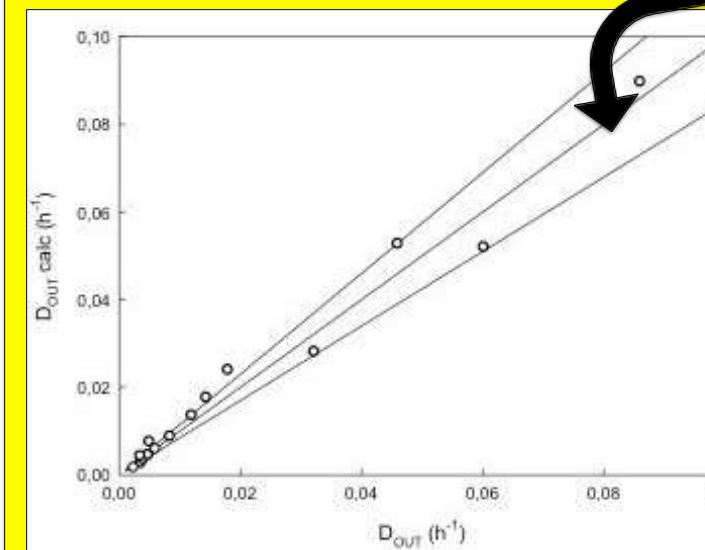


Fig. 5. Calculated vs. measured D_{OUT} . Crosshair: $\pm 10\%$ error

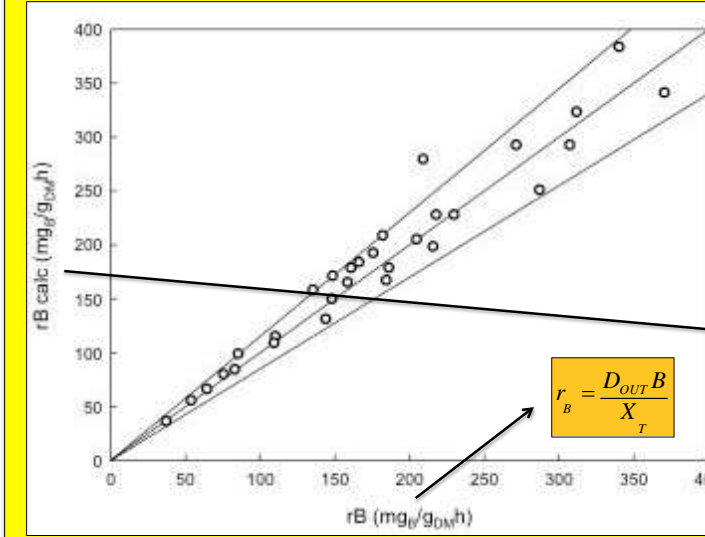


Fig. 7. r calculated vs. r measured. Crosshair: $\pm 10\%$ error.

The complete set of model parameters: r_B^{MAX} , K_B , B^{MAX} , n_B was determined by a parametric inference procedure applied to the total inhibition equation matched against data obtained in experiments carried out in continuous tests.

The concentration of acidogenic cells and solventogenic cells increased progressively with D . On the contrary the concentration of solventogenic cells decrease.

$D < 0.1 h^{-1}$: μ and μ_S increase with D

$D > 0.1 h^{-1}$: μ_S decreases and μ increases with D

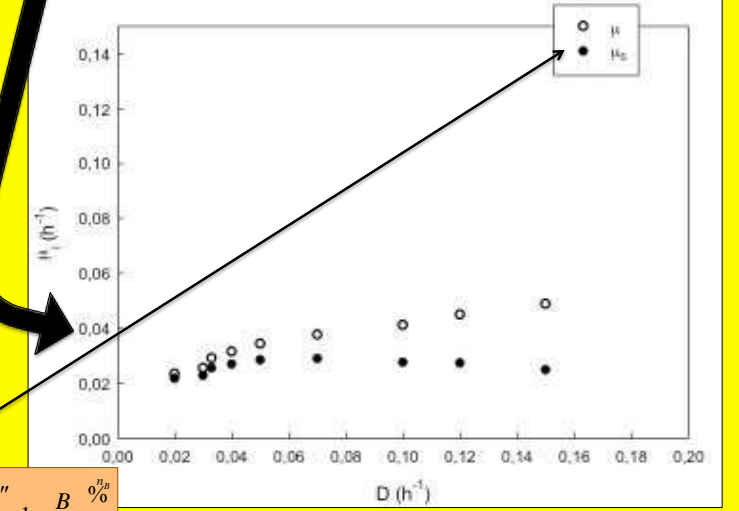


Fig. 4. μ and μ_S vs. dilution rate.

Fig. 5: a parity plot of predicted D_{OUT} vs. experimental D_{OUT} . The analysis points out that the model results are in good agreement with the experimental data points.

The productivities of butanol and ABE increased progressively with D . At $D = 0.15 h^{-1}$ and $R = 0.88$ the ABE production is $0.53 g/L h$.

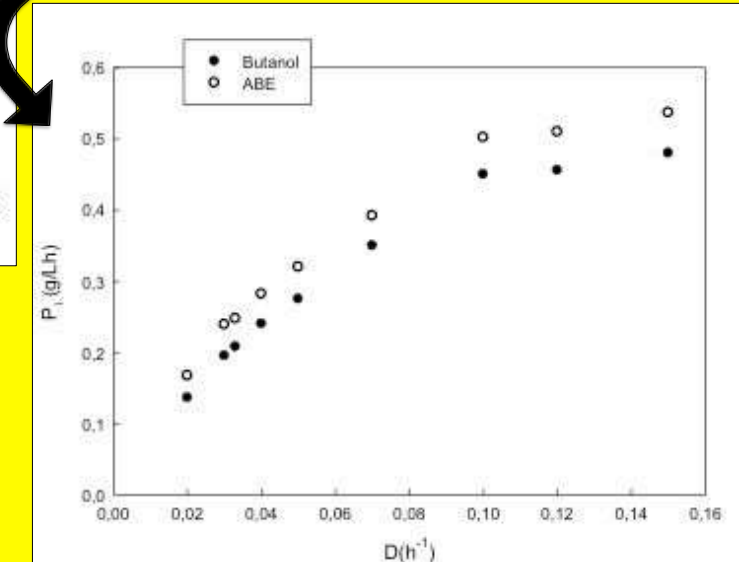


Fig. 6. Productivity vs. dilution rate

Total inhibition equation

$$r_B = r_B^{MAX} \cdot \frac{L}{L + K_B} \cdot \frac{B}{B + B^{MAX}} \cdot \frac{1}{1 + \frac{B}{K_B}^{n_B}}$$

$$\begin{aligned} r_B^{MAX} &= 300 mg / g_{DM} h \\ K_B &= 6.3 g / L \\ B^{MAX} &= 7.045 g / L \\ n_B &= 0.5395 \end{aligned}$$

Acknowledgements

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Main Remarks

- The characterization of *C. acetobutylicum* DSM 792 metabolism on lactose during solventogenic phase was carried out.
- The results have been worked out to assess the concentration of acidogenic cells (X_A), solventogenic cells (X_S), spore (X_D) and the butanol and solvent production.
- The specific butanol production rate (r_B) at different D was obtained. The experimental results were successfully correlated by a product-inhibition model.

References

- Cascone R., 2008, Chem Eng Prog., S4-S9
- Qureshi, N., 2007, Bioprocess Biosyst Eng, 30: 419-427.
- Papoutsakis, E. T., Meyer, C. L., 1989, Bioprocess Eng 4, 49-55. [4] Meyer, C.L., Papoutsakis E.T., 1989, Bioprocess Eng, 4, 1-10.
- Napoli, F., Olivieri, G., Russo, M.E., Marzocchella, A., Salatino, P., 2011, Bioprocess Technol. 102, 1608-1614.