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## QUALITATIVE PROPERTIES OF A 3-STEPS MODEL OF ANAEROBIC DIGESTION INCLUDING HYDROLYSIS OF PARTICULATE MATTER

R. Fekih-Salem<sup>*a,b,c*</sup>, N. Abdellatif,<sup>*c,d*</sup>, T. Sari<sup>*b,e*</sup>, J. Harmand<sup>*b,f*</sup>

<sup>a</sup> UMR INRA-SupAgro MISTEA, 2 p. Viala, 34060 Montpellier, France
 <sup>b</sup> EPI INRA-INRIA MODEMIC, route des Lucioles, 06902 Sophia-Antipolis, France
 <sup>c</sup> Université de Tunis El Manar, ENIT, LAMSIN, B.P. 37, 1002 Tunis Le Belvédère, Tunisie
 <sup>d</sup> Ecole Nationale des Sciences de l'Informatique, Campus Universitaire de Manouba, 2010 Manouba, Tunisie
 <sup>e</sup> Irstea, UMR Itap, 361 rue Jean-François Breton, 34196 Montpellier, France
 <sup>f</sup> Inra, UR 050, Laboratoire de Biotechnologie de l'Environnement, 11100 Narbonne, France
 Corresponding author: R. Fekih-Salem, UMR INRA-SupAgro MISTEA, 2 p. Viala

34060 Montpellier, France, fekih@supagro.inra.fr

**Introduction.** Anaerobic digestion is a biological process in which organic matter is transformed into methane and carbon dioxide (biogas) by microorganisms in the absence of oxygen. The search for models simple enough to be used for control design is of prior importance today to optimize fermentation processes and solve important problems such as the development of renewable energy from waste. Within the studies of microbiology, biochemistry and technology, the anaerobic digestion is generally considered as a three step process: hydrolysis and liquefaction of the large, insoluble organic molecules by extracellular enzymes, acid production by an acidogenic microbial consortium and a methane production stage realized by a methanogenic ecosystem. Several mathematical models describing these phenomena have been proposed in the literature. However, they are usually too complex to be used for control synthesis [1, 2, 6]. For control purposes, the most appropriate way to model the hydrolysis is still an open problem. Many chemists claim that this is a pure enzymatic phenomena while biologists emphasize the role of hydrolytic bacteria. So, there exist several choices :

- One may consider that the microbial enzymatic activity is constant without involving explicitly the biomass;

- One may divide the substrate compartment into two parts : slowly degradable substrates  $X_0$  and readily biodegradable substrate  $S_1$  which may be, for keeping the model simple enough considered to be degraded by the same biomass, *e.g.* a unique compartment grouping "hydrolitic and acidogenesis" bacteria.

The reactional scheme of the anaerobic digestion is given by :

	Macromolecules		
Hydrolysis	<i>X</i> <sub>0</sub>	$\xrightarrow{r_0}$	$k_0S_1$
	Monomers	Acidogenic B.: $X_1$	
Acidogenesis	$k_1S_1$	$\xrightarrow{r_1=\mu_1(S_1)X_1}$	$X_1 + k_2 S_2 + CO_2$
	Organic acids	Methanogenic B.: $X_2$	
Methanogenesis	$k_3S_2$	$\xrightarrow{r_2=\mu_2(S_2)X_2}$	$X_2 + CO_2 + CH_4$

where  $r_0 = k_{hyd}X_0$  or  $r_0 = \mu_0(X_0)X_1$  and  $r_i = \mu_i(S_i)X_i$ , i = 1, 2, denote the reaction rates,  $\mu_0$  is consumption rate of  $X_0$  by  $X_1$  and  $\mu_i$ , i = 1, 2, are the specific growth rates of  $X_i$  on  $S_i$ . Finally,  $k_i$ , i = 0...3, denote the pseudo-stoechiometric coefficients associated to the biological reactions.

**Model.** We consider a continuous culture, i.e the input flow rate is equal to the output flow rate [3]. The modified three step model is :

$$\begin{cases} \dot{X}_{0} = DX_{0in} - \alpha DX_{0} - r_{0}, \\ \dot{S}_{1} = D(S_{1in} - S_{1}) + k_{0}r_{0} - k_{1}\mu_{1}(S_{1})X_{1}, \\ \dot{X}_{1} = \left[\mu_{1}(S_{1}) - \alpha D\right]X_{1}, \\ \dot{S}_{2} = D(S_{2in} - S_{2}) + k_{2}\mu_{1}(S_{1})X_{1} - k_{3}\mu_{2}(S_{2})X_{2}, \\ \dot{X}_{2} = \left[\mu_{2}(S_{2}) - \alpha D\right]X_{2}, \end{cases}$$

$$(1)$$

where  $X_0(t)$  denotes the concentration of the slowly biodegradable substrate (typically the solid Chemical Oxygen Demand) at time *t*, with  $X_{0in}$  the concentration in the input,  $S_j(t)$  denote the concentrations of the substrates in the effluent, j = 1, 2 (the easily biodegradable COD and the Volatile Fatty Acids, respectively), at time *t*; with  $S_{jin}$  the input substrate concentrations *j*,  $X_i(t)$  denote the concentrations of the *i*th population of microorganisms, i = 1, 2, at time *t* (the hydrolitic and acidogens on the first side, and the methanogens on the other side). *D* denotes the dilution rate of the chemostat,  $\alpha \in [0, 1]$  represents the fraction of the biomass leaving the reactor and *V* denotes the volume of the bioreactor. According to the principle of conservation of matter within the reaction scheme we have

$$\int_{t_1}^{t_2} r_0 V d\tau \ge \int_{t_1}^{t_2} k_0 r_0 V d\tau \quad \text{i.e} \quad 1 \ge k_0,$$

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which means that, the quantity of  $X_0$  degraded is greater than or equal to the quantity of  $S_1$  produced. Similarly, we have  $k_1 \ge 1 + k_2$  and  $k_3 \ge 1$ , which means that, the quantity of  $S_1$  degraded is greater than or equal to the quantity of  $X_1$  and  $S_2$  produced. The quantity of  $S_2$  degraded is greater than or equal to the quantity of  $X_2$  produced.

**Results.** In this work, we focus on the mathematical analysis of the model of chemostat with enzymatic degradation of a substrate (organic matter) that can partly be under a solid form [5]. We study the model (1) with a constant hydrolytic kinetics,  $r_0 = k_{hyd}X_0$ , in which the growth rate  $\mu_1$  is monotonic and the growth rate  $\mu_2$  is non monotonic. The first equation of system (1) depends only on the variable  $X_0$ , and  $X_0$  globally converges towards its equilibrium

$$X_0^* = \frac{D}{k_{hyd} + \alpha D} X_{0in}.$$

If  $X_0$  is put equal to its equilibrium, then the last four equations in system (1) reduce to the following AM2 model

$$\begin{cases} S_{1} = D(S_{1in}^{*} - S_{1}) - k_{1}\mu_{1}(S_{1})X_{1}, \\ \dot{X}_{1} = \left[\mu_{1}(S_{1}) - \alpha D\right]X_{1}, \\ \dot{S}_{2} = D(S_{2in} - S_{2}) + k_{2}\mu_{1}(S_{1})X_{1} - k_{3}\mu_{2}(S_{2})X_{2}, \\ \dot{X}_{2} = \left[\mu_{2}(S_{2}) - \alpha D\right]X_{2}, \end{cases}$$

$$(2)$$

where

$$S_{1in}^* = S_{1in} + \frac{k_0 k_{hyd}}{k_{hyd} + \alpha D} X_{0in}.$$

The analysis of (2) was made in [2] and the condition of persistence of the species  $X_1$  in the AM2 model is  $\lambda_1 < S_{1in}$ . Since  $S_{1in}$  was increased (by the effect of hydrolysis) to  $S_{1in}^*$ , the condition of persistence  $\lambda_1 < S_{1in}^*$  of the species  $X_1$  in the model with hydrolysis, shows that the species are favored by the addition of the hydrolysis term.

The study of model (1) with the hydrolytic activity related to biomass,  $r_0 = \mu_0(X_0)X_1$ , was given in [4]. It is derived from a smaller order sub-model since some variables can be decoupled from the others. We study the existence and the stability of equilibrium points of the sub-model considering Monod growth rates and distinct dilution rates. In the classical chemostat model with monotonic kinetics, it is well known that only one equilibrium point attracts all solutions and that bistability never occurs [7]. In the present study, although (i) only monotonic growth rates are considered and (ii) the concentrations of input substrate concentration is less than the break-even concentration, it is shown that the considered sub-model may exhibit bistability. Hence, the importance of hydrolysis in the appearance of positive equilibrium points and the bistability is pointed out.



Existence of two positive equilibria  $E_1^*$  and  $E_1^{**}$ . The system exhibits bi-stability of the washout equilibrium  $E_0$  and a positive equilibrium  $E_1^*$ .

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### References.

- [1] G. Bastin and D. Dochain. On-line estimation and adaptive control of bioreactors, Dynamics of Microbial Competition. 1991.
- [2] B. Benyahia, T. Sari, B. Cherki, and J. Harmand. Bifurcation and stability analysis of a two step model for monitoring anaerobic digestion processes. *Journal of Process Control*, 22:1008–1019, 2012.
- [3] O. Bernard, Z. Hadj-Sadock, D. Dochain, A. Genovesi, and J.-P. Steyer. Dynamical model development and parameter identification for an anaerobic wastewater treatment process. *Biotechnology and Bioengineering*, 75:424–438, 2001.
- [4] R. Fekih-Salem, N. Abdellatif, T. Sari, and J. Harmand. On a three step model of anaerobic digestion including the hydrolysis of particulate matter. In *Proceedings of the 7th Vienna International Conference on Mathematical Modelling*, page 196, 2012.
- [5] C. Lobry and J. Harmand. A new hypothesis to explain the coexistence of n species in the presence of a single resource. CRAS série Biologie, 329:40–46, 2006.
- [6] I. Simeonov and S. Stoyanov. Modelling and dynamic compensator control of the anaerobic digestion of organic wastes. *Chem. Biochem. Eng. Q*, 17:285–292, 2003.
- [7] H.L. Smith and P. Waltman. The Theory of the Chemostat, Dynamics of Microbial Competition. Cambridge University Press, 1995.