

# STATE VARIABLES MONITORING USING A CLASS OF NONLINEAR OBSERVER BASED ESTIMATOR, APPLIED TO CONTINUOUS BIO-SYSTEM

Ricardo Aguilar-López, Ricardo Acevedo-Gómez, María Isabel Neria González\* and Alma Rosa Domínguez-Bocanegra

Departamento de Biotecnología y Bioingeniería  
CINVESTAV-IPN

Av. Instituto Politécnico Nacional No. 2508, San Pedro Zacatenco, México, D.F. 07360, MEXICO

E-mail.- [raguilar@cinvestav.mx](mailto:raguilar@cinvestav.mx)

Phone.- + 52 55 5747 3800, ext. 4307

\*División de Ingeniería Química y Bioquímica

Tecnológico de Estudios Superiores de Ecatepec

Av. Tecnológico s/n Esq. Av. Carlos Hank González Col. Valle de Anáhuac C.P. 55210, Ecatepec Estado de México

## ABSTRACT

In this work, the state estimation of key variables such as biomass and products of a sulfate reducing bacterium is predicted by using only sulfate (substrate) concentration measurements under the assumption of an unknown kinetic term. The process was developed by continuous culture, where the mathematical kinetic model for the biomass, sulfate and sulfide concentrations is presented and tuned using experimental data. The design of the nonlinear state estimator takes into account an adaptive gain. The results of the proposed estimation methodology were generated via numerical simulation; they showed a satisfactory performance.

KEYWORDS: robust estimation, bioreactor model, anaerobic biosystem

## RESUMEN

En este trabajo se predice la estimación de estados como la biomasa y la concentración de productos de una bacteria sulfato reductora empleando únicamente mediciones de la concentración de sulfato (sustrato) bajo la hipótesis de una cinética de consumo de sulfato desconocida. El proceso fue considerado como un cultivo continuo, donde el modelo cinético correspondiente para las concentraciones de biomasa, sulfato y sulfuro fue corroborado con datos experimentales. El diseño de la metodología de estimación propuesta considera una ganancia adaptable. Los resultados de estimación fueron generados por medio de simulaciones numéricas y muestran un comportamiento satisfactorio.

Palabras Clave: Estimación robusta, modelo de un birreactor, bio-sistema anaerobio

## 1. INTRODUCTION

The lack of cheap and reliable instrumentation for the online measurement of the relevant variables in many processes definitely constitutes a serious obstacle for the development of biological systems. One way to overcome this problem is to use software sensors. A software sensor can be described as the relationship between sensor hardware and estimator software. Biological processes have become widely used in the industry for the last decades, with different purposes: either to produce some chemical compounds synthesized by a microorganism to cultivate a biomass for its utilization or extraction of its metabolites or to degrade a pollutant. Therefore, bioreactors require advanced monitoring procedures to ensure the performances and efficiency bioprocesses operation [1, 2].

The estimation of the key variables such as specific substrate consumption rates, specific microbial growth rates, products and biomass concentrations are very valuable tools to analyze the performance of biotechnological systems. In particular, sulfate-reducing bacteria (SRB) are anaerobic microorganisms of great ecological importance in the global carbon and sulfur cycles, they oxidize the organic compounds coupled to the reduction of sulfate; therefore, they are used on anaerobic processes for wastewater treatment [3-5].

The sulfate-reducing bioreactors have been constructed from some biotechnological processes, as mentioned above, in which the monitoring of increases in the cell mass and products is difficult to measure given the anaerobic process conditions. Different methods for detection and enumeration of SRB in natural and industrial environments have been developed; they have been grouped in (i) direct detection methods and (ii) culture methods [6]. The direct detection involved the use of antibodies raised against SRB [7] and the use of molecular biology tools such as the 16S rRNA analysis; both techniques may be used *in situ* but required of a bigger knowledge and, in some cases, their use was not possible because of the nature of the considered sample. Culture methods for enumeration of SRB require of strict anaerobic conditions and special culture medium, experience of handling of these bacterial, the incubation times are sometimes very large, etc.[8].

Following these ideas, the estimation theory deserves an interesting research field because the estimation methodologies developed are widely employed in on-line monitoring, fault detection, control process and so on. Some of the most important estimation methodologies are related with the observers design where nonlinear Luenberger-type filters, Kalman filters, sliding-mode observers, and so on [9-13] have been presented in the open literature. On the other hand, some techniques such as neural-networks have been successfully used too [13, 14]. Previous published work on observer design also include finite time observers, algebraic estimators, observers with modeling uncertainties and so on [15-18]; however, several of these methodologies are developed on a complex mathematical frame and, consequently, the possibility of on-line implementation is hard. In this paper, it is considered the application of a class of state observer structure based on uncertainty estimator in order to estimate the key variables as biomass concentration and sulfide production in a continuous culture of *Desulfovibrio alaskensis*, where an adaptive gain based on a fractional power of the absolute value of the estimation error is considered in the observer structure. This adaptive gain coupled with a proportional term of the estimation error provides asymptotic and exponential convergence; this proposed structure looks very simple and the possible real time implementation would be feasible.

## 2. EXPERIMENTAL

*Desulfovibrio alaskensis* 6SR strain was isolated of an oil pipeline [19]. Previously, the strain was cultured in Ravot medium [20] for 15 days at 32 °C under an atmosphere of N2-CO2 (80:20, v/v). Congenital water medium (CW). A sample of congenital water was obtained from an oil pipeline located in the Mexican southeast region. Chemical determination of water: chlorides 64 000 ppm, sulfur 178 ppm, sulfate 350 to 400 ppm, pH 8.84. A 1000 mL aliquot of congenital water was saturated with N2 for 1 hour and was enriched with sodium lactate 6 mL, yeast extract 0.5 g, and reducing solution 5 mL (acid ascorbic 1 g/L, and sodium thioglycolate, 1 g/L). The pH was adjusted to 7 with KOH 1N. The CW medium was distributed in 60 mL serum bottles using the Hungate technique [21] and they were autoclaved at 120 °C for 15 min. The cultures initiate from *D. Alaskensis* in medium Ravot were used to inoculate 45 mL of CW medium. The culture was incubated for 20 days to 37 °C. This was used to inoculate three bottles with CW medium to different time: zero, 24 and 36 hours, respectively, and were incubated under same conditions. The bacterial growth was followed through Optical Density (OD). Samples from the cultures were taken anaerobically each hour. Sulphate in the medium was measured by the turbidimetric method based on the precipitation of barium [22]. Also, the production of sulfide was measured by a turbimetric method [23]. The OD reading for cell growth was transformed to dry weight (mg/mL) through a standard curve of growth.

### 3. MATHEMATICAL MODEL OF THE BIOREACTOR

Anaerobic bioreactors are large fermentation tanks provided with mechanical mixing, heating, gas collection, sludge addition and withdrawal ports, and supernatant outlets that can be considered as continuous stirred tanks for analysis purposes. However, for estimation purposes, a reduced order model which can describe the dynamic behavior of the main state variables is adequate. The kinetic parameters were determinated via standard methodology [24] in a batch culture.

If the specific growth rate follows a Monod model, i. e.:

$$\mu(S) = \mu_{\max} \frac{S}{k_S + S} \text{. The corresponding kinetic parameters can be fitted by plotting } \mu^{-1} \text{ versus } S^{-1}.$$

Therefore,

$$\mu(S) = \frac{0.035S}{0.9 + S}$$

with  $Y_{S/X} = 0.25$  and  $Y_{P/X} = 0.263$ .

Considering the above kinetic model, it is proposed the following mathematical model for a class of continuous stirred bioreactor based on classical mass balances for biomass, sulfate (substrate) and sulfide (product) concentrations:

Biomass (X).-

$$\frac{dX}{dt} = -DX + \mu(S)X \quad (1)$$

Sulfate (S).-

$$\frac{dS}{dt} = D(S_{in} - S) - \mu(s) \frac{X}{Y_{S/X}} \quad (2)$$

Sulfide (P).-

$$\frac{dP}{dt} = -DP + \mu(S) \frac{X}{Y_{P/X}} \quad (3)$$

Here  $D$  is the dilution rate,  $\mu$  is the specific growth rate,  $Y_{S/X}$  is the sulfate coefficient yield and  $Y_{P/X}$  is the sulfide coefficient yield. In accordance with the specific experimental setup, the following initial conditions are considered for the batch culture and model validation purposes:  $X_0 = 0.12 \text{ g/L}$ ,  $S_0 = 5 \text{ g/L}$ ,  $P_0 = 0.16 \text{ g/L}$ . Figure.1 shows the performance of the kinetic model considering a comparison with the experimental data which looks satisfactory.

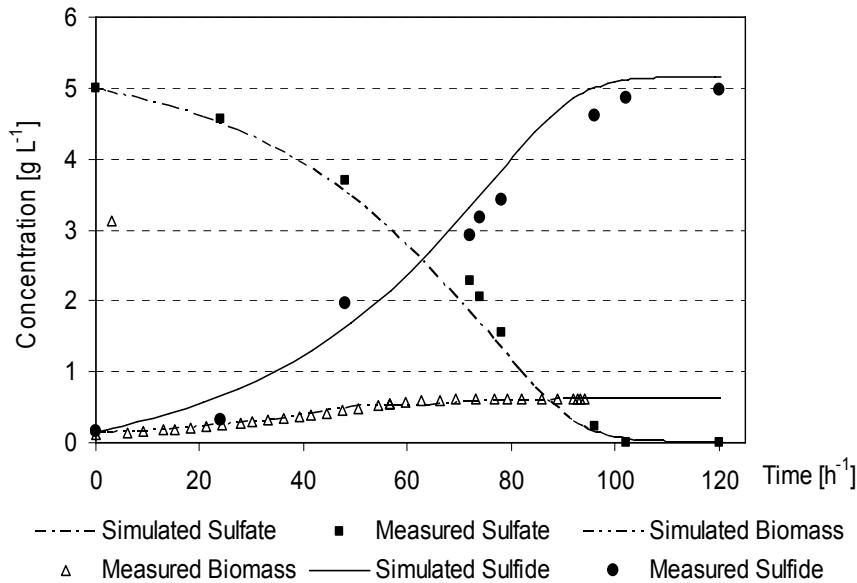


Figure 1. Kinetic model validation with experimental data.

#### 4. METHODOLOGY FOR THE OBSERVER DESIGN

Consider a canonical form of the bioreactor model:

$$\begin{aligned} \dot{x} &= f(x) + g(x)u \\ y &= h(x) = Cx \end{aligned} \tag{5}$$

Here,  $x \in \mathbb{R}^n$  is the vector of states;  $u \in \mathbb{R}^q$  is the vector control input;  $f(\cdot) : \mathbb{R}^n \rightarrow \mathbb{R}^n$  is a nonlinear, partially known vector field;  $g(\cdot) : \mathbb{R}^n \rightarrow \mathbb{R}^n$  is a linear vector of arguments and  $y \in \mathbb{R}^m$  is the system measured output. Now, consider the following assumptions:

**A1.** The system given by Equations (1a) and (1b) is locally uniformly observable, hence, for all,  $x \in \mathbb{R}^n$  and  $u \in \mathbb{R}^q$  satisfies the observability rank condition:

$$\text{rank} \left\{ \frac{\partial}{\partial x} \vartheta \right\} = n.$$

Here  $\vartheta$  is the observability vector function defined as

$\vartheta = (dL_f^0 h_1, \dots, dL_f^0 h_m, \dots, dL_f^1 h_1, \dots, dL_f^1 h_m, \dots, dL_f^{n-1} h_1, \dots, dL_f^{n-1} h_m)^T$ , being  $L_f^r h_s$  ( $s = 1, 2, \dots, m$ ) the  $r$ -order Lie derivatives.

Now, consider  $g(x) = \bar{g}(x) + \Delta g(x)$ , and functions  $f(x)$  and  $\Delta g(x)$  are model uncertainties related to the non-linear system,  $\bar{g}(x)$  is a nominal value of the control input coefficient. In the most general case, functions  $f(x)$  and  $\Delta g(x)$  are assumed to be unknown. Defining:

$$\zeta(x, u) = f(x) + \Delta g(x)u \quad (6)$$

By introducing (6) into (5), a new representation of the system is obtained (Equation 7).

$$\begin{aligned} \dot{x} &= \zeta(x, u) + \bar{g}(x)u \\ \dot{\zeta} &= \Phi(x, u) \\ y &= h(x) = Cx \end{aligned} \quad (7)$$

In order to simplify notation, this set of equations can be written in vector notation (Equation 8).

$$\begin{aligned} \dot{\Gamma} &= \Theta \\ y &= C\Gamma \end{aligned} \quad (8)$$

Here  $\Gamma = \begin{bmatrix} x \\ \zeta \end{bmatrix}$ ;  $\Theta = \begin{bmatrix} \zeta(x, u) + \bar{g}(x)u \\ \Phi(x, u) \end{bmatrix}$ . The procedure described below provides a method to estimate the

uncertainty term,  $\zeta(x, u)$ . Estimators or observers for states and uncertainties can play a key role during the early detection of hazardous and unsafe operating conditions. Following this spirit, several researches have been focused in the proposition of estimation methodologies for states and uncertainties for monitoring and control purposes [25-28].

Now, the following state observer is proposed:

$$\begin{aligned} \dot{\hat{x}} &= \hat{\zeta}(\hat{x}, u) + \bar{g}(\hat{x})u + k_1(y - \hat{y}) \\ \dot{\hat{\zeta}} &= k_2(y - \hat{y}) \\ \dot{K} &= -\lambda|y - \hat{y}|^{1/m} \end{aligned} \quad (9)$$

By defining  $\hat{\Gamma} = \begin{bmatrix} \hat{x} \\ \hat{\zeta} \end{bmatrix}$ ,  $\hat{\Theta} = \begin{bmatrix} \hat{\zeta}(\hat{x}, u) + \bar{g}(\hat{x})u \\ 0 \end{bmatrix}$  and  $K = [k_1 \ k_2]$ , Equation (9) can be rewritten as

$$\begin{aligned}\dot{\hat{\Gamma}} &= \hat{\Theta} + K(y - \hat{y}) \\ \dot{K} &= -\lambda|y - \hat{y}|^{1/m}\end{aligned}\tag{10}$$

Here the dynamic equation for  $K$  is an adaptation algorithm that updates the time-varying observer gain and  $\beta$  is a parameter design.

In order to prove the convergence of the proposed observer, let's consider the dynamic equation of the estimation errors,  $\varepsilon = \Gamma - \hat{\Gamma}$ , as follows:

$$\begin{aligned}\dot{\varepsilon} &= \mathfrak{I} - \hat{\mathfrak{I}} + K\varepsilon \\ \dot{K} &= -\lambda|\varepsilon|^{1/m}\end{aligned}\tag{11}$$

Because the error is a finite quantity, there should be a constant  $L$  that

$$\mathbf{A2.} |\Theta - \hat{\Theta}| \leq L|\Gamma - \hat{\Gamma}|$$

Taking norms to both sides of equation (11) and applying **A2**, it is obtained

$$\left| \dot{\varepsilon} \right| \leq L|\varepsilon| + K|\varepsilon| \tag{12}$$

Now, to solve the system given by Equation (11), consider function  $|\varepsilon|$  as a positive continuous function on the integration interval  $[a, b]$ ; if  $\Psi$  is the maximum of the function on the domain  $[a, b]$ , then  $|\varepsilon|$  is bounded, i.e.  $|\varepsilon| \leq \Psi \quad \forall t \in [a, b]$ , hence

$$|\varepsilon|^{1/m} \leq \Psi^{1/m} \quad m > 0 \Rightarrow \int_a^b |\varepsilon|^{1/m} \leq \Psi^{1/m}(b-a) \tag{13}$$

Here,  $m$  is restricted to be an odd number i.e.  $m = 2p + 1, p \in \mathbb{Z}^+$ . Therefore, for  $p$  large enough, the following limit is obtained:

$$\limsup \int_a^b |\varepsilon|^{1/(2p+1)} \leq \limsup \Psi^{1/(2p+1)}(b-a) \leq (b-a) \tag{14}$$

Applying the equality  $|\varepsilon| = \text{sign}(\varepsilon)\varepsilon$  to equation (12), another quota can be found

$$\text{sign}(\dot{\varepsilon})\dot{\varepsilon} \leq (L - \lambda(b-a))\text{sign}(\varepsilon)\varepsilon \tag{15}$$

By solving Equation (15), it is possible to note that the error is bounded by

$$\varepsilon \leq \varepsilon_0 \exp \left( \text{sign} \left( \frac{\cdot}{\varepsilon} \right)^{-1} \text{sign}(\varepsilon) (L - \lambda(b-a)) t \right) \quad (16)$$

Therefore, the tracking error will be asymptotically and exponentially stable if

$$\lambda > L(b-a)^{-1} \quad (17)$$

## 5. NUMERICAL EXPERIMENTS AND DISCUSSION

A mathematical model of the cellular growth of a sulphate-reducing bacterium (*Desulfovibrio alaskensis*) is presented; this model is employed as a real system for the proposed observer design where the numerical simulation of the ordinary differential equations is done via ode23s Math lab® library reproducing adequately the corresponding experimental data. The proposed estimation procedure considers the sulfate concentration (substrate) as measured output in order to infer the biomass, sulfide concentrations and the sulfate consumption rate (uncertain term), which are observable. The observer takes into account the following initial conditions: ( $X_0 = 0.1 \text{ g/L}$ ,  $S_0 = 5 \text{ g/L}$ ,  $P_0 = 0.16 \text{ g/L}$ ,  $\zeta_0 = 1.0 \text{ g/L h}$ ) and for illustration purposes, a set of nominal operation conditions is considered ( $D = 0.25 \text{ 1/hour}$  and  $S_{in} = 5 \text{ g/L}$ ) and the observer gain of  $\lambda = 0.005 \text{ 1/hr}$ . Figure 2 is related to the uncertainty (substrate consumption rate) estimation, as can be seen, a satisfactory performance is reached; from the start of the estimation procedure a small overshoot is presented, however, at 35 hours the uncertainty estimator faster converges to the real value. Figure 3 shows the states (sulfate, biomass and sulfide concentrations) estimation for the sulfate concentration estimation note an improved convergence; this is due to the fact that the sulfate concentration is the corresponding measured output. For the other concentrations, it is observed a fast convergence to the real trajectory, with a small offset which can be diminished via high values of the observer's gain.

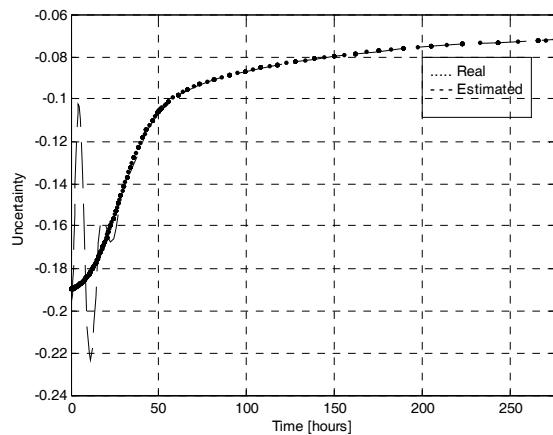


Figure 2. Uncertainty estimation performance

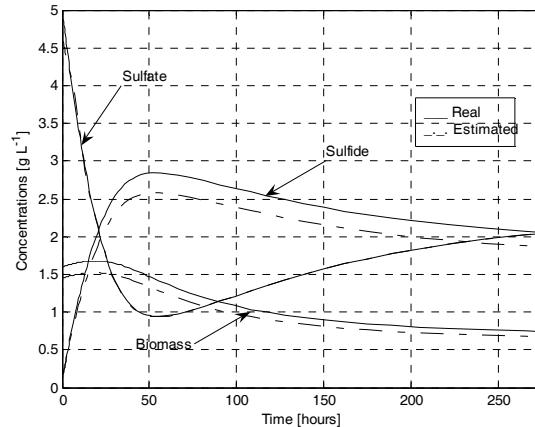


Figure 3. State estimation performance

## 6. CONCLUSIONS

In this work, the cellular growth, sulphate consumption and sulfide production are modeled for *Desulfovibrio alaskensis* using a classical mass balance and the Monod kinetic models. This model is compared with the experimental data successfully; therefore, this is employed as the *real* process for observer implementation purposes. An exponential-type state observer, coupled with an uncertainty estimator, is implemented for a continuous stirred bioreactor via numerical simulations to infer biomass and sulfide concentration and the sulfate consumption rate (uncertain term) from sulfate concentration

measurements. A theoretical frame is provided in order to show the exponential convergence properties of the proposed methodology. Numerical experiments allow observing a satisfactory estimation performance, avoiding the experimental methods for detection and enumeration of SRB and sulfide production. The mathematical analysis to show the convergence characteristics of the proposed methodology is done. The proposed nonlinear observer based estimator would be employed for monitoring, fault detection and control purposes.

## 7. ACKNOWLEDGEMENTS

R. Acevedo is very grateful to CONACyT for the financial support via a postgraduate scholarship.

## NOMENCLATURE

*D*.- Dilution rate [1/hour]

*L*.- Lipschitz constant [1/hour]

*u*.- Control input [1/hour]

*t*.- Time [hours]

*K*.- Adaptive observer vector gain [1/hour]

*X*.- Biomass concentration [g/L]

*S*.- Sulfate concentration [g/L]

*S<sub>in</sub>*.- Inlet Sulfate concentration [g/L]

*Sign*.- Discontinuous function with values [-1 1]

*m*.- Observer parameter

*P*.- Sulfide concentration [g/L]

*x*.- State variables vector

*y*.- Measured output

*Y<sub>S/X</sub>*.- Sulfate yield coefficient

*Y<sub>P/X</sub>*.- Sulfide yield coefficient

## Greek Letters :

$\varepsilon$ .- Estimation error

$\zeta$ .- Uncertain term [g/l hours]

$\lambda$ .- Observer gain [1/hour]

$\mu$ .- Specific growth rate [1/hour]

$\mu_{max}$ .- Maximum specific growth rate [1/hour]

$\Gamma$ .- Estimation matrix

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