

## BIOMASS GROWTH AND $k_{La}$ ESTIMATION USING ONLINE AND OFFLINE MEASUREMENTS

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**Abstract:** Measurement of the key process variables is essential during biopharmaceutical production. These measurements are often not available online. This work combines frequent online measurements with infrequent offline measurements to estimate the specific growth rate, biomass, and the oxygen mass transfer coefficient during continuous and fed-batch cultivations of *Bordetella pertussis* online using an Extended Kalman filter, parameter adaptation, and learning. Copyright © 2007 IFAC

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### 1. INTRODUCTION

Most biopharmaceuticals are produced in a batch or fed-batch cultivation. The quality of the product is formed in this step and is the result of the metabolic state of the micro-organisms. It is therefore essential to measure the physiological state of the process. Metabolic activity is difficult to measure directly due to the lack of sensors, but respiration can be monitored by the oxygen mass balance. The oxygen uptake rate can in turn be used to estimate the specific growth rate and biomass. The specific growth rate and biomass concentration are key parameters that define the metabolic state of micro-organisms.

In this application only dissolved oxygen measurements were available. Therefore, a software sensor based on an Extended Kalman filter (EKF) was developed to estimate specific growth rate and

biomass every minute using the oxygen uptake rate (*OUR*) as input. The choice for an EKF is in line with the process, in which the instruments and the measurement noise are known.

In bioreactors, aerated by a high air flow entering the headspace, the difference between the inlet and exhaust oxygen fraction is small and can therefore not be measured accurately. Hence *OUR* must be calculated using the oxygen balance in the liquid phase (Neeleman, 2002, Soons *et al.*, 2006, 2006a):

$$OUR \approx k_{La} \cdot (C_O^{in} - C_O^L) \quad (1)$$

Where  $k_{La}$  is the oxygen transfer coefficient. The dissolved oxygen concentration ( $C_O^L$ ) is assumed to be at pseudo-steady state i.e. accumulation of oxygen in the bioreactor is negligible. The oxygen concentration entering the bioreactor ( $C_O^{in}$ ) is

assumed to be equal to the liquid phase oxygen concentration in equilibrium with the gas phase in the headspace.

In most applications, the oxygen transfer rate  $k_{La}$  is measured in advance of the cultivation in medium and is assumed to depend on stirrer speed and volume only. However, the formation of cells, proteins, and other molecules, which absorb at gas-liquid interfaces, cause interfacial blanketing and reduce the oxygen transfer rate (Doran, 1995). Because concentrations of cells, substrates, and products change during (fed-) batch cultivation, the value of  $k_{La}$  also changes. An example of change in  $k_{La}$  due to these factors is given in Sabra *et al* (2002). Changing  $k_{La}$  causes errors in the *OUR* calculation and the estimation of the specific growth rate and biomass. It is therefore essential to deal with time-varying  $k_{La}$ .

Offline measurements are mostly considered as not suited for control and estimation purposes, because they become available with a delay and at infrequent and irregular times. These measurements however contain valuable information about the states of the system and can make the estimator more robust (Dondo and Marqués, 2003). Amongst the literature on bioprocess monitoring (e.g. Bastin and Dochain, 1990) the use of offline information for online estimation is relatively small. Myers *et al.* (1995) and Tatiraju *et al.* (1997) use offline and online measurements for state estimation, but do not estimate parameters; Lubenova *et al.* (2003); Gudi *et al.* (1995); Dondo and Marqués (2003); and Ignatova *et al.* (2003) estimate parameters in addition. In these approaches the parameters are part of the input-output equations. In this work, however, better results are obtained if the parameter  $k_{La}$  is considered as a part of the *OUR* calculations, which is done separately from the input-output equations (see figure 1).

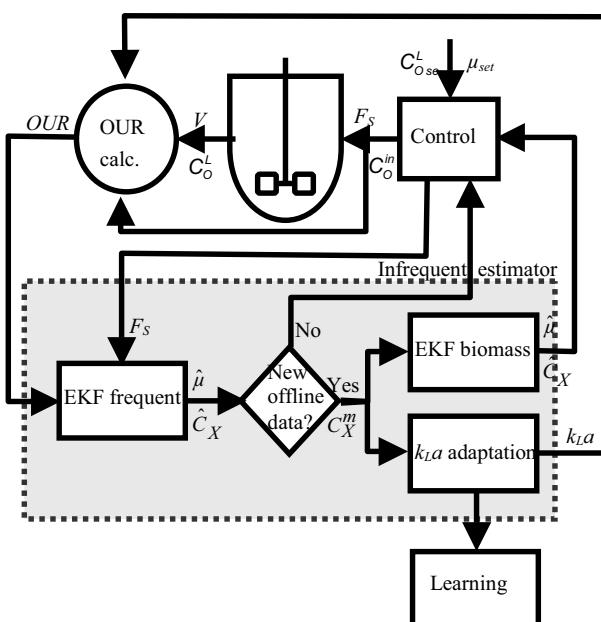


Fig. 1. System configuration (symbolic: see text or nomenclature).

This work combines frequent online measurements (oxygen uptake rate) with infrequent offline measurements (biomass) to estimate the specific growth rate and biomass accurately. Figure 1 shows an overview of the system, in which two types of estimators are involved: a *frequent* estimator using the online data and an *infrequent* estimator activated by sampled offline data. The offline measurements are also used to adapt the time-varying  $k_{La}$ . After each run improved values for  $k_{La}$  are obtained and can be used in a learning process to acquire the appropriate  $k_{La}$  time pattern.

## 2. EXTENDED KALMAN FILTER

The estimator is based on a nonlinear continuous-time model:

$$\begin{aligned} \frac{dx}{dt} &= f(x, u) \\ y &= h(x) \end{aligned} \quad (2)$$

With  $f$  a nonlinear function of the states  $x$  and inputs  $u$ , and  $y$  the output.

Lewis (1986) gives a good explanation of an Extended Kalman filter. The application in biotechnological applications is amongst others discussed by Gudi *et al.* (1995), Neleman (2002) and Keesman (2002). The structure of a discrete time EKF is shown in Fig. 2, and is based on the following equations following Lewis (1986):

$$\begin{aligned} x_{k+1} &= A_k x_k + B_k u_k + w_k \\ y_k &= C_k x_k + v_k \end{aligned} \quad (3)$$

Where  $A_k$  and  $C_k$  and  $B_k$  at each time instant follow from discretization and linearization of Eq. 2 for a time step  $\tau$ :

$$\begin{aligned} F &= \left[ \frac{\partial f}{\partial x} \right]_{\hat{x}, u}, \quad C_k = \left[ \frac{\partial h}{\partial x} \right]_{\hat{x}, u} \\ A_k &= I + F\tau, \quad B_k = \left[ \frac{\partial f}{\partial u} \right]_{\hat{x}, u} \tau \end{aligned} \quad (4)$$

and  $u_k$  is the input vector. The initial states  $x_0$  are stochastic variables with average  $\bar{x}_0$  and variance  $P_0$ :  $x_0 \sim (\bar{x}_0, P_0)$ ;  $w_k \sim (0, Q_k)$  is system noise and consists of model errors and unknown inputs; and  $v_k \sim (0, R_k)$  is measurement noise. The algorithm has two steps. The time update and the measurement update.

### 2.1 Time update

When a sample comes available at time  $k$ , first the time update  $k+1$  is calculated using the original nonlinear model.

$$\begin{aligned}\bar{x}_{k+1} &= f(\hat{x}_k, u_k) \\ \bar{y}_{k+1} &= h(\bar{x}_{k+1})\end{aligned}\quad (5)$$

giving the predicted states  $\bar{x}_{k+1}$  and the output  $\bar{y}_{k+1}$ . The prediction of the variance of the states  $\bar{P}_{k+1}$  is based on the system  $A_k$  and system noise ( $Q_k$ ).

$$\bar{P}_{k+1} = A_k \hat{P}_k A_k^T + Q_k \quad (6)$$

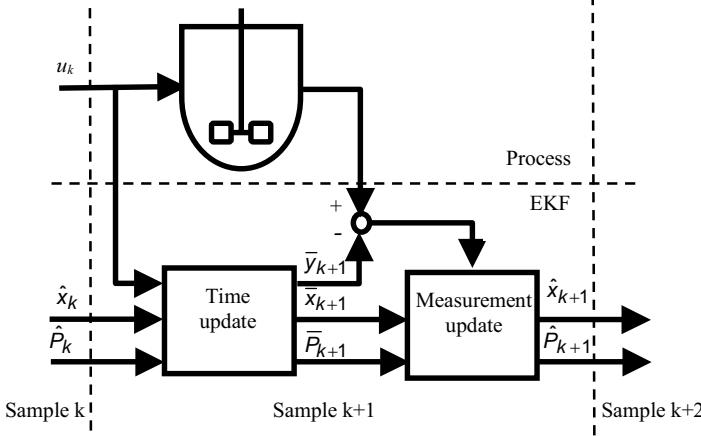


Fig. 2. Calculation scheme for the Extended Kalman filter.

## 2.2 Measurement update

Next the measurement update calculates new estimates using the predicted model states  $\bar{x}_{k+1}$  and outputs  $\bar{y}_{k+1}$ , the actual measurements  $y_{k+1}$ , and the predicted variance  $\bar{P}_{k+1}$ :

$$\begin{aligned}\hat{P}_{k+1} &= \bar{P}_{k+1} - \bar{P}_{k+1} C_{k+1}^T R_{k+1} C_{k+1} \bar{P}_{k+1} \\ K_k &= \hat{P}_{k+1} C_{k+1}^T R_{k+1}^{-1} \\ \hat{x}_{k+1} &= \bar{x}_{k+1} + K_k (y_{k+1} - \bar{y}_{k+1})\end{aligned}\quad (7)$$

Where the Kalman gain  $K_k$  is the magnitude of the correction and aims to minimize the covariance  $\hat{P}_k$ .

## 3. FREQUENT ESTIMATOR

### 3.1 Model

The frequent estimator is based on a generic model (Eqs. 8 and 9) with  $\mu$  as an additional state. The model contains only two parameters: the yield ( $Y_O$ ) and maintenance ( $m_O$ ) coefficient on oxygen (which were calculated using separate experiments):

$$\begin{aligned}\mu_{k+1} &= \mu_k \\ C_{X,k+1} &= \left(1 + \tau(\mu_k - \frac{F_{S,k}}{V_k})\right) C_{X,k}\end{aligned}\quad (8)$$

$$OUR_k = \left(\frac{\mu_k}{Y_O} + m_O\right) C_{X,k} \quad (9)$$

The states  $x$  are specific growth rate and biomass ( $\mu$  and  $C_X$ ), the output  $y$  is the oxygen uptake rate ( $OUR$ ). The volume  $V$  is assumed to be exactly known and doesn't need to be estimated.  $F_S$  is the feed rate. The prediction of the states follows every instant (one minute) by calculation of Eq. 8, the measurement update from calculation of Eq. 7. The observability matrix is full rank, which renders the linearized system observable if the oxygen uptake rate is not negligible.

### 3.2 Tuning

The parameterization of the EKF is based on simulation. Synthetic output noise is derived from real process data. System noise ( $Q$ )

$$Q = \begin{bmatrix} Q_\mu & 0 \\ 0 & Q_{C_X} \end{bmatrix} \quad (10)$$

was chosen in such a way that the deviation ( $E$ ) between estimation (EKF) and generated data by a detailed Monod model (Soons *et al.*, 2006) is minimal for a worst case scenario with a high level of noise.

$$E = \sqrt{\frac{\sum_{k=1}^N [\hat{\mu}(k) - \mu_{mod}(k)]^2}{\bar{\mu}_{mod} \cdot (N-1)}} + \sqrt{\frac{\sum_{k=1}^N [\hat{C}_X(k) - C_{X,mod}(k)]^2}{\bar{C}_{X,mod} \cdot (N-1)}} \quad (11)$$

With  $Q_\mu$  the intensity of the system noise on the specific growth rate and  $Q_{C_X}$  on the biomass. Both terms are divided by their average to make them equally important.

## 4. INFREQUENT ESTIMATOR

The infrequent estimator updates the biomass and  $k_{La}$  estimation using infrequent offline and frequent online measurements (Fig. 1). Since inaccurate values of  $k_{La}$  may cause biased estimations for the frequent estimator,  $k_{La}$  is also updated when offline samples are available.

Using a second Extended Kalman filter to estimate  $C_X$  and  $k_{La}$  for offline samples would be the logical step to incorporate the effect of product and biomass formation on  $k_{La}$ . The infrequent estimator is extended with the ten delayed biomass concentrations (analyzing biomass samples takes approximately ten sample instants, Eq. 12),  $k_{La}$ , and dissolved oxygen. Although this estimator is observable, the algorithm is complex and tuning is difficult.

As an alternative, the following three-step algorithm is designed. First the frequent estimator estimates  $\mu$  and  $C_X$  using frequent online available data (section 3). Next, if samples are available, the biomass estimator estimates biomass and finally the adaptive estimator estimates  $k_{La}$ . The main advantage

of this approach is that convergence of  $k_{La}$ -adaptation is determined by one tuning parameter which can be enforced *a priori*.

#### 4.1 EKF Biomass

The biomass estimator contains the following discrete model, in which the sample and analysis delay is assumed to be constant and is incorporated by addition of fictitious delayed states Gudi *et al.* (1995):

$$\begin{aligned} C_{X_1,k+1} &= \left(1 + \tau\left(\mu_k - \frac{F_{S,k}}{V_k}\right)\right) C_{X_1,k} \\ C_{X_2,k+1} &= C_{X_1,k} \\ . & \\ . & \\ C_{X_{11},k+1} &= C_{X_{10},k} \end{aligned} \quad (12)$$

where  $\tau$ , as before, is the frequent sampling instant. The states of the biomass estimator are the (delayed) biomass concentrations ( $C_{X_1}, C_{X_2}, \dots, C_{X_{11}}$ ); the output is biomass delayed with ten minutes ( $C_{X_{11}}$ ). The prediction of the states follows every infrequent instant (if an offline sample is available, Fig. 1) from calculation of Eq. 12, the measurement update from calculation of Eq. 7.

The observability matrix has full rank. Tuning is performed similar to the frequent estimator by optimization of the covariances of  $Q$ .

#### 4.2 Adaptive $k_{La}$ Estimator

During cultivation the microorganisms and their products may affect the physical properties of the liquid and as a result  $k_{La}$  may change. Deviations between the measured and actual biomass are assumed to be caused by errors in  $k_{La}$ . The adaptive estimator adapts  $k_{La}$  when samples are available. The relative correction  $c$  is calculated every infrequent instant and assumed constant until the next sample becomes available:

$$\begin{aligned} &\text{if } C_{X,k}^m \text{ exists} \\ k_L a_k^{Stir} &= c \cdot f(V, R^{Stir}) \\ k_L \hat{a}_k &= k_L a_k^{Stir} + \gamma(C_{X,k}^m - \hat{C}_{X,k}) \cdot T_m \\ c &= \frac{k_L \hat{a}_k}{k_L a_k^{Stir}} \end{aligned} \quad (13)$$

*else*

$$\begin{aligned} k_L a^{Stir} &= c \cdot f(V, R^{Stir}) \\ k_L \hat{a}_k &= k_L a_k^{Stir} \\ c &= c \end{aligned}$$

Depending on the oxygen demand during the cultivation, agitation speed is adjusted. The oxygen transfer coefficient  $k_{La}$  is measured in advance of the cultivation in biomass-free medium and depends on the stirrer speed and on the volume as specified in the function  $f(V, R^{Stir})$  in Eq. 13. This function is used to adjust  $k_{La}$  *a priori* for known changes stirrer speed.

The tuning parameter  $\gamma$  determines the convergence speed of the observer.  $T_m$  is the infrequent sampling instant, which can be irregular and infrequent.

#### 4.3. Learning

In some situations (e.g. production) it is not desirable to take many offline samples, because the process may be disturbed or it is not possible to take samples. The estimators discussed so far are applied online. Learning can be applied offline after each run.

In order to improve the estimation when samples are not available and  $c$  in Eq. 13 cannot be calculated, the effect of product formation on  $k_{La}$  (based on previous runs) can be used instead and reads:

$$c(C_X) = 1 + f(C_X) \quad (14)$$

where  $f(C_X)$  determines the decrease of  $k_{La}$  as a function of biomass. Batch-to-batch improvement can be obtained by estimating  $f(C_X)$  by regression after each run using data of all runs.

## 5. EXPERIMENTAL RESULTS

Fed-batch and continuous-flow stirred-tank (CSTR) cultivations with the dual substrate consuming bacterium *Bordetella pertussis* were performed with glutamate and L-lactate as the main carbon sources. Bioreactor conditions, medium, analysis, and software were applied as reported by Soons *et al.* (2006). First, batch cultivation was performed until the limiting substrates were depleted. Next, the fed-batch phase of the cultivation automatically started when the specific growth rate dropped below the set-point; the CSTR phase started when the biomass reached an optical density of one. The dynamic method was used to estimate  $k_{La}$  in medium before inoculation for a range of stirrer speeds and volumes. The CSTR experiments were performed to test the frequent EKF only, the fed-batch experiments to test the infrequent estimator (frequent EKF, infrequent EKF, and  $k_{La}$  adaptation).

Maintenance and yield coefficients were calculated on the basis of a series of four CSTR experiments with low biomass. Blanketing effects are therefore small so that  $k_{La}$  can be assumed independent of biomass and product formation, which allows calculation of proper maintenance and yield coefficients.

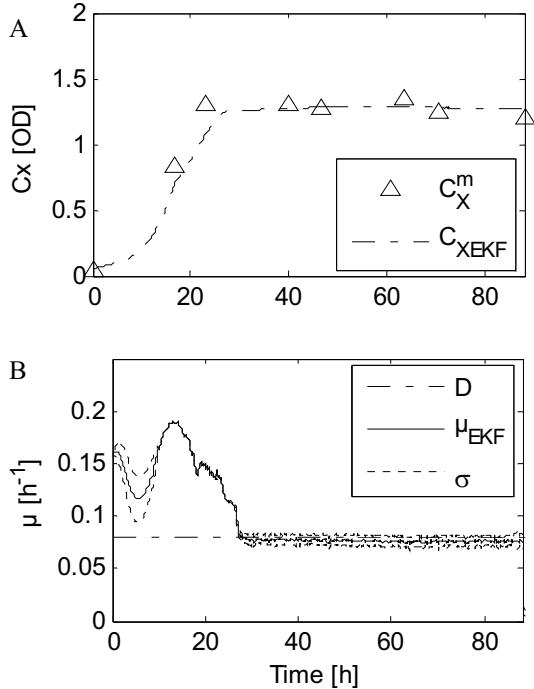


Fig. 3. Frequent estimator for CSTR cultivation  
A. Biomass. B. Estimated specific growth rate and its standard deviation ( $\sigma$ ).

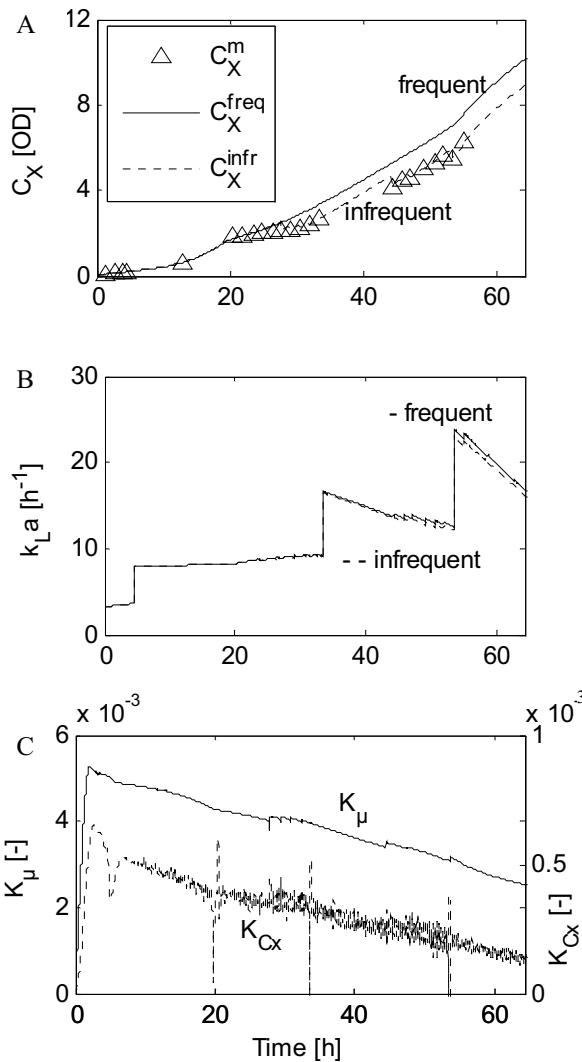


Fig. 4 Frequent and infrequent estimator for fed-batch cultivation. A. Biomass. B. Oxygen transfer coefficient. C. Kalman gains

Figure 3 shows that the estimated biomass coincides well with the offline biomass measurements during CSTR. The estimated specific growth rate converged exactly to the preset dilution rate ( $D = F_S V'$ ). Uncertainty on  $\mu$  was small throughout the cultivation. The performance of the EKF was compared to an asymptotic observer of previous work (Neeleman *et al.*, 2002), and was significantly better.

Blanketing effects however do occur during fed-batch cultivation with higher biomass concentrations, resulting in overestimation of the biomass if only the frequent estimator is used (Fig. 4). Figure 4 compares the frequent estimator with the infrequent estimator for fed-batch cultivation. An exponentially increasing feed was added to the bioreactor to keep the specific growth rate constant, resulting in an increase in volume and a decrease in  $k_{L\alpha}$ ; sampling causes a small decrease in volume and therefore slight increase in  $k_{L\alpha}$ . The abrupt changes of  $k_{L\alpha}$  were caused by changes of stirrer speed (to meet the increasing oxygen demands). Biomass was overestimated using only the frequent estimator (Eqs. 8 and 9). Biomass was estimated more accurately when the infrequent estimator (Eqs. 8, 9, 12, and 13) corrected the  $k_{L\alpha}$  and  $C_X$  estimations for biomass and product formation.

Performance of the infrequent estimator was good even when more sparse biomass measurements were available. Fewer measurements will result in larger sampling intervals  $T_m$  and larger  $k_{L\alpha}$  adaptation per offline measurement (Eq. 13). The effect of more sparse measurements on observer performance will therefore be small, however, the observer becomes more noise sensitive when the measurements contain a high level of noise. Obviously, some measurements are required to find the proper  $k_{L\alpha}$  pattern.

Figure 5 shows the values of the relative  $k_{L\alpha}$  ( $c = k_{L\hat{\alpha}} / k_{L\alpha}^{\text{Stir}}$ ) for two fed-batch cultivations at different biomass concentrations. The relative  $k_{L\alpha}$  decreased by up to 8% for a cultivation grown to 8 OD due to biomass and product formation. Using a linear relation between biomass and relative  $k_{L\alpha}$  (Eqs. 14 and 15) improves the biomass estimation ( $d=0.01$ ):

$$c(C_X) = 1 + d \cdot C_X \quad (15)$$

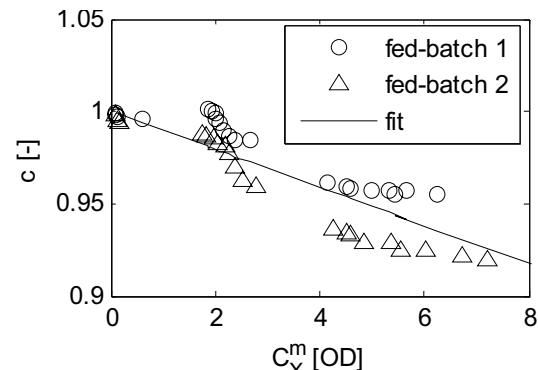


Fig. 5 Relative  $k_{L\alpha}$  ( $k_{L\alpha}$  during fed-batch compared to  $k_{L\alpha}$  in pure medium,  $c = k_{L\hat{\alpha}} / k_{L\alpha}^{\text{Stir}}$ ) as function of the biomass concentration for two fed-batches.

For high cell density cultivation (e.g. *E. coli*), in which cells can grow up to approximately 150 OD,  $k_{LA}$  will drastically decrease. It illustrates the potential of incorporating this effect in the estimation.

## 6. CONCLUSIONS

Application of the presented observer design improves bioprocess monitoring in biopharmaceutical production. The estimator consists of three parts operating on three time-scales: a *frequent* estimator that estimates specific growth rate and biomass online; an *infrequent* estimator, activated by offline measurements, that estimates the biomass and  $k_{LA}$  online; and *learning* that improves the relation between biomass and the oxygen transfer coefficient from batch-to-batch. The estimator showed convergence and fitted well to the data.

Biomass and products influence mass transfer in a bioreactor. Estimation of these  $k_{LA}$  changes is essential for proper estimation of the states during fed-batch and other high cell density cultivations that use the oxygen balance in the liquid phase.

## NOMENCLATURE

$A, B, C$	System matrices
$c$	Relative $k_{LA}$
$C_X$	Biomass concentration $OD$
$C_Q^{in}$	Oxygen concentration into reactor $mmol.l^{-1}$
$C_O^L$	Oxygen concentration in medium $mmol.l^{-1}$
$d$	Constant
$E$	Performance criterion
$F_S$	Substrate feed rate $l.h^{-1}$
$k_{LA}$	Oxygen transport coefficient $h^{-1}$
$m_O$	Maintenance on oxygen $mmol.OD^{-1}.h^{-1}$
$OTR$	Oxygen transfer rate $mol.l^{-1}.h^{-1}$
$OUR$	Oxygen uptake rate $mol.l^{-1}.h^{-1}$
$P$	Variance of the states
$Q, R$	System noise, output noise
$R_{Stir}$	Stirrer speed $rpm$
$t$	Cultivation time $h$
$T_m$	Infrequent sampling interval $h$
$u, x, y$	Inputs, states, outputs
$V$	Liquid volume $l$
$v, w$	Measurement noise, system noise
$Y_O$	Biomass yield on oxygen $OD.mmol^{-1}$
$\gamma$	Tuning parameter
$\mu$	Specific growth rate $h^{-1}$
$\tau$	Frequent sampling interval $h$
$\hat{x}, \hat{EKF}$	Estimated values
$\underline{x}, EKF$	Predicted values

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