

Modelling and Operation of a Turbidity-Meter for On-Line Monitoring of Microbial Growth in Fermenters[†]

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The PROIMI-2 prototype turbidity-meter for measuring cell concentrations in stirred tanks, has been designed and constructed using theories based on kinetic laws and the general principles of cell growth. Laboratory assays were carried out with batch cultures of Bacillus amyloliquefaciens, Zymomonas mobilis and Saccharomyces cerevisiae. The signals showed a good correlation with optical density and biomass. In the latter case up to 4.0 g per dm³ expressed as dry weight could be monitored successfully.

INTRODUCTION

Cell growth is one of the most important parameters in fermentation processes.¹ Generally, the methods used most often for quantification of microbial growth involve cell or plate counting, or dry weight.² These procedures introduce a high risk of contamination in repeated batch cultures and continuous culture fermenters.

One of the most commonly used methods for monitoring cell growth is based on spectrophotometric techniques, i.e. light scattering or light transmission. For this purpose, equipment design and operation have been developed in several laboratories with rather varied results.^{3–8} Most of the equipment installations need several opera-

tions in order to be used.^{5,7,8} This is inconvenient for automatic on-line monitoring of cell growth. A semi-automatic on-line turbidity-meter has been developed recently, but the cell concentrations measured involve dilution of the sample in special chambers outside the fermenter or alternatively the use of flowcells with different pathlengths in or outside of the fermenter vessel.^{6,8,9} Other turbidity-meters developed for the on-line monitoring of cells involve reductions in the fermenter volume and/or vessel modifications. In all cases the equipment complexity and costs had increased considerably.^{3,6–10}

The objectives of this work were the construction and operation of a simple turbidity-meter prototype, named PROIMI-2, for on-line monitoring of microbial growth in stirred tank fermenters at laboratory scale. The simple mathematical modelling of the turbidity-meter

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Fig. 1. Diagram of turbidity-meter PROIMI-2.

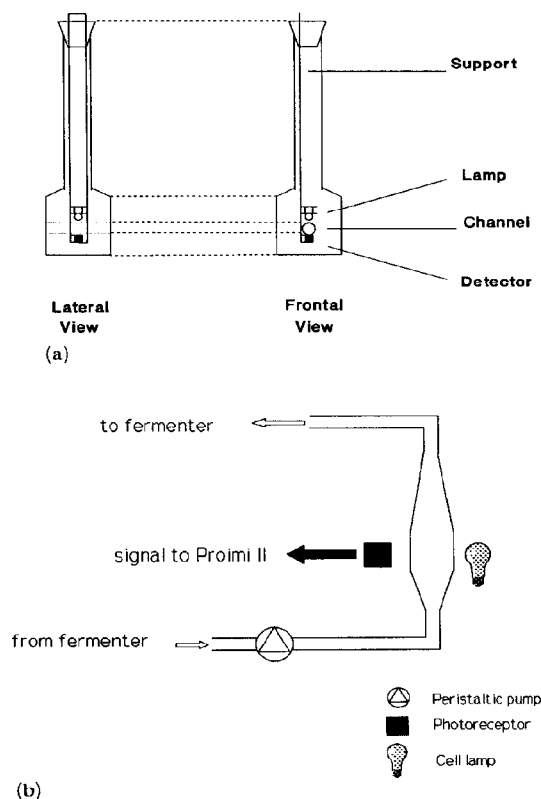


Fig. 2. Scheme of probes A and B of cell counter.

Theoretical aspects

Mathematical modelling of the turbidity-meter

Bacterial growth in the fermenter can be described by the following hypotheses:

- (1) The microorganisms are randomly distributed in the fermenter.
- (2) The growth rate of the microorganisms is constant under each of the experimental conditions.
- (3) The environmental conditions in the vessel are constant (temperature, pH, etc...).
- (4) The relative velocities of the microorganisms (particles) are constant during the fermentation process.
- (5) No cellular adhesion exists.
- (6) The volume of the microorganisms are not significant with respect to the fermenter volume. The microorganisms can be considered as point particles for kinetics considerations.

- (7) Cell growth is described by the exponential equation described previously:¹

$$x(t) = x(t_0) \cdot e^{\mu t}, \quad (1)$$

$x(t_0)$ and $x(t)$ are the cellular concentrations at zero and t times, respectively, and μ is the specific growth rate constant.

- (8) The individual velocity of the microorganisms (v_i) is random and has a maximum. The average rate for N microorganisms (α) is defined by:

$$\alpha = \sum_{d\tau} v_i / dN = 0,$$

in which N is the bacterial number in one element of volume ($d\tau$).¹³

- (9) The signal obtained (E) for the interaction of one microorganism-sensor in the solid angle (δw) is considered:¹⁴

$$\delta w = \int_r \int_\sigma [\cos \sigma / r^2] dr d\sigma = -\sin \sigma / r + Q,$$

where Q is the integration constant.

The signal E for one microorganism can be considered directly proportional to the conical shadow formed by the interposition of one bacteria to the light beam with respect to the detector. Therefore:

$$E \approx \delta w = E \approx -\sin \sigma / r + Q.$$

For n microorganisms and the distance between light source (lamp) and detector (r), the signal E can be estimated as:

$$\begin{aligned} E &\approx - \int_n \int_r [\sin \sigma / r \cdot \phi(n)] dn dr \\ &= E \approx -\sin \sigma \cdot \Omega(n) \cdot \ln(r) + K. \end{aligned}$$

E is a function of σ (conical shadow angle) and $\Omega(n)$ a function of the number of cells (n), r is a constant and K is the integration constant.

When the angle σ is very small: $\sigma \approx 0 \Rightarrow \sin \sigma \approx \sigma$,

$$\Rightarrow E \approx \sigma \cdot \Omega(n) \cdot \ln(r).$$

n is related to the cell concentration by a constant (volume of the vessel). In batch operation, the microbial growth follows the exponential equation (eqn (1)). The signal depends on the time and

can be expressed as:

$$E(t) \approx E(t_0) + \phi(\sigma, r) \cdot \mu \cdot t,$$

where ϕ is a function of angle (σ) and radius (r), and therefore of the solid angle.

RESULTS AND DISCUSSION

A diagram of the probe is shown in Fig. 2(a). Height and rotation angle of the probe can be varied in order to obtain different readings in the reactor. Changes of the probe position are useful for monitoring the homogeneity of the culture and/or the mixing times in the reactor.

The channel of probe A and the tube of probe B possibly function as a Venturi tube, because no cell growth on the channel walls was observed in submerged position in continuous culture operation for several days with high cell density cultures of the assayed microbes. On the contrary, cell growth was observed in other parts of the fermenter (vessel walls, electrodes, etc.).

In order to correlate the turbidity-meter signal with any other parameter, i.e. dry weight, optical density, etc., it was necessary to withdraw samples from a culture while the turbidity-meter was monitoring growth. The plot of the optical density (560 nm) of a batch culture of *Bacillus amyloliquefaciens* MIR-41 versus the signal of the turbidity-meter using the probe A showed a linear response (Fig. 3(a)), as was the case of *Zymomonas mobilis* PRO-910 batch cultures (Fig. 3(b)). The correlation between biomass and optical density showed some deviation from the linear relationship at high dry weights, as previously reported.¹ However, the correlation between biomass and turbidity-meter signal was linear in the range of cell density obtained (0.1–1.0 g dm⁻³) (Fig. 3(b)).

For monitoring high cell density cultures a strain of *Saccharomyces cerevisiae* was used in batch cultures. The turbidity-meter was equipped with probe B, and simultaneously the dry weight of the culture and optical density (660 nm) with appropriate dilutions were measured (Fig. 4). Linear correlation between dry weight up 4.0 g dm⁻³ and turbidity-meter-probe B signal was observed.

In order to evaluate the behaviour of the turbidity-meter *in vivo*, batch cultures of *Bacillus amyloliquefaciens* MIR-41 with starch (40 g dm⁻³) as the carbon source were used. During the

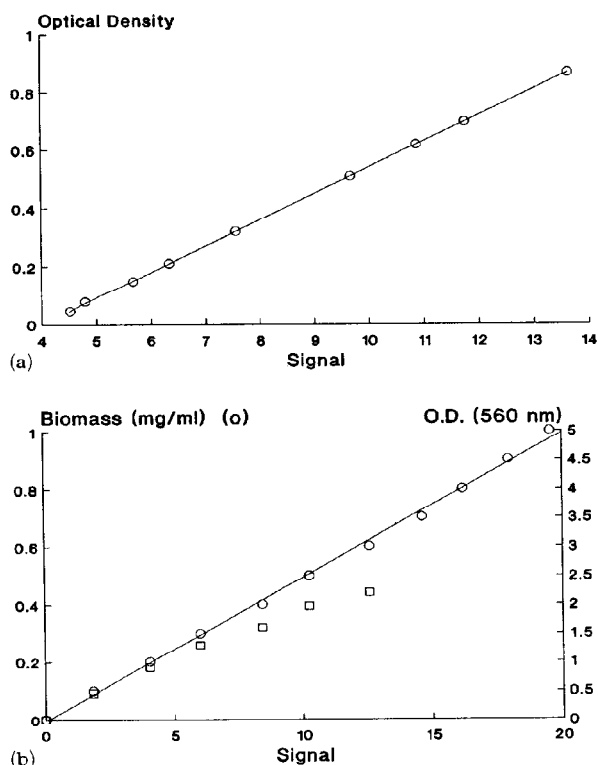


Fig. 3. (a) Correlation of the optical density (560 nm) with turbidity-meter probe A signal of *Bacillus amyloliquefaciens* MIR-41. (b) Correlation between the turbidity-meter probe A signal (O) and optical density (560 nm) (□) with biomass as dry weight (g dm⁻³) of *Zymomonas mobilis* PRO-910 batch culture.

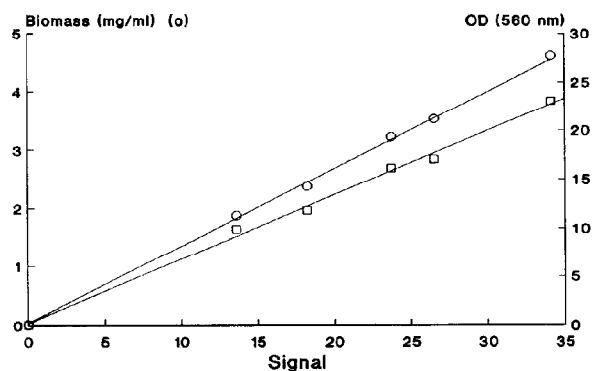


Fig. 4. Correlation between the turbidity-meter probe B signal with biomass as dry weight (g dm⁻³) of *Saccharomyces cerevisiae* batch culture (O) and optical density at 560 nm (□).

experiments the values of the turbidity-meter and the optical density were both measured and were followed in time (Fig. 5). Starch medium has a high turbidity and presents some problems for cell

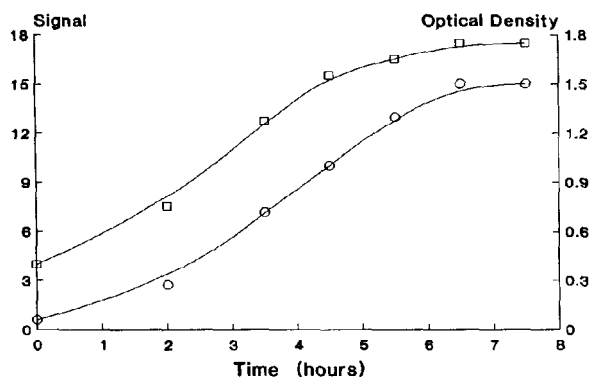


Fig. 5. Batch culture of *Bacillus amyloliquefaciens* MIR-41 in starch medium. Cellular growth was determined with the PROIMI-2 turbidity-meter probe A (□) and optical density at 560 nm (○).

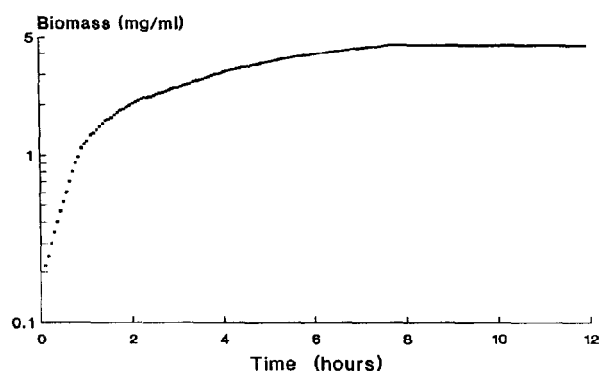


Fig. 6. Batch culture of *Saccharomyces cerevisiae* K-382. Cellular growth was determined with PROIMI-2 turbidity-meter probe B.

quantification. It is not possible to use the dry weight method, because of the content of high molecular weight compounds, which cannot be filtered out. Optical densities are usually measured in such microbial cultures. However, spectrophotometric determinations have an increased error because of the dilution of the samples in order to decrease the starch concentration and adjust the readings below one unit of absorbance. The same error occurs, when the microbial culture reaches high cellular concentrations and has to be diluted. In previous reports the determination of low and high biomass concentrations involves either the use of different flowcells or different lengths of the light paths.^{8,9} Alternatively, a system for high cell density measurements involves a circulating system and samples have to be taken outside the vessel.¹⁰ However, *Saccharomyces cerevisiae* was cultured and biomass was determined automatically with the turbidity-meter PROIMI-2 probe B according to a previous calibration, without dilution of samples and flow-cell changes (Fig. 6). With the turbidity-meter presented here, at high cell concentrations, one flow-cell was used and the upper limit of detection was approximately twice the cell concentration reported previously.⁹

In other reports, it has been suggested that gas bubbles present in the medium may interfere with the readings in turbidity-meters.³ However, the results obtained with the PROIMI-2 turbidity-meter used in aerated cultures of *Bacillus amyloliquefaciens* MIR-41 ($0.1 \text{ v v}^{-1} \text{ min}^{-1}$) did not show interference (Fig. 5). This may be due to the constant conditions under which the experiments

were carried out. Moreover, gas production by the microorganisms themselves might be negligible compared with the air influx. Similarly, no effects of aeration rates were reported in a plant cell bioreactor with a laser probe of 780 nm.¹⁵ However, the problem associated with the laser probes is the monochromatic light, which can be interfered with due to the coloured compounds or pigmented cell products.

The development of a simple mathematical model would facilitate the automation of the fermentation process. It could also be used in culture techniques where the addition of nutrients is necessary once the cellular concentration is established, such as in fed-batch cultures.¹⁶

CONCLUSIONS

The turbidity-meter PROIMI-2 designed and manufactured in our laboratory has the following characteristics and advantages: linear correlation with the readings obtained spectrophotometrically at 560 and 660 nm; similar response in low and high absorbance media; fast and reliable installation without modifications of the fermenter; no microbial growth on the walls of the detector; possibility of changing the position of the probe in the fermenter during the experiment, and low costs of construction and operation.

The turbidity-meter PROIMI-2 can be used to predict the development of the fermentation process, the on-line estimation of kinetic constants of the microorganisms, and also the automatic control of the cellular growth.

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