

ON A SPECIAL CHOICE OF NUTRIENT SUPPLY FOR CELL
GROWTH IN A CONTINUOUS BIOREACTOR. SOME
MODELING AND APPROXIMATION ASPECTS

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ABSTRACT: In this article we will consider the possibility of approximating the input function $s(t)$ (the nutrient supply for cell growth in a continuous bioreactor) with the Lindley type correction.

We prove upper and lower estimates for the one-sided Hausdorff approximation of the shifted Heaviside function $h_{t^*}(t)$ by means of the general solution of the differential equation $y'(t) = ky(t)s(t)$ with $y(t_0) = y_0$.

We will illustrate the advances of the solution $y(t)$ for approximating and modelling of: "data on the development of the *Drosophila melanogaster* population", published by biologist Raymond Pearl in 1920 (see, also Alpatov, Pearl [32]), "cancer data" (see, [33]–[34]), "data on the development of *Saccharomyces* culture in nutrient medium", published by biologist T. Carlson in 1913 (see, also [67], [69]), "growth data (mean height) of sunflower plants" [69] and "data on the growth of population of *Rhizopertha* in wheat" by Crombie in 1945 [68].

We also define a new parametric activation function based on "amendments" of "Lindley - type".

Numerical examples using *CAS Mathematica*, illustrating our results are given.

AMS Subject Classification: 41A46

Key Words: nutrient supply, prototype of model used in bioreactor modelling, "supersaturation" of the model, Heaviside function, Hausdorff distance, upper and lower bounds

Received: February 11, 2019; **Revised:** April 23, 2019;

Published (online): May 22, 2019 **doi:** 10.12732/dsa.v28i3.5

Dynamic Publishers, Inc., Acad. Publishers, Ltd.

<https://acadsol.eu/dsa>

1. INTRODUCTION

Sigmoidal functions find multiple applications to population dynamics, biostatistics, analysis of nutrient supply for cell growth in bioreactors, controllability of tumor growth, population survival functions, classical predator-prey models, neural networks, debugging and test theory and others [37]–[66].

The Verhulst model can be considered as a prototype of models used in bioreactor modelling.

In batch growth, the rate of biomass production is given by $\frac{dx}{dt} = \kappa x$, where: x = biomass concentration; κ = specific growth rate; t = time. The rate κ is a function of nutrient supply and therefore can be a function of time (i.e., if nutrient supply is changing with time.)

In general, $\kappa = F(S, P, I, X, T, \text{osmotic pressure})$; S = substrate concentration; P = product concentration; I = inhibitor concentration.

There, especially in the case of continuous bioreactor, the nutrient supply is considered as an input function $s(t)$ as follows:

$$\frac{dy(t)}{dt} = ky(t)s(t) \quad (1)$$

where s is additional specified.

To the role and choice of nutrient supply for cell growth in bioreactors are devoted to a number of studies [1]–[12].

Some concepts of multiple-nutrient-limited growth of microorganisms and its application in biotechnological processes can be found in [3].

In [13], the author consider the following hyper-logistic equation:

$$\frac{dy(t)}{dt} = ky(t) \frac{2e^{-pt}}{1 + e^{-pt}} \quad (2)$$

$$y(t_0) = y_0,$$

where $k > 0$ and $p > 0$ with general solution:

$$y(t) = y_0 e^{2k(t - t_0) + \frac{2k}{p} \ln(1 + e^{pt_0}) - \frac{2k}{p} \ln(1 + e^{pt})}.$$

For other results, see [14].

In this paper we will consider the possibility of approximating the input function $s(t)$ in the equation (1) with the Lindley type correction.

Some results for Lindley [15]–[16], power Lindley distribution, discrete Poisson-Lindley distribution, modified discrete Lindley distribution, generalized Lindley distribution, exponential modified discrete Lindley distribution, quasi Lindley distri-

bution, Kumaraswamy–Lindley distribution and transmuted Kumaraswamy quasi Lindley distribution are given in [17]–[31].

Following the ideas given in [13] we consider the following new logistic equation:

$$\frac{dy(t)}{dt} = ky(t) \frac{1 + \theta + \theta t}{1 + \theta} e^{-\theta t} \tag{3}$$

$$y(t_0) = y_0$$

where $\theta > 0$.

We prove upper and lower estimates for the one–sided Hausdorff approximation of the shifted Heaviside function $h_{t^*}(t)$ by means of the general solution of this differential equation.

We will illustrate the advances of the solution $y(t)$ for approximating and modelling of:

- "data on the development of the *Drosophila melanogaster* population", published by biologist Raymond Pearl in 1920 (see, also Alpatov, Pearl [32]);
- "cancer data" (see, [33]–[34]);
- "data on the development of *Saccharomyces* culture in nutrient medium", published by biologist T. Carlson in 1913 (see, also [67], [69]);
- "growth data (mean height) of sunflower plants" [69];
- "data on the growth of population of *Rhizapertha* in wheat" by Crombie in 1945 [68].

2. PRELIMINARIES

Definition 1. The shifted Heaviside step function is defined by

$$h_{t^*}(t) = \begin{cases} 0, & \text{if } t < t^*, \\ [0, 1], & \text{if } t = t^*, \\ 1, & \text{if } t > t^*. \end{cases} \tag{4}$$

Definition 2. [35], [36] The Hausdorff distance (the H–distance) $\rho(f, g)$ between two interval functions f, g on $\Omega \subseteq \mathbb{R}$, is the distance between their completed graphs $F(f)$ and $F(g)$ considered as closed subsets of $\Omega \times \mathbb{R}$.

More precisely,

$$\rho(f, g) = \max\left\{ \sup_{A \in F(f)} \inf_{B \in F(g)} \|A - B\|, \sup_{B \in F(g)} \inf_{A \in F(f)} \|A - B\| \right\}, \tag{5}$$

wherein $\|\cdot\|$ is any norm in \mathbb{R}^2 , e. g. the maximum norm $\|(t, x)\| = \max\{|t|, |x|\}$; hence the distance between the points $A = (t_A, x_A)$, $B = (t_B, x_B)$ in \mathbb{R}^2 is $\|A - B\| = \max(|t_A - t_B|, |x_A - x_B|)$.

3. MAIN RESULTS

3.1. A NEW MODEL

The general solution of the differential equation (3) is of the following form:

$$y(t) = y_0 e^{-\frac{k(2+\theta+\theta t)}{\theta(1+\theta)}t} + \frac{k(2+\theta+\theta t_0)}{\theta(1+\theta)} e^{-\theta t_0}. \quad (6)$$

It is important to study the characteristic - "supersaturation" of the model to the horizontal asymptote.

In this Section we prove upper and lower estimates for the one-sided Hausdorff approximation of the Heaviside step-function $h_{t^*}(t)$ by means of families (6).

Without loss of generality, we consider the following class of this family for:

$$t_0 = 0; y_0 = e^{-\frac{k(2+\theta)}{\theta(1+\theta)}}$$

$$M(t) = e^{-\frac{k(2+\theta+\theta t)}{\theta(1+\theta)}t}. \quad (7)$$

The function $M(t)$ and the "input function" $s(t)$ are visualized on Fig. 1.

Denoting by t^* the unique positive solution of the nonlinear equation:

$$(2 + \theta + \theta t^*)e^{-\theta t^*} - \frac{\theta(1 + \theta) \ln 2}{k} = 0. \quad (8)$$

Evidently, $M(t^*) = \frac{1}{2}$.

The one-sided Hausdorff distance d between the function $h_{t^*}(t)$ and the sigmoid - (7) satisfies the relation

$$M(t^* + d) = 1 - d. \quad (9)$$

The following theorem gives upper and lower bounds for d

Theorem 1. Let

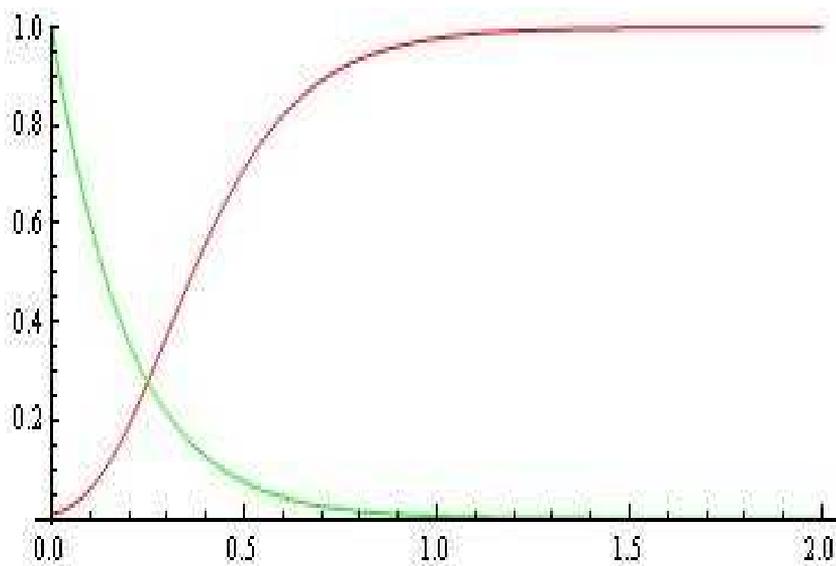


Figure 1: The functions $M(t)$ –(red) and $s(t)$ –(green) for $k = 24.3$; $\theta = 5.9$.

$$\begin{aligned} \alpha &= -\frac{1}{2}, \\ \beta &= 1 + \frac{k(1+\theta+\theta t^*)}{2(1+\theta)}e^{-\theta t^*} \\ \gamma &= 2.1\beta. \end{aligned} \tag{10}$$

For the one-sided Hausdorff distance d between $h_{t^*}(t)$ and the sigmoid (7) the following inequalities hold for the condition $\gamma > e^{1.05}$:

$$d_l = \frac{1}{\gamma} < d < \frac{\ln \gamma}{\gamma} = d_r. \tag{11}$$

Proof. Let us examine the function:

$$F(d) = M(t^* + d) - 1 + d. \tag{12}$$

From $F'(d) > 0$ we conclude that function F is increasing. Consider the function

$$G(d) = \alpha + \beta d. \tag{13}$$

From Taylor expansion we obtain $G(d) - F(d) = O(d^2)$. Hence $G(d)$ approximates $F(d)$ with $d \rightarrow 0$ as $O(d^2)$ (see Fig. 2).

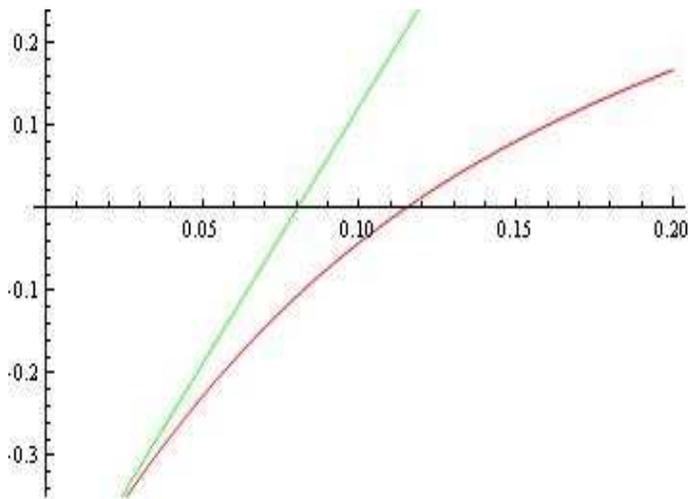


Figure 2: The functions $F(d)$ and $G(d)$ for $k = 68.3$; $\theta = 15.9$.

In addition $G'(d) > 0$.

Further, for $\gamma > e^{1.05}$ we have

$$G(d_l) < 0; \quad G(d_r) > 0.$$

This completes the proof of the theorem.

Approximations of the $h_{t^*}(t)$ by model (6) for various k and θ are visualized on Fig. 3–Fig. 4.

4. SOME APPLICATIONS

The proposed model can be successfully used to approximating data from Population Dynamics.

4.1. APPROXIMATING THE "DATA ON THE DEVELOPMENT OF THE *DROSOPHILA MELANOGASTER* POPULATION"

We will illustrate the advances of the solution $y(t)$ for approximating and modelling of "data on the development of the *Drosophila melanogaster* population", published by biologist Raymond Pearl in 1920 (see, also [32]).

We consider the following data:

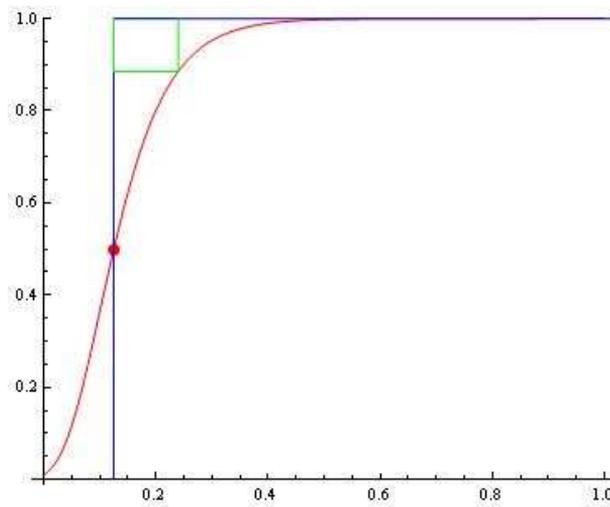


Figure 3: The model (7) for $k = 68.3$; $\theta = 15.9$; $t^* = 0.124959$; Hausdorff distance $d = 0.114808$; $d_l = 0.076393$; $d_r = 0.196472$.

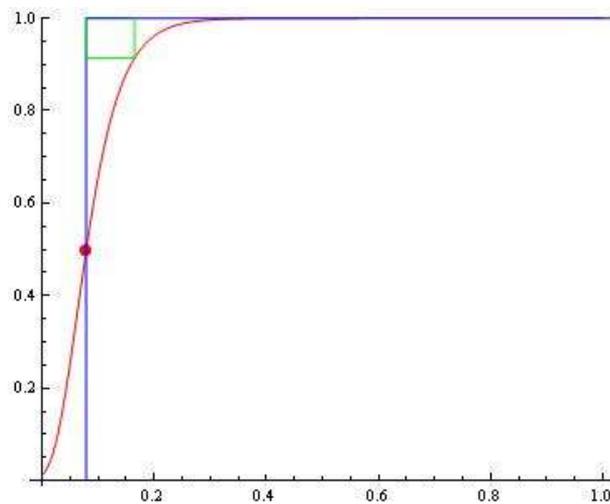


Figure 4: The model (7) for $k = 105.1$; $\theta = 24.5$; $t^* = 0.0788398$; Hausdorff distance $d = 0.0861885$; $d_l = 0.0518027$; $d_r = 0.153352$.

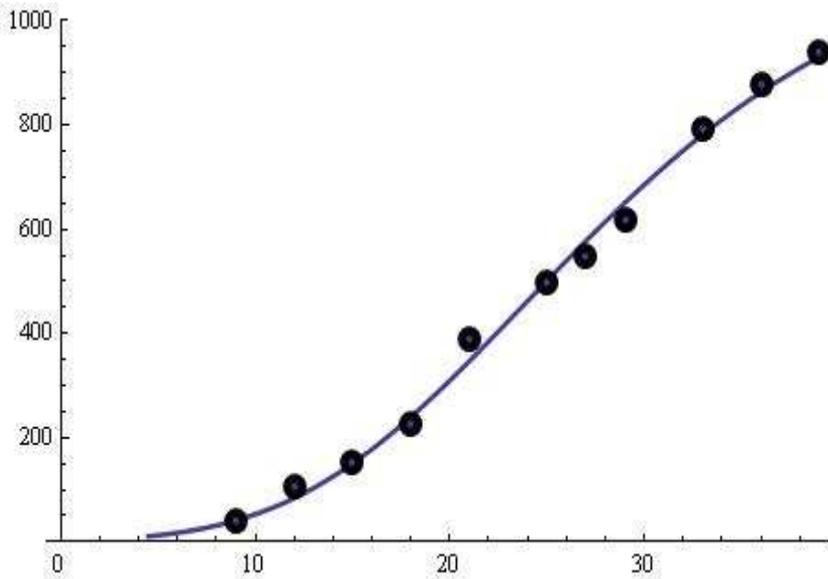


Figure 5: The fitted model $M^*(t)$.

data_Pearl

$:= \{\{9, 39\}, \{12, 105\}, \{15, 152\}, \{18, 225\}, \{21, 390\}, \{25, 547\},$
 $\{29, 618\}, \{33, 791\}, \{36, 877\}, \{39, 938\}\}.$

After that using the model

$$M^*(t) = \omega e^{-\frac{k(2+\theta+ \theta t)}{\theta(1+\theta)}} e^{-\theta t}$$

for $\omega = 1162.27$, $k = 0.383217$ and $\theta = 0.115$ we obtain the fitted model (see, Fig. 5).

4.1.1. SOME COMPARISONS BETWEEN THE NEW LOGISTIC MODEL AND THE CLASSICAL LOGISTIC MODEL OF VERHULST-PEARL.

The classic model of Verhulst -Pearl for the *data_Pearl* looks like this (see, for example, [69]):

$$M_{VP}^*(t) = \frac{1035}{1 + e^{4.27-0.17t}}$$

and is illustrated in Fig. 6

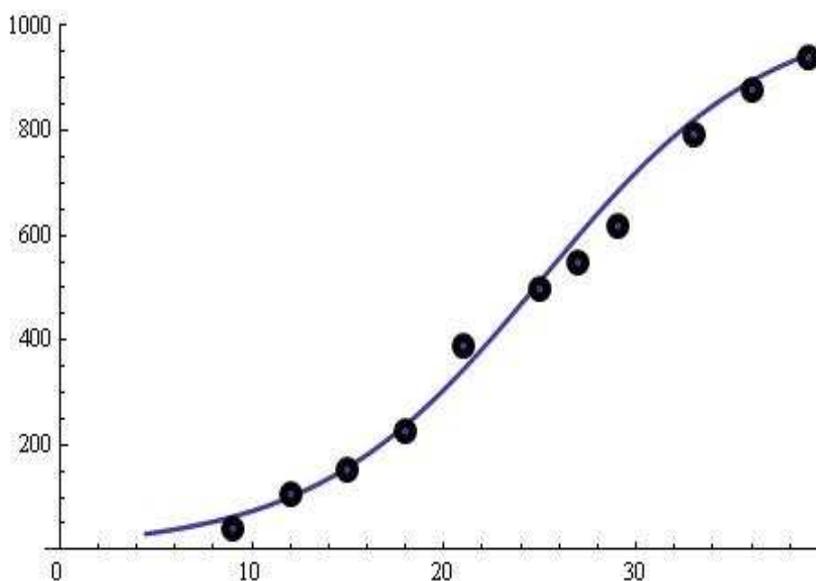


Figure 6: The fitted model $M_{VP}^*(t)$.

Of the accompanying illustrations see Fig. 5 and Fig. 6, it can be concluded that the proposed model $M^*(t)$ (7) is reliable.

4.2. APPLICATION OF THE NEW CUMULATIVE SIGMOID FOR ANALYSIS OF THE "CANCER DATA"

We will illustrate the advances of the solution $y(t)$ for approximation and modelling of "cancer data" (for some details see, [33]–[34]).

<i>days</i>	4	7	10	12	14	17	19	21
$R(t)$	0.415	0.794	1.001	1.102	1.192	1.22	1.241	1.3

Table 1: The "cancer data" [33]–[34]

Consider the model $M^*(t) = \omega e^{-\frac{k(2+\theta+\theta t)}{\theta(1+\theta)}} e^{-\theta t}$.

The model $M^*(t)$ based on the data from Table 1 for the estimated parameters:

$$\omega = 1.32845; \theta = 0.268; k = 0.306626$$

is plotted on Fig. 7.

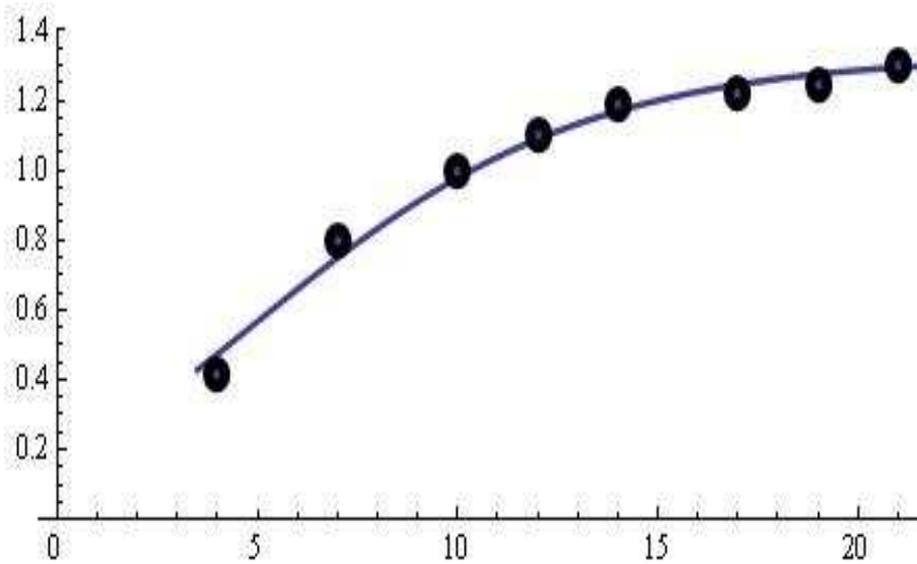


Figure 7: The model $M^*(t)$ based on the "cancer data".

4.3. APPROXIMATING THE "DATA ON THE DEVELOPMENT OF SACCHAROMYCES CULTURE IN NUTRIENT MEDIUM"

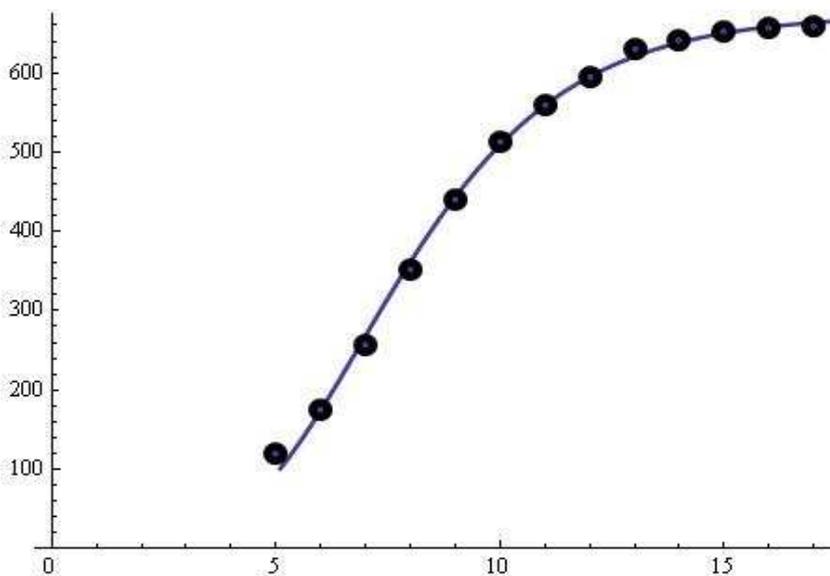
We will now analyze a sample of experimental data obtained by the biologist T. Carlson in 1913 about the development of *Saccharomyces* culture in nutrient medium (see, for example [67], [69]):

$$\begin{aligned}
 & data_Carlson \\
 & := \{ \{5, 19.1\}, \{6, 174.6\}, \{7, 257.3\}, \{8, 350.7\}, \{9, 441\}, \\
 & \quad \{10, 513.3\}, \{11, 559.7\}, \{12, 594.8\}, \{13, 629.4\}, \{14, 640.8\}, \\
 & \quad \{15, 651.1\}, \{16, 655.9\}, \{17, 659.6\} \}.
 \end{aligned}$$

After that using the model $M^*(t)$ for $\theta = 0.47$, $k = 2.9600117$ and $\omega = 673.513$ we obtain the fitted model (see, Fig. 8).

4.4. APPROXIMATING THE "GROWTH DATA (MEAN HEIGHT) OF SUNFLOWER PLANTS"

We analyze experimental growth data (mean height) of sunflower plants (DSP) (see, for example [69]):

Figure 8: The fitted model $M^*(t)$.

data_DSP

$:= \{\{14, 36.4\}, \{28, 98.1\}, \{49, 205.5\}, \{56, 228.3\}, \{70, 250.5\},$
 $\{84, 254.5\}\}.$

For $\theta = 0.08$, $k = 0.176917$ and $\omega = 262.988$ we obtain the fitted model (see, Fig. 9).

4.5. APPROXIMATING THE DATA: "THE GROWTH OF POPULATION OF *RHIZAPERTHA* IN WHEAT"

We analyze a experimental data obtained by the Crombie in 1945 [68]:

data_Crombie

$:= \{\{0, 2\}, \{14, 2\}, \{28, 2\}, \{35, 3\}, \{42, 17\}, \{49, 65\}, \{63, 119\},$
 $\{77, 130\}, \{91, 175\}, \{105, 205\}, \{119, 261\}, \{133, 302\},$
 $\{147, 330.6\}, \{161, 315\}, \{175, 333\}, \{189, 350\}, \{203, 332\},$
 $\{231, 333\}, \{245, 335\}, \{259, 330\}\}.$

After that using the model $M^*(t)$ for $\theta = 0.036$, $k = 0.113865$ and $\omega = 343.284$

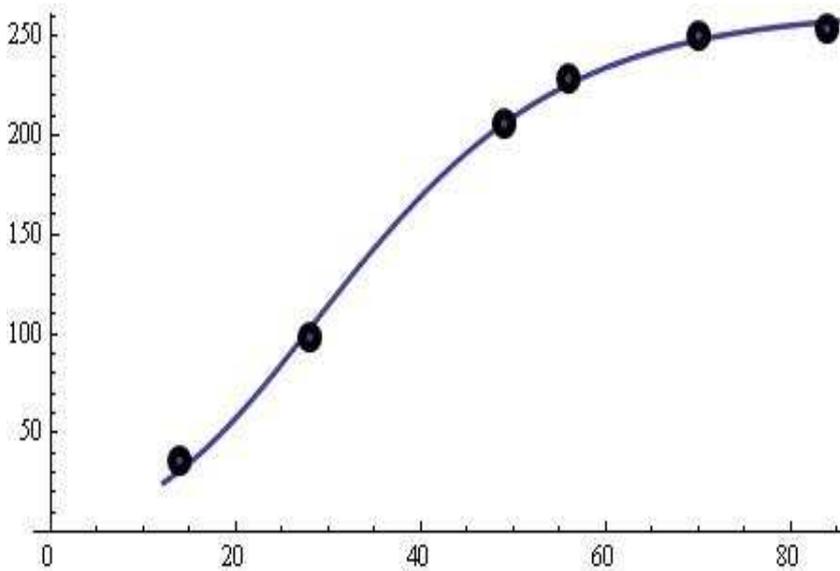


Figure 9: The fitted model $M^*(t)$.

we obtain the fitted model (see, Fig. 11).

As should be expected, the experiments conducted (see, Sections 4.1 - 4.5) show a very good approximation of data from the field of population dynamics, with suggested in this article, modified logistic model.

4.6. THE NEW ACTIVATION FUNCTION BASED ON "AMENDMENTS" OF "LINDLEY - TYPE"

Definition 3. The sign function of a real number t is defined as follows:

$$\operatorname{sgn}(t) = \begin{cases} -1, & \text{if } t < 0, \\ 0, & \text{if } t = 0, \\ 1, & \text{if } t > 0. \end{cases} \quad (14)$$

Definition 4. The new parametric activation function based on "amendments" of "Lindley - type" is defined as follows

$$A(t) = \frac{e^{-\frac{k(2+\theta+\theta t)}{\theta(1+\theta)}e^{-\theta t}} - e^{-\frac{k(2+\theta+\theta t)}{\theta(1+\theta)}e^{\theta t}}}{e^{-\frac{k(2+\theta+\theta t)}{\theta(1+\theta)}e^{-\theta t}} + e^{-\frac{k(2+\theta+\theta t)}{\theta(1+\theta)}e^{\theta t}}}. \quad (15)$$

Approximation of the $\operatorname{sgn}(t)$ by function $A(t)$ for $k = 10.1$ and $\theta = 2.5$ is visualized on Fig. 12.

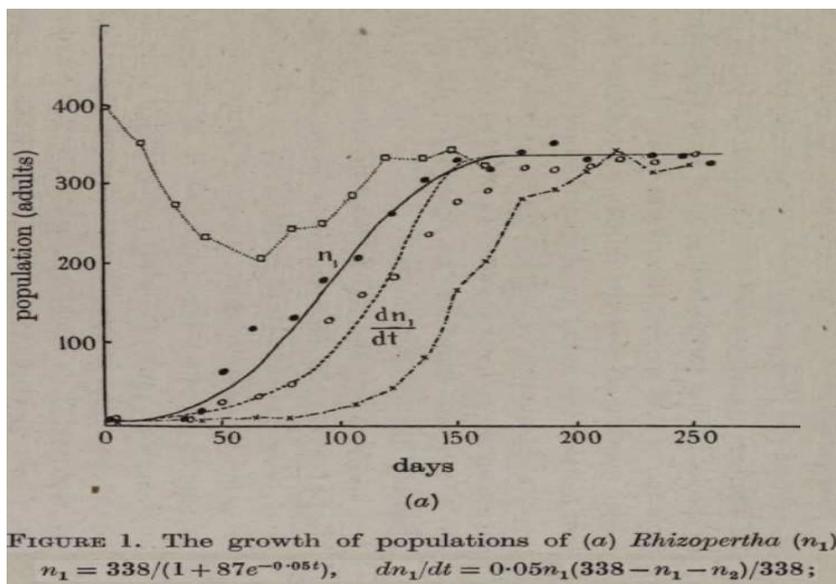


Figure 10: The fitted model by Crombie [68].

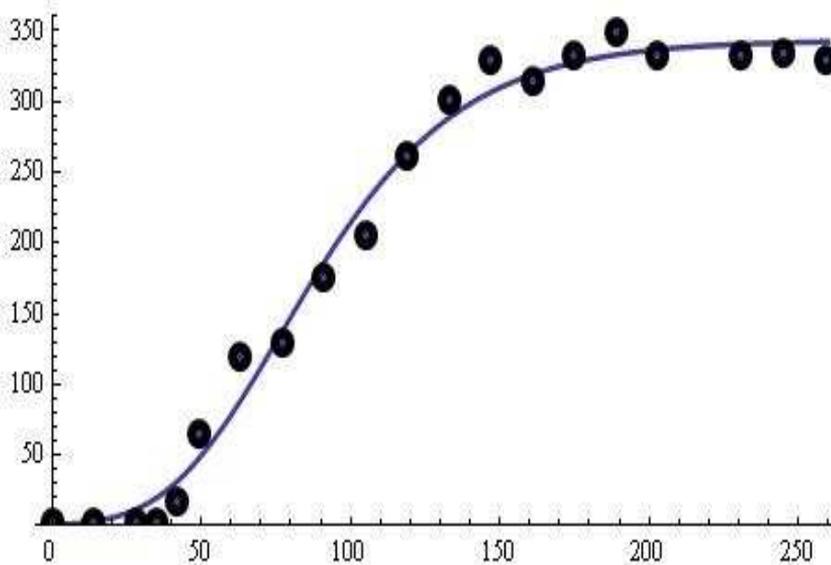


Figure 11: The fitted model $M^*(t)$.

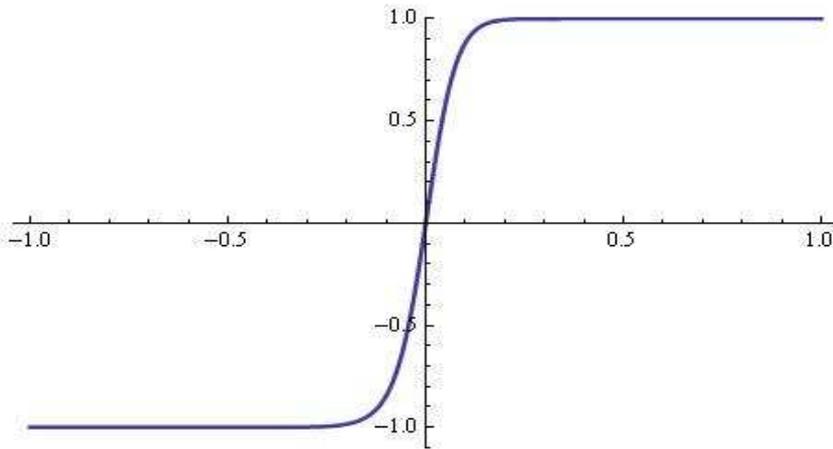


Figure 12: Approximation of the $\text{sgn}(t)$ by $A(t)$ for $k = 10.1$ and $\theta = 2.5$.

We will note that the study of the Hausdorff's approximation of the sign function by means of this new family can be done in a way given in [60] and we will omit it.

Similarly to the article cited above, recursively generable families of higher order activation functions can also be constructed.

5. CONCLUSION

A special choice of nutrient supply for cell growth in a continuous bioreactor with the Lindley type correction is introduced.

We prove upper and lower estimates for the one-sided Hausdorff approximation of the shifted Heaviside function $h_{t^*}(t)$ by means of the general solution of the differential equation $y'(t) = ky(t)s(t)$ with $y(t_0) = y_0$, where $s(t)$ is the correction of Lindley type.

We propose a software module within the programming environment *CAS Mathematica* for the analysis of the considered family of functions.

The module offers the following possibilities:

- calculation of the H-distance between the h_{t^*} and the model $M(t)$ (7);
- generation of the functions under user defined values of the parameters k and θ ;
- numerical solution of the differential model (3) and opportunities for comparison with other logistics models;
- software tools for animation and visualization.

We will explicitly note that similar approximation and modeling results associated with the use of "input function" $S(t)$ in the differential model (1) with Sen, Maiti

and Chandra-type [70]:

$$S(t) = \frac{1 + \theta + \theta t + 0.5\theta^2 t^2}{1 + \theta} e^{-\theta t}$$

and also with Yousof, Korkmaz and Sen-type [71]:

$$S(t) = \frac{1 + \theta + \theta t^b + 0.5\theta^2 t^{2b}}{1 + \theta} e^{-\theta t^b}$$

can be obtained with the mathematical apparatus outlined in this article and here we will miss them.

ACKNOWLEDGMENT

This paper is supported by the National Scientific Program "Information and Communication Technologies for a Single Digital Market in Science, Education and Security (ICTinSES)", financed by the Ministry of Education and Science.

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