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Page: [74-83]

Original research article

# A Computational Approach for Solving **Fractional Model of Malignant Tumour** Growth based on Dynamics of Cell Cycle

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## **ABSTRACT**

We have studied the fractional order mathematical model of malignant tumour growth based on cell cycle dynamics in this work. The model describes on three different tumour cell dynamics of the population; quiescent cells, interphase cells, and mitotic cells. The studied model is based on fractional order differential equations. A computational approach has been implemented to give approximate solution of this fractional model. The model can be used to describe the graphical behavior of tumour cells. The computational results have been presented graphically to show the advantage and the efficiency of the scheme for fractional order malignant tumour growth model.

Keywords: Atangana-Baleanu fractional derivative; Analytical method; Cell Cycle; Fractional differential equation; Malignant tumour

#### 1. Introduction

Fractional calculus is a branch of applied mathematics, including integrals and derivatives of real, complex and arbitaray order. In the past decade, fractional calculus has been used different science and engineering disciplines. Fractional differential equations are increasingly being used to solve and evaluate problems in the field of fluid mechanics, biology, electromagnetic, acoustics, diffusion, signal processing, and many other physical processes [1-10].

Fractional calculus is the advancement of ordinary calculus by introducing derivatives in addition integrals of fractional order. Up to the middle of the 20th century many works have been done in the area of special branch namely fractional calculus by famous mathematicians such as Caputo [11], Lutzen and Liouville [12], Miller and Ross [8] ect. The applications of fractional calculus have been studied by many famous scientist and mathematician. So to name a few, Bagley and Torvik [13] discussed a model of arbitrary order of viscoelastic

nature, cancer model is given by Ahmed et.al [14], Singh et al. [15] examined of epidemiological model for computer viruses. The Drinfeld-Sokolov-Wilson (DSW) equation is investigated by Singh et.al [16], and diabetes models of fractional order have been analyzed by Singh et. al [17] with the exponential law. Cancer is one of the most dangerous and serious disease in the history of pathology. Cancer cells grow out of control and grow abnormally in the body. Cells are the basic units of the body building and cancer grows from normal cells. More recently, mathematical biology has been contributing to the development of theoretical models for the description and understanding of tumour development. Almost all these models focus on the categorization of tumour cells in three groups: necrotic cells (which are dead and located in the most internal part of the solid tumour), quiescent cells (are not dead, but do not have sufficient nutrients for cell growth or division), proliferating cells (the active part is that it undergo mitosis). A large class of cancer development models focuses on the proliferation dynamics of cells involved in the development of the tumour mass. When examining tumour growth, the process of cell senescence and cell division can not be ignored. All eukaryotic cells undergo a cell cycle, which is a four-step sequence [18-20]. Before DNA replicates in S phase, G1 phase is required for cell growth. This is followed by a second developmental phase (G2), the mitotic phase (M) ending the cycle involving the cell nucleus and cytoplasmic division. As a result of sequence completion, two subunits enter the G1 cycle. The first three stages are usually collectively referred to as the "intermediate stage." Cells can also enter the so-called G0 state, in which they will not grow, divide or live in a quiescent state. Usually, cells lacking growth factors will stop at the checkpoint and, move from G1 to G0, and the resume circulation after a period of time. The main

focus here is the cell cycle events that occur when growth factors stimulate G0 cell proliferation.

In this article we inspired and motivated by the ongoing research in this area, we will discuss the approximate method for solving nonlinear equations by homotopy analysis sumudu transform method (HASTM). The HASTM is a novel and efficient mixture of the homotopy analysis method (HAM) [21-23] and sumudu transform method (ST) [24]. Research study of the linear fractional malignant tumour equation explained by the homotopy analysis sumudu transforms method [25-27]. It provides the solutions in terms of convergent series with easily computable components in a direct way without using, perturbation, linearization or restrictive assumptions. The advantage of this method is that it combines two powerful methods to obtain accurate and approximate analytical solutions of nonlinear equations. The HASTM solution contains auxiliary parameters  $\hbar$  that can be used to manage convergence of solution.

The advancement of this paper is given as follows: In section 2, the integrals and sumudu transformation method of Atangana -Baleanu derivative and fractional order are discussed. In section 3, the fundamental plan of HASTM is presented. In section 4, fractional order for malignant tumour assoicated with non-singular kernel is discussed. In section 5, stability is verified. In section 6, solution of HASTM solution for malignant tumour associated with AB derivative is investigated. In section 7, numerical results and discussion are elaborated. In section 8, closing remarks are discussed.

## 2. Preliminaries

**Definition 1.** Let  $f \in H^1(a,b), b > a$ , then for  $\lambda \in (0,1]$  the Atangana-Baleanu fractional (AB) derivative [28] in Caputo sense is given below

$${}^{ABC}_{a}D_{t}^{\lambda}(\varphi(\tau)) = \frac{B(\lambda)}{1-\lambda} \int_{a}^{\tau} \varphi'(\delta) E_{\lambda} \left[ -\lambda \frac{(\tau-\delta)^{\lambda}}{1-\lambda} \right] d\delta$$
(2.1)

where  $B(\lambda)$  is a normalization function under the conditions B(0) = B(1) = 1. The related fractional integral is defined by

$$I^{\lambda}(\varphi(\tau)) = \frac{1-\lambda}{B(\lambda)}\varphi(\tau) + \frac{\lambda}{B(\lambda)\Gamma(\lambda)} \int_{0}^{\tau} \varphi(\delta)(\tau-\delta)^{\lambda-1} d\delta$$
(2.2)

**Definition 2.** The sumudu transform [24] is obtained over the set of function

$$A = \{ \varphi(\tau) : \ni N, \rho_1, \rho_2 > 0, |\varphi(\tau)| < N \exp\left(\frac{|\tau|}{\rho_i}\right),$$
if  $\tau \in (-1)^i \times [0, \infty) \}$ ,
as
$$S[\varphi(\tau)] = G(u) = \int_0^\infty \varphi(u\tau) \exp(-\tau) dt, \quad u \in (-\rho_1, \rho_2)$$
(2.3)

**Definition 3.** The sumudu transform of AB fractional derivative in Caputo sense [29] is given as follows

$$S\left({}^{ABC}_{0}D^{\lambda}_{t}\left(\varphi(\tau)\right)\right) = \frac{B(\lambda)}{1-\lambda+\lambda u^{\lambda}}\left(G(u)-\varphi(0)\right). \tag{2.4}$$

# 3. Fundamental plan of method

To discuss the initial idea of HASTM, let us take a non-linear partial differential equation of fractional order,

$${}^{ABC}_{0}D^{\eta}_{\theta}\upsilon(\varsigma,\theta) + R\upsilon(\varsigma,\theta) + N\upsilon(\varsigma,\theta) = \varphi(\varsigma,\theta),$$
  
0 < \eta \le 1,

(3.1)

where  $\upsilon(\varsigma, \vartheta)$  is a function of two variable  $\varsigma$  and  $\vartheta$ ,  ${}^{ABC}_{\phantom{ABC}}D^{\eta}_{\vartheta}$  denotes a fractional operator of order  $\eta$  defined in terms of AB fractional derivative,  $k \in N, R$  represents a linear operator bounded in two variables  $\varsigma$  and  $\vartheta$  i.e. we can have the number  $\alpha > 0$ 

s.t.  $\|R\upsilon\| \le \alpha \|\upsilon\|$  and N denote the general nonlinear differential operator in  $\varsigma$  and  $\vartheta$ , in Lipschitz continous with  $\delta > 0$  holding the condition  $|N\upsilon - N\varphi| \le \tau |\upsilon - \varphi|$  and  $\varphi(\varsigma,\vartheta)$  is the source term. Applying ST on Eq. (3.1), we have

$$S\left[ ABC \atop 0 \right] O_{\theta}^{\eta} v + S\left[ Rv + Nv \right] = S\left[ \varphi(\varsigma, \theta) \right].$$
 (3.2)

By explaining the differential characteristics of ST on Eq. (3.2), we obtain the following result

$$\frac{B(\eta)}{1-\eta+\eta u^{\eta}}S[\upsilon(u)-\upsilon(0)]+S[R\upsilon+N\upsilon]=S[\varphi(\varsigma,\vartheta)].$$
(3.3)

According to the Eq. (3.3), the nonlinear operator is expressed as follows

$$N[\rho(\varsigma,\vartheta;p)] = S[\rho(\varsigma,\vartheta;p)] - \upsilon(0) + \frac{1 - \eta + \eta u^{n}}{B(\eta)} \{S[R\rho(\varsigma,\vartheta;p) + N\rho(\varsigma,\vartheta;p)] - S[\varphi(\varsigma,\vartheta;p)]\},$$
(3.4)

where  $0 \le p \le 1$  is a parameter known as the embedding parameter and  $\rho(\varsigma, \vartheta; p)$  is denoting a function  $\varsigma, \vartheta$  and p.

Next ,we propose the homotopy represented by Eq. (3.5)

$$(1-p)S[\rho(\varsigma,\vartheta;p)-\nu_0] = \bar{h}N[\nu(\varsigma,\vartheta)],$$
(3.5)

where S denotes the sumudu transform operator,  $\overline{h} \neq 0$  is an auxiliary parameter,  $\upsilon_0(\varsigma, \vartheta)$  is an initial approximation of  $\upsilon(\varsigma, \vartheta)$  and  $\rho(\varsigma, \vartheta; p)$  is the unknown function. If the embedded parameters are p=0 and p=1, it is easy to explain

$$\rho(\varsigma, \vartheta; 0) = \upsilon_0(\varsigma, \vartheta), \rho(\varsigma, \vartheta; 1) = \upsilon(\varsigma, \vartheta).$$
(3.6)

Therefore, when p values from 0 to 1, the solution  $\rho(\varsigma, \vartheta; p)$  changes from the initial approximation  $\upsilon_0(\varsigma, \vartheta)$  to the solution  $\upsilon(\varsigma, \vartheta)$ . Enlarging the function  $\rho(\varsigma, \vartheta; p)$  in the form of Taylor series, we get the subsequent equation about p

$$\rho(\varsigma,\vartheta;p) = \nu_0(\varsigma,\vartheta) + \sum_{n=1}^{\infty} \nu_n(\varsigma,\vartheta)p^n,$$
(3.7)

where,

$$\upsilon_{n}(\varsigma,\vartheta) = \frac{1}{n!} \frac{\partial^{n}}{\partial q^{n}} \left\{ \rho(\varsigma,\vartheta;p) \right\} \Big|_{p=0}.$$
(3.8)

If the initial approximation  $v_0(\varsigma, \theta)$  and the convergence control parameter  $\overline{h}$  are suitably specified, then Eq.(3.7) converges at p=1, we have

$$\upsilon(\varsigma,\vartheta) = \upsilon_0(\varsigma,\vartheta) + \sum_{n=1}^{\infty} \upsilon_n(\varsigma,\vartheta).$$
 (3.9)

The result expressed by the Eq. (3.9) must be one of the solutions of the given sequence of nonlinear differential equations. By using the result (3.9), then the governing equation can be obtained by using the Eq. (3.5). We express the vector as follows

$$\vec{\upsilon}_n = \{\upsilon_0(\varsigma, \vartheta), \upsilon_1(\varsigma, \vartheta), \upsilon_2(\varsigma, \vartheta), \dots, \upsilon_n(\varsigma, \vartheta)\}.$$
(3.10)

Now, we differentiate the Eq. (3.5) n-times w.r.t. p, then divide by n! Finally, in the case of p=0, we get the following equation

$$S\left[\upsilon_{n}(\varsigma,\vartheta)-\chi_{n}\upsilon_{n-1}(\varsigma,\vartheta)\right]=\overline{h}\,\Re_{n}\left(\vec{\upsilon}_{n-1}\right).$$
(3.11)

Now using the inverse of sumudu transform on Eq. (3.11), we arrived at the subsequent result

$$\upsilon_{n}(\varsigma,\vartheta) = \chi_{n}\upsilon_{n-1}(\varsigma,\vartheta) + \overline{h}S^{-1} \Big[ \mathfrak{R}_{n}(\vec{\upsilon}_{n-1}) \Big],$$
(3.12)

where  $\chi_n$  is given as

$$\chi_n = \begin{cases} 0, & n \le 1, \\ 1, & n > 1. \end{cases}$$
 (3.13)

And we define the value of  $\mathfrak{R}_n(\nu_{n-1})$  in an improved manner as

$$\Re_{n}(\vec{v}_{n-1}) = S\left[\upsilon_{n-1}(\varsigma,\vartheta)\right] - \left\{\upsilon(0) + \frac{1 - \eta + \eta u^{n}}{B(\eta)}S\left[\varphi(\varsigma,\vartheta)\right]\right\} (1 - \chi_{n}) + \frac{1 - \eta + \eta u^{n}}{B(\eta)}S\left[R\upsilon_{n-1} + p_{n-1}\right].$$
(3.14)

In Eq. (3.14),

 $P_n$  is the homotopy polynomial [30] and is presented as

$$P_{n} = \frac{1}{\Gamma(\eta)} \left[ \frac{\partial^{n}}{\partial q^{n}} N \rho(\varsigma, \theta; p) \right]_{p=0}, \quad (3.15)$$

and

$$\rho(\varsigma, \theta; p) = \rho_0 + p\rho_1 + p^2\rho_2 + \cdots$$
 (3.16)

Using Eq. (3.14) in Eq. (3.12), we have

$$\upsilon_{n}(\varsigma,\vartheta) = \left(\chi_{n} + \overline{h}\right)\upsilon_{n-1}(\varsigma,\vartheta) - \\
\overline{h}\left(1 - \chi_{n}\right)S^{-1}\left\{\upsilon(0) + \frac{1 - \eta + \eta u^{n}}{B(\eta)}S\left[\varphi(\varsigma,\vartheta)\right]\right\} + \\
\frac{1 - \eta + \eta u^{n}}{B(\eta)}\overline{h}S^{-1}\left[S\left[R\upsilon_{n-1} + p_{n-1}\right]\right].$$
(3.17)

An improvement in the scheme used is that a novel correction function (3.17) has been introduced with the aid of homotopy polynomials. Therefore, using Eq. (3.17), we can determine  $\upsilon_n(\varsigma,\vartheta)$  for  $n \ge 1$  different components and the solution is represented by the following equation

$$\upsilon(\varsigma,\vartheta) = \sum_{n=0}^{\infty} \upsilon_n(\varsigma,\vartheta). \tag{3.18}$$

# 4. Fractional Model for Malignant Tumour Growth Assoicated with Non-Singular Kernel

When  $q_h(t)$  be the quiescent cells number (G0) at a time t,  $u_h(t)$  be the mitotic cells number at a time t and  $v_h(t)$  be the interphase cells number (G1;s and G2-phase) at a time t, then the mathematical model describing the tumour cell cycle as follows [31]

$$\frac{dq_h}{dt} = \alpha_Q \upsilon_h(t) - \left(\beta_Q + \gamma_{G_0}\right) q_h(t),$$

$$\frac{du_h}{dt} = 2\sigma_1 u_h(t) + \beta_Q q_h(t) - \delta_1 u_h \frac{d\upsilon_h}{dt}$$

$$= -\left(\omega_0 + \alpha_Q\right) \upsilon_h(t).$$
(4.1)

With the following intial condition

$$q_h(0) = c_1, u_h(0) = c_2, v_h(0) = c_3,$$

where  $\gamma_{G_0}$ ,  $\alpha_Q$ ,  $\beta_Q$ ,  $\sigma_1$ ,  $\omega_0$  and  $\delta_1$  represent death rate G0 cells, transition rate from G1 to G0, transition rate from G0 to G1, M cells divison rate, death rate G1 cells and death rate M cells respectively. This mathematical model for tumour cell cycle assocated with intger order derivative was studied by using homotopy perturbation scheme [32] and modified differential transform technique [33].

The classical derivatives especially Riemann and Caputo fractional derivatives, have restrictions beacuase their kernel is singular. Because the kernel is used by the physical system to explain the memory effect. It is clear that because of this weakness, mutually derived memory may not accurately reflect the full effect. We use the fractional derivative given by Atangana and Baleanu [28] to study the malignant tumour Growth model and to explain in a better and more efficient manner. The main advantages of this kind of opreator is that the singular power-law kernel is now replaced by non singular kernel ,which is easier to use in theoretical analysis, numerical calculations and real world applications.

We modify the model (4.1) by empolying the Atangana-Baleanu fractional derivative as follows

$$^{ABC}_{\phantom{ABC}0}D^{\lambda}q_{h}(t) = \alpha_{Q}\upsilon_{h}(t) - \left(\beta_{Q} + \gamma_{G_{0}}\right)q_{h}(t),$$

$$^{ABC}_{\phantom{ABC}0}D^{\lambda}_{t}u_{h}(t) = 2\sigma_{1}u_{h}(t) + \beta_{Q}q_{h}(t) - \delta_{1}u_{h}(t),$$

$$^{ABC}_{\phantom{ABC}0}D^{\lambda}_{t}\upsilon_{h}(t) = -\left(\omega_{0} + \alpha_{Q}\right)\upsilon_{h}(t).$$

$$(4.2)$$

With the following intial condition

$$q_h(0) = c_1, u_h(0) = c_2, v_h(0) = c_3,$$
 (4.3)

# 5. Stability analysis

In this section we discuss the steadystate stability analysis without cancer and establish the conditions of model parameters for tumour growth or eradication. The fractional order mathematical model of malignant tumour growth based on cell cycle dynamics (4.2) gives a linear system with the only stationary state  $R^* = (0,0,0)$ . Calculate the eigen values of the coefficient matrix to analyze the stability.

$$A = \begin{vmatrix} -\left(\beta_{\mathcal{Q}} + \gamma_{G_0}\right) & 0 & 0\\ \beta_{\mathcal{Q}} & 2\sigma_1 - \delta_1 & 0\\ 0 & 0 & -\left(\omega_0 + \alpha_{\mathcal{Q}}\right) \end{vmatrix}.$$

Characteristic equation is given as

$$\lambda^3 + b_1 \lambda^2 + b_2 \lambda - b_3 = 0,$$

where

$$\begin{aligned} b_{1} &= \delta_{1} - 2\sigma_{1} - \left(\beta_{Q} + \gamma_{G_{0}} + \omega_{0} + \alpha_{Q}\right), \\ b_{2} &= \left(\omega_{0} + \alpha_{Q}\right) \left(\beta_{Q} + \gamma_{G_{0}}\right) + \left(\omega_{0} + \alpha_{Q}\right) \left(2\sigma_{1} - \delta_{1}\right) + \\ \left(\beta_{Q} + \gamma_{G_{0}}\right) \left(2\sigma_{1} - \delta_{1}\right), \\ b_{3} &= \left(2\sigma_{1} - \delta_{1}\right) \left(\beta_{Q} + \gamma_{G_{0}}\right) \left(\omega_{0} + \alpha_{Q}\right). \end{aligned} \tag{5.1}$$

The charactersitic values are,

$$\begin{split} &\lambda_{1} = -\left(\beta_{\mathcal{Q}} + \gamma_{G_{0}}\right), \lambda_{2} = \left(2\sigma_{1} - \delta_{1}\right), \\ &\lambda_{3} = -\left(\omega_{0} + \alpha_{\mathcal{Q}}\right). \end{split}$$

All roots are real,

$$\lambda_1 < 0, \lambda_3 < 0.$$

The stationary state  $R^*$  is locally asymptotically stable, if  $\sigma_1 < \frac{\delta_1}{2}$ .

$$\sigma_1 < \frac{\delta_1}{2}$$
 means that the death rate of

mitotic cells exceeds the division rate, so cancer cells will not grow and the tumour will disappear at one point [31].

# 6. HASTM Solution for Malignant Tumour Growth Associated with AB Fractional Derivative

This section represents HASTM solution to analyze the numerical results of malignant tumour related to AB derivative. Firstly ,we apply the Sumudu transform method to malignant tumour model (4.2) ,it yields

$$\frac{B(\lambda)}{1-\lambda+\lambda u^{\lambda}} \Big[ S(q_{h}(t)) - c_{1} \Big] 
= S\Big[ \alpha_{Q} \upsilon_{h}(t) - (\beta_{Q} + \gamma_{G_{0}}) q_{h}(t) \Big], 
\frac{B(\lambda)}{1-\lambda+\lambda u^{\lambda}} \Big[ S(u_{h}(t)) - c_{2} \Big] 
= S\Big[ 2\sigma_{1}u_{h}(t) + \beta_{Q}q_{h}(t) - \delta_{1}u_{h}(t) \Big], 
\frac{B(\lambda)}{1-\lambda+\lambda u^{\lambda}} \Big[ S(\upsilon_{h}(t)) - c_{3} \Big] 
= S\Big[ -(\omega_{0} + \alpha_{Q})\upsilon_{h}(t) \Big].$$
(6.1)

On simplification we have,

$$S(q_{h}(t))-c_{1}-\frac{B(\lambda)}{1-\lambda+\lambda u^{\lambda}}S\left[\alpha_{Q}\nu_{h}(t)-\left(\beta_{Q}+\gamma_{G_{0}}\right)q_{h}(t)\right]=0,$$

$$S(u_{h}(t))-c_{2}-\frac{B(\lambda)}{1-\lambda+\lambda u^{\lambda}}S\left[2\sigma_{1}u_{h}(t)+\beta_{Q}q_{h}(t)-\delta_{1}u_{h}(t)\right]=0,$$

$$S(\nu_{h}(t))-c_{3}-\frac{B(\lambda)}{1-\lambda+\lambda u^{\lambda}}S\left[-\left(\omega_{0}+\alpha_{Q}\right)\nu_{h}(t)\right]=0.$$

$$(6.2)$$

We present the nonlinear operators as  $N_{1}\left[\xi_{1}(t;x)\right] = S\left[\xi_{1}(t;x)\right] - c_{1} - \frac{B(\lambda)}{1 - \lambda + \lambda u^{\lambda}} S\left[\alpha_{Q}\xi_{3}(t;x) - \left(\beta_{Q} + \gamma_{G_{0}}\right)\xi_{1}(t;x)\right] = 0,$   $N_{2}\left[\xi_{2}(t;x)\right] = S\left[\xi_{2}(t;x)\right] - c_{2} - \frac{B(\lambda)}{1 - \lambda + \lambda u^{\lambda}} S\left[2\sigma_{1}\xi_{2}(t;x) + \beta_{Q}\xi_{1}(t;x) - \delta_{1}\xi_{2}(t;x)\right] = 0,$   $N_{3}\left[\xi_{3}(t;x)\right] = S\left[\xi_{3}(t;x)\right] - c_{3} - \frac{B(\lambda)}{1 - \lambda + \lambda u^{\lambda}} S\left[-\left(\omega_{0} + \alpha_{Q}\right)\xi_{3}(t;x)\right] = 0.$  (6.3)

Hence we have,

$$\mathfrak{R}_{1,n}\left(q_{h(n-1)}\right) = S \left\lceil q_{h(n-1)} \right\rceil - c_1 \left(1 - \chi_n\right) - \frac{1 - \lambda + \lambda u^{\lambda}}{B(\lambda)} S \left[\alpha_{\mathcal{Q}} \upsilon_{h(n-1)} - \left(\beta_{\mathcal{Q}} + \gamma_{G_0}\right) q_{h(n-1)}\right], \\
\mathfrak{R}_{2,n}\left(u_{h(n-1)}\right) = S \left\lceil u_{h(n-1)} \right\rceil - c_2 \left(1 - \chi_n\right) - \frac{1 - \lambda + \lambda u^{\lambda}}{B(\lambda)} S \left[2\sigma_1 u_{h(n-1)} + \beta_{\mathcal{Q}} q_{h(n-1)} - \delta_1 u_{h(n-1)}\right], \\
\mathfrak{R}_{3,n}\left(\upsilon_{h(n-1)}\right) = S \left\lceil \upsilon_{h(n-1)} \right\rceil - c_3 \left(1 - \chi_n\right) - \frac{1 - \lambda + \lambda u^{\lambda}}{B(\lambda)} S \left[-\left(\omega_0 + \alpha_{\mathcal{Q}}\right) \upsilon_{h(n-1)}\right]. \tag{6.4}$$

The  $n^{th}$  order deformation equation is expressed as follows

$$S \left\lceil q_{hn}(t) - \chi_{n}\left(q_{h(n-1)}\right)(t) \right\rceil = \overline{h} \,\mathfrak{R}_{1,n}\left(q_{h(n-1)}\right),$$

$$S \left\lceil u_{hn}(t) - \chi_{n}\left(u_{h(n-1)}\right)(t) \right\rceil = \overline{h} \,\mathfrak{R}_{2,n}\left(u_{h(n-1)}\right),$$

$$S \left\lceil v_{hn}(t) - \chi_{n}\left(v_{h(n-1)}\right)(t) \right\rceil = \overline{h} \,\mathfrak{R}_{3,n}\left(v_{h(n-1)}\right).$$

$$(6.5)$$

Applying the inverse sumudu transform to Eq. (6.5), it yields

$$q_{hn}(t) = \chi_{n}(q_{h(n-1)})(t) + \overline{h}S^{-1} \left[ \mathfrak{R}_{1,n}(q_{h(n-1)}) \right],$$

$$u_{hn}(t) = \chi_{n}(u_{h(n-1)})(t) = \overline{h}S^{-1} \left[ \mathfrak{R}_{2,n}(u_{h(n-1)}) \right],$$

$$v_{hn}(t) = \chi_{n}(v_{h(n-1)})(t) = \overline{h}S^{-1} \left[ \mathfrak{R}_{3,n}(v_{h(n-1)}) \right].$$
(6.6)

Taking the initial condition

 $q_h(0) = 2 \times 10^5$ ,  $u_h(0) = 10^5$ ,  $v_h(0) = 4 \times 10^5$ , and solving Eq.(6.6) for  $n = 1, 2, 3, \dots$ . We obtain the values of  $q_{hn}(t)$ ,  $u_{hn}(t)$ ,  $v_{hn}(t)$ . So, the solution of fractional order malignant tumour cell model (4.2) is given as,

$$q_h(t) = q_{h0} + q_{h1} + q_{h3} + \cdots,$$
  

$$u_h(t) = u_{h0} + u_{h1} + u_{h3} + \cdots,$$
  

$$v_h(t) = v_{h0} + v_{h1} + v_{h3} + \cdots,$$

## 7. Numerical Results and Discussion

This section elaborate the numerical simulation of malignant tumour. The numerical results for the model (4.1) are calculated using HASTM. We have taken the values of parameter  $\alpha_Q = 0.02$ ,  $\beta_Q = 0.2$ ,  $\gamma_{G_0} = 0.1/10^4$ ,  $\sigma_1 = 1$ ,  $\delta_1 = 0.28$ ,  $\omega_0 = 0.11$  and the initial conditions are given as  $q_h(0) = c_1 = 2 \times 10^5$ ,  $u_h(0) = c_2 = 10^5$ ,  $v_h(0) = c_3 = 4 \times 10^5$ . In this model, the dynamics of three different tumour cell population (quiescent cells, interphase cells, and mitotic cells) are being studied. From Fig. 1(a), it is easy to see that the number of quiescent cells increases as the value of  $\lambda$ 

increases after some times is nature is

opposite. Similarly, the effect of  $\lambda$  on

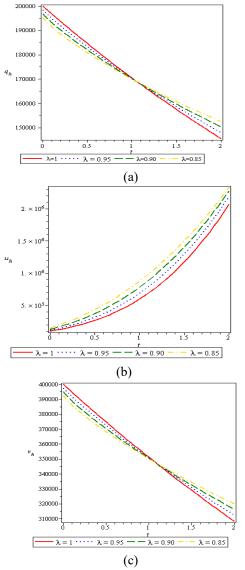
mitotic cells and interphase cells is shown in Figs. 1(b)-(c). From Figs. 2(a)-(b), it can

be seen that as values of  $\alpha_{\scriptscriptstyle O}$  (transition rate

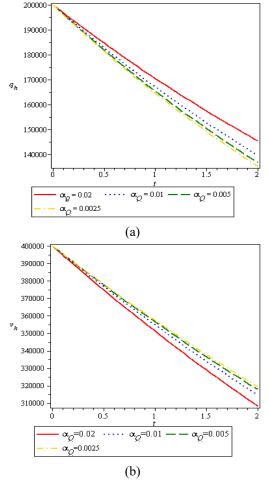
G1 to G0) increases, the number of

quiescent cells increases and as a values of

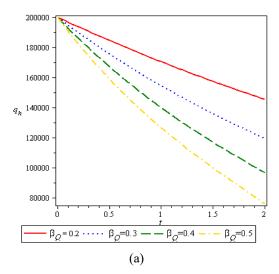
 $\alpha_{\mathcal{Q}}$  increase the number of interphase cells decrease. It can be seen that from Fig. 3 (a)-(b) as the the value of  $\beta_{\mathcal{Q}}$  (transition rate from G0 to G1) increases, the number of quiescent cells decreases and as the value of  $\beta_{\mathcal{Q}}$  (transition rate from G0 to G1) increases, the number of mitotic cells increases. Fig. 4 shows that as the values of death rate of M cells  $\left(\delta_{1}\right)$  increases the number of mitotic cells decreases.

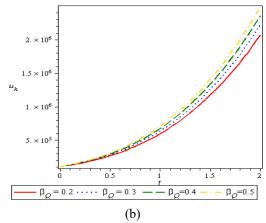


**Fig. 1.** Effect of order of AB fractional derivative on population (a) Quiescent cells (b) Mitotic cells (c) Interphase cells.



**Fig. 2.** Effect of transition rate from G1 to G0 on population when  $\lambda = 1$  (a) Quiescent cells (b) Interphase cells.





**Fig. 3.** Effect of transition rate from G0 to G1 on population when  $\lambda = 1$  (a) Quiescent cells (b) Mitotic cells.

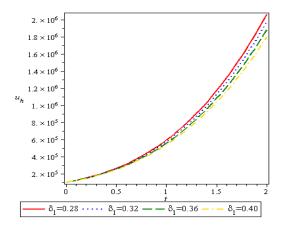


Fig. 4. Effect of death rate of M cells on population when  $\lambda = 1$  for Mitotic cells.

## 8. Conclusion

In this article, we studied the cell cycle model of malignant tumour associated with non-singular kernel. It can be seen that the growth of cancer mainly depends on the death rate of mitotic cells and the rate at which mitotic cells enter G1 without treatment. When the rate of mitotic is greater than the death rate of mitotic cells, cancer begins to grow. We use homotopy analysis sumudu transform method to study the fractional order mathematical model of malignant tumour growth based on cell cycle dynamics. The main benefit of this method is that it provides an analytical

approximate solution. The effects of order of fractional derivative and other parameters

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