

Article

## Time lag analysis of novel arithmetic modeling in breast cancer

Kalyan Das<sup>1</sup>, Ranjith Kumar<sup>2</sup>, Pankaj Taneja<sup>3</sup>, M Lutfor Rahman<sup>4</sup>

<sup>1</sup>National Institute of Food Technology Entrepreneurship and Management, HSIIDC Industrial Estate, Kundli, Haryana, India

<sup>2</sup>Anurag University, Hyderabad, Telangana, India

<sup>3</sup>Sharda University, Knowledge Park III, Greater Noida, Uttar Pradesh, India

<sup>4</sup>Institute of Statistical Research and Training, University of Dhaka, Dhaka 1000, Bangladesh

E-mail: daskalyan27@gmail.com, ranjithreddy1982@gmail.com, pankaj12750@rediffmail.com, lutfor@isrt.ac.bd

Received 13 January 2021; Accepted 20 February 2021; Published 1 June 2021



### Abstract

In this paper, a mathematical model which considers population dynamics among infected and uninfected cancer tumor cells has been proposed. Delay differential equations have been utilized to demonstrate the framework to consider the periods of the cell cycle. We examine the steadiness of the framework and demonstrate a hypothesis dependent on the contention standard to decide the dependability of a fixed point and show that the solidness may rely upon the delay. We show hypothetically as well as through numerical results that periodic oscillations may arise through Hopf bifurcations. In this paper we study a stochastic model for the conduct of malignancy tumors, depicted by a stochastic differential condition with multiplicative noise term. We study the existence of the solution process, as well as its behavior in the framework of stochastic inclusion problems and long time behavior.

**Keywords** breast cancer; time lag; stability; hopf-bifurcation; stochasticity.

Computational Ecology and Software  
ISSN 2220-721X  
URL: <http://www.iaees.org/publications/journals/ces/online-version.asp>  
RSS: <http://www.iaees.org/publications/journals/ces/rss.xml>  
E-mail: [ces@iaees.org](mailto:ces@iaees.org)  
Editor-in-Chief: WenJun Zhang  
Publisher: International Academy of Ecology and Environmental Sciences

### 1 Introduction

Breast cancer is a frequently occurred cancer among women, which is affecting 2.1 million women each year. It caused approximately 15% of all cancer deaths among women in 2018 (WHO, 2020). Symptoms of breast cancer include breast lump, bloody discharge from the nipple, and changes in the nipple or breast shape or texture. The treatment of cancer depends on type of cancer. For instance, chemotherapy, radiation, hormone therapy, and surgery some options to treat breast cancer (Diaby et al., 2015). The target organ breast cancer is a collection of malignancies in breast epithelial cells. Histological subtypes include ductal (70-80% of diagnosed cases), lobular (10-15% of diagnosed cases) or medular (3-5% of diagnosed cases). About 65% of













To establish the Hopf bifurcation at  $\tau = \bar{\tau}$ , we have to show that  $\frac{d\mu}{d\tau} \neq 0$ . For this differentiating (14) and (15) with respect to  $\tau$  and  $\tau = \bar{\tau}, \mu = 0$  and  $v = \bar{v}$  we get

$$\begin{aligned} L \frac{d\mu}{d\tau} + M \frac{dv}{d\tau} &= U \\ -M \frac{d\mu}{d\tau} + L \frac{dv}{d\tau} &= V \end{aligned} \quad (17)$$

where

$$\begin{aligned} L &= P_1 - Q_2 \tau \cos v \tau + Q_1 \cos v \tau - Q_1 v \tau \sin v \tau; \quad M = -2v - Q_2 \tau \sin v \tau + Q_1 \sin v \tau + Q_1 v \tau \cos v \tau; \\ U &= Q_2 v \sin v \tau - Q_1 v^2 \cos v \tau; \quad V = Q_2 v \cos v \tau + Q_1 v^2 \sin v \tau; \end{aligned}$$

Solving (17) we get

$$\frac{d\mu}{d\tau}(\bar{\tau}) = \frac{LU - MV}{L^2 + M^2} \text{ and the sign of } \frac{d\mu}{d\tau}(\bar{\tau}) \text{ is same as } LU - MV.$$

$$\begin{aligned} LU - MV &= (P_1 Q_2 v + 2Q_1 v^3) \sin v \tau + (2Q_2 v^2 - P_1 Q_1 v^2) \cos v \tau - Q_1^2 v^2 \\ &= v \left[ 2v(Q_1 v \sin v \tau + Q_2 \cos v \tau) - P_1(Q_1 v \cos v \tau - Q_2 \sin v \tau) - Q_1^2 v^2 \right] \end{aligned}$$

$$\text{Let } \phi(h) = h^2 + A_1 h + A_2$$

Clearly  $\phi(h)$  is the L.H.S. of (16) with  $v^2 = h$  and  $\phi(v^2) = 0$  then  $v$  is the first positive root of (16) and

$$\frac{d\mu}{d\tau} > 0 \text{ at } \tau = \bar{\tau} \text{ and stability cannot take the place these values of } \tau \text{ and}$$

$$\frac{d\mu}{d\tau}(\bar{\tau}) = \frac{v^2}{L^2 + M^2} \frac{d\phi}{dh}(v^2) \text{ where } \bar{\tau} = \frac{1}{\bar{v}} \arctan \frac{Q_1 \bar{v}^3 - (P_2 Q_1 + P_1 Q_2) \bar{v}}{\bar{v}^2 (Q_2 - P_1 Q_1) - P_2 Q_2} + \frac{n\pi}{\bar{v}}, n = 0, 1, 2, 3, \dots$$

By Hopf bifurcation theorem we can state that if  $M_1 > 0$ , and  $M_2 < 0$ , then there exist  $\bar{\tau} > 0$  such that  $E^*$  possesses a supercritical type Hopf-bifurcation as  $\tau$  increases and passes through  $\bar{\tau}$ .

#### 4 Stochastic Stability Analysis

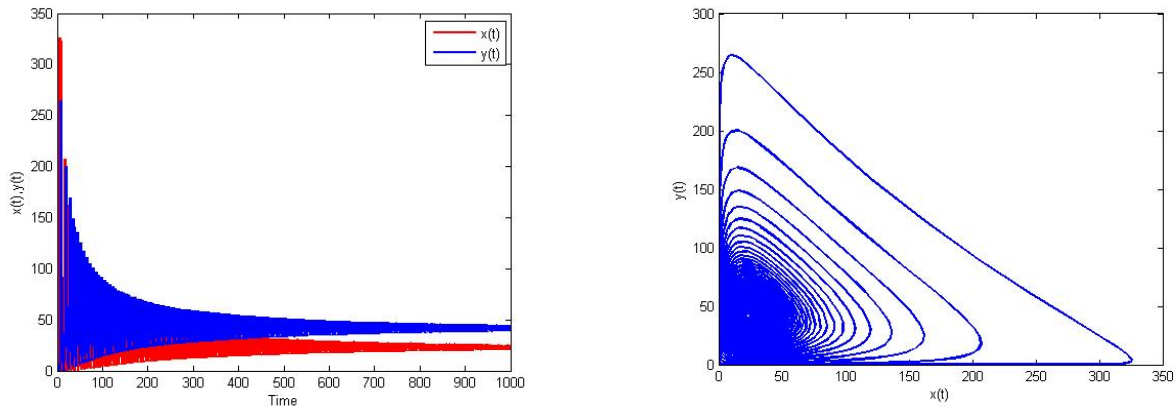
In this section we presented the environmental disturbances on the system (1) by white noise theory. These results are discussed at positive equilibrium point. To discuss the stability of the stochastic system, we consider the linearized model with the perturbations  $x_1$  and  $x_2$ . By using mean-square fluctuations we characterized the stochastic stability of the system. The stochastic perturbed system with delay is given by

$$\frac{dx(t)}{dt} = \left[ rx \left( 1 - \frac{x+y}{k} \right) - dx - \beta xy \right] dt + q_1 \xi_1(t) \quad (18)$$

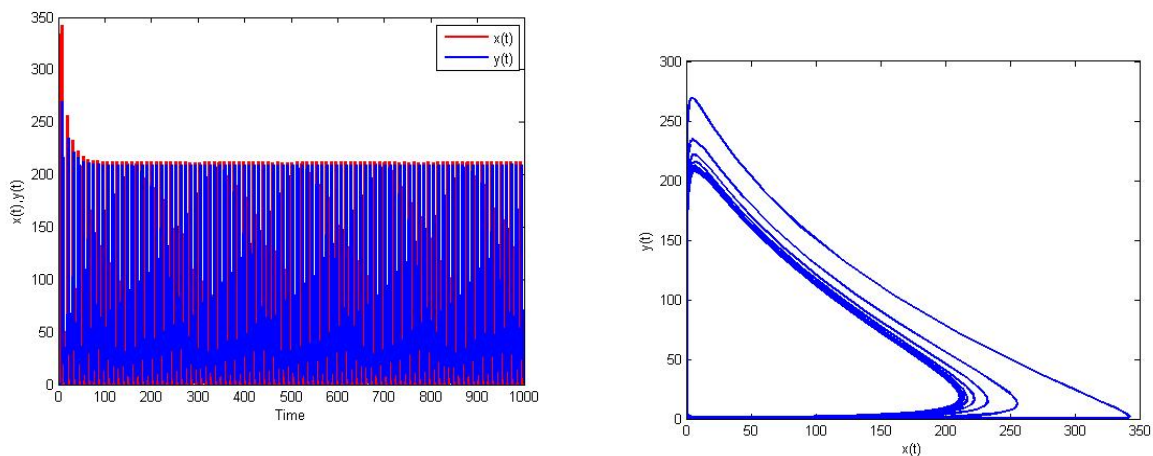


$$\frac{dy(t)}{dt} = \left[ \beta x(t-\tau)y(t-\tau) + sy \left( 1 - \frac{x+y}{k} \right) - \alpha y \right] dt + q_2 \xi_2(t)$$

Linearising above system with the perturbations  $x_1(t)$  and  $x_2(t)$  i.e.,  $x = x_1 + x^*$ ,  $y = x_2 + y^*$  then



**Fig. 2** The trajectories and phase graphs of system (1) with the parameter values as  $r = 2; k = 400; d = 0.211; \beta = 0.035; s = 0.01; \alpha = 0.814;$  and  $\tau = 0.09$  is stable.



**Fig. 3** The trajectories and phase graphs of system (1) with the parameter values as  $r = 2; k = 400; d = 0.211; \beta = 0.035; s = 0.01; \alpha = 0.814;$  and  $\tau = 0.2$  is unstable and a periodic orbit bifurcate from  $E^*$

$$\frac{dx_1(t)}{dt} = \frac{-2rx^*x_1}{k} - \frac{-rx^*x_2}{k} - \beta x^*x_2 + q_1 \xi_1(t) \tag{19}$$

$$\frac{dx_2(t)}{dt} = \beta x_1(t-\tau) - \frac{sy^*x_1}{k} - \frac{2sy^*x_2}{k} + q_2 \xi_2(t)$$

Applying Fourier transforms both sides and we obtain,

$$q_1 \xi_1(t) = \left( i\omega + \frac{2rx^*}{k} \right) x_1(\omega) + \left( \frac{rx^*}{k} + \beta x^* \right) x_2(\omega)$$

$$q_2 \xi_2(t) = \left( \frac{sy^*}{k} - \beta e^{-i\omega\tau} \right) x_1(\omega) + \left( i\omega + \frac{2sy^*}{k} \right) x_2(\omega)$$

The matrix form of above system is given by

$$\bar{\xi}(\omega) = M(\omega) X(\omega) \quad (20)$$

and denoting the elements of  $M(\omega)$  are  $m_{11}, m_{12}, m_{21}$  &  $m_{22}$  (row wise) then

$$M(\omega) = \begin{bmatrix} m_{11} & m_{12} \\ m_{21} & m_{22} \end{bmatrix} \text{ where}$$

$$\bar{\xi}(\omega) = \begin{bmatrix} q_1 \xi_1(t) \\ q_2 \xi_2(t) \end{bmatrix}; X(\omega) = \begin{bmatrix} x_1(\omega) \\ x_2(\omega) \end{bmatrix}; m_{11} = i\omega + \frac{2rx^*}{k}; m_{12} = \frac{rx^*}{k} + \beta x^*;$$

$$m_{21} = \frac{sy^*}{k} - \beta e^{-i\omega\tau}; m_{22} = i\omega + \frac{2sy^*}{k}.$$

Here  $M(\omega)$  is non-singular matrix then inverse of this matrix exist, therefore from (20)

$$X(\omega) = M^{-1}(\omega) \bar{\xi}(\omega) = N(\omega) \bar{\xi}(\omega) \quad (21)$$

$$\text{Where } N(\omega) = \begin{bmatrix} n_{11} & n_{12} \\ n_{21} & n_{22} \end{bmatrix} = M^{-1}(\omega) = \begin{bmatrix} \frac{m_{22}(\omega)}{|M(\omega)|} & -\frac{m_{12}(\omega)}{|M(\omega)|} \\ -\frac{m_{21}(\omega)}{|M(\omega)|} & \frac{m_{11}(\omega)}{|M(\omega)|} \end{bmatrix}$$

Now from the spectral density, we define

$$S_g(\omega) d\omega = \lim_{T \rightarrow \infty} \frac{|\bar{g}(\omega)|^2}{T}$$

Where  $g(t)$  a random function with is mean zero and  $S_g(\omega)$  represents the variance of the elements of  $g(t)$  within the interval  $[\omega, \omega + d\omega]$ .

The inverse transform of  $S_g(\omega)$  is the auto covariance function is given by

$$C_g(\tau') = \frac{1}{2\pi} \int_{-\infty}^{\infty} S_g(\omega) e^{i\omega\tau'} d\omega$$

and the variance function  $g(t)$  is given by

$$\sigma_g^2 = C_g(0) = \frac{1}{2\pi} \int_{-\infty}^{\infty} S_g d\omega$$

From (21), the mean value of the population is  $\bar{x}_i = \sum_{j=1}^2 n_{ij} \xi_j(\omega)$  where  $b_{ij}, i, j = 1, 2$

Therefore,  $S_{x_i} = \sum_{j=1}^2 q_j |n_{ij}(\omega)|^2, (i = 1, 2)$

The fluctuations of  $x_i (i = 1, 2)$  are given by

$$\sigma_{x_i}^2 = \frac{1}{2\pi} \int_{-\infty}^{\infty} S_{x_i} d\omega = \frac{1}{2\pi} \sum_{j=1}^2 \int_{-\infty}^{\infty} q_j |n_{ij}(\omega)|^2 d\omega$$

Therefore, from above variances and from system (18), we can find

$$\sigma_{x_1}^2 = \frac{1}{2\pi} \left[ q_1 \int_{-\infty}^{\infty} \frac{|m_{22}(\omega)|^2}{|M(\omega)|} d\omega + q_2 \int_{-\infty}^{\infty} \frac{|-m_{12}(\omega)|^2}{|M(\omega)|} d\omega \right]$$

$$\sigma_{x_2}^2 = \frac{1}{2\pi} \left[ q_1 \int_{-\infty}^{\infty} \frac{|-m_{21}(\omega)|^2}{|M(\omega)|} d\omega + q_2 \int_{-\infty}^{\infty} \frac{|m_{11}(\omega)|^2}{|M(\omega)|} d\omega \right]$$

Where  $|M(\omega)| = M_1(\omega) + iM_2(\omega)$  and

$$M_1(\omega) = -\left( \omega^2 - \frac{3rsx^*y^*}{k^2} + \frac{s\beta x^*y^*}{k} - \frac{r\beta x^*}{k} \cos \omega\tau - \beta^2 x^* \cos \omega\tau \right)$$

$$M_2(\omega) = -\left( -\frac{2sy^*\omega}{k} - \frac{2rx^*\omega}{k} + \frac{r\beta x^*}{k} \sin \omega\tau + \beta^2 x^* \sin \omega\tau \right)$$

The variances of  $x_i (i = 1, 2)$  are given by

$$\sigma_{x_1}^2 = \frac{1}{2\pi} \left[ q_1 \int_{-\infty}^{\infty} \frac{|m_{22}(\omega)|^2}{M_1^2(\omega) + M_2^2(\omega)} d\omega + q_2 \int_{-\infty}^{\infty} \frac{|m_{12}(\omega)|^2}{M_1^2(\omega) + M_2^2(\omega)} d\omega \right]$$

$$\sigma_{x_2}^2 = \frac{1}{2\pi} \left[ q_1 \int_{-\infty}^{\infty} \frac{|m_{21}(\omega)|^2}{M_1^2(\omega) + M_2^2(\omega)} d\omega + q_2 \int_{-\infty}^{\infty} \frac{|m_{11}(\omega)|^2}{M_1^2(\omega) + M_2^2(\omega)} d\omega \right] \tag{22}$$

Where  $|m_{11}|^2 = \omega^2 + \left(\frac{2rx^*}{k}\right)^2$ ;  $|m_{12}|^2 = \left(\frac{rx^*}{k} + \beta x^*\right)^2$

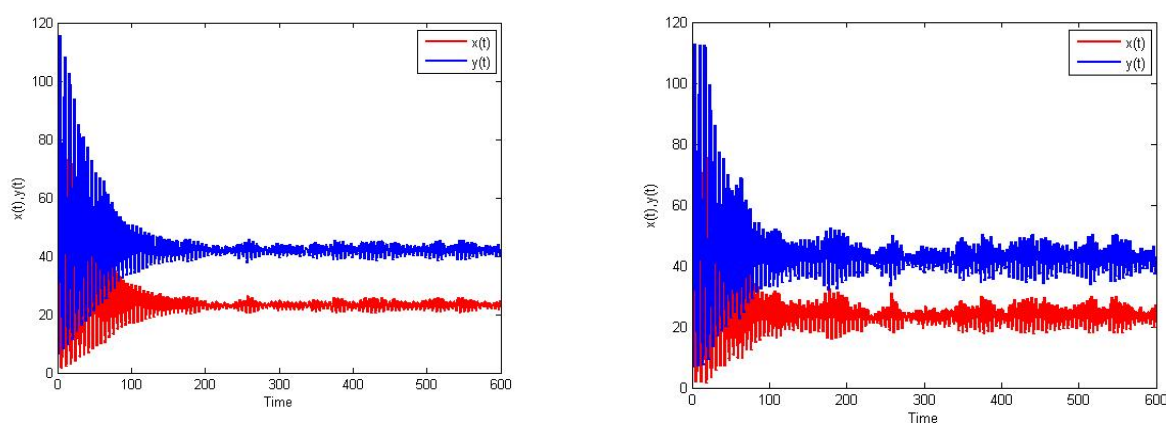
$$|m_{21}|^2 = \left( \frac{sy^*}{k} - \beta \cos \omega \tau \right)^2 + (\beta \sin \omega \tau)^2; |m_{22}|^2 = \omega^2 + \left( \frac{2sy^*}{k} \right)^2$$

The results in (22) gives the variances of the system (18) populations  $x$  and  $y$ . Generally, to find these integrals it is very difficult. Therefore, by using numerical simulations we can explain these results easily. Taking different parameter values and for the some time lag we can calculate variance and this is small then the corresponding population is stable, otherwise unstable.

## 5 Conclusion and Discussions

In this paper we have formulated a delayed cancer model with infected cells and antiseptic tumor cells. Breast cancer risk assessments using statistical data based on epidemiologic data are valid; however, no single model includes family history, estrogen-dependent mechanisms, and abnormal breast disease. For this reason it is helpful to use different models in a specialized risk assessment clinic, but this requires a full understanding of the strengths and powers of a particular relationship.

From Fig. 2, we have a tendency to look at how quickly the positive balance of program (1) is stabilized as at home. Modeling within this state of affairs produces sustainable energy. With the same set of parameter values, we find that when the system is stable, it can be seen in Fig. 2 and when it is unstable Fig3 with constant stability. That means the system changes its behavior from stable to unstable when time delay crosses its critical value, which gives Hopf-bifurcation. From the biological point of view delay has influence in spread of virus in tumor cells. Conjointly one should bear in mind the pause to allow for the management of the germ cell growth. The existence of time-resolved solutions is consistent in cancer models. It means that the tumor level can get around a difficult and immediate purpose despite no other treatment. We have accomplished numerical simulation of the random model quickly so as to add the increasing white noise of the same amount of fluctuations present in the system and that we often look at the fluctuations in the amplitude of the sounds (Fig. 4).



**Fig. 4** The trajectories of system (18) with the parameter values as

$r = 2; k = 400; d = 0.211; \beta = 0.035; s = 0.01; \alpha = 0.814;$  and  
 $q_1 = 0.1, q_2 = 0.4$  (left figure) &  $q_1 = 0.7, q_2 = 0.9$  (right figure)

## References

- Adam J, Bellomo N. 1997. A Survey of Models for Tumor Immune Dynamics. Birkhauser, Boston, USA
- Araujo RP, Elwaina DLSM. 2004. A history of the study of solid tumour growth: The contribution of mathematical modelling. *Journal of Bulletin of Mathematical Biology*, 66: 1039-1091
- Byrne HM. 1997. The effect of time delays on the dynamics of avascular tumor growth. *Mathematical Biosciences*, 144(2): 83-117
- Criscitiello C, Azim HA, Schouten PC, Linn SC, Sotiriou C. 2012. Understanding the biology of triple negative breast cancer. *Ann Oncol*, 23(Suppl 6): vi13-8
- de Boer Hoogeweg RP, Dullens H, de Weger R, Den Otter W. 1985. Macrophage T lymphocyte interactions in the anti-tumor immune response: a mathematical model. *Journal of Immunology*, 134(4): 2748-2758.
- Diaby V, Tawk R, Sanogo V, Xiao H, Montero AJ 2015. A review of systematic reviews of the cost-effectiveness of hormone therapy, chemotherapy, and targeted therapy for breast cancer. *Breast Cancer Research and Treatment*, 151(1): 27-40
- El-Gohary A. 2008. Chaos and optimal control of cancer self-remission and tumor system steady states. *Chaos, Solitons and Fractals*, 37(5): 1305-1316
- Forys U, Bodnar M, Poleszczuk J. 2011. Negativity of delayed induced oscillations in a simple linear DDE. *Applied Mathematics Letters*, 24(6): 982-986
- Forys U, Kolev M. 2002. Time delays in proliferation and apoptosis for solid avascular tumor. *Prep Institute of Applied Mathematics and Mechanics, rw02-10 (110)*. Warsaw University, Poland
- Freedman, H.I., Erbe, L.H., Rao, V.S.H., 1986. Three species food chain model with mutual interference and time delays. *Mathematical Biosciences*, 80: 57-80
- Goldstein B, Faeder JR, Hlavacek WS. 2004. Mathematical and computational models of immune-receptor signaling. *Nature Reviews Immunology*, 4(6): 445-456
- Gopalsamy K. 1992. *Stability and Oscillations in Delay Differential Equations of Population Dynamics*. Kluwer Academic, USA
- Khajanchi S. 2018. Modeling the dynamics of glioma-immune surveillance. *Chaos Solitons and Fractals*, 114: 108-118
- Kirschner D, Panetta JC. 1998. Modeling the immunotherapy of tumor - immune interaction. *Journal of Mathematical Biology*, 37(3): 235-252
- Koscielny S, Tubiana M, Valleron AJ. 1985. A simulation model of the natural history of human breast cancer. *British Journal of Cancer*, 52(4): 515-524
- Kronik N, Kogan Y, Vainstein V, Agur Z. 2008. Improving alloreactive CTL immunotherapy for malignant gliomas using a simulation model of their interactive dynamics. *Cancer Immunology, Immunotherapy*, 57(3): 425-439
- Kuznetsov VA, Makalkin IA, Taylor MA, Perelson AS. 1994. Non-linear dynamics of immunogenic tumors: Parameter estimation and global bifurcation analysis. *Bulletin of Mathematical Biology*, 56(2): 295-321
- Lehmann BD, Bauer JA, Chen X, Sanders ME, Chakravarthy AB, Shyr Y, Pietenpol JA 2011. Identification of human triple-negative breast cancer subtypes and preclinical models for selection of targeted therapies. *Journal of Clinical Investigation*, 121(7): 2750-2767
- Miękisz J, Poleszczuk J, Bodnar M, Forys U. 2011. Stochastic models of gene expression with delayed degradation. *Bulletin of Mathematical Biology*, 73(9): 2231-2247
- Network CGA. 2012. Comprehensive molecular portraits of human breast tumours. *Nature*, 490(7418): 61-70
- Norton L. 1988. A Gompertzian model of human breast cancer growth. *Cancer Research*, 48(24 Pt 1): 7067-7071

- Rejniak KA, Anderson ARA. 2011. Hybrid models of tumor growth. *Wiley Interdisciplinary Reviews: Systems Biology and Medicine*, 3(1): 115-125
- Preziosi L. 2003. *Cancer Modelling and Simulation (Mathematical Biology and Medicine Series, Vol. 3. Chapman & Hall/CRC), USA*
- Oncology NCCNCPGi 2015. *Clinical Practice Guidelines in Oncology: Genetic/Familial High-Risk Assessment: Breast and Ovarian. USA*
- Parise CA, Caggiano V 2014. Breast cancer survival defined by the ER/PR/HER2 subtypes and a surrogate classification according to tumor grade and immunohisto chemical biomarkers. *Journal of Cancer Epidemiology*, 469251
- Petrucelli N, Daly MB, Feldman GL. 2010. Hereditary breast and ovarian cancer due to mutations in BRCA1 and BRCA2. *Genetics in Medicine*, 12(5): 245-259
- Piotrowska MJ, Foryś U. 2011. Analysis of the Hopf bifurcation for the family of angiogenesis models, *Journal of Mathematical Analysis and Applications*, 382(1): 180-203
- Piotrowska M, Forys U. 2011. The nature of Hopf bifurcation for the Gompertz model with delays, *Mathematical and Computer Modelling*, 54: 9-10
- Retsky M, Swartzendruber D, Wardwell R, Bame P, Petrosky V. Re: Larry Norton 1989. A Gompertzian model of human breast cancer growth. *Cancer Research*, 49(22): 6443-6444
- Ribba B, Holford NH, Magni P, Trocóniz I, Gueorguieva I, Girard P, Sarr C, Elishmereni M, KloftC, Friberg LE. 2014. A review of mixed-effects models of tumor growth and effects of anticancer drugtreatment used in population analysis. *CPT: Pharmacometrics and Systems Pharmacology*, 3: e113
- Sarkar RR, Banerjee S. 2005. Cancer self remission and tumor stability—a stochastic approach. *Mathematical Biosciences*, 196(1): 65-81
- Starkov KE, Bunimovich-Mendrazitsky S. 2016. Dynamical properties and tumor clearance conditions for a nine-dimensional model of bladder cancer immunotherapy. *Mathematical Biosciences and Engineering*, 13(5): 59-1075
- Sorlie T, Tibshirani R, Parker J, Hastie T, Marron JS, Nobel A, Deng S, Johnsen H, Pesich R, Geisler S. 2003. Repeated observation of breast tumor subtypes in independent gene expression data sets. *Proceedings of the Natural Academy of Sciences of USA*, 100(14): 8418-8423
- Speer JF, Petrosky VE, Retsky MW, Wardwell RH 1984. A stochastic numerical model of breast cancer growth that simulates clinical data. *Cancer Research*, 44(9): 4124-4130
- Spratt JA, von Fournier D, Spratt JS, Weber EE. 1993. Mammographic assessment of human breast cancer growth and duration. *Cancer*, 71: 2020-2026
- Villasana M, Radunskaya A. 2004. A delay differential equation model for tumor growth. *Journal of Mathematical Biology*, 47: 270-294
- Wong-Brown MW, Meldrum CJ, Carpenter JE, Clarke CL, Narod SA, Jakubowska A, Rudnicka H, Lubinski J, Scott RJ. 2015. Prevalence of BRCA1 and BRCA2 germline mutations in patients with triple-negative breast cancer. *Breast Cancer Research and Treatment*, 150(1): 71-80
- Yafia R. 2006. Dynamics analysis and limit cycle in a delayed model for tumor growth with quiescence. *Nonlinear Analysis: Modelling and Control*, 11(1): 95-110