

Computational Modelling of Dynamical System and the Type of Stability

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Received: 17 May 2023/Accepted 14 July 2023/Published 17 July 2023

Abstract: *The study of computational modelling of a dynamical system and the type of stability was investigated using ODE45 numerical techniques. Due to the decrease and increase of the growth rates of yeast species 1 and 2 otherwise called environmental perturbation on the prediction of the extent of the proportion decrease and increase in biodiversity. A biodiversity gain was observed when the growth rates increased together from 101% - 150%. When growth rates are decreased together by 50%, it was also found that, there is a biodiversity loss of yeast species. Finally, the region of instability was found since the pairs of eigenvalues are positive.*

Keywords: *Mathematical model; ODE45; Delay, Dynamical System, and stability*

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1.0 Introduction

A dynamic system is a system in which motion takes place under the influence of some forces. Dynamical systems are key foundation of every evolving real-world situation and the system can be used to describe the asymptotic behavior of a natural or man-made system.

In recent years, the dynamical system has had many applications to science and engineering

from many researchers (Bertoin, 2016; Chellaboina *et al.*, 2003; Edward and Ford, 2003; Eli and Abanum, 2020; Godspower *et al.*, 2020; Hale, 1969; Yan, Y. & Ekakaa, 2011) in Mathematical Modeling and Ecological Modeling. Some of which have gone under the related headings of chaos theory or nonlinear theory. The dynamical systems have two types' continuous and discrete dynamical systems. If the time in the equation is implicit then is called an autonomous equation. Further, if the time is explicit then it is called non-autonomous. Mathematics has always benefitted from its involvement with developing sciences. Each successive interaction revitalizes and enhances the field. Biomedical sciences are the premier science of the near future. Mathematical biology is a fast-growing well recognized and the most exciting modern application of mathematics. The increasing use of mathematics in biology is inevitable as biology becomes quantitative.

The biological mathematical model becomes the theme for the dynamical systems. Therefore, the biological model was studied theoretically and numerically. The complexity of biological sciences makes interdisciplinary involvement essential. For the mathematician, biology opens up new and exciting branches while for the biologists' mathematical modelling offers another research tool.

Eli and Abanum (2020) presented a comparison between the Analytical and Numerical results of the stability analysis of a dynamical system. They formulated the system of ordinary Differential Equations involving Sickle Cells, HIV and T-Cells with the aid of a biological mathematical model. The eigenvalues were obtained to test for the trivial steady-state solution or points using a characteristic equation which is analytical.

Finally, they carried out a numerical simulation to test the level of reliability of the result. Solomonovich *et al.* (1998) studied the stability analysis problem for a new class of discrete-time recurrent neural networks with mixed time delays. The mixed time delays that consist of both the discrete and distributed time delays are addressed, for the first time, when analyzing the asymptotic stability for discrete-time neural networks. The activation functions are not required to be differentiable or strictly monotonic. The existence of the equilibrium point was first proved under mild conditions by constructing a new Lyapunov-Krasovskii functional, a linear matrix inequality (LMI) approach is developed to establish sufficient conditions for the discrete-time neural networks to be globally asymptotically stable. As an extension, they further consider the stability analysis problem for the same class of neural networks but with state-dependent stochastic disturbances. All the conditions obtained are expressed in terms of LMIs whose feasibility can be easily checked by using the numerically efficient Matlab LMI Toolbox. A simulation example is presented to show the usefulness of the derived LMI-based stability condition.

Eli and Ekakaa, (2021) studied the effect of discrete time delays on the stability of a dynamical system using ODE45 numerical simulation techniques. It was shown from the result that the dynamical system is dominantly unstable.

Edward and Ford, (2003) studied the boundness and stability of differential equations. Their paper discusses the qualitative behaviour of solutions to different equations, focusing on the boundedness and stability of solutions. Examples demonstrate how the use of Lipschitz constants can provide insights into the qualitative behaviour of solutions to some nonlinear problems. Manchester Centre for Computational Mathematics their findings, and their conclusions. Chellaboina *et al.*, (2003),

discussed a dissipative dynamical systems approach to stability analysis of time-delay systems. In their paper, the concepts of dissipativity and exponential dissipativity are used to provide sufficient conditions for guaranteeing the asymptotic stability of a time delay dynamical system. Specifically, representing a time delay dynamical system as a negative feedback interconnection of a finite-dimensional linear dynamical system and an infinite-dimensional time delay operator, they show that the time delay operator is dissipative concerning a quadratic supply rate and with a storage functional involving an integral term identical to the integral term appearing in standard Lyapunov-Krasovskii functionals. Finally, using the stability of feedback interconnection results for dissipative systems, they develop enough conditions for asymptotic stability of time delay dynamical systems. The overall approach provides a dissipativity theoretic interpretation of Lyapunov-Krasovskii functionals for asymptotically stable dynamical systems with arbitrary time delay.

Rajendra (2021) analysed the standard approach of dynamical systems towards biomedical science. The purpose of their study is to meet the current and future needs for the interaction between various science and technology areas on the one hand and dynamical systems on the other hand. They discuss various models for interacting populations like Predator-Prey model, Lotka - Volterra system, the Realistic Predator-Prey model, Continuous Population models for single species, Discrete Population models for single species etc. They develop models which capture the essence of various interactions allowing the outcomes to be more fully understood. It has a very broad perspective to analyse biomedical issues using one of the most important branches of mathematics. The main objective of the study is to elaborate dynamical tools and this type of study will act as the bridge between



biomedical sciences and mathematical sciences. Anand and Melba Mary (2016), also investigated an Improved Dynamic Response of DC to DC Converter Using Hybrid PSO Tuned Fuzzy Sliding Mode Controller. According to them, DC/DC switching converters are widely used in numerous appliances in modern existence. In their paper, the dynamic and transient response of phase shift series resonant DC/DC converter is improved using hybrid particle swarm optimization tuned fuzzy sliding mode controller under starting and load step change conditions. The aim of the control is to regulate the output voltage beneath the load change. The model of the hybrid particle swarm optimization tuned fuzzy sliding mode controller is implemented using the Sim Power Systems toolbox of MATLAB SIMULINK. The performance of the proposed dynamic novel control under step load change conditions is investigated.

Theory of Dynamic Interactions: Innovations was examined by Alejandro Álvarez (2016). The theory of dynamic interactions suggests a new paradigm of mechanics and initiates them into a new area of knowledge, hitherto undeveloped. In their paper, they describe the innovations that this theory brings to physics, and in particular, the ideas expressed in a new book by Doctor Barceló: New Paradigm in Physics. It is necessary to analyse the incorporation into mechanics not only of knowledge about bodies with inertial movement but also that of non-inertial systems. It is necessary for a new structure of knowledge that can incorporate both inertial and accelerated systems. In this paper, we referred to the main innovations and novel ideas proposed by Doctor Barceló in his new book, concerning the rotational dynamics

1.1 Model assumptions

For this study, we shall consider the following assumptions:

- (i) The growth of yeast species 1 and yeast species 2 depends on the

difference between the survival rate and the death rate.

- (ii) The growth of these two species can also be affected by the interaction within each species otherwise called the self-interaction coefficient of the intra-competition process.
- (iii) The growth of these two species can also be affected by the inter-competition coefficient which specified the contribution of each species to inhibit the growth of each species.
- (iv) The growth of these two competing species can also be affected by the initial data values of yeast species 1 and yeast species 2 when under the length of the growing season in the unit of weeks.

1.2 Mathematical formulations

For this study, we have considered the following multi-parameter continuous dynamical system of a nonlinear first Order Ordinary Differential Equation from Eli and Ekakaa, 2021.

$$\frac{dx}{dt} = \alpha_1 x - \beta_1 x^2 - \gamma_1 xy \quad (1)$$

$$\frac{dy}{dt} = \alpha_2 y - \beta_2 y^2 - \gamma_2 xy \quad (2)$$

$x(t)$ denotes the biomass of yeast specy 1 (candida albican) at time t in the unit of weeks.

$y(t)$ denotes the biomass of yeast specy 2 (candida parapsilosis) at time t in the unit of weeks.

α_1 and α_2 specifies the growth rate of yeast species 1 and 2 respectively.

β_1 and β_2 specifies the intra-competition coefficient of yeast species 1 and 2 respectively.

γ_1 and γ_2 denotes the competition of yeast species 1 and yeast 2 respectively where γ_1 is the contribution of the yeast species to inhibit the growth of species 2 as γ_2 is the contribution of the yeast specy 2 to inhibit the growth of the specy 1



2.0 Linearization of the Dynamical System

To obtain different eigenvalues to test for the stability of the system, it is important to linearize the system by letting the function F_1 and F_2 represent equations (1) and (2) respectively as

$$F_1 = \alpha_1 x - \beta_1 x^2 - \gamma_1 xy \tag{3}$$

$$F_2 = \alpha_2 x - \beta_2 y^2 - \gamma_2 xy \tag{4}$$

Differentiating (3) and (4) partially w.r. to x and y we get

$$J_{11} = \frac{\delta F_1}{\delta x} = \alpha_1 - 2\beta_1 x - \gamma_1 y \tag{5}$$

$$J_{12} = \frac{\delta F_1}{\delta y} = -\gamma_1 x \tag{6}$$

$$J_{21} = \frac{\delta F_2}{\delta x} = -\gamma_2 y \tag{7}$$

$$J_{22} = \frac{\delta F_2}{\delta y} = \alpha_2 - 2\beta_2 y - \gamma_2 x \tag{8}$$

With the following model parameters values, in Eli and Ekakaa, 2021.

$$\alpha_1 = 0.1, \alpha_2 = 0.08, \beta_1 = 0.0014, \beta_2 = 0.001, \gamma_1 = 0.0012, \gamma_2 = 0.0009,$$

3.0 Results and Discussion

3.1 Results

Given the characteristics equation

$$|J - \lambda I|$$

Applying the ODE 45 numerical method where I is the identity matrix of order 2×2 matrix and λ is a scalar.

Here a numerical method is applied to obtain the eigenvalues using the Matlab ODE 45. The possible steady state is the point $(x, y) = (0, 0)$, applying the numerical method, we have obtained two eigenvalues $\lambda_1 = 0.0800$, $\lambda_2 = 0.100$.

To check for the stability type of this steady-state solution for this interacting problem, a Jacobian Matrix was defined from which two eigenvalues were calculated, the characteristics equation being $|J - \lambda I| = 0$.

Table 1: Quantifying the effect of decreasing the model parameter $\alpha_1 = 0.1$ and $\alpha_2 = 0.08$ on the type of stability of a trivial steady state solution using Matlab Algorithm.

| α_1 & α_2 Variation | α_1 | α_2 | x | y | λ_1 | λ_2 | TOS |
|--------------------------------------|------------|------------|---|---|-------------|-------------|----------|
| 100% | 0.1000 | 0.0800 | 0 | 0 | 0.0800 | 0.1000 | Unstable |
| 10% | 0.0100 | 0.0080 | 0 | 0 | 0.0080 | 0.0100 | Unstable |
| 20% | 0.0200 | 0.0160 | 0 | 0 | 0.0160 | 0.0200 | Unstable |
| 90% | 0.0900 | 0.0720 | 0 | 0 | 0.0720 | 0.0900 | Unstable |
| 99% | 0.0990 | 0.0792 | 0 | 0 | 0.0792 | 0.0990 | Unstable |
| 101% | 0.1010 | 0.0808 | 0 | 0 | 0.0808 | 0.1010 | Unstable |
| 110% | 0.1100 | 0.0880 | 0 | 0 | 0.880 | 0.1100 | Unstable |
| 120% | 0.1200 | 0.0960 | 0 | 0 | 0.0960 | 0.1200 | Unstable |
| 130% | 0.1300 | 0.1040 | 0 | 0 | 0.1040 | 0.1300 | Unstable |
| 140% | 0.1400 | 0.1120 | 0 | 0 | 0.1120 | 0.1400 | Unstable |
| 150% | 0.1500 | 0.1200 | 0 | 0 | 0.1200 | 0.1500 | Unstable |

**TOS = Type of Stability

Table 2: Quantifying the effect of decreasing the α_1 and α_2 together by 10% on the biodiversity loss: using ODE44

| LGS | $x(old)$ | $x(new)$ | BL | $y(old)$ | $y(new)$ | BL |
|-----|----------|----------|---------|----------|----------|---------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 16.6859 | 44.3329 | 38.3574 | 27.5848 | 28.0849 |
| 15 | 41.5737 | 14.1675 | 65.9220 | 48.3189 | 25.7551 | 46.6976 |



| | | | | | | |
|-----|---------|---------|---------|---------|---------|---------|
| 22 | 52.8941 | 12.1988 | 76.9373 | 58.4827 | 24.3292 | 58.3993 |
| 29 | 62.0864 | 10.6148 | 82.9032 | 67.4179 | 23.1850 | 65.6100 |
| 36 | 68.1802 | 9.3161 | 86.3360 | 74.2993 | 22.2493 | 70.0545 |
| 43 | 71.2927 | 8.2356 | 88.4482 | 79.1076 | 21.4726 | 72.8564 |
| 50 | 72.1270 | 7.3237 | 89.8462 | 82.2717 | 20.8186 | 74.6953 |
| 57 | 71.4373 | 6.5459 | 90.8369 | 84.2948 | 20.2619 | 75.9630 |
| 64 | 69.7796 | 5.8762 | 91.5789 | 85.5525 | 19.7835 | 76.8756 |
| 71 | 67.5880 | 5.2947 | 92.1662 | 86.3374 | 19.3689 | 77.5661 |
| 78 | 65.1155 | 4.7863 | 92.6495 | 86.8294 | 19.0070 | 78.1100 |
| 85 | 62.5255 | 4.3391 | 93.0603 | 87.1433 | 18.6891 | 78.5536 |
| 92 | 59.9118 | 3.9436 | 93.4177 | 87.3469 | 18.4084 | 78.9250 |
| 99 | 57.3254 | 3.5920 | 93.7339 | 87.4793 | 18.1592 | 79.2417 |
| 106 | 54.8017 | 3.2783 | 94.0178 | 87.5682 | 54.8017 | 37.4183 |
| 113 | 52.3567 | 2.9973 | 94.2752 | 87.6282 | 17.7385 | 79.7571 |
| 120 | 49.9996 | 2.7447 | 94.5106 | 87.6693 | 17.5601 | 79.9701 |
| 127 | 47.7351 | 2.5170 | 94.7272 | 87.6981 | 17.3994 | 80.1599 |
| 134 | 45.5637 | 2.3111 | 94.9278 | 87.7184 | 17.2543 | 80.3299 |
| 141 | 43.4849 | 2.1245 | 95.1144 | 87.7329 | 17.1229 | 80.4830 |

Table 3: Quantifying the effect of decreasing the α_1 and α_2 together by 50% on the biodiversity loss: using ODE45

| LGS | $x(old)$ | $x(new)$ | BL | $y(old)$ | $y(new)$ | BL |
|-----|----------|----------|---------|----------|----------|---------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 21.6928 | 27.6293 | 38.3574 | 31.6419 | 17.5077 |
| 15 | 41.5737 | 23.2156 | 44.1580 | 48.3189 | 33.1851 | 31.3207 |
| 22 | 52.8941 | 24.5379 | 53.6093 | 58.4827 | 34.6034 | 40.8315 |
| 29 | 62.0864 | 25.6433 | 58.6973 | 67.4179 | 35.8806 | 46.7788 |
| 36 | 68.1802 | 26.5290 | 61.0899 | 74.2993 | 37.0108 | 50.1868 |
| 43 | 71.2927 | 27.2026 | 61.8439 | 79.1076 | 37.9959 | 51.9694 |
| 50 | 72.1270 | 27.6795 | 61.6239 | 82.2717 | 38.8437 | 52.7861 |
| 57 | 71.4373 | 27.9800 | 60.8328 | 84.2948 | 39.5661 | 53.0623 |
| 64 | 69.7796 | 28.1261 | 59.6930 | 85.5525 | 40.1766 | 53.0387 |
| 71 | 67.5880 | 28.1402 | 58.3652 | 86.3374 | 40.6895 | 52.8715 |
| 78 | 65.1155 | 28.0433 | 56.9330 | 86.8294 | 41.1184 | 52.6446 |
| 85 | 62.5255 | 27.8548 | 55.4504 | 87.1433 | 41.4761 | 52.4047 |
| 92 | 59.9118 | 27.5918 | 53.9459 | 87.3469 | 41.7738 | 52.1748 |
| 99 | 57.3254 | 27.2689 | 52.4313 | 87.4793 | 42.0213 | 51.9643 |
| 106 | 54.8017 | 26.8987 | 50.9163 | 87.5682 | 54.8017 | 37.4183 |
| 113 | 52.3567 | 26.4916 | 49.4017 | 87.6282 | 42.3984 | 51.6156 |
| 120 | 49.9996 | 26.0564 | 47.8868 | 87.6693 | 42.5410 | 51.4756 |
| 127 | 47.7351 | 25.6003 | 46.3700 | 87.6981 | 42.6600 | 51.3558 |
| 134 | 45.5637 | 25.1292 | 44.8482 | 87.7184 | 42.7595 | 51.2537 |
| 141 | 43.4849 | 24.6479 | 43.3185 | 87.7329 | 42.8428 | 51.1668 |



Table 4: Quantifying the effect of decreasing the α_1 and α_2 together by 90% on the biodiversity loss: using ODE45.

| LGS | $x(old)$ | $x(new)$ | BL | $y(old)$ | $y(old)$ | BL |
|-----|----------|----------|---------|----------|----------|---------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 28.1100 | 6.2204 | 38.3574 | 36.8449 | 3.9431 |
| 15 | 41.5737 | 37.1350 | 10.6768 | 48.3189 | 44.6518 | 7.5895 |
| 22 | 52.8941 | 45.8815 | 13.2578 | 58.4827 | 52.5360 | 10.1684 |
| 29 | 62.0864 | 53.2195 | 14.2816 | 67.4179 | 59.6228 | 11.5624 |
| 36 | 68.1802 | 58.4830 | 14.2230 | 74.2993 | 65.3549 | 12.0384 |
| 43 | 71.2927 | 61.6247 | 13.5611 | 79.1076 | 69.6293 | 11.9815 |
| 50 | 72.1270 | 62.9925 | 12.6645 | 82.2717 | 72.6429 | 11.7037 |
| 57 | 71.4373 | 63.0556 | 11.7330 | 84.2948 | 74.7001 | 11.3824 |
| 64 | 69.7796 | 62.2176 | 10.8369 | 85.5525 | 76.0697 | 11.0842 |
| 71 | 67.5880 | 60.8160 | 10.0195 | 86.3374 | 76.9785 | 10.8399 |
| 78 | 65.1155 | 59.0715 | 9.2819 | 86.8294 | 77.5802 | 10.6522 |
| 85 | 62.5255 | 57.1376 | 8.6170 | 87.1433 | 77.9810 | 10.5140 |
| 92 | 59.9118 | 55.1155 | 8.0056 | 87.3469 | 78.2521 | 10.4123 |
| 99 | 57.3254 | 53.0630 | 7.4354 | 87.4793 | 78.4356 | 10.3381 |
| 106 | 54.8017 | 51.0219 | 6.8972 | 87.5682 | 54.8017 | 37.4183 |
| 113 | 52.3567 | 49.0158 | 6.3812 | 87.6282 | 78.6506 | 10.2451 |
| 120 | 49.9996 | 47.0587 | 5.8819 | 87.6693 | 78.7127 | 10.2164 |
| 127 | 47.7351 | 45.1601 | 5.3944 | 87.6981 | 78.7572 | 10.1951 |
| 134 | 45.5637 | 43.3242 | 4.9152 | 87.7184 | 78.7892 | 10.1794 |
| 141 | 43.4849 | 41.5533 | 4.4419 | 87.7329 | 78.8126 | 10.1676 |

Table 5: Quantifying the effect of decreasing the α_1 and α_2 together by 95% on the biodiversity loss: using ODE45

| LGS | $x(old)$ | $x(new)$ | BL | $y(old)$ | $y(old)$ | BL |
|-----|----------|----------|--------|----------|----------|---------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 29.0281 | 3.1573 | 38.3574 | 37.5897 | 2.0015 |
| 15 | 41.5737 | 39.3015 | 5.4656 | 48.3189 | 46.4410 | 3.8865 |
| 22 | 52.8941 | 49.3011 | 6.7928 | 58.4827 | 55.4322 | 5.2162 |
| 29 | 62.0864 | 57.5671 | 7.2790 | 67.4179 | 63.4338 | 5.9096 |
| 36 | 68.1802 | 63.2806 | 7.1863 | 74.2993 | 69.7562 | 6.1146 |
| 43 | 71.2927 | 66.4562 | 6.7840 | 79.1076 | 74.3244 | 6.0464 |
| 50 | 72.1270 | 67.5988 | 6.2781 | 82.2717 | 77.4369 | 5.8767 |
| 57 | 71.4373 | 67.3144 | 5.7714 | 84.2948 | 79.4929 | 5.6966 |
| 64 | 69.7796 | 66.0850 | 5.2946 | 85.5525 | 80.8160 | 5.5364 |
| 71 | 67.5880 | 64.2990 | 4.8662 | 86.3374 | 81.6674 | 5.4090 |
| 78 | 65.1155 | 62.1954 | 4.4846 | 86.8294 | 82.2157 | 5.3136 |
| 85 | 62.5255 | 59.9341 | 4.1446 | 87.1433 | 82.5721 | 5.2456 |
| 92 | 59.9118 | 57.6161 | 3.8318 | 87.3469 | 82.8084 | 5.1960 |
| 99 | 57.3254 | 55.2956 | 3.5407 | 87.4793 | 82.9650 | 5.1604 |
| 106 | 54.8017 | 53.0116 | 3.2665 | 87.5682 | 54.8017 | 37.4183 |



| | | | | | | |
|-----|---------|---------|--------|---------|---------|--------|
| 113 | 52.3567 | 50.7843 | 3.0033 | 87.6282 | 83.1445 | 5.1167 |
| 120 | 49.9996 | 48.6252 | 2.7488 | 87.6693 | 83.1952 | 5.1035 |
| 127 | 47.7351 | 46.5415 | 2.5003 | 87.6981 | 83.2310 | 5.0937 |
| 134 | 45.5637 | 44.5357 | 2.2562 | 87.7184 | 83.2565 | 5.0866 |
| 141 | 43.4849 | 42.6085 | 2.0152 | 87.7329 | 83.2749 | 5.0813 |

Table 6: Quantifying the effect of decreasing the α_1 and α_2 together by 99.9% on the biodiversity loss: using ODE45

| LGS | $x(old)$ | $x(new)$ | BL | $y(old)$ | $y(old)$ | BL |
|-----|----------|----------|--------|----------|----------|---------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 29.9553 | 0.0641 | 38.3574 | 38.3418 | 0.0406 |
| 15 | 41.5737 | 41.5274 | 0.1115 | 48.3189 | 48.2806 | 0.0793 |
| 22 | 52.8941 | 52.8204 | 0.1393 | 58.4827 | 58.4201 | 0.1071 |
| 29 | 62.0864 | 61.9945 | 0.1480 | 67.4179 | 67.3367 | 0.1205 |
| 36 | 68.1802 | 68.0814 | 0.1449 | 74.2993 | 74.2072 | 0.1240 |
| 43 | 71.2927 | 71.1963 | 0.1352 | 79.1076 | 79.0113 | 0.1217 |
| 50 | 72.1270 | 72.0373 | 0.1244 | 82.2717 | 82.1747 | 0.1179 |
| 57 | 71.4373 | 71.3562 | 0.1135 | 84.2948 | 84.1987 | 0.1140 |
| 64 | 69.7796 | 69.7075 | 0.1034 | 85.5525 | 85.4580 | 0.1105 |
| 71 | 67.5880 | 67.5241 | 0.0946 | 86.3374 | 86.2442 | 0.1079 |
| 78 | 65.1155 | 65.0591 | 0.0867 | 86.8294 | 86.7374 | 0.1060 |
| 85 | 62.5255 | 62.4756 | 0.0798 | 87.1433 | 87.0520 | 0.1047 |
| 92 | 59.9118 | 59.8678 | 0.0734 | 87.3469 | 87.2563 | 0.1037 |
| 99 | 57.3254 | 57.2867 | 0.0675 | 87.4793 | 87.3892 | 0.1031 |
| 106 | 54.8017 | 54.7678 | 0.0619 | 87.5682 | 54.8017 | 37.4183 |
| 113 | 52.3567 | 52.3271 | 0.0566 | 87.6282 | 87.5387 | 0.1022 |
| 120 | 49.9996 | 49.9739 | 0.0514 | 87.6693 | 87.5799 | 0.1020 |
| 127 | 47.7351 | 47.7130 | 0.0464 | 87.6981 | 87.6088 | 0.1018 |
| 134 | 45.5637 | 45.5449 | 0.0414 | 87.7184 | 87.6292 | 0.1017 |
| 141 | 43.4849 | 43.4690 | 0.0365 | 87.7329 | 87.6438 | 0.1016 |

Table 7: Quantifying the effect of increasing the α_1 and α_2 together by 101% on the biodiversity gain: using ODE45

| LGS | $x(old)$ | $x(new)$ | BG | $y(old)$ | $y(old)$ | BG |
|-----|----------|----------|--------|----------|----------|--------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 30.1671 | 0.6427 | 38.3574 | 38.5137 | 0.4074 |
| 15 | 41.5737 | 42.0397 | 1.1209 | 48.3189 | 48.7043 | 0.7976 |
| 22 | 52.8941 | 53.6346 | 1.4000 | 58.4827 | 59.1123 | 1.0765 |
| 29 | 62.0864 | 63.0087 | 1.4856 | 67.4179 | 68.2341 | 1.2106 |
| 36 | 68.1802 | 69.1698 | 1.4514 | 74.2993 | 75.2232 | 1.2435 |
| 43 | 71.2927 | 72.2561 | 1.3513 | 79.1076 | 80.0715 | 1.2185 |
| 50 | 72.1270 | 73.0219 | 1.2408 | 82.2717 | 83.2423 | 1.1797 |
| 57 | 71.4373 | 72.2448 | 1.1304 | 84.2948 | 85.2556 | 1.1398 |
| 64 | 69.7796 | 70.4971 | 1.0283 | 85.5525 | 86.4979 | 1.1050 |
| 71 | 67.5880 | 68.2228 | 0.9393 | 86.3374 | 87.2689 | 1.0789 |



| | | | | | | |
|-----|---------|---------|--------|---------|---------|---------|
| 78 | 65.1155 | 65.6757 | 0.8602 | 86.8294 | 87.7495 | 1.0597 |
| 85 | 62.5255 | 63.0202 | 0.7912 | 87.1433 | 88.0553 | 1.0466 |
| 92 | 59.9118 | 60.3474 | 0.7270 | 87.3469 | 88.2527 | 1.0370 |
| 99 | 57.3254 | 57.7081 | 0.6676 | 87.4793 | 88.3806 | 1.0302 |
| 106 | 54.8017 | 55.1370 | 0.6118 | 87.5682 | 54.8017 | 37.4183 |
| 113 | 52.3567 | 52.6489 | 0.5580 | 87.6282 | 88.5239 | 1.0222 |
| 120 | 49.9996 | 50.2527 | 0.5062 | 87.6693 | 88.5633 | 1.0198 |
| 127 | 47.7351 | 47.9525 | 0.4555 | 87.6981 | 88.5909 | 1.0180 |
| 134 | 45.5637 | 45.7486 | 0.4058 | 87.7184 | 88.6102 | 1.0167 |
| 141 | 43.4849 | 43.6400 | 0.3567 | 87.7329 | 88.6241 | 1.0158 |

Table 8: Quantifying the effect of increasing the α_1 and α_2 together by 105% on the biodiversity gain: using ODE45

| LGS | $x(old)$ | $x(new)$ | BG | $y(old)$ | $y(old)$ | BG |
|-----|----------|----------|--------|----------|----------|----------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 30.9493 | 3.2522 | 38.3574 | 39.1483 | 2.0619 |
| 15 | 41.5737 | 43.9474 | 5.7095 | 48.3189 | 50.2827 | 4.0642 |
| 22 | 52.8941 | 56.6660 | 7.1310 | 58.4827 | 61.6931 | 5.4894 |
| 29 | 62.0864 | 66.7579 | 7.5242 | 67.4179 | 71.5625 | 6.1475 |
| 36 | 68.1802 | 73.1522 | 7.2924 | 74.2993 | 78.9642 | 6.2786 |
| 43 | 71.2927 | 76.0943 | 6.7351 | 79.1076 | 83.9497 | 6.1210 |
| 50 | 72.1270 | 76.5603 | 6.1465 | 82.2717 | 87.1326 | 5.9084 |
| 57 | 71.4373 | 75.4136 | 5.5662 | 84.2948 | 89.0967 | 5.6965 |
| 64 | 69.7796 | 73.2958 | 5.0391 | 85.5525 | 90.2723 | 5.5168 |
| 71 | 67.5880 | 70.6855 | 4.5829 | 86.3374 | 90.9860 | 5.3843 |
| 78 | 65.1155 | 67.8394 | 4.1832 | 86.8294 | 91.4221 | 5.2893 |
| 85 | 62.5255 | 64.9220 | 3.8329 | 87.1433 | 91.6966 | 5.2251 |
| 92 | 59.9118 | 62.0136 | 3.5081 | 87.3469 | 91.8698 | 5.1781 |
| 99 | 57.3254 | 59.1647 | 3.2087 | 87.4793 | 91.9810 | 5.1460 |
| 106 | 54.8017 | 56.4052 | 2.9260 | 87.5682 | 54.8017 | -37.4183 |
| 113 | 52.3567 | 53.7464 | 2.6542 | 87.6282 | 92.1037 | 5.1074 |
| 120 | 49.9996 | 51.1957 | 2.3921 | 87.6693 | 92.1370 | 5.0961 |
| 127 | 47.7351 | 48.7547 | 2.1360 | 87.6981 | 92.1601 | 5.0879 |
| 134 | 45.5637 | 46.4225 | 1.8848 | 87.7184 | 92.1761 | 5.0819 |
| 141 | 43.4849 | 44.1969 | 1.6375 | 87.7329 | 92.1876 | 5.0776 |

Table 9: Quantifying the effect of increasing the α_1 and α_2 together by 110% on the biodiversity gain: using ODE45

| LGS | $x(old)$ | $x(new)$ | BG | $y(old)$ | $y(old)$ | BG |
|-----|----------|----------|---------|----------|----------|---------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 31.9537 | 6.6031 | 38.3574 | 39.9633 | 4.1867 |
| 15 | 41.5737 | 46.4320 | 11.6859 | 48.3189 | 52.3401 | 8.3221 |
| 22 | 52.8941 | 60.6086 | 14.5848 | 58.4827 | 65.0583 | 11.2437 |
| 29 | 62.0864 | 71.5689 | 15.2732 | 67.4179 | 75.8600 | 12.5219 |



| | | | | | | |
|-----|---------|---------|---------|---------|---------|----------|
| 36 | 68.1802 | 78.1682 | 14.6493 | 74.2993 | 83.7312 | 12.6944 |
| 43 | 71.2927 | 80.8459 | 13.3998 | 79.1076 | 88.8396 | 12.3022 |
| 50 | 72.1270 | 80.8818 | 12.1381 | 82.2717 | 92.0076 | 11.8339 |
| 57 | 71.4373 | 79.2319 | 10.9110 | 84.2948 | 93.8886 | 11.3812 |
| 64 | 69.7796 | 76.6322 | 9.8204 | 85.5525 | 94.9719 | 11.0100 |
| 71 | 67.5880 | 73.5943 | 8.8866 | 86.3374 | 95.6123 | 10.7427 |
| 78 | 65.1155 | 70.3762 | 8.0790 | 86.8294 | 95.9960 | 10.5570 |
| 85 | 62.5255 | 67.1317 | 7.3669 | 87.1433 | 96.2333 | 10.4311 |
| 92 | 59.9118 | 63.9315 | 6.7093 | 87.3469 | 96.3790 | 10.3405 |
| 99 | 57.3254 | 60.8249 | 6.1047 | 87.4793 | 96.4723 | 10.2801 |
| 106 | 54.8017 | 57.8332 | 5.5318 | 87.5682 | 54.8017 | -37.4183 |
| 113 | 52.3567 | 54.9652 | 4.9821 | 87.6282 | 96.5728 | 10.2073 |
| 120 | 49.9996 | 52.2255 | 4.4517 | 87.6693 | 96.5997 | 10.1864 |
| 127 | 47.7351 | 49.6129 | 3.9338 | 87.6981 | 96.6181 | 10.1712 |
| 134 | 45.5637 | 47.1249 | 3.4263 | 87.7184 | 96.6308 | 10.1603 |
| 141 | 43.4849 | 44.7577 | 2.9270 | 87.7329 | 96.6398 | 10.1523 |

Table 10: Quantifying the effect of increasing the α_1 and α_2 together by 120% on the biodiversity gain: using ODE45

| LGS | $x(old)$ | $x(new)$ | BG | $y(old)$ | $y(old)$ | BG |
|-----|----------|----------|---------|----------|----------|----------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 34.0545 | 13.6117 | 38.3574 | 41.6683 | 8.6317 |
| 15 | 41.5737 | 51.7457 | 24.4674 | 48.3189 | 56.7464 | 17.4413 |
| 22 | 52.8941 | 68.9815 | 30.4144 | 58.4827 | 72.2405 | 23.5245 |
| 29 | 62.0864 | 81.5480 | 31.3461 | 67.4179 | 84.8743 | 25.8928 |
| 36 | 68.1802 | 88.2206 | 29.3932 | 74.2993 | 93.4924 | 25.8321 |
| 43 | 71.2927 | 90.1036 | 26.3854 | 79.1076 | 98.6929 | 24.7579 |
| 50 | 72.1270 | 88.9970 | 23.3894 | 82.2717 | 101.6643 | 23.5714 |
| 57 | 71.4373 | 86.2511 | 20.7367 | 84.2948 | 103.3194 | 22.5692 |
| 64 | 69.7796 | 82.7078 | 18.5273 | 85.5525 | 104.2420 | 21.8457 |
| 71 | 67.5880 | 78.8502 | 16.6630 | 86.3374 | 104.7697 | 21.3492 |
| 78 | 65.1155 | 74.9115 | 15.0440 | 86.8294 | 105.0717 | 21.0094 |
| 85 | 62.5255 | 71.0265 | 13.5961 | 87.1433 | 105.2484 | 20.7763 |
| 92 | 59.9118 | 67.2684 | 12.2791 | 87.3469 | 105.3576 | 20.6196 |
| 99 | 57.3254 | 63.6609 | 11.0520 | 87.4793 | 105.4239 | 20.5130 |
| 106 | 54.8017 | 60.2206 | 9.8882 | 87.5682 | 54.8017 | -37.4183 |
| 113 | 52.3567 | 56.9501 | 8.7732 | 87.6282 | 105.4939 | 20.3880 |
| 120 | 49.9996 | 53.8467 | 7.6943 | 87.6693 | 105.5120 | 20.3522 |
| 127 | 47.7351 | 50.9063 | 6.6434 | 87.6981 | 105.5242 | 20.3267 |
| 134 | 45.5637 | 48.1223 | 5.6154 | 87.7184 | 105.5326 | 20.3085 |
| 141 | 43.4849 | 45.4878 | 4.6062 | 87.7329 | 105.5385 | 20.2952 |



Table 11: Quantifying the effect of increasing the α_1 and α_2 together by 130% on the biodiversity gain: using ODE45

| LGS | $x(old)$ | $x(new)$ | BG | $y(old)$ | $y(old)$ | BG |
|-----|----------|----------|---------|----------|----------|----------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 36.2833 | 21.0474 | 38.3574 | 43.4777 | 13.3490 |
| 15 | 41.5737 | 57.5359 | 38.3950 | 48.3189 | 61.5577 | 27.3987 |
| 22 | 52.8941 | 77.9715 | 47.4106 | 58.4827 | 80.0090 | 36.8080 |
| 29 | 62.0864 | 91.8830 | 47.9922 | 67.4179 | 94.3666 | 39.9726 |
| 36 | 68.1802 | 98.1581 | 43.9686 | 74.2993 | 103.4549 | 39.2407 |
| 43 | 71.2927 | 98.9256 | 38.7597 | 79.1076 | 108.5634 | 37.2352 |
| 50 | 72.1270 | 96.5533 | 33.8657 | 82.2717 | 111.2748 | 35.2528 |
| 57 | 71.4373 | 92.6643 | 29.7141 | 84.2948 | 112.6998 | 33.6973 |
| 64 | 69.7796 | 88.1519 | 26.3290 | 85.5525 | 113.4614 | 32.6219 |
| 71 | 67.5880 | 83.4533 | 23.4736 | 86.3374 | 113.8720 | 31.8919 |
| 78 | 65.1155 | 78.7983 | 21.0130 | 86.8294 | 114.1003 | 31.4074 |
| 85 | 62.5255 | 74.2973 | 18.8272 | 87.1433 | 114.2332 | 31.0867 |
| 92 | 59.9118 | 69.9885 | 16.8192 | 87.3469 | 114.3090 | 30.8677 |
| 99 | 57.3254 | 65.8988 | 14.9558 | 87.4793 | 114.3561 | 30.7236 |
| 106 | 54.8017 | 62.0287 | 13.1875 | 87.5682 | 54.8017 | -37.4183 |
| 113 | 52.3567 | 58.3740 | 11.4929 | 87.6282 | 114.4039 | 30.5559 |
| 120 | 49.9996 | 54.9281 | 9.8570 | 87.6693 | 114.4161 | 30.5087 |
| 127 | 47.7351 | 51.6811 | 8.2666 | 87.6981 | 114.4242 | 30.4751 |
| 134 | 45.5637 | 48.6234 | 6.7152 | 87.7184 | 114.4296 | 30.4512 |
| 141 | 43.4849 | 45.7448 | 5.1972 | 87.7329 | 114.4335 | 30.4339 |

Table 12: Quantifying the effect of increasing the α_1 and α_2 together by 140% on the biodiversity gain: using ODE45

| LGS | $x(old)$ | $x(new)$ | BG | $y(old)$ | $y(old)$ | BG |
|-----|----------|----------|---------|----------|----------|----------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 38.6468 | 28.9324 | 38.3574 | 45.3971 | 18.3528 |
| 15 | 41.5737 | 63.8197 | 53.5097 | 48.3189 | 66.7910 | 38.2294 |
| 22 | 52.8941 | 87.5347 | 65.4906 | 58.4827 | 88.3405 | 51.0540 |
| 29 | 62.0864 | 102.4011 | 64.9333 | 67.4179 | 104.2176 | 54.5844 |
| 36 | 68.1802 | 107.8518 | 58.1863 | 74.2993 | 113.5240 | 52.7929 |
| 43 | 71.2927 | 107.2154 | 50.3875 | 79.1076 | 118.3732 | 49.6358 |
| 50 | 72.1270 | 103.4914 | 43.4850 | 82.2717 | 120.7869 | 46.8146 |
| 57 | 71.4373 | 98.4433 | 37.8038 | 84.2948 | 121.9870 | 44.7147 |
| 64 | 69.7796 | 92.9624 | 33.2229 | 85.5525 | 122.5996 | 43.3033 |
| 71 | 67.5880 | 87.4485 | 29.3847 | 86.3374 | 122.9161 | 42.3672 |
| 78 | 65.1155 | 82.1058 | 26.0926 | 86.8294 | 123.0914 | 41.7623 |
| 85 | 62.5255 | 76.9963 | 23.1438 | 87.1433 | 123.1846 | 41.3587 |
| 92 | 59.9118 | 72.1645 | 20.4513 | 87.3469 | 123.2402 | 41.0927 |
| 99 | 57.3254 | 67.6118 | 17.9439 | 87.4793 | 123.2734 | 40.9172 |
| 106 | 54.8017 | 63.3317 | 15.5651 | 87.5682 | 54.8017 | -37.4183 |



| | | | | | | |
|-----|---------|---------|---------|---------|----------|---------|
| 113 | 52.3567 | 59.3151 | 13.2904 | 87.6282 | 123.3062 | 40.7151 |
| 120 | 49.9996 | 55.5484 | 11.0976 | 87.6693 | 123.3144 | 40.6586 |
| 127 | 47.7351 | 52.0179 | 8.9721 | 87.6981 | 123.3198 | 40.6186 |
| 134 | 45.5637 | 48.7101 | 6.9054 | 87.7184 | 123.3235 | 40.5903 |
| 141 | 43.4849 | 45.6114 | 4.8902 | 87.7329 | 123.3260 | 40.5699 |

Table 13: Quantifying the effect of increasing the α_1 and α_2 together by 150% on the biodiversity gain: using ODE45

| LGS | $x(old)$ | $x(new)$ | BG | $y(old)$ | $y(new)$ | BG |
|-----|----------|----------|---------|----------|----------|----------|
| 1 | 20.0000 | 20.0000 | 0 | 30.0000 | 30.0000 | 0 |
| 8 | 29.9745 | 41.1519 | 37.2897 | 38.3574 | 47.4320 | 23.6581 |
| 15 | 41.5737 | 70.6097 | 69.8421 | 48.3189 | 72.4603 | 49.9626 |
| 22 | 52.8941 | 97.6041 | 84.5274 | 58.4827 | 97.1934 | 66.1917 |
| 29 | 62.0864 | 112.9470 | 81.9192 | 67.4179 | 114.3188 | 69.5674 |
| 36 | 68.1802 | 117.1731 | 71.8578 | 74.2993 | 123.6055 | 66.3616 |
| 43 | 71.2927 | 114.9486 | 61.2347 | 79.1076 | 128.0916 | 61.9208 |
| 50 | 72.1270 | 109.8164 | 52.2542 | 82.2717 | 130.1902 | 58.2442 |
| 57 | 71.4373 | 103.6292 | 45.0631 | 84.2948 | 131.1913 | 55.6339 |
| 64 | 69.7796 | 97.1923 | 39.2846 | 85.5525 | 131.6714 | 53.9071 |
| 71 | 67.5880 | 90.9004 | 34.4920 | 86.3374 | 131.9211 | 52.7973 |
| 78 | 65.1155 | 84.8762 | 30.3471 | 86.8294 | 132.0471 | 52.0764 |
| 85 | 62.5255 | 79.1915 | 26.6547 | 87.1433 | 132.1175 | 51.6096 |
| 92 | 59.9118 | 73.8549 | 23.2728 | 87.3469 | 132.1578 | 51.3021 |
| 99 | 57.3254 | 68.8587 | 20.1192 | 87.4793 | 132.1805 | 51.0992 |
| 106 | 54.8017 | 64.1919 | 17.1349 | 87.5682 | 54.8017 | -37.4183 |
| 113 | 52.3567 | 59.8355 | 14.2843 | 87.6282 | 132.2037 | 50.8688 |
| 120 | 49.9996 | 55.7715 | 11.5438 | 87.6693 | 132.2093 | 50.8045 |
| 127 | 47.7351 | 51.9815 | 8.8958 | 87.6981 | 132.2130 | 50.7593 |
| 134 | 45.5637 | 48.4478 | 6.3297 | 87.7184 | 132.2155 | 50.7273 |
| 141 | 43.4849 | 45.1535 | 3.8374 | 87.7329 | 132.2172 | 50.7043 |

LGS = Length of growing season, $x(old)$ = Measures the biomass of yeast specie 1 when all model parameters are fixed at 100%, $y(new)$ = Measures the biomass of yeast specie 2 when α_1 and α_2 only are varied, BL = Biodiversity Loss, BG = Biodiversity Gain



4.2 Discussion of Results

Table 2 to Table 13 are the results of the effect of decreasing and increasing α_1 and α_2 together. Table 2 shows the extent of the percentage of biodiversity loss (BL) due to the low environmental perturbation value by 10% (0.10) concerning yeast species. A close look at the first-row result of example 1, we observed that the numerically simulated data of yeast species 1 and 2 when the model parameters are fixed at 100% and when α_1 and α_2 together are varied at 10% that is $x(\text{old})$ and $x(\text{new})$, $y(\text{old})$ and $y(\text{new})$ gives the same value of 20kg and 30kg which is consistent with the notion of population modelling and prediction which shows that the population size of the growing population does not change. Applying the method of ODE45 numerical simulation, we observed that in the same first row in example 1, the estimated population is approximately 0.00 because the $x(\text{old})$ and $x(\text{new})$, $y(\text{old})$ and $y(\text{new})$ are equal. Observing from example 2 to

example 21 of the yeast species 1 and 2 when model parameter values are fixed at 100% and α_1 and α_2 together only are varied at 10%, the solution map or solution trajectories follow a random pattern, from this prediction $x(\text{old})$ and $y(\text{old})$ in table 2 to table 3 remain the same because all the model parameters are fixed at 100% while $x(\text{new})$ and $y(\text{new})$ varies because of the variation of α_1 and α_2 together. Now the biomass of yeast species 1 and 2, $x(\text{new})$ and $y(\text{new})$ in the unit of the kilogram (kg) has lost biomass over that of $x(\text{old})$ and $y(\text{old})$ showing evidence of biodiversity loss. From Table 2 to Table 13, we varied α_1 and α_2 together called the environmental perturbation from 10% to 150% in (new) and (new) while in (old) and (old) all the model parameters are fixed at 100%. As we increased the decrease from 10% to 99.99%, the value of the first scenario is less the value of the second scenario and third is less than the fourth.

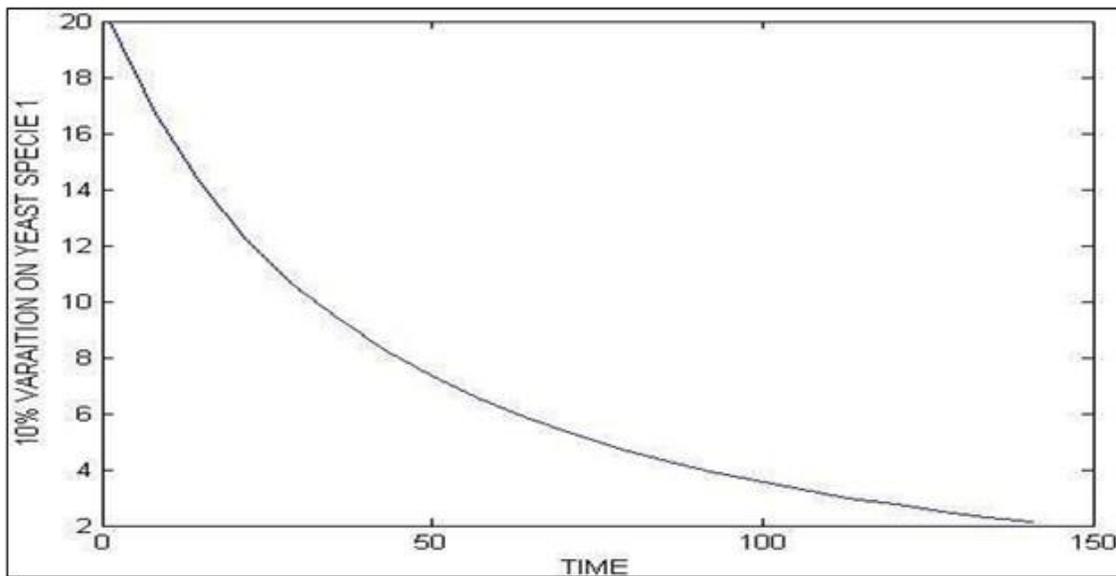


Fig. 1. Graphical Simulation on the effects of decreasing α_1 and α_2 together by 10% on biodiversity loss of yeast specie 1



As the percentage increased, there was an improved biomass of yeast species 1 and 2 called $x(\text{new})$ and $y(\text{new})$ which gets closer to the biomass of yeast species 1 and 2 at fixed model parameters called $x(\text{old})$ and $y(\text{old})$. The result obtained shows that as the percentage increases, the loss in biodiversity from 80 (approximately) to 0.00.

From example 1 to example 21, we consistently observed an increase on yeast species 1 and 2, and the remains after varying together α_1 and α_2 from 101% to 150% are greater than that is

when all the model parameters are fixed at 100%.

The values of the first scenario are less than the values of the second scenario and the second is less than the third and so on as we increased α_1 and α_2 together, the values get bigger which means the biomass of yeast species 1 and 2 in the unit of a kilogram (kg) has maintained an improved (new) and (new) over that of (old) and (old) showing evidence of biodiversity gain.

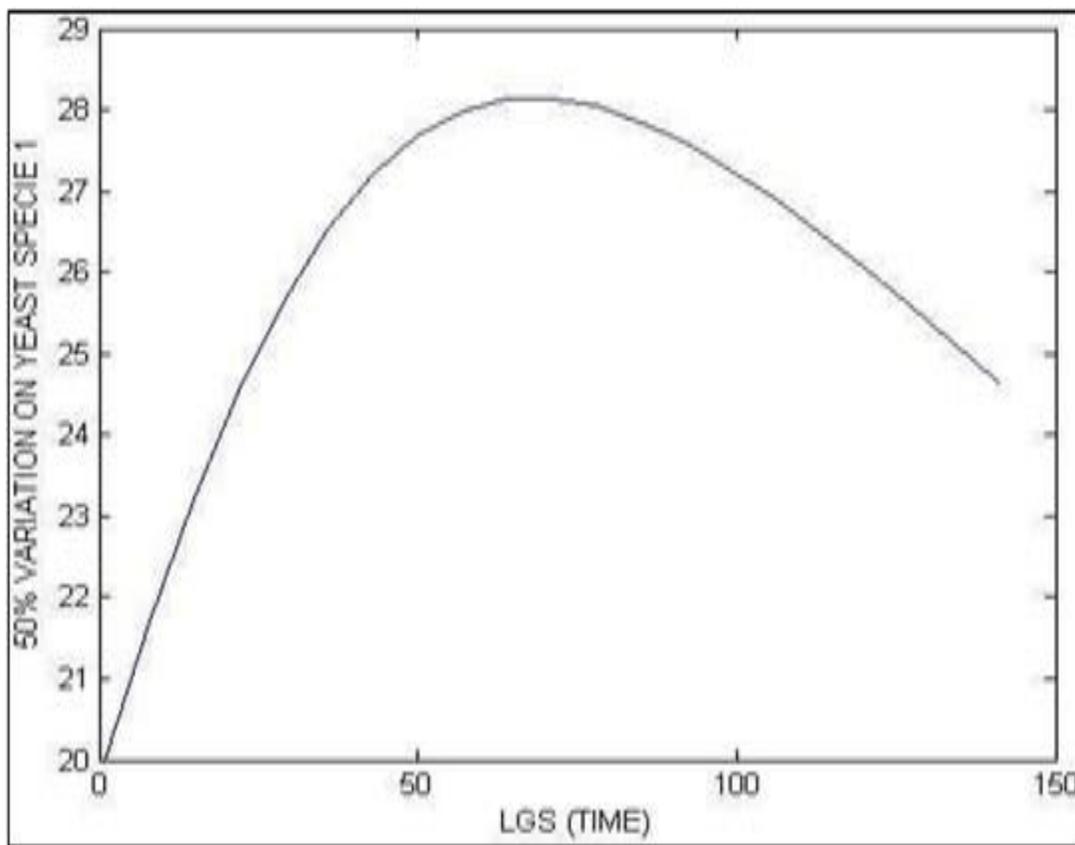


Fig 2. Graphical Simulation on the effects of decreasing α_1 and α_2 together by 50% on biodiversity loss of yeast specie 1.



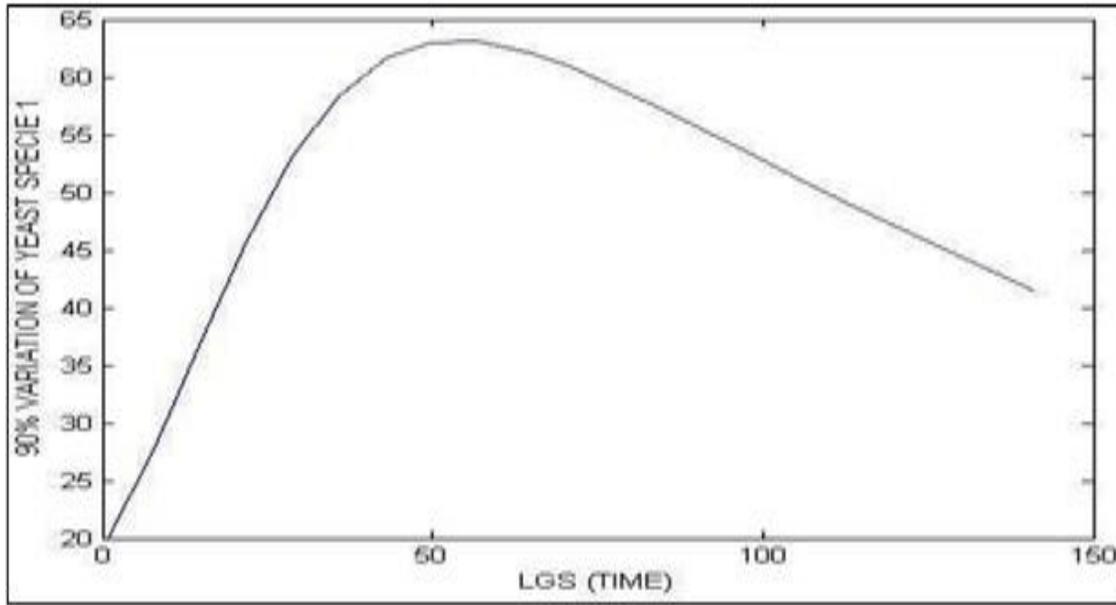


Fig. 3. Graphical Simulation on the effects of decreasing α_1 and α_2 together by 90% on biodiversity loss of yeast specie 1.

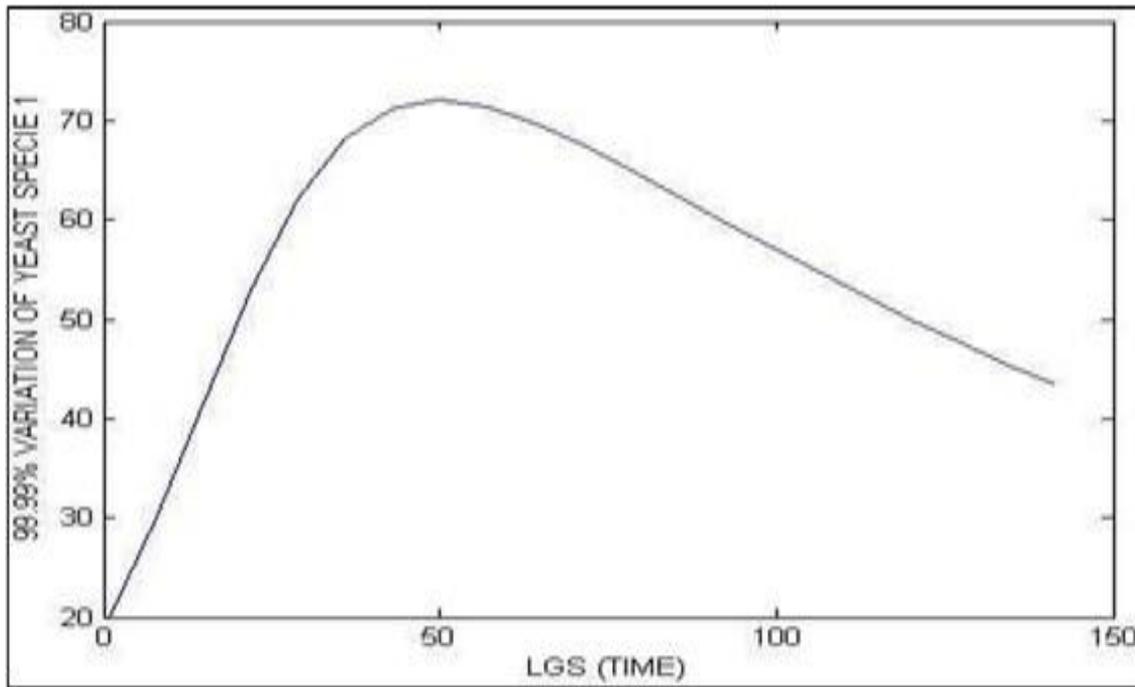


Fig. 4. Graphical Simulation on the effects of decreasing α_1 and α_2 together by 99.99% on biodiversity loss of yeast specie 1.



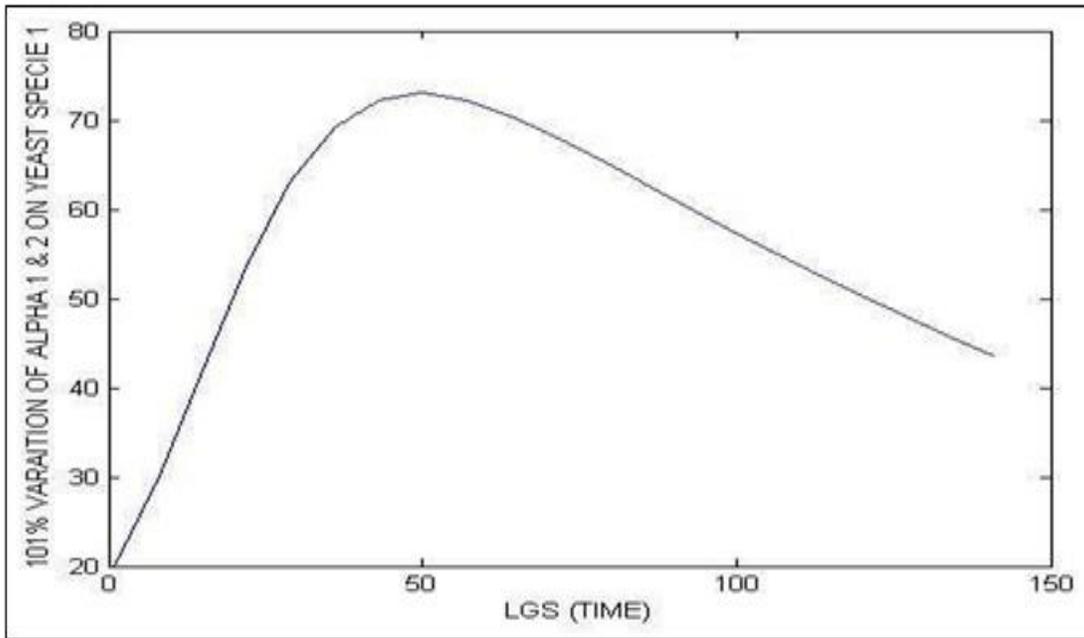


Fig. 5. Graphical Simulation on the effects of increasing α_1 and α_2 together by 101% on biodiversity loss of yeast specie 1.

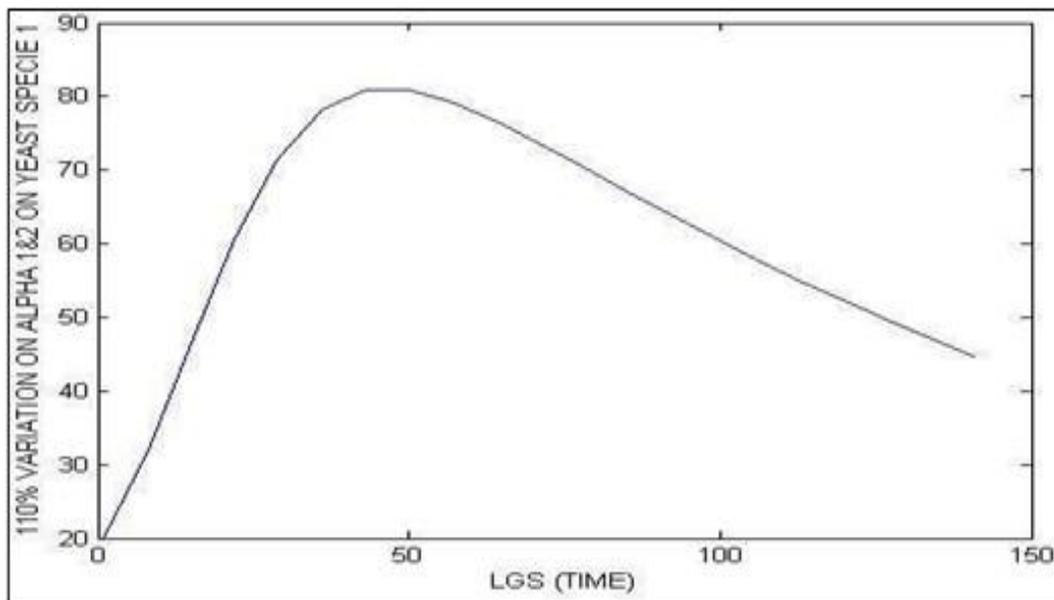


Fig. 6. Graphical Simulation on the effects of increasing α_1 and α_2 together by 110% on biodiversity loss of yeast specie 1.



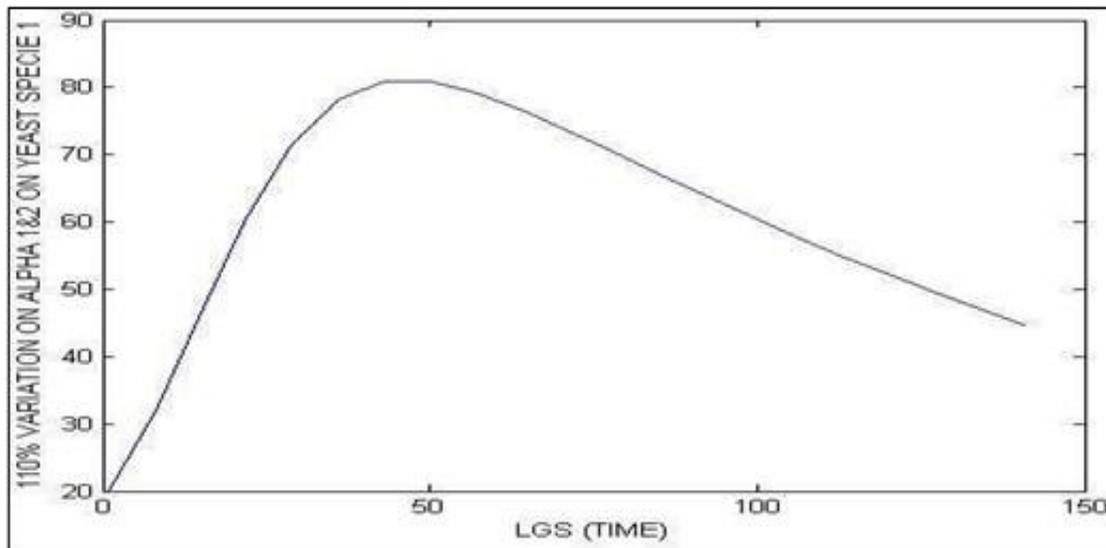


Fig. 7. Graphical Simulation on the effects of increasing α_1 and α_2 together by 150% on biodiversity loss of yeast specie 1

4.0 Conclusion

On the implementation of ODE45 computational approach, a region of instability was found. From example 1 to example 21, we consistently observed an increase in yeast species 1 and 2. The remains after varying α_1 and α_2 together from 101% to 150% are greater the remains when all model parameters are fixed at 100%. It was also found that the values of the first scenario are less than the values of the second scenario and the second is less than the third and so on as we increased α_1 and α_2 together, the values get bigger which means the biomass of yeast species 1 and 2 in the unit of a kilogram (Kg) has maintained an improved $x(\text{new})$ and $y(\text{new})$ over that of $x(\text{old})$ and $y(\text{old})$ showing evidence of biodiversity gain.

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- Declarations**
The authors declare that they have no conflict of interest.
- Data availability**
All data used in this study will be readily available to the public.
- Consent for publication**
Not Applicable
- Availability of data and materials**
The publisher has the right to make the data Public.
- Competing interests**
The authors declared no conflict of interest.
- Funding**
There is no source of external funding
- Authors' contributions**
Both authors contributed equally to the work

