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| Book Name: | [**Microbiology and Biotechnology Research: An Overview**](https://www.bookpi.org/bookstore/product/microbiology-and-biotechnology-research-an-overview-vol-1/) |
| Manuscript Number: | **Ms\_BPR\_4726** |
| Title of the Manuscript: | **DNA Tandem Repeats as Iterable Objects to Count Cell Divisions: A Computational Model** |
| Type of the Article | **Book Chapter** |

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**A research paper already published in a journal can be published as a Book Chapter in an expanded form with proper copyright approval.**

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**This chapter is an extended version of the article published by the same author(s) in the following journal.**

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| PART 1: Comments | | |
|  | Reviewer’s comment | Author’s Feedback *(Please correct the manuscript and highlight that part in the manuscript. It is mandatory that authors should write his/her feedback here)* |
| **Please write a few sentences regarding the importance of this manuscript for the scientific community. A minimum of 3-4 sentences may be required for this part.** | Yes. **This article is important for scientific community since this article focuses on on an issue that has not been adequately studied** |  |
| **Is the title of the article suitable?**  **(If not please suggest an alternative title)** | **Yes. It is clear, concise, and informative. It accurately reflects the content of the paper** |  |
| Is the abstract of the article comprehensive? Do you suggest the addition (or deletion) of some points in this section? Please write your suggestions here. | **Yes. The abstract is well-written and easy to read.** |  |
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| **Are the references sufficient and recent? If you have suggestions of additional references, please mention them in the review form.**  **-** | **Yes** |  |
| Is the language/English quality of the article suitable for scholarly communications? | **Yes** |  |
| Optional/General comments |  |  |

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| **PART 2:** | | |
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| **Are there ethical issues in this manuscript?** | *(If yes, Kindly please write down the ethical issues here in detail)* |  |

**Reviewer details:**

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