## Accepted Manuscript

Mathematical modeling and numerical simulation of the mitotic spindle orientation system

Bashar Ibrahim

 PII:
 S0025-5564(16)30334-0

 DOI:
 10.1016/j.mbs.2018.03.030

 Reference:
 MBS 8078

To appear in:

Mathematical Biosciences

Received date:22 November 2016Revised date:23 March 2018Accepted date:24 March 2018

Please cite this article as: Bashar Ibrahim, Mathematical modeling and numerical simulation of the mitotic spindle orientation system, *Mathematical Biosciences* (2018), doi: 10.1016/j.mbs.2018.03.030

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.



## Highlights

- A mathematical model was developed to describe the spindle position checkpoint activation and silencing in budding yeast.
- The numerical simulations of the nonlinear ordinary differential equations model reproduce the experimental features of SPOC mechanism.
- Bifurcation analysis reveals the orientation dependency on spindle pole bodies, and how this dependence is altered by parameter values.
- Partial differential equations based model and linear stability show the effects of diffusion coefficient on SPOC mechanism.

A CERTIN

Download English Version:

## https://daneshyari.com/en/article/8876950

Download Persian Version:

https://daneshyari.com/article/8876950

Daneshyari.com