

# Mathematical modeling *Pseudomonas* spp. growth and microflora composition variation in *Agaricus bisporus* fruiting bodies during chilled storage

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## Abstract

The objective of this study was to determine the dominant microorganism in *Agaricus bisporus* fruiting bodies (ABFB) and to develop kinetic models to describe its growth. The variation in microflora composition in ABFB stored at 4 °C during storage was studied using 16S rDNA sequence analyses. ABFB presented complex microbial communities at the initial stage of storage (day 0), with the dominant microorganism being *Pseudomonas* spp. along with small proportions of *Pedobacter*, *Sphingobacterium*, *Bacillus*, *Corynebacterium*, *Lactobacillus*, *Sphingomonas*, and *Staphylococcus*. On day 12, an increase of *Pseudomonas* spp. and a significant decrease of *Pedobacter* were observed. As the dominant spoilage microorganism during the storage, the relative abundance of *Pseudomonas* spp. showed an increasing trend. Additionally, samples inoculated with a cocktail of *Pseudomonas* spp., i.e., *P. fluorescens*, *P. migulae*, *P. tolaasii*, and *P. agarici*, were incubated at temperatures of 4, 10, 16, 20, 25, and 32 °C to assess their growth kinetics. Three primary models (Huang, Baranyi, and Reparameterized Gompertz model) and three secondary models (Huang square-root, Ratkowsky square-root, and Arrhenius-type model) were compared to evaluate the effect of temperature on bacterial growth using the Integrated Pathogen Modeling Program. The Reparameterized Gompertz model was a better fit than the Huang and Baranyi models to describe the growth of *Pseudomonas* spp., and had the lowest mean square error (MSE) and sum of squared errors (SSE) values, ranging from 0.012 to 0.291 and 0.059–1.783, respectively. Huang square-root model MSE and SSE values were 0.002 and 0.007, respectively, and its predicted minimum growth temperature was -0.72 °C. Therefore, the Huang square-root model was more suitable for describing the effect of temperature on growth of *Pseudomonas* spp. on ABFB. The models developed in this study can be used to evaluate the growth behavior of *Pseudomonas* spp. on ABFB and assess their shelf-life.