Modeling the growth behavior of foodborne pathogens by real-time PCR

- Predicting the influence of temperature abuse -

成果の特徴

- Develop a rapid and specific genome-based method to collect numerous bacterial growth data from various food samples.
- Construct predictive models that enable the estimation of bacterial response to fluctuating temperature conditions.
- Evaluate the difference between the prediction results from real-time PCR data and traditional agar plate count method.

Growth pattern in the temperature abuse scenario 8 20 Plate count data <u>Real-time PCR data</u> 7 15 6 Log CFU/mL emperature 5 10 4 3 5 2 1 0 16 32 64 80 96 112 128 144 0 48 Incubation time (h)

成果の内容

成果の活用

Genome-based analysis method such as real-time PCR enables accurate quantification of bacterial number despite the influence of stress conditions caused by processing treatments, food properties, temperature, and storage conditions that the traditional agar method cannot do. Real-time PCR is a promising data collection tool for customized prediction models, which is important for food companies to estimate the behavior of different pathogens in specific food types subject to food company conditions.

参考

<u>Noviyanti, F.</u>, Shimizu, S., Hosotani, Y., Koseki, S., Inatsu, Y., Kawasaki, S. (2020) *Foodborne Pathogens and Disease* **17**, 693-700.



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