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ONLINE ISSN : 1882-5982 PRINT ISSN : 1340-8267

Japanese Journal of Food Microbiology

Vol. 25 (2008), No. 3 pp.109-119

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Studies on a Rapid Method for Detecting *Salmonella* in Animal Feed Using the QUALIBAXTM System

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(Received March 12, 2008) (Accepted June 20, 2008)

Abstract

Performance of the QUALIBAXTM System (the BAX Method), a rapid PCR-based qualitative method for detecting pathogens in food, was compared to the Official Method in Japan for detecting low levels of *Salmonella* in feed ingredients and formula feed. A total of 78 samples, which included 17 naturally contaminated lots, was tested. A pre-enrichment step, where 25 g samples were grown in 250 m*l* of buffered peptone water with 0.6% Tween, allowed for BAX Method results in 32 hours that were consistent with the Official Method. Analysis determined 97.4% agreement between the two methods, with a false positive rate of 1.3% and an indeterminate rate of 1.3% for the BAX Method. The detection limit was determined by MPN to be approximately 2 cfu/100 g sample. Additional samples were taken from an Official Method selective enrichment in either selenite cystine broth or Hajna tetrathionate broth instead of BAX Method, the time to

result extended beyond 32 hours but all of the BAX Method results agreed with the Official Method, demonstrating advanced precision. These studies showed that the QUALIBAXTM System can be used by feed manufacturers as a fast and accurate method for detecting *Salmonella*.

Key words:

Salmonella, QUALIBAXTM System, PCR, Feed

[PDF (1145K)] [References]

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To cite this article:

Tetsuo CHIHARA, Yoshihiro SEKIGUCHI, Yoshiyasu HASHIMOTO, Shinji OSHIMA, Yasutoshi SUGIMOTO, Hiroaki MOTO, Koichi KIMURA and Kenzo UEHASHI, "Studies on a Rapid Method for Detecting *Salmonella* in Animal Feed Using the QUALIBAXTM System", Japanese Journal of Food Microbiology: **25**: 109-119 (2008).

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