Accepted Manuscript

Multiple-locus variable-number tandem repeat analysis using multiplex polymerase chain reaction and next-generation sequencing - a novel high-throughput method for subtyping *Listeria* strains



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PII:	S0956-7135(18)30289-5
DOI:	10.1016/j.foodcont.2018.06.006
Reference:	JFCO 6180
To appear in:	Food Control
Received Date:	09 November 2017
Accepted Date:	02 June 2018

Please cite this article as: Chirapiphat Phraephaisarn, Yurika Kitai, Rabuesak Khumthong, Hajime Takahashi, Chihiro Ohshima, Punnida Techaruvichit, Mongkol Vesaratchavest, Rutjawate Taharnklaew, Suwimon Keeratipibul, Multiple-locus variable-number tandem repeat analysis using multiplex polymerase chain reaction and next-generation sequencing - a novel high-throughput method for subtyping *Listeria* strains, *Food Control* (2018), doi: 10.1016/j.foodcont.2018.06.006

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ACCEPTED MANUSCRIPT

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3	subtyping <i>Listeria</i> strains
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