

MOLECULAR EPIDEMIOLOGY, INC.



## **PFGE Subtyping Report**

F04/MB454/04

Customer ID #	<b>XXXX-XXXX</b>	Job ID #	>	xxxxx-xx-xxxx-xxxx	MEI Report #	<b>XXXXXXXXX</b>
Customer Sample #	xxxxx-x		٦	Requested Service(s)	Genetic S	ubtyping
Sample Type	NA			Test Method	MB454	51 0
Sample Source	NA			Sample Received	00/00/000	0
<b>Customer Comments</b>	NA			Date Reported	00/00/000	0
MEI Lab Sample #	XXXX-X			Page Number	1 of 1	
% RFLP Pattern Si 응 章	milarity			F	Pattern #	Organism
					BC001	Bacillus cereus
		1 111	Ī		BC002	Bacillus cereus
	1 11	11	1		BC003	Bacillus cereus
	1000				BC004	Bacillus cereus
		11-1-1100			BC005	Bacillus cereus
	1	11 1 1 100	I		BC006	Bacillus cereus
	11		1		BC007	Bacillus cereus
		11111			BC008	Bacillus cereus
	11	111	I		BC009	Bacillus cereus
Deviations	None					
PFGE ConclusionEleven isolates, identified as Bacillus cereus, were analyzed by Pulsed Field Gel Electrophoresis (PFGE). The resulting Restriction Fragment Length Polymorphism (RFLP) patterns were compared to each other and assigned a pattern number. The isolates exhibiting indistinguishable patterns have the same pattern number.						
PFGE Review					Date:	00/00/0000
QC Approval					Date:	00/00/0000
QA Approval	<u> </u>				Date:	00/00/0000
REPORTED BY		•		Molecular Epiden 15300 Bothell Way Lake Forest Park, <u>www.molecularepi</u>	/ NE, WA 98155	

This laboratory report relates only to the portion of the sample which was tested in this report. Interpretation of these results is the sole responsibility of the customer.

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