

**Table S1.** PCR primer pairs used to detect enterotoxin genes in *S. aureus* strains and their predicted amplification size.

Target gene	Primer Sequence (5'-3')	Product size (bp)	Reference
sea	GCAGGGAACAGCTTAGGC GTTCTGTAGAACTATGAAACACG	520	
seb	ATGTAATTGATATCGCAGTG TGCAGGCATCATATCATACCA	643	
sec	CTTGTATGTATGGAGGAATAACA A TGCAGGCATCATATCATACCA	283	Monday and Bohach (1999)
sed	GTGGTGAAATAGATAGGACTGC ATATGAAGGTGCTCTGTGG	384	
see	TACCAATTAACTTGTGGATAGAC CTCTTGACCTTACCGC	170	
seg	CGTCTCCACCTGTTGAAGG CCAAGTGATTGTCTATTGTCG	327	

#### Reference

28. Monday, SR; Bohach, GA. Use of multiplex PCR to detect classical and newly described pyrogenic toxin genes in Staphylococcal isolates. *Journal of Clinical Microbiology*. 1999, 37(10), 3411-3414.