

CORRECTION

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Correction: Molecular analysis of subtilase cytotoxin genes of food-borne Shiga toxin-producing *Escherichia coli* reveals a new allelic *subAB* variant

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Correction

After publication of this work [1], we noticed that the sequences of primers subA-L and subAB5'-OEP in Table two (Table 1 here) are not correct. We therefore provide the correct primer sequences in a new Table two (Table 1 here).

Table 1 Designations, targets, and positions of primers for PCR analysis and Southern blot hybridization

Primer	Target	Sequence (5' – 3')	GenBank accession number	Reference
saaDF	<i>saa</i>	5'-CGT GAT GAA CAG GCT ATT GC-3'	AF399919	[26]
saaDR	<i>saa</i>	5'-ATG GAC ATG CCT GTG GCA AC-3'	AF399919	[26]
subAB-V-for	<i>subAB</i>	5'-CTT CCC TCA TTG CCT CAC G-3'	AY258503	This study
subAB-V-rev	<i>subAB</i>	5'-GGC TGG CCT GTT GTG TAA A-3'	AY258503	This study
tia_lo	<i>tia</i>	5'-TCC ATG CGA AGT TGT TAT CA-3'	U20318	[15]
tia_sense	<i>tia</i>	5'-TTC TCT TTT TAC CCT GCT TTT TGC-3'	FJ664545	[15]
subAB-for5	<i>subAB</i> ₁	5'-CGT ATC TGC GCC ATA TCC TG-3'	AY258503	This study
subAB-rev5	<i>subAB</i> ₁	5'-CTG TTC CGA GCA GCC ATA TC-3'	AY258503	This study
subAB3'tia	<i>subAB</i> ₂₋₁	5'-ACT GGC TGT TCT AAC CG-3'	AEZO02000028.1	This study
subA_out	<i>subA</i> ₂₋₂	5'-GAA TCA ACA ACA GAT ACG AC-3'	AEZO02000020.1	This study
subA-L	Linker ^a	5'-GGG AGG ATT AAC CAT CG-3'	AEZO02000020.1	This study
subAB5'OEP	<i>subAB</i> ₂₋₂	5'-ATG AAT GAG AGC ATC CCT-3'	AEZO02000020.1	This study
subAB2-3'out	<i>subAB</i> ₂₋₂	5'-AGG TCG GCT CAG TGT TC-3'	AEZO02000020.1	This study

^aIntergenic linker between the OEP-locus and *subA*₂₋₂.

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Reference

1. Funk J, Stoeber H, Hauser E, Schmidt H: Molecular analysis of subtilase cytotoxin genes of food-borne shiga toxin-producing *escherichia coli* reveals a new allelic *subAB* variant. *BMC Microbiol* 2013, 13:230.

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