

Table I. Primers used in conventional and AMOS-PCR for identification of *Brucella* in samples collected from seropositive cattle and buffaloes in the Delta region of Egypt.

Conventional PCR primer	Nucleotide sequence 5'-3 '	Amplicon size
B4 primer	TGG CTC GGT TGC CAA TAT CAA	223 bp
B5 primer	CGC GCT TGC CTT TCA GGT CTG	
AMOS-PCR		-
<i>B. abortus</i>	GAC-GAA-CGG-AAT-TTT-TCC-AAT-CCC	498 bp
<i>B. melitensis</i>	AAA-TCG-CGT-CCT-TGC-TGG-TCT-GA	731 bp
<i>B. ovis</i>	CGG-GTT-CTG-GCA-CCA-TCG-TCG	976 bp
<i>B. suis</i>	GCG-CGG-TTT-TCT-GAA-GGT-TCA-GG	285 bp
IS711	TGC-CGA-TCA-CTT-AAG-GGC-CTT-CAT	-