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Corrigendum to “Unravelling the diversity of *Anaplasma* species circulating in selected African wildlife hosts by targeted 16S microbiome analysis” [Current Research in Microbial Sciences Volume 5 (2023) 100198]

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The authors regret that the forward primer and probe sequences of the *Anaplasma*/*Ehrlichia* group-specific assay in the paper are incorrect.

CORRECT SEQUENCES:

Primer, Ma16SF: (5'-ACA GAA GAA GTC CCG GCA AA-3'), Ma16SR: (5'-TTG CCC CCT CCG TAT TAC C-3') (Inqaba Biotech, South Africa) and a TaqMan MGBTM probe, Ma16SP: (FAM-5'-CCG TGC CAG CAG C-3'-MGB) (Thermo Fisher Scientific, South Africa) were designed to target a 64 bp fragment in the V3 hypervariable region that is conserved between *Anaplasma* and *Ehrlichia* species.

IN THE PAPER (INCORRECT):

Primers, Ma16SF: (5'-ACA GAA GTC CCG GCA AA-3'), Ma16SR: (5'-TTG CCC CCT CCG TAT TAC C-3') (Inqaba Biotech, South Africa) and a TaqMan MGBTM probe, Ma16SP: (FAM-5'-CCG TGC CAG C-3'-MGB) (Thermo Fisher Scientific, South Africa) were designed to target a 64 bp fragment in the V3 hypervariable region that is conserved between *Anaplasma* and *Ehrlichia* species.

The authors would like to apologise for any inconvenience caused.

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