

**S2 File:** Alignment of 184 amino acids of partial *Culicoides* mtDNA COI gene in fasta+gap format for used in phylogenetic analysis.

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>Cbrev-01 [GenBank:KP201844]
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>Anopheles quadriannulatus [GenBank:DQ792581]

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>Anopheles dirus\_D [GenBank:AJ877572]

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>Anopheles janconnae [GenBank:HQ335348]

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