

<https://doi.org/10.11646/zootaxa.4920.4.10>

<http://zoobank.org/urn:lsid:zoobank.org:pub:F889E84A-58C4-4CB5-AB9C-97E80FFBAFCA>

**MIREILLE BELLE MBOU OKASSA, DOLLON MBAMA NTABI & ARSENE LENGA (2020)  
Morphological and molecular identification of specimens in the genus *Euseius*  
(Acarı: Phytoseiidae) from the Republic of Congo. *Zootaxa*, 4768: 479–498.**

A table of data was accidentally omitted from the recent paper by Belle Mbou Okassa *et al.* (2020). That omission is corrected here.

**TABLE 5.** Mean genetic Tamura 3 parameter and K2P distances (minimum and maximum values) within and between *Euseius sp* and *E. fustis*, with outgroup species *Typhlodromus pyri* for the two molecular markers used.

A. 12S rRNA	<i>Euseius sp</i>	<i>Euseius fustis</i>	<i>Typhlodromus pyri</i>
<i>Euseius sp</i>	0 (0–0.1)		
<i>Euseius fustis</i>	0.12 (0.12–0.13)	0 (0–0)	
<i>Typhlodromus pyri</i>	0.34 (0.34–0.36)	0.36 (0.36–0.36)	0 (0–0)
B. ITSS	<i>Euseius sp</i>	<i>Euseius fustis</i>	<i>Typhlodromus pyri</i>
<i>Euseius sp</i>	0 (0–0)		
<i>Euseius fustis</i>	0.2 (0.2–0.3)	0 (0–0)	
<i>Typhlodromus pyri</i>	0.16 (0.16–0.16)	0.17 (0.17–0.17)	0 (0–0.01)