

Correction

Correction: Wilson, N.G., et al. Tropical Range Extension for the Temperate, Endemic South-Eastern Australian Nudibranch *Goniobranchus splendidus* (Angas, 1864). *Diversity* 2016, 8, 16

Nerida G. Wilson ^{1,2,*} , Anne E. Winters ³ and Karen L. Cheney ³

¹ Western Australian Museum, 49 Kew Street, Welshpool, WA 6106, Australia

² School of Animal Biology, University of Western Australia, Crawley, WA 6009, Australia

³ School of Biological Sciences, The University of Queensland, St Lucia, QLD 4072, Australia; e.anne06@gmail.com (A.E.W.); k.cheney@uq.edu.au (K.L.C.)

* Correspondence: nerida.wilson@museum.wa.gov.au; Tel.: +61-08-9212-3844

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The authors wish to make the following corrections to this paper [1]. The details are as follows:

1. In Table 1, the pending Genbank numbers should be MH018011, MH017998, MH018010, MH01797. Please, replace Table 1

Table 1. Species of *Goniobranchus* included in the phylogenetic analysis.

Species	Voucher	Locality	COI	16 S
<i>G. albopustulosus</i>	CASIZ 142953	Maui, Hawaii	JQ 727828	JQ 727701
<i>G. daphne</i>	UQ collection	Gold Coast Seaway, QLD	KJ 001297	KJ 018921
<i>G. epicurea</i>	SAM D 19285	Tasmania, Australia	EF 535114	AY 458804
<i>G. heatherae</i>	CASIZ 175546	Cape Peninsula, South Africa	JQ 727844	JQ 727720
<i>G. kuniei</i>	SAM D 19261	Heron Is., GBR, QLD	EF 535112	AY 458807
<i>G. splendidus</i>	SAM D 19292	Mooloolaba, QLD	EF 535115	AY 458815
<i>G. splendidus</i>	CASIZ 146039	Mooloolaba, QLD	EU 982738	EU 982789
<i>G. cf. splendidus</i>	UQ 1102	Mackay, QLD	pending	pending
<i>G. tasmaniensis</i>	SAM 19295	Triabunna, Tasmania	EF 535113	AY 458817
<i>G. tinctorius</i>	UQ 1097	Mackay, QLD	pending	pending
<i>G. verrieri</i>	CASIZ 158796	Batangas, Philippines	JQ 727858	JQ 727740

with the following

Table 1. Species of *Goniobranchus* included in the phylogenetic analysis.

Species	Voucher	Locality	COI	16 S
<i>G. albopustulosus</i>	CASIZ 142953	Maui, Hawaii	JQ 727828	JQ 727701
<i>G. daphne</i>	UQ collection	Gold Coast Seaway, QLD	KJ 001297	KJ 018921
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<i>G. cf. splendidus</i>	UQ 1102	Mackay, QLD	MH018011	MH017998
<i>G. tasmaniensis</i>	SAM 19295	Triabunna, Tasmania	EF 535113	AY 458817
<i>G. tinctorius</i>	UQ 1097	Mackay, QLD	MH018010	MH01797
<i>G. verrieri</i>	CASIZ 158796	Batangas, Philippines	JQ 727858	JQ 727740

2. Please note that Figure 2 contains a mistake, *splendidus* Mackay (UQ1097) should be *splendidus* Mackay (UQ1102). Please replace Figure 2

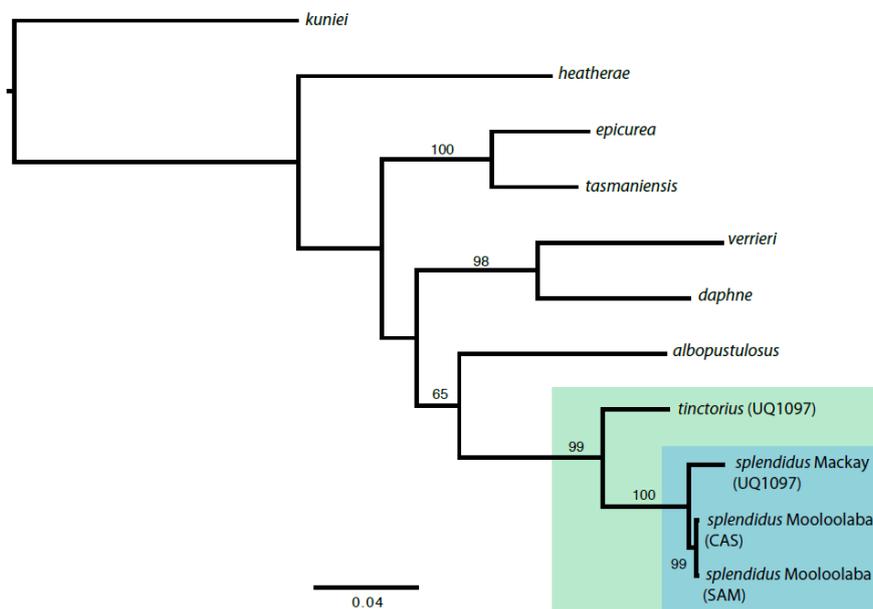


Figure 2. Maximum-likelihood topology for Cytochrome Oxidase I gene (COI) + 16S DNA from selected species of *Goniobranchus* nudibranchs. Specimens of *G. splendidus* shown in blue box, and sister species *G. tinctorius* shown in green box. Bootstrap values less than 60 are not shown.

with the following

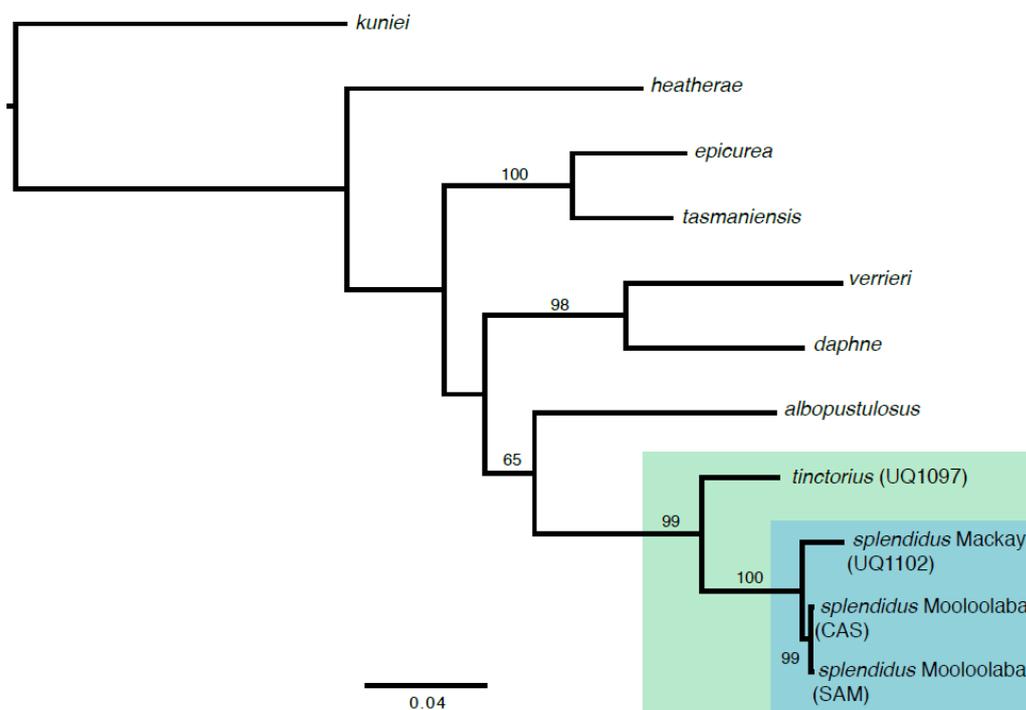


Figure 2. Maximum-likelihood topology for Cytochrome Oxidase I gene (COI) + 16S DNA from selected species of *Goniobranchus* nudibranchs. Specimens of *G. splendidus* shown in blue box, and sister species *G. tinctorius* shown in green box. Bootstrap values less than 60 are not shown.

We apologize for any inconvenience caused to the readers by these changes. The manuscript will be updated, and the original publication will remain available from the article webpage.

Reference

1. Wilson, N.G.; Winters, A.E.; Cheney, K.L. Tropical Range Extension for the Temperate, Endemic South-Eastern Australian Nudibranch *Goniobranchus splendidus* (Angas, 1864). *Diversity* **2016**, *8*, 16. [[CrossRef](#)]



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