

AUTHOR’S CORRECTION

High-Level Diversity of Dinoflagellates in the Natural Environment, Revealed by Assessment of Mitochondrial *cox1* and *cob* Genes for Dinoflagellate DNA Barcoding

Senjie Lin, Huan Zhang, Yubo Hou, Yunyun Zhuang, and Lilibeth Miranda

Department of Marine Sciences, University of Connecticut, Groton, Connecticut 06340

Volume 75, no. 5, p. 1279–1290, 2009. Page 1280, column 2, line 14 from the bottom: “2008” should read “2007.”
 Page 1285: Table 4 should appear as shown below.

TABLE 4. Distances between species in the same genus and between strains within species derived from *cob*, based on the TVM+G model

Group	Range	Mean	SD	No. of sequences	No. of taxa
Congeneric, interspecific					
<i>Alexandrium</i>	0–0.00299	0.0009	0.0014	5	21
<i>Amphidinium</i>	0.23652	0.2365	0.0000	2	1
<i>Ceratium</i>	0.02096	0.0210	0.0000	2	1
<i>Heterocapsa</i>	0.11976–0.25150	0.1986	0.0696	3	3
<i>Karenia</i>	0	0.0000	0.0000	2	1
<i>Prorocentrum</i> (<i>P. lima</i> and <i>P. micans</i> vs others)	0.00898–0.06587	0.0355	0.0242	7	35
<i>Pyrocystis</i>	0.00898	0.0090	0.0000	2	1
<i>Scrippsiella</i>	0.00898–0.04790	0.0339	0.0217	3	11
<i>Symbiodinium</i>	0.02695–0.04491	0.0339	0.0096	3	3
Intraspecific					
<i>Peridinium aciculiferum</i>	0	0.0000	0.0000	5	10
<i>Polarella glacialis</i>	0	0	0	2	1
<i>Prorocentrum dentatum/P. donghaiense</i>	0	0	0	2	1
<i>Prorocentrum minimum</i>	0.00599–0.00898	0.006487	0.0035	4	6
<i>Scrippsiella hangoei</i>	0–0.00299	0.0018	0.0015	5	10