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PG-7 Two dinoflagellate taxa, *Durinskia* (Peridiniales, Dinophyceae) and *Goniodoma* (Gonyaulacales, Dinophyceae) from Okinawa, Japan

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Dinoflagellates have attracted much attention in research as they are often associated with ciguatera poisoning and other related toxic phenomena in subtropical to tropical coastal areas. Hence, the complete taxonomic description of different potential toxic and nontoxic dinoflagellates is important. The present research has been conducted with the aim to describe the taxonomy of different dinoflagellate taxa from Okinawa, Japan. Among our isolated strains, we have been focusing on the two taxa, Durinskia and Goniodoma to confirm their taxonomic and phylogenetic status as both of genera have not been well described previously. Based on light microscopy, Durinskia strain's cells are round to ovoid, measuring 15 to 24.4 μ m length, 13.1 to 23.8 μ m width with a conspicuous, bright red, large and rectangular eyespot. The thecal plate arrangement (Po. x, 4', 2a, 6'', 5c, 4s, 5'', 2''') of this species is similar to that of D. baltica and D. capensis. Based on cell shape, size, eyespot, cingular displacement, nucleus (eukaryotic and dinokaryon), and thecal plate arrangements, the present strain is placed within the genus Durinskia. The genus Durinskia currently has two species, D. baltica and D. capensis. SSU rDNA sequences indicate that this strain is distinct from D. baltica and D. capensis by 26 and 53 basepair substitutions, respectively. In case of rbcL gene sequence analysis, the present Durinskia strain formed same clade with other endosymbionts and showed genetic differences compared to D. baltica and D. capensis. These results suggest that this strain is distinct from the two known species and constitute a potentially undescribed species of *Durinskia*.

The Goniodoma strain's cells consist of thick thecal plates with many trichocysts pours. Cells are spherical to hemispherical in shape and slightly shorter than wide. Cell size ranges from 48 to 64 μ m in length and 51to 67 μ m in wide. The cell size and shape agree with G. polyedricum. However, no Goniodoma sequence data is available so far for SSU rDNA or other DNA sequences. The preliminary analysis of SSUrDNA sequences shows that the present strain of Goniodoma is closely related to the sequence from Alexandrium tamutum. Alexandrium is very similar to Goniodoma but differs in size, thickness, size and shape of thecal plates. Further detailed studies to confirm the species and taxonomic position of the present Goniodoma strain are being carried out based on morphology and molecular phylogeny.