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Thriving populations with low genetic diversity in giant clam species, *Tridacna maxima* and *T. noae*, at Dongsha Atoll, South China Sea

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ACCEPTED MANUSCRIPT

1	Thriving populations with low genetic diversity in giant clam species, <i>Tridacna maxima</i> and
2	T. noae, at Dongsha Atoll, South China Sea
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14	ABSTRACT
15	Giant clams provide valuable functions to the coral reefs and traditional fisheries in the South
16	China Sea (SCS), but many populations face escalating threats of overexploitation and habitat
17	loss. To provide critical biodiversity data for development of protection and sustainable
18	utilisation strategies, we examine the status of wild tridacnine population at Dongsha Atoll,
19	the largest northernmost atoll of the SCS. This study also examines the genetic patterns of
20	Tridacna maxima and T. noae, based on partial mitochondrial COI gene sequences. We
21	found four species at Dongsha Atoll with an overall density of 3.14 per 100 m ² . Over-
22	harvesting may have depleted populations of <i>T. squamosa</i> and <i>Hippopus</i> hippopus, and they
23	may no longer be reproductively viable. Populations of <i>T. maxima</i> and <i>T. noae</i> appear to be
24	thriving and replenished by recruits, but species showed low levels of mitochondrial genetic
25	diversity that could reduce its adaptability and may become further impacted by exploitation

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