

GENETIC COMPARISON OF *DONAX STRIATUS* POPULATIONS ALONG THEIR GEOGRAPHICAL DISTRIBUTION

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ABSTRACT

Shells of *Donax striatus* sampled from a population at Cocal Beach, Trinidad (Wade 1967), and from a population at Praia do Tibau, Brazil (own observations), show strong variation in shell shape, size and colour. In order to estimate the genetic relatedness among geographically distant *D. striatus* populations across its geographical distribution, genetic analyses will be carried out.

Keywords: population genetics, bivalves, Brazil

INTRODUCTION

Even though the genetic structure among populations represents the effect of the interaction among gene flow, genetic drift, selection and mutation, the relative importance of each of these factors may often be difficult to separate (Balloux & Lugon-Moulin 2002). For many species, their geographical distribution exceeds the distances that can be covered by any single dispersing individual. This can lead to a genetic structure that is characterized by a correlation between genetic and geographical distance between populations (isolation by distance; Wright 1943). However, in some organisms empirical findings of genetic differentiation may be difficult to reconcile with knowledge on the biology of the species. For most marine organisms, including marine bivalves, the mechanisms affecting genetic structure still remain largely unknown. Exposed intertidal sandy beaches are commonly dominated by surf clams of the genus *Donax* (Ansell 1983). Being the main primary consumers in soft bottom communities they can contribute up to 95 % of the total biomass (McLachlan *et al.* 1981; Arntz & Fahrback 1991). Despite the key role of surf clams in sandy beach ecosystems and their importance for commercial and recreational fishery, the population genetic of most donacids remains still unknown. *Donax striatus* Linnaeus, 1767 (Bivalvia: Donacidae) inhabits the intertidal zone along the Caribbean (south of Gibara, Cuba) and western Atlantic coast (north of Areia Branca, Brazil) (WoRMS 2010). The taxonomy of this tropical surf clam is controversially discussed (Coan 1983). The high variability in shell shape, size and colour, confirmed by our pilot survey in May 2007 at the Brazilian beach Tibau, raised doubts concerning the taxonomic identity of the species (Wade 1967). As morphometric comparisons revealed significant differences, the present project aims (1) to establish a marker for molecular barcoding, (2) clarify the taxonomy of the species and (3) estimate the genetic relatedness among geographically distant populations of *D. striatus*.

MATERIALS AND METHODS

Specimens of the *D. striatus* were sampled at seven beaches of the Brazilian state Ceará (from 2°52'S, 40°57'W to 4°38'S, 37°28'W). *D. hanleyanus* was sampled as the out-group from the Argentinean beach Santa Teresita (36°32'S, 56°41'W). In order to facilitate the diffusion of the preservative into the tissue, the shells were broken or sheared immediately before conservation in 96 % ethanol. DNA was extracted with the Qiagen DNeasy kit according to the manufacturer's recommendations. 10-100 ng DNA were used for PCR-amplification of a fragment of the cytochrome oxidase I (Cox1) using the primers HCO (5'-TAACTTCAGGTGACCAAAAATCA-3') and LCO (5'-GGTCAACAAATCATAAAGATATTGG-3') (Folmer *et al.* 1994) in 25 µl reactions. PCR products were purified using the Qiagen DNA purification kit according to the manufacturer's recommendations. Sequencing was conducted on an ABI 3730xl automated sequencer. Sequence data were processed and aligned using

ClustalW. Phylogenetic analysis was performed using the software package for inference of evolutionary trees PAUP* version 4.0beta10.

PRELIMINARY RESULTS

Sequence data from the Cox1 proved useful for species discrimination within the genus *Donax*. There is no molecular evidence for differentiation at the species level within *D. striatus*. By the deadline of the abstract submission the present project is already in process. Therefore, results of the genetic comparison of *D. striatus* across its geographical distribution will be presented at the author's poster.

REFERENCES

- ANSELL, A.D. 1983. The biology of the genus *Donax*. In: McLachlan, A., Erasmus, T. & Junk, W. (eds.) *Developments in Hydrobiology Vol. 19. Sandy Beaches as Ecosystems*. Dr. W. Junk Publishers, The Hague, The Netherlands, The Netherlands:607-635.
- ARNTZ, W.E. & FAHRBACH, E. 1991. El Niño - Klimaexperiment der Natur: Die physikalischen Ursachen und biologischen Folgen. Birkhäuser, Basel, Boston, Berlin. p. 264.
- BALLOUX, F. & LUGON-MOULIN, N. 2002. The estimation of population differentiation with microsatellite markers. *Molecular Ecology* 11: 155-165.
- COAN, E. 1983. The Eastern Pacific Donacidae. *The Veliger* 25 (4): 273-298.
- FOLMER, O., BLACK, M., HOEH, W., LUTZ, R. & VRIJENHOEK, R. 1994. DNA primers for amplification of mitochondrial cytochrome c oxidase subunit I from diverse metazoan invertebrates. *Molecular Marine Biology and Biotechnology* 3 (5): 294-299.
- MCLACHLAN, A., ERASMUS, T., DYE, A.H., WOOLRIDGE, T., VAN DER HORST, G., ROSSOUW, G., LASIAK, T.A. & MCGWYNNE, L.E. 1981. Sand beach energetics: an ecosystem approach towards a high energy interface. *Estuarine, Coastal and Shelf Science* 13: 11-25.
- WADE, B.A. 1967. On the taxonomy, morphology, and ecology of the beach clam, *Donax striatus* Linné. *Bulletin of Marine Science* 17: 723-740.
- WORMS. 2010. *Donax striatus* Linnaeus, 1767. In: Bouchet, P., Gofas, S. & Rosenberg, G. (eds.) *World Marine Mollusca database*. Accessed through: World Register of Marine Species at <http://www.marinespecies.org/aphia.php?p=taxdetails&id=494732>.
- WRIGHT, S. 1943. Isolation by distance. *Genetics* 28: 114-138.