

Weak connectivity and population cohesiveness in rudderfish *Kyphosus sandwichensis* (Teleostei: Kyphosidae) inhabiting remote oceanic islands

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[Ver número de ResearchID y ORCID de Web of Science](#)

AQUATIC CONSERVATION-MARINE AND FRESHWATER ECOSYSTEMS

DOI: 10.1002/aqc.3438



Acceso anticipado: AUG 2020

Tipo de documento: Article; Early Access

[Ver impacto de la revista](#)

Abstract

Population connectivity has a fundamental role in metapopulation dynamics, with important implications in conservation. Easter Island (EI) and Salas y Gomez Island (SG) in the Pacific Ocean are ideal for the study of population connectivity because they are separated by 415 km and isolated from other islands in the Pacific Ocean by >2,000 km. Considering that dispersal processes could play a critical role in the persistence of its populations, the connectivity pattern of the rudderfish *Kyphosus sandwichensis* was evaluated between EI and SG using both a population genetics and a biophysical modelling approach. The variability in the control region of the mitochondrial DNA did not show a significant phylogeographical pattern, and the variability in 16 microsatellite loci suggested that individuals of *K. sandwichensis* located at EI and SG belong to the same genetic population. However, historical migration showed that 0.2% of the recruits at EI come from SG and that 0.15% at SG come from EI per year. Using simulated larval release during September and a larval development of 30 days in the plankton, biophysical modelling did not detect migration between the islands. Furthermore, self-recruitment shows interannual variation ranging from 5 to 10% of the total released larvae. Whereas the genetic data showed a lack of population genetic structure but low connectivity of *K. sandwichensis* between EI and SG, the biophysical modelling showed null movement of particles between the islands. Stochastic movement of larvae or adults could explain the pattern observed, with rafting as an example. These low-frequency and stochastic movements may be important in maintaining the cohesiveness between EI and SG.

Palabras clave

Palabras clave de autor: [marine protected areas](#); [population connectivity](#); [Rapa Nui ecoregion](#); [reef-associated fish](#)

KeyWords Plus: [PLANKTONIC LARVAL DURATION](#); [MARINE PROTECTED AREAS](#); [EASTER-ISLAND](#); [TELEOSTEI](#)

[ATHERINOPSIDAE](#); [DISPERSAL](#); [DNA](#); [SOFTWARE](#); [GENETICS](#); [PATTERNS](#); [PACIFIC](#)

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Financiación

Entidad financiadora Mostrar más información	Número de concesión
Chilean Millennium Initiative grant ESMOI	
Conicyt-Fondequip	EQM150077
NLHPC	ECM-02
Comision Nacional de Investigacion Cientifica y Tecnologica (CONICYT)	
Comision Nacional de Investigacion Cientifica y Tecnologica (CONICYT) CONICYT FONDECYT	3150419
CONICYT PIA Apoyo CCTE	AFB170008

[Ver texto de financiación](#)

Editorial

WILEY, 111 RIVER ST, HOBOKEN 07030-5774, NJ USA

Información de la revista

- **Impact Factor:** [Journal Citation Reports](#)

Categorías / Clasificación

Áreas de investigación: Environmental Sciences & Ecology; Marine & Freshwater Biology; Water Resources

Categorías de Web of Science:Environmental Sciences; Marine & Freshwater Biology; Water Resources

Información del documento

Idioma:English

Número de acceso: WOS:000559475400001

ISSN: 1052-7613

eISSN: 1099-0755