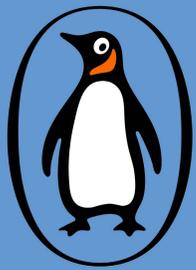


# Spatial Heterogeneity as a Genetic mixing mechanism in highly philopatric Colonial Seabirds

R CRISTOFARI<sup>1,2,3,\*</sup>, E TRUCCHI<sup>3</sup>, JD WHITTINGTON<sup>3,1,2</sup>, S VIGETTA<sup>1</sup>, H GACHOT-NEVEU<sup>1</sup>, NC STENSETH<sup>3</sup>, Y LE MAHO<sup>1,2</sup>, C LE BOHEC<sup>2,1</sup>

<sup>1</sup>Institut Pluridisciplinaire Hubert Curien, LEA-647 BioSensib, Centre National de la Recherche Scientifique - University of Strasbourg, Strasbourg, France (robin.cristofari@iphc.cnrs.fr), <sup>2</sup>Centre Scientifique de Monaco, LEA-647 BioSensib, Monaco, Principality of Monaco, <sup>3</sup>Centre of Ecological and Evolutionary Synthesis, University of Oslo, Oslo, Norway



## Background.

- How is genetic diversity maintained in philopatric colonial systems such as in dense penguin colonies?
- Philopatric behaviour has several selective advantages, yet its possible consequence, inbreeding depression, may cause entire populations to crash.
- In the **King penguin *Aptenodytes patagonicus***, return rates of chicks to their natal sub-colony are remarkably high. And when starting to breed in an area, adults tend to return year after year to their previous breeding territories.

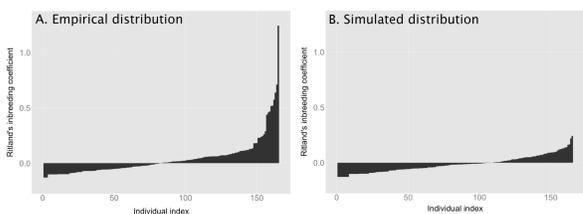
● In order to assess the importance and consequences of this phenomenon, we present the first fine-scale study of the genetic structure in a king penguin colony.

## Methods.

- 175 chicks were sampled and the nests precisely geolocated, during the early 2010 breeding season, on Possession Island, Crozet Archipelago (*fig. 1*).
- Tick infestation, chick survival rate, and site occupancy chronology were assessed for several years.
- Samples were genotyped at 8 microsatellite loci.
- Heterogeneous patterns were investigated through spatial distribution of individual inbreeding and pairwise relatedness, both using continuous, and clustered analysis methods.

## Results.

● The Colony is in equilibrium, genetically diverse, and no global processes (such as colony-wide autocorrelation) are visible - yet **no global process does not mean no structure!**

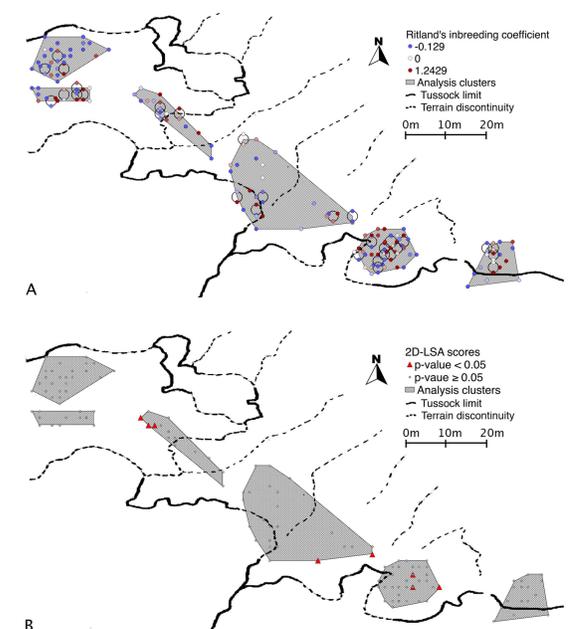


**Fig. 2. Individual inbreeding distribution deviates from expectations.** A. Observed individual inbreeding distribution (Ritland's coefficient). B. Simulated distribution for a population of non-related individuals.

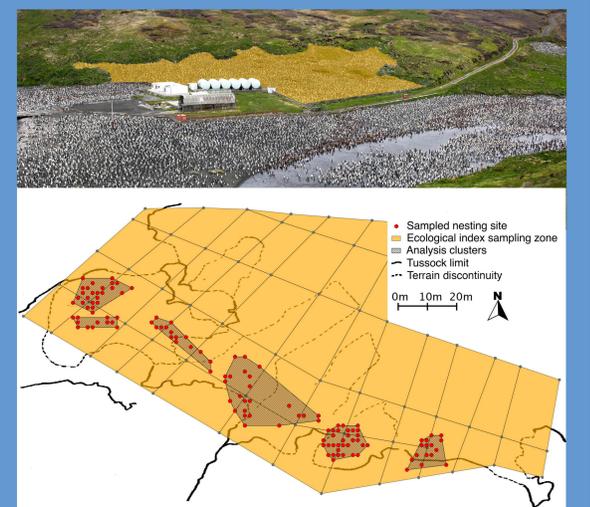
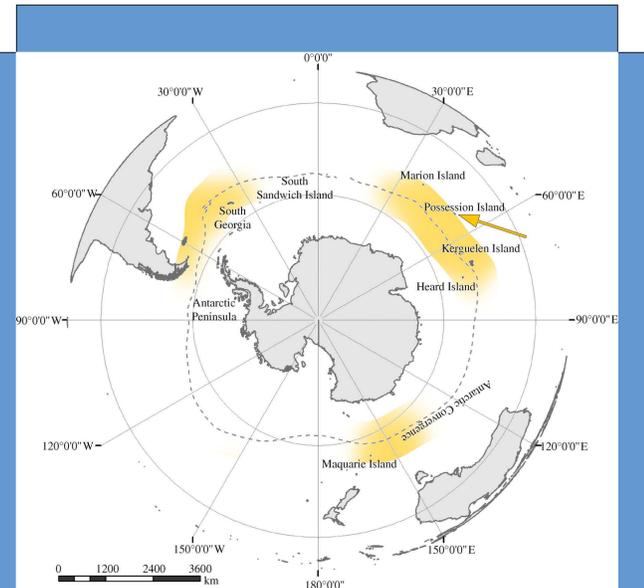
- High mean individual inbreeding level (near half-sib), and some individuals are more related than expected by mere chance (*fig. 2*).
- Higher or lower inbreeding and pairwise relatedness appear to be organized in patches (*fig. 3*).
- These patches correlate with ecological site-quality descriptors: better patches attract early and more successful breeders. They bring about higher individual inbreeding levels in offsprings. Lower-quality sites on the other hand promote outbreeding (*fig. 3 and 4*).

## Conclusions.

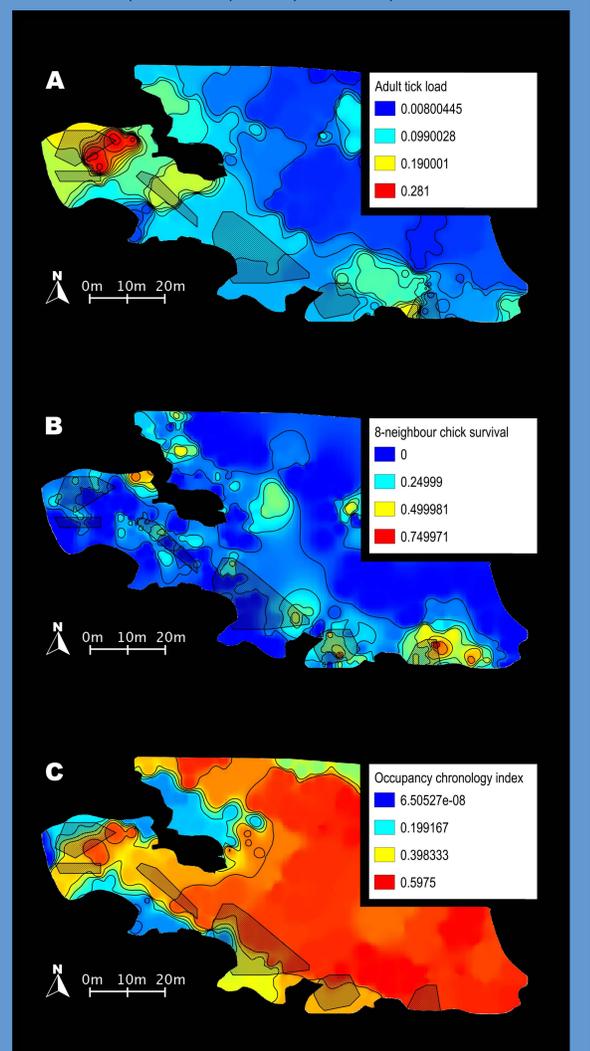
- Despite highly philopatric behaviour, colonial seabirds such as king penguins manage to keep high levels of genetic diversity and mixing within colonies.
- Our results stress the importance of understanding intra-colonial dispersal and genetic mixing mechanisms in order to better estimate species-wide gene flow and population dynamics.
- **Heterogeneity in nesting-site quality may be an important inbreeding avoidance and genetic mixing driver in highly philopatric species.**



**Fig. 3. Individual inbreeding and nearest-neighbour-relatedness tend to cluster.** A. Distribution of Ritland's individual inbreeding coefficient along the sampling area. Shaded zones: clusters C1 to C6. B. 2D-LSA scores. Red triangles represent individuals that are significantly more related to their 9 nearest neighbours than to random individuals.



**Fig. 1. Sampling design.** Sampling was restricted to the periphery of the colony. Orange zones boundaries are marked on the ground for remote parameter assessment. Shaded clusters run from C1 (north-west) to C6 (south-east).



**Fig. 4. Ecological descriptors of breeding-site quality exhibit a strongly heterogeneous distribution across the colony.** A. Adult tick load, averaged for years 2005-2012. B. 8-neighbour chick survival, averaged for years 2010-2012. C. Site occupancy chronology, averaged for years 2006-2013. Ratio of brooding birds amongst 50 randomly selected breeders.

