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► To cite this version:

Meritxell Genovart, Olivier O. Gimenez, Albert Bertolero, Rémi Choquet, Daniel Oro, et al.. Decrease in social cohesion in a colonial seabird under a perturbation regime. Scientific Reports, 2020, 10 (18720), 10.1038/s41598-020-75259-3 . hal-03020416

HAL Id: hal-03020416

<https://hal.science/hal-03020416>

Submitted on 24 Nov 2020

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**Decrease in social cohesion in a long-lived species under a perturbation
regime**

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Abstract

1. Environmental perturbations may have a strong impact on the dynamics of a population and their understanding may help to mitigate the effects of global change. In social animals, social interactions can influence behavioural processes and can play an important role on populations' resilience. However little is known about the effects of perturbations on the strength of social cohesion that keep group-living animals connected.
2. To explore the strength of social cohesion and its dynamics under perturbations, we studied an ecological system involving a colonial, long-lived species living in a site experiencing a shift to a perturbed regime. This regime, caused by the invasion of predators, led an Audouin's gull *Larus audouinii* colony to hold from 70% to only 3% of the total world population in only one decade (32% mean annual decline). Birds breed aggregated in discrete and annually changing patches within colonies, which allowed us to disentangle whether annual aggregations were random or resulted from social ties among individuals. Our goals were 1) to uncover if there were long-term social ties between individuals and 2) to examine whether the perturbation regime affected social cohesion.
3. We explored social cohesion by means of contingency tables and by modeling interdependencies among observations within the Social Network Analysis framework, using additive and multiplicative effects (AME) and accounting for missing data. We analysed 25 years of monitoring with an individual capture-recapture database of more than 3,500 individuals.
4. We showed that there are social ties between individuals over the years. Furthermore, social cohesion strongly decreased after entering the perturbation regime. We propose that sociality and individual behavioural heterogeneity play

a major role driving dispersal between sites and thus population dynamics in social animals.

5. Perturbations may lead not only to changes in individuals' behaviour and fitness but also to a change in populations' social cohesion. The demographic consequences of the breaking down of social ties are still not well understood, but they can be critical for population dynamics of social species. Further studies considering individual heterogeneity, sociality and different types of perturbations should be carried out to improve our understanding on the resilience of social species.

Keywords: Audouin's gull, colonial species, decision-making, non-linear response, perturbations, resilience, social cohesion, social network analyses.

Introduction

Ecosystems are subject to perturbations, both natural and human induced, affecting individuals, populations and communities. When they are strong or are maintained through time, these perturbations may cause a shift between stable states at the level of both individual and population and even lead to population collapses and extinctions^{1,2}. Understanding how individuals and populations will respond to these perturbations is critical both from a ‘pure’ ecological standpoint and also from an applied point of view to mitigate the effects of global change^{3–5}. Population dynamics may be directly affected by these perturbations through a decrease in demographic parameters such as survival or fecundity, or by a change, immediate or delayed, on individual behaviour, such as an increase in dispersal⁶. We define population resilience as the maximal pulse perturbation a population can tolerate or absorb without going extinct^{1,7}. In social animals, social behavioural processes, such as information sharing and decision-making, add another dimension to understanding the resilience of populations facing perturbations. For instance, the amount of social information can be enhanced not only by positive density-dependence, but also by social cohesion^{8–11}. Social cohesion favors the exchange of private information and consequently reduce uncertainty in resource acquisition (e.g. shelter against predators, food, mates) or in decision-making in the face of disturbances, such as dispersal to non-perturbed or less perturbed sites^{6,12–14}. Thus, the structure of a group may affect social interactions, information transfer, and collective decisions¹⁵. Some recent studies also show that spatial cohesion may be risk sensitive, and that responses may differ depending on the risk and the social group¹⁶. However, little is known about the effects of environmental perturbations on the cohesion of social groups in empirical studies of social animals¹⁷.

The analysis of social relationships in animal populations may include a range of social dynamics, from simple and ephemeral contacts, to permanent and strong bonds between individuals^{11,18–20}. Coloniality is a life-history strategy where individuals show a social link among conspecifics by breeding in large and dense groups^{21,22}. However, many colonial species are philopatric, thus this link may not necessarily reflect individual social ties but a shared tendency to breed in the same birthplace. This tendency may result from the need to share information about resources, especially when they are patchy and more unpredictable, or it may result from the advantages of social defence against predators^{23,24}. A challenge lies in disentangling whether annual association between individuals is only due to philopatry, or also due to the existence of social ties within groups of individuals over time²⁵. Social ties between neighbouring pairs in breeding colonies are rarely considered in behavioural and ecological studies²⁶ and, if true, such associations may suggest the evolution of social cohesion for exploiting the evolutionary advantages of social living (including social information sharing) for individual fitness prospects.

Social network theory, which originated in sociology to study human relationships and social organization^{9,27,28} now provides both a conceptual framework and the analytical tools to explore social cohesion and social processes in animal populations^{29–33}.

Network theory is now being simultaneously developed in a number of fields, including statistical physics, sociology, molecular biology, and computer science. As a result, the field is changing at a rapid pace. While not all developments can or should be applied toward the study of animal societies³⁴, this rush of novel ideas from outside disciplines is enriching behavioural ecology³⁵.

To assess the existence of social cohesion and its dynamics under perturbations, we studied an ecological system involving a colonial, social vertebrate (the Audouin's gull

Larus audouinii) living in a site experiencing a shift to a perturbed regime. Interestingly from a social point of view, the species breeds aggregated in spatially-discrete patches within large colonies. Each breeding season, some patches go extinct and some are colonized³⁶, forcing individuals to breed in patches different from the ones they were born in or they bred in the previous year. These colonization-extinction processes may allow us to disentangle whether social aggregation among individuals is an annual random association, or it rather results from social cohesion among individuals over the years.

An extensive long-term individual monitoring program has been carried out since 1988 at the Ebro Delta, including the main breeding site for the species, the Punta de la Banya³⁷. At the study site, population dynamics has undergone different phases: an initial growing phase after site colonization, a stable phase of dynamic equilibrium, and a final transition phase to population collapse^{17,38} (Figure 1). This collapse was due to the arrival of terrestrial predators, and led this colony to hold from 70% to only 3% of the total world population in only a decade (32% mean annual decline)^{36,37} (Figure 1). Most predators were foxes, but also badgers and other mesocarnivores. Predators invaded the site likely due to their increasing densities in recent decades³⁹ and the attractiveness of the site in terms of food availability and lack of competition. The perturbation regime caused changes in the spatial distribution of patches at the site, changes in age structure, decrease in fecundity and a progressively increase of dispersal to other sites^{6,36}. The response of this population to predators has been not immediate probably due to strong philopatry, high site-suitability inertia and social behavioural processes, such as conspecific attraction¹⁷. This raise in dispersal was caused by social processes, as social copying¹⁷, however it remains to assess how social cohesion among individuals, if occurred, was affected by dispersal processes. One possibility is that

dispersal would broke social cohesion by individual heterogeneity in the willingness to
 disperse⁴⁰. In contrast, social cohesion can be maintained over time when dispersal
 occurs collectively at the scale of social groups to the same sites^{25,41}.
 Taking advantage of the long-term monitoring of this long-lived species, the knowledge
 of its population dynamics, and the use of tools recently developed in the Social
 Network Analysis (SNA) framework, we specifically addressed the following
 questions: 1) is there any long-term social ties between individuals breeding in the same
 patch? and 2) have perturbations, in this case a perturbation regime, affected social
 cohesion? We finally discuss the role and consequences of social cohesion in population
 dynamics and resilience in social species.

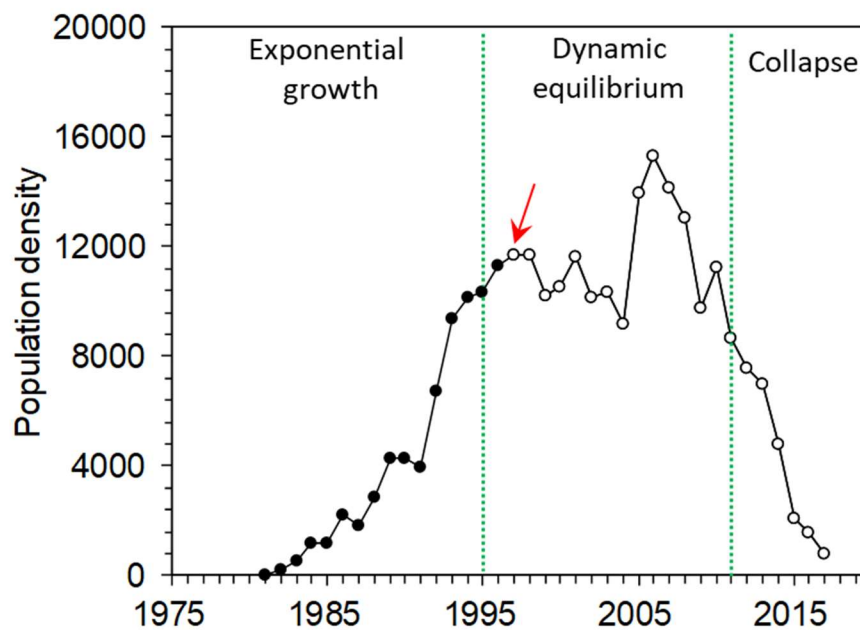


Figure 1. Number of breeding pairs in the Punta de la Banya colony from colonization
 in 1981 to 2017. The observed phases in the population dynamics (growth, dynamic
 equilibrium and collapse) are separated by green lines, which are identified by
 chronological clustering analysis³⁸. Red arrow indicates the arrival of predators to the

colony (open dots for the time series). We compared the period 2002-2010 with the period 2011-2017 corresponding to the dynamic equilibrium and collapse phases respectively.

Results

We analysed a total of 1,610,922 dyadic interactions during the first period (2002-2011) and 368,142 during the second period (2012-2017).

When assessing the social ties with the contingency table approach during the period of stability, the assumption that breeding aggregations in Audouin's gull were at random was rejected, and those individuals that bred together during the sub-period 2002-2006 had a higher probability of breeding together during the sub-period 2007-2011 ($\chi^2_1 = 64.685$, $P < 0.0001$). When randomly reducing sample size of the data set, results were still statistically significant in more than 95% of the cases (1000 randomizations).

Accordingly, when assessing the social ties with dependent regression terms in the AME function, we showed that the probability of breeding together during the second sub-period (2007-2011) depended on whether they have bred previously together in the first sub-period (2002-2006), with a statistically significant coefficient of regression parameter (Table 1; Figure 2). When analysing this data set with permuted data, we concluded that there was indeed a non-random association of individuals within the patches, with our statistic being among the 5% extreme values.

When we analysed the social ties during the transition to collapse phase, we observed that the probability of breeding together during the period 2012-2017 did not depend on whether they have bred together the five previous years ($\chi^2 = 1.957$, p-value = 0.162) and we could not reject the hypothesis of a random association between individuals. In addition, the SNA approach showed that breeding aggregations in Audouin's gull

during the transition phase did not depend on whether they have bred together the five previous years (Table 1; Figure 2).

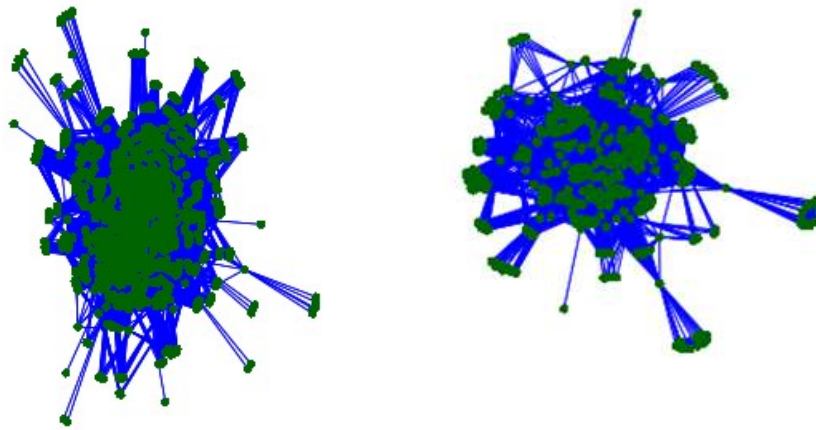
	Stable period				Transition to collapse period			
	pmean	psd	z-stat	p-value	pmean	psd	z-stat	p-value
intercept	-1.467	0.087	-16.808	0.000	-0.231	0.018	-12.629	0.000
.dyad	6.420	0.328	19.576	0.000	-0.004	0.022	-0.188	0.851

Table 1. Results of the AME regression function to test if there were social ties between individuals while breeding during the stable period and during the transition phase to collapse period. The alternative hypothesis is that individuals aggregate annually at random for breeding. “.dyad”: coefficient of the dependent regression term considering the previous dyadic relationship between individuals; “pmean”: posterior mean estimate; “psd”: posterior standard deviation; “z-stat “: nominal z-score.

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a)

b)



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b) **Figure 2.** Graphical representation of social networks by the association

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between individuals of Audouin's gulls in breeding patches comparing a) the

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stability phase (2002-2011) and b) the transition phase to colony collapse (2012-

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2017) (see Figure 1). Each node represents an individual and each edge links

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those individuals that have bred together in the same patch. We used the half-

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weight association index (HWI) to estimate the strength of relationship between

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pairs of individuals an index more suitable when not all individuals within each

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group have been identified ^{42,43}.

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192 **Discussion**

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By studying a particular ecological system of a colonial long-lived species that

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experiences a perturbation regime, we showed that social ties among individuals persist

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over unperturbed years and that perturbations may decrease social cohesion in animal

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populations.

The characteristic breeding behaviour of the study species that aggregates in patches that change annually, allowed us to show that individuals do not annually breed aggregated at random but rather there is some group stability, with individuals establishing social ties that persist over time. Our study system resembles what it was recorded for Slender-billed gulls (*Chroicocephalus genei*), a colonial breeder with weak inter-annual breeding-site fidelity: some individuals bred together across breeding seasons and some social groups showed tenacity despite the colony often moving every year⁴¹. Group stability can emerge as a product of network self-organization, but may provide the necessary conditions for the evolution of other social processes^{44,45}. Our results would support the idea that social aggregation during breeding would provide other advantages than the mere defence against predators^{46,47}, such as social information sharing (e.g.^{48,49}). Social information sharing is crucial for decision-making in risky behaviours, such as dispersal, and previous studies showed that the perturbed regime in our study site caused dispersal to other sites, including colonization of new habitats^{36,50}. In our case study, sociality may have played a major role driving dispersal and thus population dynamics, both during the exponential growth after colonization and the collapse after the perturbation regime. This idea is also reinforced with a mechanistic dynamical model that shows that population dynamics of Audouin's gulls at the study site can only be explained by dispersal runaway caused by social copying¹⁷.

The importance of social information compared to private information increases under perturbations, even when the quality of social information does not increase compared to a non-perturbed regime^{51,52}. For instance, Maldonado et al.⁵³ show that experimental disturbances applied to a social bird may impact its foraging efficiency by social instability caused by the split of social groups. In colonial birds, breeding failure, which

is a proxy of environmental stress, may trigger splitting of the social groups (e.g.⁴¹). At demographic level, the alteration of social network structure may affect the behaviour of populations. For instance, under stress conditions, sociality may operate through feedback loops such as social copying for dispersal, causing non-linear population dynamics and playing a critical role on the resilience of populations (e.g.¹⁷). We showed here that after a perturbation, not only the number of individuals in the population may decrease (by increased mortality or dispersal) but also its social cohesion, likely reducing but also altering the information transfer within the social network composed by those individuals that remain in the site where perturbation occurs. Among other demographic processes, dispersal may alter social connections of both individuals remaining and those dispersing, with consequences for social network structure⁴⁰. The perturbation regime suffered by the study population has likely triggered a social transition⁵⁴ in collective behaviour from philopatric to dispersal and with the fast diffusion of innovations such as the colonization of harbours by large number of individuals, a habitat safe from predators never occupied before⁵⁰. Previous studies have shown that responses of populations to perturbations may also depend on individual personalities in the population⁵⁵⁻⁵⁷. For example, dispersers are different from non-dispersing individuals for a suite of phenotypic traits, including their behavioural profile⁵⁸⁻⁶⁰. Heterogeneities in personalities for dispersal decision-making may have also played a role in our studied population, with most individuals dispersing to other sites after a period of disturbance, while some individuals remaining philopatric. This change may have also further consequences for social network stability, as performance in social groups may improve with heterogeneity in individual personalities^{60,61}.

Our study opens new research avenues about resilience of social populations under perturbations; if perturbations affect social cohesion and heterogeneity in personalities in the population, we may wonder whether this population would be equally resilient to future perturbations. Additionally, in our study population, sociality seemed to operate not right after the first perturbation episode but after cumulative maintained perturbation⁵⁰; would the type of perturbation, either pulse or in regime⁶² influence the response of social groups? We have shown here that a regime perturbation may decrease social cohesion in animal populations, but further studies should be carried out to improve our understanding on the demographic consequences of the breaking down of social ties under perturbations for population dynamics and resilience in social species.

Material and Methods

Study species and study area

The Audouin's gull is a long-lived seabird with more than 80% of the global population breeding in the western Mediterranean (<http://www.iucnredlist.org/details/22694313/0>;³⁷. The species was critically endangered until the early 80's, when it colonized a new site, the Punta de la Banya in the Ebro Delta (Figure 2). Here, the large availability of both suitable breeding habitat and food resulted in a rapid and exponential growth, ending with the site holding more than 70% of the total world population in 2006. The global population dynamics was mainly driven by this colony and after the exponential growth, the species was downgraded to a conservation category of "least concern"⁶³. However, the Punta de la Banya colony is now collapsing and even if the species is colonizing new sites, the global population is decreasing at a 5% annual rate³⁷. In 1997, first carnivores (mainly

foxes, but also badgers, beech martens and least weasels) arrived at Punta de la Banya, and since then the site has been perturbed by the presence of carnivores.

Annual censuses of breeding pairs at every patch within colonies at the Ebro Delta area have been carried out since colonization in 1981 to 2017 (Figure 1, 3 and Table S1). In the Ebro Delta there are three colonies: Punta de la Banya, colonized in 1981 and occupied throughout the study period, Sant Carles de la Ràpita harbour, occupied since 2011 to 2015, and Sant Antoni, occupied from 2013 up to now (Figure 3, Figure S1).

Within colonies, individuals are distributed in patches⁶⁴. As patch location may change from one year to another, we annually geolocalized, mapped and defined the breeding patches (Figure 3 and S2).

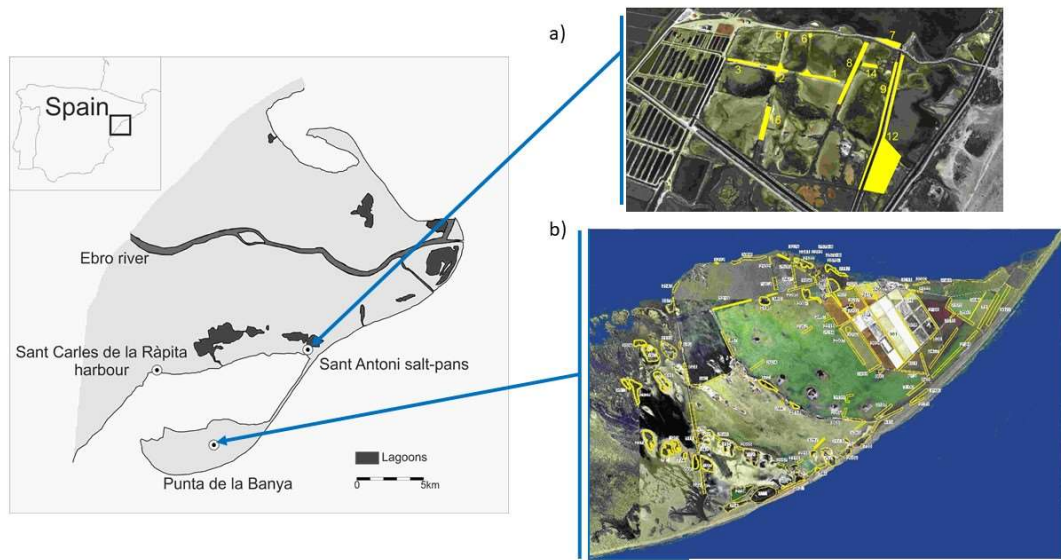


Figure 3. Map of the study area comprising the 3 main colonies and the distribution of patches within colonies during the study period at a) Sant Antoni and b) Punta de la Banya. Sant Carles de la Ràpita colony is considered to have only one patch.

The species is monogamous, there is assortative mating by age, and from an evolutionary point of view is a bet-hedger, laying commonly 3 eggs, although few

chicks survive, except in years with high food availability, when the strength of density-dependence is low ⁶⁵.

Individual data

During 1988-2017 a total of 30,290 individuals were captured and ringed as chicks at the Punta de la Banya ^{66,67}. From 2002 to 2017, resightings were made using spotting scopes from a distance all over the western Mediterranean with a total of 63,106 resights in the study area and 5,593 different individuals resighted. Each year we recorded the breeding patch for each individual. To make sure that individuals were breeding and that they did so in a particular patch, we only selected those individuals seen during the breeding season in a particular patch showing unequivocal breeding behaviour. Specifically, individuals making alarm calls, incubating eggs or with chicks. After this selective filter, our final database included 3,548 individuals.

SNA framework

Our social network, defined as the observed pattern of breeding association, was constructed taking individuals (N=3548) as the nodes of the network and each edge dyad (i.e. pair of individuals) representing the fact that individuals breed in the same patch. We ended with a global sociomatrix, i.e. the matrix representation of the dyadic relationships among individuals, of 3,548* 3,548. Edges showed if two individuals bred in the same patch at least once in a certain period (see below). The network was not directional. Based on previous results on population dynamics, and on the population size of this species and colony ^{37,50}, we divided our dataset in two main periods: a period defined as “stable phase” from 2002 to 2011, and a period of “transition phase to collapse”, from 2012 to 2017 (Figure 1).

We used the recently developed AME function from the AMEN package ^{68,69}, that can be applied to binary, ordinal, and continuous network data. This new approach is a

random-effects regression model; uses an iterative Markov chain Monte Carlo (MCMC) algorithm that provides Bayesian inference of the parameters in the social relations regression model (SRM;⁷⁰) using additive and multiplicative effects and combining the linear regression model with the covariance structure of the SRM⁶⁹. The AME method, not currently used in research on animal social networks is also able to cope with missing and censored data, our data set complying with the assumption that individuals are missing at random. Coping with missing data is highly relevant when analysing sociality on wild populations, as detection rate for individuals is almost always imperfect, and properly controlling for missed observations is a very important step in social network analysis^{71,72}. To create and visualize our networks we used the packages Amen⁶⁸, Asnipe⁷³, gdata⁷⁴ and igraph in R⁷⁵.

Are there social ties that persist over time?

We investigated if individuals create social ties that persist over time longer than one breeding occasion by means of two approaches: i) contingency tables and ii) the inclusion of time dependent regression terms in the AME modelling framework⁶⁸ (see previous section). We used both methods because this is the first application of the AME approach in an ecological context. We analysed data of the period of stability, from 2002 to 2011, dividing this period into two sub-periods of five years (2002-2006 and 2007-2011). In the contingency table approach, we tested if the probability of breeding together at least once during the second sub-period was independent of having bred together at least once in the first period. We built a 3x3 table of frequencies, showing the frequencies of two individuals breeding or not together at least once during the second sub-period depending on whether they bred together or not at least once during the first sub-period, and pulling apart those dyads with missing data. We then tested for deviation of random frequencies by Chi Square test.

With the SNA approach, we analysed the social ties between individuals using the AME function provided in the Amen package in R and including data from the first period (five previous years) as predictors of association during the second sub-period. We considered that this time window was not too large to include important death events, but large enough to account for the imperfect detection of individuals. To achieve convergence, we increased the number of iterations to 100,000 from the default value of 10,000 and lengthened the burn-in period to 500.

Had perturbations affected social cohesion in this species?

To assess if perturbations affected social structure in this species, we analysed as previously, with both the contingency table approach and the SNA approach, the social ties during the period of “transition phase to collapse” (2012-2017). To do so, we tested if the probability of breeding together in this phase (2012-2017) was independent of having bred together during five previous years (2007-2011). We then compared these results from those previously observed during the “stability phase”.

A potential concern was the reduced power during the collapse period because the number of individuals decreased from the stability period. In order to have a similar power in both analyses, we performed the contingency table analysis during the stability period by drawing at random a number of observed associations equal to the number of observed associations during the collapse. We did this repeatedly (1000 times) and calculated which percentage of times the resulting chi-square was significant at the 5% level.

Regarding the SNA approach, it is advised ⁷⁶ to do permutations on the raw data prior to the analysis and compare the result of some relevant statistic obtained with the original data to the distribution of the same statistic over the permutations. We chose the regression coefficient of the association of a dyad on the previous association of the

same dyad as our statistic of social cohesion. This statistic is provided by the function
ame of the package `amen` ⁶⁸. We calculated this statistic on the original data. Then,
within each year, we rearranged randomly the individuals among the patches, keeping
the same number of individuals within each patch. We did this 200 times and calculated
each time the regression coefficient. Then, we situated the value of the regression
coefficient from the original data among the distribution of regression coefficients from
the permutated data and examined how extreme it was. If it was among the 5% extreme
values, we concluded that there was indeed a non-random association of individuals
within the patches.

Acknowledgements

We would like to thank all the people who have helped with the fieldwork in the Ebro
delta over the years, particularly Julia Piccardo, Toni Curcó and the technical staff and
volunteers at the Ebro Delta Natural Park. We would also like to thank Peter Hoff, for
his advices and solving an analytical problem we encountered while using the AMEN R
package. We also thank the Regional Government of Catalonia, for permits to access
the study sites. Elisabeth Rochon corrected the English. Funding came from the Spanish
Ministry of Science (CGL2017-85210), grant PICS INTERACT n°07699 (2016, CSIC-
CNRS). MG was partially supported by the European Union (MINOUW Project,
H2020-634495) and the Spanish Ministry of Science (CGL2017-85210). We have no
conflict of interest to declare.

Authors' contributions

MG conceived the idea; MG, OG, RP and RC designed methodology; MG, DO and AB collected the data; MG analysed the data; MG led the writing of the manuscript. All authors critically contributed to the drafts and gave final approval for publication.

Data accessibility

Data is available via CSIC repository.

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