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Interpretation of interannual variability in long-term aquatic ecological surveys

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Abstract :

Long-term ecological surveys (LTES) often exhibit strong variability among sampling dates. The use and interpretation of such interannual variability is challenging due to the combination of multiple processes involved and sampling uncertainty. Here, we analysed the interannual variability in ~30 years of 150 species-density (fish and invertebrate) and environmental-observation time series in four aquatic-systems (stream, river, estuary, and marine continental shelf) with different sampling efforts to identify the information provided by this variability. We tested, using two empirical methods, whether we could observe simultaneous fluctuation between detrended time series corresponding to widely acknowledged assumptions about aquatic population dynamics: spatial effects, cohort effects, and environmental effects), suggesting that sampling uncertainty overrode the effects of biological processes. Our study does not question the relevance of LTES for detecting important trends, but clearly indicated that the statistical power to interpret interannual variations in aquatic-species densities is low, especially in large systems where the degree of sampling effort is always limited.

Keywords : long-term ecological surveys (LTES), aquatic species (stream -river-estuary- marine), interannual variability, symbolic analyses, sampling efficiency

36 Introduction

Long-term ecological surveys (LTES) are essential to examine the dynamics of ecosystems 37 and identify the processes involved (Hobbie et al. 2003; Magurran et al. 2010; Kuebbing et al. 38 39 2018). Indeed, LTES are key to assess ecological responses (gradual or abrupt) to slow continuous environmental changes (e.g. climatic change; Sommer 2012) and sudden infrequent 40 disturbance events (e.g., hurricanes, fires; Scheffer et al. 2001). This is particularly relevant in 41 the Anthropocene, in which all ecosystems are threatened by a combination of multiple stressors 42 such as climatic shifts, over-exploitation of resources, pollution, habitat degradation, and 43 44 invasion by exotic species (Dudgeon et al. 2006; Halpern et al. 2008; Tilman et al. 2017; Belliard et al. 2018). In this context, LTES are of particular interest for quantifying the 45 46 ecological impacts of current environmental changes and for reliably predicting future impacts 47 (Kuebbing et al. 2018). LTES are thus crucial to provide the knowledge necessary for developing appropriate management tools to mitigate the impacts of anthropogenic pressures 48 on ecosystems (Geheber and Piller 2012; Robertson et al. 2012) and to evaluate the 49 effectiveness of restoration measures (Morandi et al. 2014). Nevertheless, their establishment 50 and maintenance remains highly challenging, partly due to their cost in a context of reduced 51 long-term funding (Lindenmayer 2017), but also due to a bias against the publication of long-52 term research results in favour of short-term "newsy" articles (Kuebbing et al. 2018), and a 53 focus on funding equipment rather than funding long-term research contracts (Lindenmayer 54 55 2018).

Beyond revealing ecological trends or abrupt shifts and associated drivers, most long-term
ecological time-series exhibit strong variability among sampling dates (Knapp and Smith 2001;
Kratz et al. 2003). Understanding this variability is essential to identify the processes governing
population and community dynamics (Lepš et al. 2019). However, the use and interpretation of
such temporal (usually annual) variability is challenging due do the combination of multiple

processes involved and sampling uncertainty. This temporal variability might be related to 61 temporal variability in environmental conditions (e.g., interannual variations in phytoplankton 62 linked to nutrient concentration; Salmaso 2005; 2010), change in biomass of trophic-related 63 64 species (e.g., amphipod oscillation linked to diatom biovolume; Johnson and Wiederholm 1992). Variability could be also linked to variation in data collection (changes in sampling 65 techniques, samplers, sampling timing, sampling area; Trenkel et al. 2004), and sampling 66 inefficiency, i.e. when the study organisms being sampled do not reflect characteristics of the 67 whole population of interest (Peterson et al. 2004; Trenkel and Cotter 2009). Sampling 68 inefficiency depends on sampling techniques, design, and effort but also on the study 69 organisms' dispersal capacity and their distribution area compared to the finite study area 70 71 (Koenig et al. 1996). To analyse and interpret temporal variability in LTES, two approaches 72 could be considered: an empirical approach based only on observational data or a numerical simulation including specific hypotheses concerning population dynamics. When the 73 underlying processes of population dynamics are well identified and sampling noise is reduced. 74 numerical simulation can allow testing alternative hypotheses about the ecological processes 75 driving temporal variability and the effects of sampling uncertainty (Bret et al. 2017; Rivot et 76 al. 2004). However, in the case of LTES influenced by multiple processes and with unknown 77 sampling noise, such as LTES in large aquatic systems (e.g., river, estuary or marine continental 78 shelf), an empirical approach is more suitable and may help identify the most relevant 79 ecological processes involved. 80

In this study, we used an empirical approach to identify and interpret the information provided by the interannual variability in a unique data set grouping four independent LTES of aquatic species in France: 1) trout populations in small streams, 2) fish populations in a large river, 3) fish and invertebrate populations in an estuary, and 4) fish and invertebrate marine populations on the continental shelf (Fig. 1). These four LTES include relatively long-term (~ 86 30 years) annual observations of both multiannual aquatic species density with annual reproduction cycles (with an identification of age classes) and environmental measurements at 87 various reaches Moreover, the datasets were characterized by various degrees of sampling 88 89 effort, defined here as the percentage of the study area sampled (sampled area divided by the total study area): low for the three large aquatic systems (river, estuary and marine continental 90 shelf) and high for streams. Previous analyses of these LTES revealed trends and abrupt shifts 91 in community composition and certain aquatic species related to environmental changes. For 92 example, Daufresne et al. 2015 showed a positive effect of restored flows on the proportion of 93 midstream fish in the large river; Chaalali et al. (2013) showed negative effect of water 94 warming on shrimp reproduction in the estuary; and Poulard and Blanchard 2005 reported a 95 96 positive effect of sea temperature increase on subtropical species (Table 1). Here, we did not examine trends but rather focused on understanding interannual variability and potential 97 98 drivers.

We thus examined the shape of the time series after removing the long-term trend (Lepš et al. 99 2019) and compared the shapes among time series to identify whether we could detect similarity 100 in shape among time series; i.e. simultaneous year-to-year fluctuations. For the four LTES, 101 characterized by a range of sampling efforts, we tested whether we could observe simultaneous 102 fluctuations between time series corresponding to widely acknowledged assumptions about 103 aquatic population dynamics: 1) spatial effects, spatial correlation between densities of 104 individuals (especially for fish juveniles) between pairs of neighbouring reaches, i.e. reaches 105 106 with similar environmental conditions and close enough to allow dispersal between them 2) cohort effects, temporal correlation between the densities of juveniles and the densities of older 107 individuals with a one-year time lag (only for fish) and 3) environmental effects, correlation 108 109 between aquatic species (or guild) densities and the main environmental drivers (identified in previous studies). For spatial effect, we expected a positive association between the interannual 110

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111 variability of aquatic-species densities from neighbouring reaches; time series should fluctuate simultaneously, rising and falling together. For cohort effects, we expected a positive 112 association between the interannual variability of fish density of juveniles at year t and older 113 individuals at year t+1; time series of fish densities of juveniles and older individuals should 114 fluctuate together in the same direction with a one-year time lag. For environmental effects, we 115 expected a positive association between the interannual variability of environmental and 116 ecological time series when the sign of the relationship between species (or guild) densities and 117 environmental parameters was positive, and negative otherwise; time series should fluctuate 118 119 together in the same direction when the sign of the relationship was positive, and antagonistically (one rising and the other falling at the same time) otherwise. For this, we 120 121 applied two empirical methods: a residual Pearson correlation analysis and a symbolic analysis 122 based on Information Theory.

123 Methods

Data description

Trout populations in small streams

This dataset included 5-18 years of brown trout (Salmo trutta) time series sampled between 126 1990 and 2014 in 42 stream reaches distributed across 23 streams in France (Fig. 1). At each 127 stream reach (~100 m²), trout density (number of individuals per m²) was estimated by counting 128 all individuals sampled using a two-pass removal electrofishing technique, without block nets 129 (Fig. 1). Each year, the whole reach ($\sim 100\%$ of the study reach) was electrofished in September 130 by various operators following the European Committee for Standardization guidelines (CEN 131 2003; Bret et al. 2015). Based on their size, trout individuals were separated in the field into 132 three age classes: young-of-the-year (age-0), one-year-old fish (age-1), and 2 years and older 133 (age-2⁺). This dataset also included daily water temperature and flow velocity time series for 134 each stream reach (see Bret et al. 2015 for more details on the environmental data). Spatial 135

effect was examined between the closest stream reaches (< 5 km apart) not disconnected bydams.

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Fish populations in a large river

This dataset included 18-34 years of freshwater fish time series sampled between 1979 and 140 2015 in six river reaches situated along the Rhône River (Fig. 1). At each river reach (~1 km²), 141 fish density (number of individuals per m²) was estimated based on electrofishing using either 142 point abundance sampling (between 20 and 200 sampling points with an average area of 7m² 143 or 32m² depending on the study reach) or continuous sampling along the banks during 30 min 144 (Fig. 1). Each year, around 0.1% of the study reach was sampled in fall by various academic 145 146 teams (see Daufresne et al. 2015; Lamouroux et al. 2015 for more details on the dataset). Based on their size, fish individuals (for each species) were separated in the field into two age classes: 147 young-of-the-year (age-0), and one year and older (age-1⁺). Among the 41 species sampled 148 within the six river reaches, we removed species that occurred in less than 30% of all samples 149 150 (reaches and years combined), thereby preserving 16 species. For the analysis of both spatial effect and cohort effect, we only considered, for each river reach, species occurring in more than 151 30% of all sampling dates (see Table S1 for the resulting number of studied species per 152 analysis). Spatial effect was analysed for the closest river reaches (< 35 km apart) with similar 153 environment and not disconnected by dams. For the analysis of *environmental effects*, the 16 154 selected species were grouped into three guilds, according to their habitat preferences following 155 156 Daufresne et al. (2015), among which we examined lithophilic, midstream (species preferring deep and fast water), and southern guilds. This dataset also included daily water temperature 157 and discharge time series for each river reach (see Daufresne et al. 2015; Maire et al. 21019 for 158 159 more details on the environmental data).

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Fish and invertebrate populations in an estuary

This dataset included 24-30 years of estuarine-species time series sampled between 1979 162 and 2014 in four estuary reaches situated along the Gironde estuary (Fig. 1). At each estuary 163 reach (~5 km²), estuarine fish and invertebrate density (number of individuals per m³) were 164 estimated based on point abundance sampling (three sampling points along one transect; 165 166 Pasquaud et al. 2012). At the three sampling points (one close to each bank and one on the middle axis of the estuary), surface and bottom samples were taken simultaneously using, 167 respectively, two 4.0 x 1.0 m rectangular frame nets, fitted at either side of the boat, and a 2.0 168 169 x 1.2 m frame trawl, kept at 0.2 m above the bottom by skates. Sampling lasted around 7 min and was performed during daytime, between the mid-discharge and the slack period of high 170 171 tide, with the gear being towed against the current (Fig. 1). Each year, around 0.01% of the 172 study reach was sampled every month by academic teams (see Lobry et al. 2006 for more details on the dataset). For each species, we calculated a mean autumn density by averaging the 173 densities of the September, October, and November samples. Based on their size, fish 174 175 individuals (for each species) were separated in the field into two age classes: young-of-theyear (age-0), and one year and older (age-1⁺). Among the 59 species sampled within the four 176 estuary reaches, we removed species that occurred in less than 30% of all samples (reaches and 177 years combined), thereby preserving 16 species. For the analysis of both *spatial* and *cohort* 178 *effects*, we only considered, for each estuary reach, species occurring in more than 30% of all 179 sampling dates (see Table S1 for the resulting number of studied estuarine species per analyses). 180 Spatial effect was examined between each pair of the four estuary reaches (< 20km apart). For 181 the analysis of environmental effects, the 16 selected species were grouped into four guilds, 182 according to their habitat preference following Lobry et al. (2006), among which we examined 183 184 freshwater and marine guilds. This dataset also included daily water temperature, salinity, and

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Marine fish and invertebrate populations on continental shelf

This dataset included 26 years of marine-species time series sampled between 1987 and 189 2014 on the eastern continental shelf of the Bay of Biscay (Fig. 1). The study area extended 190 from 47°53'12.01"N - 5°38'35.99"O to 43°41'15.00"N - 1°13'14.99"O, covering around 75, 574 191 km² between 15 and 600m depth. Fish and invertebrate density (number of individuals per km²) 192 and biomass (kg per km²) were estimated based on a stratified groundfish survey, using a 36/47 193 GOV trawl with a 20 mm mesh codend liner. Each year, between 56 and 154 hauls spread over 194 195 the study area (30 min samples at a towing speed of 4 knots), i.e. around 0.007% of the study 196 area, were collected in November by Ifremer (see Poulard and Trenkel 2007 for more details on the dataset; EVHOE survey, Fig. 1). We computed density and biomass time series of the 197 entire Bay of Biscay by averaging across all hauls. Among the 296 marine species sampled 198 including both fish and invertebrates, we conserved only 32 species, i.e. the species occurring 199 at more than 30% of all sampling dates. For the analysis of environmental effects, the marine 200 selected species were separated into three functional groups according to their biogeographical 201 affinity (temperature preference) following Quéro (1984), among which we examined boreal 202 and subtropical groups. This dataset also included daily sea surface temperature and annual 203 fishing pressure time series, estimated using commercial catch data obtained from the data base 204 205 held by the International Council for the Exploration of the Sea (http://www.ices.dk/marinedata/dataset-collections/Pages/Fish-catch-and-stock-assessment.aspx). For the analysis of 206 spatial effect, as the environmental fluctuations were relatively homogeneous within the Bay 207 208 of Biscay and most studied marine species exhibit large scale dispersal, we split the Bay of Biscay into two zones (northern and southern) separated perpendicularly to the coast to display 209

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similar bathymetric features (Poulard and Blanchard 2015). For both zones, we computed marine species density time series by averaging the densities recorded in all hauls from each zone (equivalent to reach in the three other LTES). Note that zone limits varied among species according to their spatial distribution to maintain a similar number of occurrences in each zone (see Fig. S1 for the zone delimitation by species).

Data analyses

We used two different methods to examine the existence of environmental effects, cohort 217 effects, and spatial effects in the detrended time series. The first method (described below) 218 consisted in describing the rhythm of each time series using symbolic transformation (Fig. 2 a) 219 220 and techniques from Information Theory, and quantifying the association between the rhythms of the different time series, i.e. the degree to which time series tended to oscillate together, 221 rising and falling with the same frequency (Cazelles 2004; Descamps et al. 2010). Previous 222 studies already tested the application of this method on ecological data, compared their results 223 with cross-correlation techniques, and concluded this method improved the detection of weak 224 relationships between time series, even short ones and in the presence of noise (Cazelles 2003; 225 2004). We compared the results of this method with a second more conventional approach, 226 which consisted in fitting a LOESS model to each time series and then calculating the Pearson 227 correlation among detrended LOESS residuals (linear model, Fig. 2 b). 228

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Symbolic transformation

To describe the rhythm of continuous time series, we discretized them into a sequence of a finite number of symbols (Cazelles 2004). As our objective was to identify the main drivers of interannual variability in ecological time series, we used a simple discretization based on three symbols ("Flat", "Increase", and "Decrease") reflecting changes between consecutive values (Fig. 2 a). Therefore, continuous time series *X* of length *n* were transformed into discrete sequences *S* of length *n*-1. First, for each species, guild or environmental parameter, we computed the series Δ of length *n*-1 defined as $\Delta_t = X_t - X_{t-1}$, and computed two thresholds *I* (for increase) and *D* (for decrease) calculated as the standard deviation of positive values of Δ , and the standard deviation of negative values of Δ , respectively.

240 Then, given a time series *X*, and the corresponding thresholds *I* and *D*:

if $X_{t-1} < X_t$ and $X_t - X_{t-1} > I$, S_t was coded 'Increase'

242 if $X_{t-1} > X_t$ and $|X_t - X_{t-1}| > D$, S_t was coded 'Decrease'

243 if $|X_t - X_{t-1}| \le D$ or $|X_t - X_{t-1}| \le I$, S_t was coded 'Flat'.

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Mutual information

To quantify the degree of association between two time series, we calculated the mutual information between the corresponding symbolic sequences (Pompe 1993; Prichard and Theiler 1995). The mutual information I_{SIS2} between the symbolic sequences *S1* and *S2* is defined as: $I_{S1S2} = H_{S1} + H_{S2} - H_{S1S2}$

where H_{SI} and H_{S2} are the entropy of the symbolic sequences *S1* and *S2*, and H_{SIS2} is their joint entropy. The entropy of a symbolic sequence S is defined as:

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$$H_S = -\sum_{i=1}^{k} P(S_i) \log (P(S_i))$$

where $P(S_i)$ is the probability of observing the symbol $S_i \in$ ("Increase", "Decrease", "Flat"), estimated as the proportion of the symbol in the sequence, and k is the number of different symbols in the sequence (k = 3 in our case). The joint entropy H_{SIS2} between two symbolic sequences *S1* and *S2* is defined as:

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$$H_{S1S2} = -\sum_{i=1}^{k} \sum_{j=1}^{k} P(S1_i, S2_j) \log (P(S1_i, S2_j))$$

where $P(S1_i,S2_j)$ is the probability of observing simultaneously the symbols SI_i and $S2_j$, estimated as the proportion of pairs $(SI_i, S2_j)$ observed in the sequence.

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Statistical significance level

To quantify the statistical significance of this measure of common information between 262 two time series, we compared the observed mutual information with those obtained with n =263 1000 couples of randomly-generated time series (Cazelles 2004). These randomly-generated 264 time series were built under the null hypothesis of no association between two observed time 265 series. The randomly-generated time series Z were obtained by resampling the observed time 266 series based on a Markov process scheme that preserved the short-term temporal correlations 267 268 of the series (Cazelles et al. 2014). The randomly-generated time series were thus computed in the following way (Cazelles and Stone 2003; Descamps et al. 2010): 269

1. A frequency histogram with n_b equal-sized intervals was created from the values of the observed time series *X* to estimate its distribution ($n_b = 3$ for time series < 30 years);

272 2. A transition matrix describing the time evolution of the series was created based on the 273 frequencies of the data contained within each interval. The transition matrix gives the 274 probability of moving from values that are within interval b_i at time *t* to values that are within 275 interval b_i at time t+1;

276 3. An initial value Z_0 was randomly selected from the observed time series X;

4. To determine Z_{t+1} , we randomly selected the interval b_j to which Z_{t+1} belonged based on the interval b_i of Z_t , and the probabilities from the transition matrix. The value of Z_{t+1} was then randomly selected among all the elements of interval b_i ;

5. The previous step was iterated to obtain a randomly-generated time series *Z* of the samelength as the observed time series *X*.

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282 We then transformed the 1000 randomly-generated time series Z into 1000 randomlygenerated symbolic sequences, and computed the distribution of the 1000 randomly-generated 283 mutual information values under the Null hypothesis of independence. We then compared the 284 285 mutual information of the observed time series (I_{obs}) with the 95th percentiles of the distribution of the 1000 randomly-generated mutual information (I_{sur}) . We determined a p-value for the 286 287 association using a procedure based on ranks; i.e. p-value was defined as 1 minus the proportion of randomly-generated mutual information values exceeded by the observed mutual 288 information value (the rank of Iobs relative to all randomly-generated mutual information 289 values). All analyses were performed in R using the package synchrony and infotheo. 290

Mutual information allowed quantifying the degree of association between two time series, but without any assumption regarding the sign of the relationship between the two time series. Thus, for significant associations, we verified that joint probabilities contributing to the mutual information corresponded to our ecological hypotheses. Similarly, we verified whether the signs of the linear models between residuals corresponded to our ecological hypotheses (see the hypotheses section and Table 1).

For comparative purposes, all ecological time series were standardized by dividing all 297 values by the respective mean prior to symbolic transformation. For each dataset, our 298 hypotheses were generic across all reaches for both *environmental* and *cohort effects*, and across 299 all pairs of neighbouring reaches for *spatial effects*. For example, for a given species, a *cohort* 300 *effect* should occur at all reaches. To increase the statistical power of our tests for shorter time 301 series, we concatenated, separately for each dataset, the time series of all reaches (for the 302 Stream, River and Estuary datasets) to test both cohort and environmental effects; and we 303 concatenated the symbolic sequences of all pairs of neighbouring reaches for the Stream and 304 305 *River* datasets to test *spatial effect* (see the description of hypotheses for details). Note that randomly-generated time series were independently computed for each reach, independently 306

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transformed into randomly-generated symbolic sequences, and concatenated afterwards. For both *spatial* and *environmental effects*, association between time series was tested with no time lag, except for one hypothesis for the *Marine* dataset (see the hypotheses description and Table 1). In our analyses, significance thresholds were not corrected for multiple tests. Instead, we considered the number of positive tests as evidence for the validity of the tested hypothesis. In results, we provided $I_{obs} - I_{sur}$ and the associated P-value for the symbolic analysis and R² and the P-value for the residual approach.

Description of hypotheses

Trout populations in small streams

We tested spatial, cohort, and environmental effects for the trout populations. For the 316 317 spatial effect, we concatenated the time series of age-0 trout density of all pairs of neighbouring reaches. For cohort and environmental effects, we concatenated all (42 reaches) time series of 318 trout density (for both age-0 and age-1) and environmental conditions. For environmental effect, 319 we expected annual median of mean-daily water-temperature and age-0 trout-density time 320 321 series to fluctuate simultaneously in the same direction, while annual median of mean-daily water-temperature and age-1 trout-density time series, and flow velocity during emergence 322 period and age-0 trout-density time series should fluctuate antagonistically (Table 1). 323

Fish populations in a large river

We tested *spatial*, *cohort*, and *environmental effects* for various fish populations. For the *spatial effect*, we concatenated the time series of age-0 fish density for six species (Table S1) of all pairs of neighbouring reaches. For *cohort effect*, we concatenated all (six reaches) time series of age-0 and age-1⁺ fish density for 16 species (Table S1). For *environmental effect*, we concatenated all time series of age-0 fish, age-0 lithophilic-fish, southern-fish, midstream-fish densities, and environmental conditions. We expected the time series of the number of days with high flows during cyprinid reproductive period to fluctuate antagonistically with age-0

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fish and age-0 lithophilic-fish densities (Table 1). We expected southern-fish densities to fluctuate simultaneously with time series of annual maximum of mean daily water temperature and time series of mean water temperature during reproductive and growth periods. Finally, we expected midstream-fish densities and annual low-flow hydraulic conditions time series to fluctuate simultaneously.

Fish and invertebrate populations in an estuary

We tested *spatial*, *cohort*, and *environmental effects* for both fish and invertebrate 338 populations. For the spatial effect, we examined the fluctuation of density time series among 339 each pair of estuary reaches for 13 species (Table S1). For cohort effect, we concatenated all 340 (four reaches) time series of age-0 and age-1⁺ fish density for four species (Table S1). For 341 342 environmental effect, we concatenated all time series of environmental conditions, and expected mean annual salinity time series to fluctuate simultaneously with marine-species densities, and 343 antagonistically with freshwater-species and Palaemon longirostris densities (Table 1). We also 344 expected mean annual salinity time series to fluctuate simultaneously with Argyrosomus regius, 345 Engraulis encrasicolus, and Dicentrarchus labrax densities, and antagonistically with 346 Palaemon longiros densities and the percentage of shrimp ovigerous female. 347

Marine fish and invertebrate populations on continental shelf

For this dataset, we could only test the spatial and environmental effects for the marine 349 populations. For the *spatial effect*, we examined the fluctuation of density time series between 350 the northern and the southern zones for 32 species (Table S1, Fig. S1). For environmental effect, 351 we expected mean annual sea surface temperature time series to fluctuate simultaneously with 352 subtropical-species and antagonistically with boreal-species densities (Table 1). We also 353 expected a simultaneous fluctuation between the annual commercial catch and scientific survey 354 355 biomass. Finally, we expected an antagonist fluctuation between the annual commercial catch biomass at year t and the scientific survey biomass at year t+1. 356

358 **Results**

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Trout populations in small streams

360 Using the residual Pearson correlation approach and the symbolic analysis, we found significant spatial effect between interannual variation of age-0 trout densities of neighbouring 361 reaches (Table 2). Indeed, using the symbolic analysis, we found a significant association 362 between variations in age-0 trout time series; i.e. more than what could be expected by chance 363 (I_{obs} - I_{sur} = 0.30, P < 0.001, n = 35), with joint probabilities contributing to the mutual 364 information corresponding to our hypothesis (age-0 trout density in neighbouring reaches 365 oscillated together, rising and falling simultaneously). Similarly, using the residual analysis, we 366 367 found a positive significant relationship between residuals of age-0 in neighbouring reaches (R² = 0.48, P < 0.001). We also found a significant *cohort effect* (association between variation in 368 age-0 trout density at year t and age-1 trout density at year t+1) using both methods ($I_{obs} - I_{sur}$) 369 = 0.2, P < 0.001, n = 394, using the symbolic analysis with the joint probabilities contributing 370 to the mutual information for analogous symbols; and $R^2 = 0.44$, P < 0.001 using the residual 371 analysis). However, we found no significant environmental effects as interannual variations in 372 environmental drivers were not linked with interannual variations of trout densities (I_{obs} - I_{sur} < 373 0 for the symbolic analysis, and $R^2 < 0.025$ for the residual analysis, n = [405-521] for three 374 environmental drivers, see table 1). 375

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Fish populations in a large river

Among the five tests of *spatial effect* performed for the dominant species (*Alburnus alburnus*, *Squalius cephalus*, *Rutilus rutilus*, *Gobio gobio*, and *Chondrostoma nasus*), we found a significant relationship between temporal variations of age-0 densities of neighbouring reaches only for gudgeon (*Gobio gobio*) based on both the mutual information approach (I_{obs} -

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I_{sur} = 0.029, P = 0.023, n = 32) and the residuals approach ($R^2 = 0.22$, P = 0.04, Table 2). For the 16 tested species, we only found a significant *cohort effect* for pumpkinseed (*Lepomis gibbosus*) based on both the mutual information approach ($I_{obs} - I_{sur} = 0.011$, P = 0.038, n = 104) and the residuals approach ($R^2 = 0.36$, P < 0.001). The results for the *environmental effects* hypotheses showed no significant associations between variations in environmental drivers and variations in guild densities with either method ($I_{obs} - I_{sur} < 0$, P > 0.05, n=1130 for the symbolic analysis; $R^2 < 0.2$ for the residual analysis).

Fish and invertebrate populations in an estuary

Among the 78 tests performed (13 selected species among 6 pairs of reaches), we found 28 391 significant spatial effects for interannual variations of age-0 densities of neighbouring reaches 392 based on the mutual information approach (I_{obs} - $I_{sur} > 0$, P < 0.05, n = [24-30], Table 2). Note 393 that we obtained 35 significant associations, but for seven tests, the joint probabilities 394 contributing to the mutual information did not correspond to our hypothesis of positive 395 associations. In particular, results showed significant spatial effects among most pairs of 396 reaches for Alosa alosa, Alosa fallax, Anguilla anguilla, and Sprattus sprattus. With the residual 397 approach, we obtained 56 significant positive relationships ($R^2 > 0.2$, P < 0.05), among which 398 35 with $R^2 > 0.4$. We found no significant *cohort effect* at time lag 1 (I_{obs} - $I_{sur} < 0$, P > 0.05, n 399 = 98 for the symbolic analysis; $R^2 < 0.2$, P > 0.05 for the residual analysis) and no *environmental* 400 effects (I_{obs} - I_{sur} < 0, P > 0.05, n = [72-96] for the symbolic analysis; R² < 0.2, P > 0.05 for the 401 residual analysis). 402

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Marine fish and invertebrate populations on continental shelf

Among the 32 dominant species, we found 12 significant *spatial effects* ($I_{obs} - I_{sur} > 0$, P < 0.05, n = 26 for the symbolic analysis, Table 2). For four other species, the association was

407 significant but the joint probabilities contributing to the mutual information did not correspond to our hypothesis. Using the residual method, we found significant positive correlations 408 between the residuals of both zones with R² value higher than 0.2 for 18 species, and higher 409 than 0.4 for 10 species. We found no significant correlation between interannual variations in 410 sea surface temperature and interannual variations in subtropical or boreal guild densities using 411 both approaches. Among the 17 dominant commercially exploited species, we found no 412 significant association between interannual variations in commercial catches and interannual 413 variations scientific survey biomass at time lag 0. In contrast, we found a significant negative 414 relationship between the residuals of both commercial catches and scientific biomass at lag 1 415 for *Capros aper* ($R^2 = 0.22$, P = 0.027 for the residual analysis). 416

418 Discussion

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A relatively low number of significant associations

In this study, we analysed more than 150 aquatic-ecological time series from four 420 independent datasets to examine the information provided by the short term (interannual) 421 temporal variability in long-term aquatic ecological time series. Overall, we identified few 422 significant associations between the interannual variability of the different time series, both 423 environmental and ecological, and thus did not find all expected patterns resulting from 424 population dynamics. For all datasets and using two analytical methods, we found no significant 425 association between the short-term variations in environmental parameters and those of species 426 427 or guilds densities, although we tested environmental parameters that have been previously identified as long-term drivers of aquatic population dynamics (e.g., Bret et al. 2015; Daufresne 428 et al. 2015; Maire et al. 2019). Similarly, we found only few significant tests for cohort effects 429 430 that is an association between the interannual variations in densities of juveniles and older individuals at a one-year lag: only 2 of the 21 tests performed (9%) were significant, while 431

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cohort tracking has been commonly observed in fish populations (e.g., Cattanéo et al. 2002;
Lobón-Cerviá et al. 2012; Bret et al. 2015). Nevertheless, for *spatial effects*, among the 116
tests performed, 36% were significant based on the symbolic approach (39% based on the
residuals approach). The higher number of significant tests for *spatial effect* seems consistent
as the densities of individuals sampled in neighbouring sites, with similar environmental
conditions and located close enough to allow dispersal between them, should oscillate
simultaneously.

Interannual variability in aquatic ecological observations due to uncertainty in density estimates

442 We hypothesize that the overall low number of significant results was mainly linked to uncertainty in density estimates due to high sampling inefficiency and variable bias between 443 sampling dates; i.e. variable proportion of the population sampled on each survey. More 444 generally, because the entire population is rarely sampled in ecological studies, the proportion 445 of the population captured (the catchability or capture probability) strongly depends on 446 sampling effort, i.e. the area (or volume) sampled compared to the area (or volume) over which 447 the population is distributed (Hilborn and Walters 2013). This might explain why we found 448 significant spatial effects and cohort effects for the Stream dataset for which the whole reach 449 was sampled, while little for the other datasets for which less than 0.1% of the study area was 450 451 sampled.

In addition, catchability can also vary over time and space although it is often assumed to be constant (Korman and Yard 2017). Variation in catchability depends on the temporal and spatial variation in the efficiency of the sampling gear used to capture individuals, and how sampling points are distributed relative to the distribution of individuals (Trenkel and Cotter 2009). Aquatic populations are usually distributed unevenly over different habitats within the Can. J. Fish. Aguat. Sci. Downloaded from www.nrcresearchpress.com by IFREMER BIBLIOTHEQUE LA PEROUSE on 01/31/20 For personal use only. This Just-IN manuscript is the accepted manuscript prior to copy editing and page composition. It may differ from the final official version of record.

457 study area (Ahmadi-Nedushan et al. 2006; Mérigoux et al. 2009), and the spatial distribution of 458 habitats varies over time according to environmental conditions (Korman and Campana 2009; Dunbar et al. 2012; Capra et al. 2017). Thus, catchability could also be affected by the 459 460 environmental conditions prior and during the surveys. For example, previous studies showed that spatial distribution of aquatic organisms was influenced by high wind velocity and 461 462 associated marine currents in the ocean (Poulard and Trenkel 2007), high turbidity in rivers as the associated reduction in predation risk modified fish behaviour (Dodrill et al. 2016; Ward et 463 al. 2016) or changes in flow and associated modifications in the proportion and spatial 464 465 distribution of habitats (Girard et al. 2014). These environmental conditions could have also affected the sampling capacity; e.g., high wind velocity modifies ship motion and trawl 466 467 geometry (Poulard and Trenkel 2007), and high discharge increases the size of the study area 468 (or volume), and consequently reduces sampling effort.

Moreover, catchability strongly depends on species traits and dispersal capacity (Trenkel 469 and Cotter 2009). In particular, many aquatic species are gregarious (e.g., especially cyprinid 470 species as Alburnus alburnus, Chondrostoma nasus; Vaudor et al. 2011), thereby not evenly 471 distributed over the study area (spatial aggregation). Spatial distribution and behaviour of 472 aquatic species can also vary with age and size; e.g. larger fish individuals are often distributed 473 at greater depths and further from the shore compared to smaller individuals (Vehanen et al. 474 1999; Woillez et al. 2007). This variation of catchability with age and size might explain the 475 low number of significant *cohort effect* detected. The efficiency of a sampling design (gear. 476 477 timing, spatial distribution of effort) to capture individuals also varies with organism sizes and ages (Borgstroem and Skaala 1993; Korman et al. 2009). For example, the EVHOE survey 478 (Marine dataset) was designed for catching young (small) organisms and hence was less reliable 479 480 for estimating the density of larger individuals (Trenkel and Cotter 2009). This might explain 481 why we found almost no significant association between scientific and commercial catch482 biomass as commercial fishing mainly targeted larger fish.

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Complex interacting processes prevent the understanding of population dynamics at the annual time scale, even using 30-years of annual surveys

In addition to the uncertainty in density estimates, there are additional explanations for the 486 overall low number of significant results. First, the lack of significant environmental effects 487 might be because the environmental drivers selected for this study were not appropriate to 488 explain the observed interannual variations although relevant for long-term trends, e.g., 489 significant increase in southern fish density with global warming (Daufresne et al. 2015), and 490 491 decrease in shrimp density with estuary salinization (Béguer et al. 2012). Moreover, the mechanisms driving interannual variations were probably more complex than those driving 492 long-term trends and might involve a combination of interacting environmental parameters 493 (e.g., substrate quality and clogging, temperature rapid variations, short-term flow variations) 494 at different phases of the life cycle, in particular during the first developmental stages of larvae 495 (Schiemer et al. 2003) and recruitment of juvenile fish (Myers 1998; Grenouillet et al. 2001) 496 but also during the period of reproduction and incubation (Korman et al. 2011). Second, the 497 lack of consistency between the densities of juveniles and older individuals the following year 498 might be due to high levels of dispersion, and/or high levels of mortality, linked to unknown or 499 unmeasured environmental drivers, biotic interactions between and within species (predation, 500 competition, facilitation) or endogenous regulatory processes such as density-dependent 501 regulation (Grenouillet et al. 2001; Daufresne and Renault 2006; Lobón-Cerviá 2013). The lack 502 of significant cohort effects could also be due to an inappropriate assignment of age classes. 503 For both *River* and *Estuary* datasets, and for long-lived fish, the age class age-1⁺ aggregated 504 various cohorts, potentially affected by different environmental conditions. Third, non-505

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significant *spatial effect* tests could be also due to the definition of neighbouring reaches, probably not similar and/or close enough to host the same population. Indeed, for the *Estuary* dataset, we found more significant results among the closest stations (< 5 km), than among the most distant ones (>20 km). Similarly, Bret et al. (2015) found a decrease in *spatial effect* among age-0 trout densities with increasing distance between streams.

Methodological limitations

We could also question the relevance of symbolic analyses to assess the relationships 513 between the interannual variability in environmental and ecological time series, as well as the 514 relevance of the selected rules at the different steps of the analysis (i.e. symbolic transformation 515 516 and statistical significance level). However, we repeated our analyses using different number 517 of symbols (from 3 to 5), different threshold values (I and D, see methods), different transformation rules (coding peaks and troughs instead of increases and decreases, Cazelles et 518 al. 2004; or coding low, medium, and high density values); and using different procedures for 519 520 randomly-generated time series computation (e.g. preserving or not temporal dependencies; see Theiler et al. 1991). These complementary analyses led to similar conclusions (results not 521 shown). Consistently, we obtained comparable results with the symbolic and residuals 522 approaches, suggesting that our results are largely independent of the analytical method of our 523 empirical approach. Besides, numerical simulations of aquatic-species density time series based 524 on known ecological patterns, associated with tests of the statistical power of our two analytic 525 methods to detect association between interannual variability of the different time series, could 526 be an alternative approach. However, such numerical simulations are not straightforward, as 527 illustrated for the River dataset by Vaudor et al. (2015), but rather require to make strong and 528 529 questionable assumptions about numerous ecological patterns and processes (e.g., dispersal ability of the different species, strength and direction of environmental effects, and density-530

536 Conclusion

Our study clearly indicates that the statistical power to identify and interpret significant 537 interannual variations in aquatic species densities is low, especially in large systems where a 538 complete sampling of populations is unrealistic and sampling effort is limited. Although the 539 significant spatial effects observed in our study suggests that interannual variations contains 540 541 some valuable information of similarities in population dynamics, whose environmental drivers are probably multiple and not clearly identified yet; our results point out that moderate and/or 542 isolated annual ecological changes cannot be revealed by the analysis of interannualvariations 543 in such LTES. 544

Our study is focused on interannual variations and does not question the relevance of 545 LTES for detecting important trends or shifts as demonstrated in previous studies. However, 546 interannual variability linked to sampling inefficiency lowers the statistical power to identify 547 trends in large systems, supporting that long time series are required for detecting trends (Hatch 548 2003; Nicholson and Jennings 2004). Consistently, Vaudor et al. (2015) showed, when studying 549 the effects of major flow restoration measures with the *River* dataset, that the probability of 550 551 detecting moderate changes (50-200 %) in population density after a restoration event averaged 16%, compared to 61% for large changes (500-1000 %), and that the number of surveys well 552 balanced before and after restoration increased this probability. 553

554 Improved analyses of the drivers of short-term population dynamics would require more 555 exhaustive and specific monitoring methods, involving intensive sampling at a finer temporal

scale. Alternative approaches would be to combine LTES with intensive continuous sampling 556 during specific periods corresponding to particular life stages (e.g., fish recruitment), or 557 measure population dynamic through individual tracking using telemetry devices or mark-558 recapture techniques to estimate the probability of detection of species, thereby assessing the 559 observation error of the LTES for a given population (Capra et al. 2018; Radinger et al. 2019). 560 Finally, developing and improving population dynamic models, also using LTES (and 561 combining spatial replications of LTES; Clark et al. 2015) to test the relevance of simulations, 562 should also be encouraged; however, the requirement of continuous sampling and/or specific 563 experiments under controlled conditions to calibrate the model drivers should not be 564 overlooked. 565

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Table 1: Summary of the hypotheses for environmental effects. This table summarize the relationship (including the sign) between

aquatic species (or guild) and environmental parameters observed in previous studies examining the same datasets.

Datasets	Environmental parameters	Ecological data	Expected sign of the relationship	References		
Stream	Annual median of mean daily water temperature	age-0 trout d.	+			
	Annual median of mean daily water temperature	age-1 trout d.	-	Bret et al. 2015		
	Flow velocity during emergence period (March-April)	-				
	Nbr. of days with high flows during cyprinid reproductive period (March-June)	age-0 fish d.	-			
	Nbr. of days with high flows during cyprinid reproductive period (March-June)	age-0 lithophilic-fish d.	-	Development of al 2015		
River	Annual maximum of mean daily water temperature	southern-fish d.	+	Daulfesne et al. 2015		
	Mean water temperature during reproductive and growth periods (April-August)	southern-fish d.	+	Lamouroux et al. 2015		
	Annual low-flow hydraulic conditions	midstream-fish d.	+			
	Mean annual salinity	marine-species d.	+			
	Mean annual salinity	freshwater-species d.	-			
	Mean annual temperature	Argyrosomus regius d.	+			
F (Mean annual temperature	Engraulis encrasicolus d.	+	D(1 2012		
Estuary	Mean annual temperature	Dicentrarchus labrax d.	+	Beguer et al. 2012		
	Mean annual temperature	Palaemon longirostris d.	-	Pasquaud et al. 2012		
	Mean annual salinity	Palaemon longirostris d.	-			
	Mean spring temperature	Shrimp reproduction (%	-			
		ovigerous female)				
	Mean annual sea surface temperature	subtropical-species d.	+			
16	Mean annual sea surface temperature	boreal-species d.	-	Poulard and Blanchard 2005		
Marine	Annual commercial catch b. (year t)	scientific survey b. (year t)	+			
	Annual commercial catch b. (year t)	scientific survey b. (year t+1)	-			

d. = density, b. = biomass; nbr. = number

Table 2: Summary of our results: number of significant associations per number of test performed for each hypothesis and dataset. We considered a significant synchrony, for the symbolic analysis, when the observed mutual information was higher than the 95th percentiles of the distribution of the 1000 surrogate mutual information ($I_{obs} - I_{sur} > 0$,) and the joint probabilities contributing to the mutual formation corresponded to our hypothesis. For the residual analysis, we considered a significant synchrony when the correlation between residuals was significant (P < 0.05) with R^2 >0.4, and the sign of the relationship corresponded to our hypothesis.

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	Spatial effects			Cohort effects			Environmental effects		
Datasets	Symbolic	Residual		Symbolic	Residual		Symbolic	Residual	
		$R^2 > 0.4$	$R^2 > 0.2$		$R^2 > 0.4$	$R^2 > 0.2$		$R^2 > 0.4$	$R^2 > 0.2$
Stream	1/1	1/1	1/1	1/1	1/1	1/1	0/3	0/3	0/3
River	1/5	0/5	1/5	1/16	0/16	1/16	0/5	0/5	0/5
Estuary	28/78	35/78	56/78	0/4	0/4	0/4	0/8	0/8	0/8
Marine	12/32	10/32	18/32	-	-	-	0/36	0/36	1/36

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791 Figure legends:

Figure 1: Site localisation and sampling methods of the four datasets: 1) Trout populations in small streams (in green), 2) Fish populations in a large river (in red), 3) Fish and invertebrate populations in an estuary (in purple), and 4) Marine fish and invertebrate populations on continental shelf (in orange). The theoretical hydrographical network (RHT, Pella et al. 2012) was used to represented the French stream network (Strahler order >2).

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Figure 2: Illustration of the first step of both approaches used to evaluate the synchrony between time series: a) the symbolic transformation, and b) the residual computation. In both panels, open dots represent a standardized fish density time series. In panel a, the continuous time series is transformed into a discrete sequence of three symbols (F for "Flat", I for "Increase", and D for "Decrease"). This step precedes the mutual information calculation between two symbolic sequences. In panel b, residuals of a LOESS model are extracted. This step precedes the linear model among residuals from two time series.



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