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## MASS EFFECTS MEDIATE COEXISTENCE IN COMPETING SHREWS

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**Abstract.** Recent developments in metacommunity theory have raised awareness that processes occurring at regional scales might interfere with local dynamics and affect conditions for the local coexistence of competing species. Four main paradigms are recognized in this context (namely neutral, patch-dynamics, species-sorting, and mass-effect), that differ according to the role assigned to ecological or life-history differences among competing species, as well as to the relative time scale of regional vs. local dynamics. We investigated over four generations the patterns of regional and local coexistence of two species of shrews (*Crocidura russula* and *Sorex coronatus*) sharing a similar diet (generalist insectivores), in a spatially structured habitat at the altitudinal limit of their distributions. Local populations were small and regional dynamics were strong, with high rates of extinction and recolonization. Niche analysis revealed significant habitat differentiation on a few important variables, including temperature and availability of winter resting sites. In sites suitable for both species, we found instances of local coexistence with no evidence of competitive exclusion. Patterns of temporal succession did not differ from random, with no suggestion of a colonization–competition trade-off. Altogether, our data provide support for the mass-effect paradigm, where regional coexistence is mediated by specialization on different habitat types, and local coexistence by rescue effects from source sites. The strong regional dynamics and demographic stochasticity, together with high dispersal rates, presumably contributed to mass effects by overriding local differences in specific competitive abilities.

**Key words:** *coexistence; colonization; competition; Crocidura russula; habitat differentiation; habitat suitability model; mass effect; regional vs. local dynamics; shrews; Sorex coronatus; source–sink dynamics; stochasticity.*

### INTRODUCTION

Much of community theory to date has focused primarily on processes affecting species interactions at a local scale, as modeled by classical population-dynamics equations (Lotka–Volterra type and extensions; Lotka 1924, Volterra 1926). Under these settings, and notwithstanding interactions with higher trophic levels, species coexistence is largely mediated by resource partitioning. Competitive exclusion can only be avoided if niche displacement depresses interspecific competition below intraspecific levels (Chesson 2000a). However, recent empirical and theoretical studies have raised awareness that processes occurring at larger spatial scales might interfere with local dynamics and affect conditions for local coexistence (reviewed in Amarasekare 2003, Leibold et al. 2004). The concept of metacommunity is now emerging as a relevant tool to formalize interactions among regional and local scales, shedding new light on reasons why competitors coexist more often than expected (Holyoak et al. 2005).

Four main lines of research are to be distinguished within this framework (Leibold et al. 2004). The neutral

(NE) paradigm (Hubbell 2001) assumes that interacting species differ neither in ecology nor in life history. In the absence of extrinsic processes (speciation or immigration from outside the metacommunity), competitive exclusion should eventually drive to extinction all species but one. However, as transient dynamics can be very long, low rates of speciation or immigration are enough to maintain some diversity at equilibrium. Whether diversity accrues within or among sites depends on the relative forces of drift (which accelerates the local loss of species) and dispersal (which homogenizes species distributions and thus promotes local coexistence).

The patch-dynamics (PD) paradigm (e.g., Levins and Culver 1971, Tilman 1994) considers multiple identical patches that undergo stochastic or deterministic extinctions counterbalanced by dispersal. It differs from the null (neutral) model by assuming that regional coexistence is mediated by life-history differences among species (namely a trade-off between competitive ability and dispersal). Models along this line often assume that local dynamics occur on a shorter time scale than extinction–colonization dynamics (patch-occupancy models), which limits the scope for local coexistence.

The two other paradigms assume that patches differ in conditions (which allows regional coexistence through specialization on different patch types), but assign different roles to dispersal. In the species-sorting (SS)

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paradigm (Leibold 1998, Chase and Leibold 2003), intra-patch population dynamics occur on a shorter time scale than colonization–extinction processes, so that locally superior species have ample time to outcompete rivals. The scope for local coexistence is thereby limited, and species distributions closely match local conditions (Cottenie et al. 2003, Leibold and Norberg 2004). In the mass-effect (ME) paradigm (Levin 1974, Shmida and Wilson 1985, Holt 1993, Pulliam 2000, Mouquet and Loreau 2002), by contrast, local and regional processes occur at the same time scale, which allows dispersal to quantitatively affect local dynamics and promote local coexistence (extinction rates of competitively inferior species are reduced by source–sink dynamics and rescue effects).

Stochastic processes may obviously affect population dynamics in all four paradigms. On the one hand, stochasticity is expected to accelerate the local loss of diversity through increased drift, in the same way that genetic drift accelerates allele fixation. On the other hand, it makes the outcome of competitive interactions less predictable. Random events in small populations may actually override possible differences in specific competitive abilities (SS or ME models), in the same way that genetic drift may override possible differences in allelic fitness (Orrock and Fletcher 2005).

To sum up, ME and SS differ from NE and PD by assuming that intrinsic differences among local sites affect the regional distribution of competing species; SS differs from ME by assuming different time scales for local and regional processes; and PD differs from NE by assuming a trade-off between colonization and competitive abilities. Even though real metacommunities are not expected to conform to one single paradigm, the classification and schematization just discussed provide a useful framework to delineate processes and address relevant empirical issues (Leibold et al. 2004).

In the present study, we focus on two species of shrews sharing a similar diet (generalist insectivores), and investigate the temporal patterns of local and regional coexistence in a spatially structured habitat. In order to delineate which of the four metacommunity paradigms best explains the observed patterns, we address the following questions. Do the species under study occupy sites that differ according to ecologically relevant factors (as expected from SS and ME, against PD and NE)? Do they differ in colonization or competitive abilities (as expected from PD, against NE)? Are regional processes (immigration–emigration) strong enough to prevent competitive exclusion from sites otherwise suitable for both species (as expected from ME, against SS)? As it turns out, responses to these questions point to the mass effect (ME) paradigm as the most appropriate model to account for observed patterns of local and regional coexistence in the system under study.

## METHODS

### *Study species*

The two soricid shrews, *Crocidura russula* (Hermann, 1780) and *Sorex coronatus* (Millet, 1828), closely resemble each other in terms of feeding ecology, morphology, and life history. The two species display very similar body mass (*C. russula*, 9–12 g, *S. coronatus*, 9–11 g), the clearest discriminating morphological character being tooth coloration (the enamel is reddish in *S. coronatus* and whitish in *C. russula*). Both are ground-foraging insectivores, sharing a generalist diet consisting mostly of arthropods, worms, and mollusks (Bever 1983, Castien and Gosalbez 1995, Hausser 1995). Shrews are characterized by a highly elevated rate of metabolism, which imposes very important energetic needs (Genoud 1985). Owing to the seasonality in prey availability, winter food shortage constitutes the major source of mortality that limits natural populations (Genoud and Hausser 1979, Bouteiller and Perrin 2000, Butet et al. 2006). Competition for food is thus likely to play a crucial role in species interactions. Other potential competitors (i.e., other shrews of the genus *Sorex* and *Neomys*) are virtually absent from the study area (see *Results*). Predators are few and similar for both species (mainly domestic cats and nocturnal raptors such as the Common Barn Owl, *Tyto alba*; Indelicato [2000]). Finally, both species are annual breeders with juvenile dispersal (Churchfield 1990, Favre et al. 1997).

Widespread in western Europe, these shrews have a lowland distribution in Switzerland, usually found below 1000 m above sea level, a.s.l. (Hausser 1995). *Crocidura russula* is preferentially anthropophilic below 600 m and strictly so at higher altitudes (Genoud 1985), relying on thermally favorable sites (farms, stables, and compost heaps) to meet the energetic needs of the cold season (Genoud and Hausser 1979). *Sorex coronatus* is found in more diverse habitat types including forests, hedges, and meadows (Genoud 1982). However, despite slightly divergent preferences, many habitats are suitable for both species (Genoud 1982, Butet et al. 2006). Given the similarities in the trophic niche and the high energetic requirements, strong competition is expected at the local scale. Some authors (Croin Michielsen 1966, Genoud 1982; J. Hausser, *unpublished data*) indeed suggested that *S. coronatus* might be excluded from anthropophilic areas by *C. russula*, based on the observations that the species are only rarely found in syntopy, despite being sympatric over a large part of western Europe (Genoud 1985) and that *S. coronatus* is found next to human settlements in northern Europe, where *C. russula* is absent (Genoud 1982). Competitive dominance by *C. russula* might point to a patch-dynamics component in specific interactions (colonization vs. competition trade-off). To what extent dispersal suffices to counteract competitive exclusion and allow local coexistence remains an open empirical question.

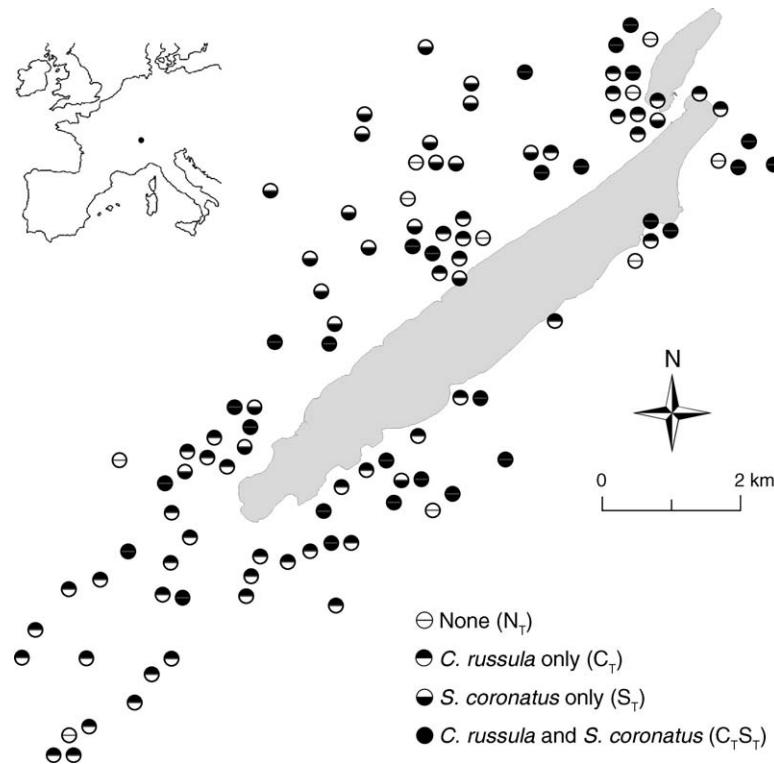


FIG. 1. Map of the sampled sites. Open circles represent sites that remained unoccupied throughout the study ( $N_T$ ). Solid circles ( $C_T S_T$ ) indicate sites where both shrew species (*Crocidura russula*, C; *Sorex coronatus*, S) occurred at least once (but not necessarily within the same year). Half-solid, half-open circles are sites where only one of the two species was caught throughout the study ( $C_T$ , *C. russula* only;  $S_T$ , *S. coronatus* only). Subscript  $T$  refers to the site occupancy pattern over the four-year survey. The lake, Lac de Joux, is light gray.

#### Field sampling

The study area was located in the Vallée de Joux ( $6^{\circ}15' E$ ,  $46^{\circ}37' N$ , 1000–1300 m a.s.l.), Switzerland, at the upper altitudinal limit of both species. *C. russula* is strictly anthropophilic at these altitudes (Genoud 1985). Harsh environmental conditions might simultaneously increase energetic needs and decrease prey abundance, thus enhancing competition for food. Demographic and environmental stochasticity are also a priori likely to play a role in their distribution patterns. Abundances were monitored in summer (August to September) in 106 sites (62 in 2003; 84 in 2004, 2005, and 2006), over a  $6 \times 14$  km area (Fig. 1). Sampling sites consisted of  $20 \times 20$  m areas within private gardens. These were always adjacent to human habitations, and usually comprised some lawn, a vegetable garden, a compost pile, wooden piles, stone walls, and hedges. Of these sites, 42 were sampled over all four years, 40 sites over three consecutive years, two sites over two consecutive years, and 22 sites only once. It should be noted that our choice of sampling sites (gardens) preferentially focused on *C. russula*, and that *S. coronatus* may occupy localities outside human settlements. Individuals were live-trapped using 20 Longworth small-mammal traps per site (Longworth Scientific Instruments, Abingdon, UK),

baited with *Tenebrio molitor* (yellow mealworm) larvae. After a pre-baiting period of 2–4 days, each site was visited four times during two consecutive days. The traps were opened daily at around 06:30 hours, checked at 10:00, and again at 13:30 hours, before being closed for the night. All caught individuals were sexed, aged, individually marked by toe clipping, and immediately released.

A site was considered as occupied by species A if at least one individual was captured. Site colonization rate ( $\text{yr}^{-1}$ ) was estimated as the number of annual transitions from an empty to an occupied state (i.e., the sum of cases in which a site was empty in year  $t$  and occupied in year  $t + 1$ ), divided by the total number of transitions from an empty state (i.e., the sum of cases in which a site was empty in year  $t$  and monitored in year  $t + 1$ ). A site extinction rate was similarly estimated as the number of transitions from an occupied to an empty state, divided by the total number of transitions from an occupied state. For both rates, analysis was restricted to the set of sites that were occupied at least once over the four years.

#### Habitat modeling

Habitat preferences of both species were modeled with the Ecological Niche Factor Analysis (ENFA, Biomapper 3.2; Hirzel et al. 2002, 2004a) at 25-m resolution.

TABLE 1. Scores of the environmental variables on the first four axes of the Ecological Niche Factor Analysis (ENFA) for the shrews *Sorex coronatus* (*S.c.*) and *Crocidura russula* (*C.r.*).

Variable	Marginality		Specialization 1		Specialization 2		Specialization 3		Range (units)
	<i>S.c.</i>	<i>C.r.</i>	<i>S.c.</i>	<i>C.r.</i>	<i>S.c.</i>	<i>C.r.</i>	<i>S.c.</i>	<i>C.r.</i>	
BUILD†	0.74	0.76	-0.25	0.01	0.04	-0.10	-0.22	0.16	0.7–33.5 (no. buildings)
ELEV†	-0.34	-0.21	-0.87	-0.73	0.56	0.08	-0.03	0.67	1008–1299 (m)
EAST†	0.25	-0.01	-0.06	-0.02	0.20	-0.32	-0.13	0.07	-0.99 to 1
FROST‡	0.24	-0.05	-0.28	-0.18	-0.41	0.02	0.39	-0.38	9–24 (days)
NDVI§	-0.05	-0.24	0.23	0.33	0.48	-0.34	0.11	0.55	-0.02 to 0.5
NORTH†	-0.14	-0.01	0.10	0.23	-0.14	0.14	0.43	0.05	-1 to 1
WHAB†	0.46	0.56	-0.02	-0.17	0.47	-0.11	0.27	0.24	0–27.2 (building/m <sup>2</sup> )
WRAD‡	0.04	0.00	0.19	0.49	0.09	0.87	0.72	0.14	3612–10346 (kJ/d)

Notes: A positive marginality coefficient indicates that the species is found at values higher than average. Only absolute values matter for the specialization axes (see Hirzel et al. 2002). BUILD, number of habitations ( $f$ ) within a 100 m radius, weighted by distance ( $d$ ) to the closest building,  $f(\exp[-d/100])$ ; ELEV, elevation; EAST, easterly aspect; FROST, number of frost days per year; NDVI, normalized difference vegetation index (correlated with vegetation biomass); NORTH, northerly aspect; WHAB, number of buildings occupied in winter within a 100 m radius; WRAD, average solar radiations in winter (December to February). Units are given when applicable.

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‡ Provided by N. E. Zimmermann and F. Kienast from the Swiss Federal Institute for Forest, Snow and Landscape Research, 8903 Birmensdorf, Switzerland.

§ Advanced Spaceborne Thermal Emission and Reflection Radiometer, NASA, Jet Propulsion Laboratory, California Institute of Technology, Pasadena, California, USA. (<http://asterweb.jpl.nasa.gov/index.asp>)

This multivariate analysis extracts a series of independent factors (linear combinations of environmental variables) that maximize the marginality and specialization of a focal species, relative to a reference area. Marginality measures how much the environmental conditions in “presence” sites depart from average, and specialization measures the narrowness of their distributions, relative to that of reference sites. ENFA only requires presence data, and thus is often applied when species absences do not necessarily reflect habitat unsuitability (Hirzel et al. 2001, 2004b, Reutter et al. 2003, Brotons et al. 2004, Engler et al. 2004, Chefaoui et al. 2005). It is well adapted to our case, where absences may originate from stochastic demographic processes or competitive exclusions rather than environmental incompatibilities.

To reflect the sampling pattern, the study area was restricted to a 200-m buffer around human habitations. Eight environmental variables (Table 1) were chosen in accordance with the ecology of our study species (Hausser 1995): three topographic variables (altitude and northerly and easterly aspects); two climatic variables (winter solar radiation and number of frost days per year); two anthropogenic variables (number of winter-inhabited buildings within a 100 m radius, and number of buildings within a 100 m radius weighted by distance); and one biotic variable measuring plant productivity (normalized difference vegetation index, NDVI; Rouse et al. 1973). Each site where *C. russula* (respectively, *S. coronatus*) occurred at least once during the survey was considered as a presence point in the *C. russula* (respectively, *S. coronatus*) ENFA analysis. The marginality factor plus all factors explaining more than 10% of specialization were used to build habitat suitability maps. This provided an index of habitat

suitability to *C. russula* ( $H_C$ ) and to *S. coronatus* ( $H_S$ ) for each site.

The quality and robustness of the models were evaluated by the continuous Boyce index as implemented in Biomapper 3.2, using  $k$ -fold cross-validation (Fielding and Bell 1997). This index measures how better than chance expectation a model is (for more details, see Hirzel et al. 2006). The presence data set was split into  $k$  partitions (for *C. russula*,  $k = 6$ ; for *S. coronatus*,  $k = 4$ ); then a new model was built with  $k - 1$  partitions (calibration data set) and was validated with the omitted one (evaluation data set). We repeated this procedure for the  $k$  independent partitions to get mean and standard deviation of the Boyce index (Hastie et al. 2001).

#### Site differentiation and spatial exclusion

The overall degree of differentiation or exclusion between *C. russula* and *S. coronatus* was estimated by the kappa statistic,  $\kappa$  (Cohen 1960), which compares the agreement between two categorical variables (here, species occurrences, 0 or 1) in the same way as correlation coefficients do for quantitative variables

$$\kappa = \frac{N \sum_{i=0}^1 x_{ii} - \sum_{i=0}^1 x_{i.} x_{.i}}{N^2 - \sum_{i=0}^1 x_{i.} x_{.i}}$$

where  $N$  represents the total number of observations,  $x_{11}$  is the number of co-occurrences,  $x_{00}$  is the number of times both species were absent,  $x_{1.}$  (respectively,  $x_{.1}$ ) stands for the number of times *C. russula* (respectively, *S. coronatus*) was present, and  $x_{0.}$  (respectively,  $x_{.0}$ ) is the number of times *C. russula* (respectively, *S. coronatus*) was absent. Kappa varies from  $-1$  (the two

TABLE 2. Patterns of site occupancy by shrew species (*Crocicidura russula*, C; *Sorex coronatus*, S) over the four study years.

Site occupation	Number of sites				Site code	All years
	2003	2004	2005	2006		
C <sub>0</sub> S <sub>0</sub>	13	29	24	30	N <sub>T</sub>	10
C <sub>1</sub> S <sub>0</sub>	28	35	36	28	C <sub>T</sub>	46
C <sub>0</sub> S <sub>1</sub>	16	16	17	24	S <sub>T</sub>	22
C <sub>1</sub> S <sub>1</sub>	5	4	7	2	C <sub>T</sub> S <sub>T</sub>	28
Total	62	84	84	84	Total	106

Notes: C<sub>0</sub>S<sub>0</sub>, sites with no species during the year; C<sub>1</sub>S<sub>0</sub> (respectively, C<sub>0</sub>S<sub>1</sub>), sites with *C. russula* (respectively, *S. coronatus*) only; C<sub>1</sub>S<sub>1</sub>, sites with both species present within the same year. N<sub>T</sub>, number of sites that remained unoccupied throughout the study; C<sub>T</sub> (respectively, S<sub>T</sub>), number of sites where *C. russula* only (respectively, *S. coronatus* only) were caught throughout the study; C<sub>T</sub>S<sub>T</sub>, number of sites where both species occurred at least once (but not necessarily within the same year). Subscript T refers to the site occupancy pattern over the four-year survey.

species are never found together), through 0 (independence), to 1 (always together). Kappa values were computed both over the four-year study period and for each year separately. To test significance, yearly observations at a site were bootstrapped (10 000 replicates), and observed kappa values were considered significant if the 95% confidence interval of the bootstrap distribution did not include zero.

To investigate habitat differentiation, we classified sites into four groups (Table 2). The group C<sub>T</sub>S<sub>T</sub> comprised sites where both species occurred at least once, but not necessarily within the same year. The groups C<sub>T</sub> and S<sub>T</sub> comprised sites where only *C. russula* (respectively, *S. coronatus*) was found at least once over the four years. Finally, the group N<sub>T</sub> comprised sites where none of the species were found during the whole survey. Subscript T refers to the site occupancy pattern over the four-year survey. Environmental variables were compared between sites S<sub>T</sub>, C<sub>T</sub>, and C<sub>T</sub>S<sub>T</sub> using one-way MANOVA. Significance of pairwise comparisons between groups was evaluated with Tukey hsd post hoc tests.

To quantify possible habitat-specific differences in demographic parameters, we also contrasted *C. russula* colonization rate, extinction rate, occupancy, and density in sites where *S. coronatus* was either present (C<sub>T</sub>S<sub>T</sub>) or absent (C<sub>T</sub>), using bilateral Mann-Whitney tests. Symmetric analyses were performed on *S. coronatus* (S<sub>T</sub> vs. C<sub>T</sub>S<sub>T</sub> sites).

To test for possible interactions between species, sites were classified according to yearly occupation: C<sub>1</sub>S<sub>1</sub> if both species co-occurred, C<sub>1</sub>S<sub>0</sub> (respectively, C<sub>0</sub>S<sub>1</sub>) if only *C. russula* (respectively, *S. coronatus*) was found, and C<sub>0</sub>S<sub>0</sub> if both species were absent in the year being considered (Table 2). Exclusion was investigated by contrasting, for a given species, the suitability of unoccupied sites with the other species present, against unoccupied sites with the other species absent (i.e., C<sub>0</sub>S<sub>0</sub>

vs. C<sub>0</sub>S<sub>1</sub> for *C. russula*, and C<sub>0</sub>S<sub>0</sub> vs. C<sub>1</sub>S<sub>0</sub> for *S. coronatus*). The rationale for this was that, if species A were excluding species B from otherwise favorable sites, then the suitability to B of sites inhabited only by A should exceed the suitability of unoccupied sites. The same analyses were also performed on the subsample of sites where both species occurred at least once (C<sub>T</sub>S<sub>T</sub>,  $n = 28$ ).

### Temporal patterns

To detect temporal exclusion, we also restricted analysis to those sites where both species were found in at least one year over the study period (C<sub>T</sub>S<sub>T</sub>,  $n = 28$ ). Kappa statistics and significance levels were first calculated, with years considered as independent sampling units. We then tested whether the observed number of simultaneous co-occurrences differed from random expectation. Significance levels were tested through 10 000 permutations of presence-absence data (independently for each species) within sites sampled for at least two consecutive years. Permutations, rather than bootstrapping, were used in this case because expected co-occurrences have to be calculated given observed frequencies. Co-occurrences were considered to differ significantly from random expectations if observed values were outside the 95% CI of the permutation distribution.

To detect specific differences in colonizing or competitive abilities, we tested in the same way whether successions differed from random (namely, whether colonization events by species A occurred preferentially in the absence of species B, and whether extinction of species A occurred preferentially in the presence of species B).

All bootstrapping and permutation statistics were performed in R (R Development Core Team 2007).

## RESULTS

### Field sampling

Over the four years of the study, *C. russula* was found in 74 sites, with about one-half of the sites being occupied every year (33/62 in 2003, 40/84 in 2004, 44/84 in 2005, and 30/84 in 2006) (see Appendix for specific information). Altogether, 545 individuals were caught, with an average of  $3.8 \pm 3.3$  individuals per occupied site (mean  $\pm$  SD). Each marked individual was captured 1.70 times, on average, providing a rough estimation of capture probability per trapping session of  $1.70/4 = 0.43$  (assuming equal trappability for all individuals within a garden). The probability of missing one individual over four trapping sessions was thus  $(1 - 0.43)^4 = 0.11$ , and that of missing both individuals of a breeding pair was 1% ( $0.11^2 = 0.011$ ). No individual was recaptured over different sites or different years. Twenty-four colonizations (site empty in one year but occupied in the following year) out of 51 possible such events, and 40 extinctions (site occupied in one year but empty in the following year) out of 102 possible such events were

TABLE 3. MANOVA  $P$  levels for differences in environmental variables between sites inhabited exclusively by *C. russula* or *S. coronatus* (respectively,  $C_T$  and  $S_T$ ) and by both species ( $C_T S_T$ ).

Name	Group effect	$C_T$ vs. $S_T$	$C_T$ vs. $C_T S_T$	$S_T$ vs. $C_T S_T$
BUILD	***	$C_T > S_T$ ***	$C_T > C_T S_T$ ***	ns
ELEV	***	$C_T < S_T$ ***	$C_T < C_T S_T$ *	ns
EAST	**	$C_T < S_T$ **	ns	ns
FROST	***	$C_T < S_T$ ***	$C_T < C_T S_T$ **	ns
NDVI	***	$C_T < S_T$ ***	$C_T < C_T S_T$ ***	ns
NORTH	ns	ns	ns	ns
WHAB	***	$C_T > S_T$ ***	$C_T > C_T S_T$ ***	ns
WRAD	ns	ns	ns	ns
Global effect (MANOVA)	***			

Notes: Significance levels of pairwise comparisons between groups were tested with Tukey hsd post hoc tests. MANOVA was used to test only the group effect. Levels of significance are: \*  $P < 0.05$ ; \*\*  $P < 0.01$ ; \*\*\*  $P < 0.001$ . Variables for which *C. russula* prefers higher values than *S. coronatus* are indicated by  $C_T > S_T$ , and those where lower values are preferred by  $C_T < S_T$ .

detected during the survey, so that colonization and extinction rates can be estimated as  $0.47 \text{ yr}^{-1}$  and  $0.39 \text{ yr}^{-1}$ , respectively.

*Sorex coronatus* was less abundant, with 50 sites occupied at least once over the four years (21/62 in 2003, 21/84 in 2004, 24/84 in 2005, and 26/84 in 2006). In total, 182 individuals were caught (average  $2.0 \pm 1.3$  individuals per occupied site). Each marked individual was captured 1.58 times, on average, providing a rough estimation of capture probability per trapping session of  $1.58/4 = 0.39$ . The probability of missing one individual over four trapping sessions was thus  $(1 - 0.39)^4 = 0.13$ , and that of missing both individuals of a breeding pair was less than 2% ( $0.13^2 = 0.017$ ). During the study, 29 colonizations (out of 60 possible events) and 31 extinctions (out of 63 possible events) occurred, so that colonization and extinction rates amounted to  $0.48 \text{ yr}^{-1}$  and  $0.49 \text{ yr}^{-1}$ , respectively.

Neither colonization nor extinction rates differed significantly between the two species ( $P$  values = 1.00 for colonization and 0.27 for extinction, using binomial tests to compare two proportions; Crawley [2005:84]). Out of the 106 sites investigated, 28 were occupied by both species, and simultaneously so in 18 instances (Table 2). Other potential competitors can be safely neglected in the analysis, since only five individual shrews from two other species (four *Neomys fodiens* and one *Sorex minutus*) were captured over the four years of the study.

#### Habitat analyses

The ENFA habitat model for *C. russula* was built with four axes, explaining 100% of the marginality and 82% of the specialization. The scores for environmental variables (Table 1) indicate a strong preference for human settlements (density of buildings and heated habitations). Solar radiation and altitude also play a role in niche specialization. The fit was excellent, as quantified by a very high Boyce index with a low variance ( $0.88 \pm 0.07$ , mean  $\pm$  SD).

The habitat model for *S. coronatus* was built with five axes, explaining 100% of the marginality and 87% of the specialization. The scores for environmental variables

(Table 1) point to similar requirements, with some preferences for human settlements and specialization on altitude. The fit was also excellent, with a continuous Boyce index amounting to  $0.86 \pm 0.07$  (mean  $\pm$  SD).

#### Habitat differentiation and exclusion

The kappa index estimated over all sites pointed to a significant discrimination between the two species ( $\kappa = -0.27$ ; 95% CL  $[-0.39, -0.16]$ ) (kappa values were also significant for each year when analyzed separately). Environmental variables appear to globally differ between sites occupied exclusively by *C. russula* ( $C_T$ ) or *S. coronatus* ( $S_T$ ) and by both species ( $C_T S_T$ ) (MANOVA,  $P < 0.0001$ ; Table 3). A comparison between sites occupied exclusively by *C. russula* or *S. coronatus* ( $C_T$  vs.  $S_T$ ) revealed habitat differentiation on six environmental variables, with *C. russula* showing a stronger preference for human settlements, lower elevation, and more open and warmer sites.

Comparison of *C. russula* sites with vs. without *S. coronatus* ( $C_T S_T$  vs.  $C_T$ ) similarly revealed a preference for human settlements and warmer sites at lower elevations. In contrast, a comparison of *S. coronatus* sites with vs. without *C. russula* ( $C_T S_T$  vs.  $S_T$ ) revealed no significant differences. Furthermore, *C. russula*-free sites were not more suitable for *C. russula* in the presence of *S. coronatus* (Mann-Whitney tests: for all sites,  $P = 0.91$ ; restricted to sites where both species occurred,  $P = 0.69$ ). *S. coronatus*-free sites were not more suitable for *S. coronatus* in the presence of *C. russula* (Mann-Whitney tests: for all sites,  $P = 0.98$ ; restricted to sites where both species occurred,  $P = 0.80$ ).

The density and occupancy rate of *C. russula* (respectively, *S. coronatus*) were significantly higher in sites unoccupied by the other species (respectively,  $C_T$  or  $S_T$ ) than in sites favorable for both species ( $C_T S_T$ ) (Table 4;  $P < 0.0002$  for the four comparisons). Extinction rates were also significantly higher in sites favorable to both species ( $C_T S_T$ ) than in sites inhabited only by the focal species (Table 4; *C. russula*,  $P = 0.005$ ; *S. coronatus*,  $P = 0.01$ ). Reciprocally, colonization rate for *C. russula* was significantly higher in sites unoccupied by *S. coronatus* ( $C_T$ ) than in sites favorable for both

TABLE 4. Mean values of demographic variables (density, occupancy, colonization, and extinction rates) generally differ between sites occupied either by the focal species (*C. russula*,  $C_T$ ; or *S. coronatus*,  $S_T$ ) or by both species ( $C_T S_T$ ).

Variable	<i>C. russula</i> mean for sites		<i>P</i>	<i>S. coronatus</i> mean for sites		<i>P</i>
	$C_T$	$C_T S_T$		$S_T$	$C_T S_T$	
Occupancy	0.80	0.48	0.001	0.73	0.42	0.0001
Density	3.3	1.54	<0.0001	1.58	0.70	0.0002
Colonization rate	0.70	0.39	0.037	0.50	0.52	0.89
Extinction rate	0.37	0.69	0.005	0.42	0.77	0.01

Note: Given are *P* values for the differences among groups (bilateral Mann-Whitney tests).

species ( $C_T S_T$ ;  $P = 0.037$ ), but there was no such effect for *S. coronatus*.

Considering only the sites where both species occurred ( $C_T S_T$ ), we found no suggestion of temporal exclusion. Kappa values were significant neither over all years ( $\kappa = -0.04$ ; 95% CL  $[-0.23, 0.16]$ ) nor for each year separately, and simultaneous occupation did not occur significantly less than random expectation. The observed value was 18 times when both species were found in the same site within the same year over the study period; 95% CL under random expectation: [15, 24]; the maximum value would be 314 (total sites samples in four years) if both species were always present in each of the sites. Similarly, the patterns of temporal succession in sites where both species co-occurred did not deviate significantly from random: neither of the two species was more likely to settle in the absence of the other species (Fisher tests: *C. russula*,  $P = 0.70$ ,  $n = 44$ ; *S. coronatus*,  $P = 0.34$ ,  $n = 70$ ), or more likely to become extinct in the presence of the other species (Fisher tests: *C. russula*,  $P = 1$ ,  $n = 28$ ; *S. coronatus*,  $P = 0.61$ ,  $n = 42$ ). Here,  $n$  represents the number of annual transitions from an occupied to an unoccupied site (for extinction) and from an empty to an occupied site (for colonization). These transitions were recorded only on sites where both species were found over the four-year study (but not necessarily within the same year, i.e.,  $C_T S_T$ ). Results remained unchanged when considering the presence of the other species in the year before the transition (*P* values ranged from 0.45 to 1.0).

#### DISCUSSION

The two shrew species co-occurred in 28 of the 106 localities investigated, pointing to an important habitat overlap and a large scope for competition. However, this number was less than that expected by chance, obtained as the product of specific occurrence probabilities ( $74 \times 50/106 = 35$ ). Accordingly, the kappa index was significantly negative.

From our analyses, this deficit in co-occurrences mostly stemmed from a habitat differentiation. Although preferences seem broadly similar and distributions seem to be affected by the same environmental variables, the two species were specializing on slightly different values. A comparison of sites occupied exclusively by one or the other species ( $C_T$  vs.  $S_T$ ) revealed that *C. russula* preferred warmer sites at lower

elevation, with less vegetation and closer to winter-heated human habitations. The same results emerged when comparing *C. russula* sites with and without *S. coronatus* ( $C_T S_T$  vs.  $C_T$ ): *C. russula* sites were warmer, at lower elevation, with less vegetation and closer to human habitations. The reverse comparison ( $C_T S_T$  vs.  $S_T$ ) revealed no significant differences, possibly because our sampling design focused on localities a priori favorable to *C. russula* (human habitations), thereby ignoring a fraction of *S. coronatus* habitat. These differences in habitat quality translated into differences in demography: *C. russula* displayed higher density, occupation, and colonization rates, as well as a lower extinction rate, in pure *C. russula* sites ( $C_T$ ) than in sites also favorable to *S. coronatus* ( $C_T S_T$ ). The reverse also applied to *S. coronatus*, except for colonization rate. Our data thus provide evidence that habitat differentiation (even though incomplete) mediates regional coexistence in these competing species. This argues in favor of the species-sorting and mass-effect paradigms, but against the neutral and patch-dynamics paradigms, which assume undifferentiated habitat use.

Furthermore, we argue that the patterns of local coexistence and temporal occupation tend to favor the mass-effect against the species-sorting paradigm. Indeed, the latter assumes that local dynamics occur on shorter time scales than regional dynamics, so that locally favored species have ample time to exclude competitors (Leibold et al. 2004), although competitive exclusion is not necessarily required for species sorting. This should leave little scope for local coexistence, and should induce a close match between species distributions and habitat conditions. Under the mass-effect perspective, by contrast, immigration and emigration affect local population dynamics, so that species can be rescued from local competitive exclusion in communities where they are bad competitors by immigrants from communities where they are good competitors (Leibold et al. 2004). Accordingly, we found substantial numbers of local co-occurrences (28 mixed sites out of 106, with simultaneous occurrence in 18 instances) and no support for competitive exclusion. No species excluded its competitor from otherwise suitable sites, and the sites occupied by both species showed no sign of temporal exclusion (the patterns of temporal occupation and succession did not differ from random). Furthermore, the rates of extinction and colonization documented

here point to similar time scales for regional and local processes. The characteristic times to extinction of occupied sites (obtained as the inverse of extinction rate) were indeed very short, less than three years in both species (2.55 years for *C. russula* and 2.03 for *S. coronatus*). In addition, a parallel analysis of *C. russula* dynamics in this same metapopulation (Jaquiéry et al., *in press*) showed that local population sizes were significantly affected by the connectivity to potential source populations and by the quality of these sources, confirming important mass effects in the system under study.

Because our sampling design focused a priori on *C. russula* sites, the niche of *S. coronatus* might have remained partly unsampled. Assuming that better quality *S. coronatus* habitat exists outside human habitations, sampling this habitat would have resulted in a lower kappa index (because species would then co-occur in a lower proportion of sites), a stronger niche differentiation, and possibly lower extinction rate and higher colonization rates for this species. However, our main conclusion (namely that mass effects mediate coexistence in our study system) would have remained unchanged. The additional data would have argued more strongly against the patch-dynamics and neutral paradigms (because these models assume no habitat differentiation), and the patterns of local coexistence documented here would still support the mass effect (against the species-sorting) paradigm.

The landscape under study displays spatial autocorrelation of environmental variables (e.g., elevation, number of frost days), so that the sites favorable to one species tend to be somewhat aggregated. Such spatial structures are expected to affect local dynamics (be it only because dispersal range is limited), but there is no simple way to analyze such effects. However, we note that this feature only reinforces our main conclusion (namely that coexistence is mediated by mass effects). First, the neutral and patch-dynamics models assume no habitat differentiation, and therefore no landscape structure. Habitat heterogeneity and spatial structure are, by contrast, constitutive parts of the species-sorting and mass-effect paradigms. Second, the main effect of spatial structure is to lower the opportunity for local coexistence (because dispersing individuals are more likely to settle in sites already occupied by conspecifics), and thereby the power to detect mass effect. Neglecting spatial aspects in our analyses is thus conservative regarding our main conclusion.

Observations by previous authors (Croin-Michielsen 1966, Genoud 1985; see *Methods*) suggested a patch-dynamics component, in which one species (*S. coronatus*) would benefit from its ability to colonize empty sites, and the other (*C. russula*) from its ability to exclude competitors. We found no evidence for such a component. Extinction and colonization rates were similar (although estimations for *S. coronatus* might be

biased due to our sampling design), and the patterns of temporal successions in sites suitable for both species did not differ from random. In particular, *S. coronatus* was not more likely than *C. russula* to settle in empty sites or to be excluded from occupied sites. The only asymmetry noticed (namely, sparser and smaller populations in *S. coronatus*) was more likely to result from the asymmetry in the sampling design, which focused on sites a priori favorable to *C. russula*.

Altogether, spatial and temporal patterns point to a mass-effect scenario in which regional coexistence is mediated by specialization on different habitat types, while local coexistence is maintained by recurrent dispersal from source habitats. A general role of habitat heterogeneity in maintaining regional coexistence of competing species seems well supported both theoretically (Chesson 2000a, b, Amarasekare 2003, Mouquet and Loreau 2003) and empirically (Codeco and Grover 2001, Yu et al. 2001). By contrast, the role of mass effects in maintaining local diversity seems less documented. A recent meta-analysis by Cottenie (2005) suggests that a majority of published metacommunity data sets are structured by species-sorting processes (44%), followed by a combination of SS and mass-effects (29%), vs. less than 10% for neutral or patch-dynamics processes. This meta-analysis also points to significant effects of dispersal abilities and spatial scale of the study area on metacommunity structuring.

In our own study, the high dispersal abilities of both species, combined with the relatively small spatial scale involved ( $6 \times 14$  km) and the high regional dynamics, certainly concurred to generate important mass effects. Extinction and colonization rates were particularly high because both species were living here in marginal conditions, at the altitudinal limit of their distributions. Consequences are twofold. First, environmental stochasticity certainly exceeded the levels normally experienced in core areas (Guo et al. 2005, Holt et al. 2005). At this altitude, a strong winter might easily wipe out a series of local populations. Second, local populations were small ( $3.8 \pm 3.3$  individuals captured per site for *C. russula* and  $2.0 \pm 1.3$  for *S. coronatus*), normally not exceeding one pair with their progeny, which further induces high levels of demographic stochasticity through the random nature of individual survival, fecundity, and dispersal.

Frequent extinctions may not only select for high dispersal, but also maintain local populations below their carrying capacities for substantial periods of time. Competitive exclusions should not occur as long as populations are not limited by resources. Demographic stochasticity, furthermore, makes the outcome of competitive interactions less predictable, because random events may override possible differences in competitive ability, rendering interactions effectively neutral (Orrock and Fletcher 2005). These blurring effects should lower the match between species distri-

bution and environmental conditions, and enhance the scope for spatial effects.

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#### APPENDIX

A table showing coordinates, number of trapped *Crocidura russula* and *Sorex coronatus* individuals, and habitat suitability for each site, 2003–2006 (*Ecological Archives* XXXXXXXXXX).