Migratory connectivity analysis

by EURING Migration Atlas

Sylvia melanocephala (EURING code 12670)

1.1 Connectivity between individuals

The analysis evaluated 436 individuals (872 encounters) filtered from a total of 65061 records in the EURING databank which were considered for the Atlas. The species shows a significant connectivity from clustering, with a number of first-level clusters = 9 (Table 12670-1; Figure 12670-1).

Table 12670-1. Results from the migratory connectivity analysis. For each cluster, the degree of connectivity (r_M) , its statistical significance (p-value) and 95% confidence interval limits are shown. When the p-value is less than or equal to 0.1, the degree of clustering structure (oasw) and the best number of clusters identified are reported.

CI.	T 1 2	2.7	Migratory		Lower 95%	Upper 95%	Best	
Cluster	Level of		connectivity	p-	confidence	confidence	number of	
name	clustering	individuals	$(r_{\rm M})$	value	limit	limit	clusters	oasw
0	0	436	0.999	0.001	0.996	1.000	9	0.671
1	1	80	1.000	0.001	1.000	1.000	8	0.959
2	1	28	0.991	0.001	0.897	0.997	2	0.840
3	1	45	1.000	0.001	0.999	1.000	3	0.679
4	1	106	0.999	0.001	0.997	1.000	4	0.711
5	1	33	0.796	0.001	0.551	1.000	7	0.640
6	1	50	0.990	0.001	0.962	1.000	2	0.670
7	1	41	1.000	0.001	1.000	1.000	9	0.833
8	1	35	1.000	0.001	1.000	1.000	3	0.992
9	1	18	-	-	-	-	-	-
11	2	1	-	-	-	-	-	-
12	2	3	-	-	-	-	-	-
13	2	1	-	-	-	-	-	-
14	2	4	-	-	-	-	-	-
15	2	12	-	-	-	-	-	-
16	2	1	-	-	-	-	=	-
17	2	56	-	-	-	-	=	-
18	2	2	-	-	-	-	-	-
21	2	25	0.882	0.001	0.794	0.958	2	0.725
22	2	3	-	-	-	-	-	-
31	2	33	1.000	0.001	1.000	1.000	2	0.654
32	2	8	-	-	-	-	-	-
33	2	4	-	-	-	-	-	-
41	2	13	-	-	-	-	=	-
42	2	59	0.996	0.001	0.992	1.000	9	0.769
43	2	31	0.999	0.001	0.995	1.000	6	0.908
44	2	3	-	-	-	-	-	-

Cluster	Level of	N c	Migratory onnectivity		Lower 95% confidence	Upper 95% confidence	Best number of	
name	clustering	individuals	$(r_{ m M})$	p- value	limit	limit	clusters	oasw
51	2	8	-	-	-	-	-	_
52	2	3	_	-	-	-	_	-
53	2	7	_	-	_	_	_	-
54	2	1	-	_	_	_	-	_
55	2	5	_	_	_	_	_	_
56	2	5	_	_	_	_	_	_
57	$\frac{1}{2}$	$\overset{\circ}{4}$	_	_	_	_	_	_
61	2	45	0.967	0.001	0.900	1.000	6	0.647
62	2	5	-	-	-	-	-	-
71	2	$\overset{\circ}{2}$	_	_	_	_	_	_
72	2	5	_	_	_	_	_	_
73	2	13	_	_	_	_	_	_
74	2	4						
75	$\frac{2}{2}$	9	_		_	_		_
76	$\frac{2}{2}$	$\frac{3}{4}$	_	_	_	_	_	_
77	$\frac{2}{2}$	1	_	-	_	_	_	_
78	$\frac{2}{2}$	$\overset{1}{2}$	_	-	-	_	-	_
79	$\frac{2}{2}$	1	-	-	-	-	-	-
81	$\frac{2}{2}$	$\frac{1}{22}$	1 000	0.053	1.000	1.000	2	0.955
	$\frac{2}{2}$	$\frac{22}{2}$	1.000	0.055	1.000	1.000	Δ	0.955
82			-	-	-	-	-	-
83	2	11	-	-	-	-	-	-
211	3	18	-	-	-	-	-	-
212	3	7	-	-	-	-	-	-
311	3	27	1.000	0.001	1.000	1.000	9	0.660
312	3	6	-	-	-	-	-	-
421	3	3	-	-	-	-	-	-
422	3	10	-	-	-	-	-	-
423	3	8	-	-	-	-	-	-
424	3	5	-	-	-	-	-	-
425	3	3	-	-	-	-	-	-
426	3	18	-	-	-	-	-	-
427	3	3	-		-	-	-	-
428	3	4	-	-	-	-	-	-
429	3	5	-	-	-	-	-	-
431	3	3	-	-	-	-	-	-
432	3	7	-	-	-	-	-	-
433	3	2	-	-	-	-	-	-
434	3	13	-	-	-	-	-	-
435	3	1	-	-	-	-	-	-
436	3	5		-	-	-	-	-
611	3	9	-	-	-	-	-	-
612	3	8	_	-	-	-	-	-
613	3	5	_	-	-	-	-	-
614	3	10	_	-	_	_	-	_
615	3	3	_	_	-	-	-	_
616	3	10	-	_	-	-	-	_
811	3	21	_	_	-	-	_	_
812	3	1	_	_	_	_	_	_

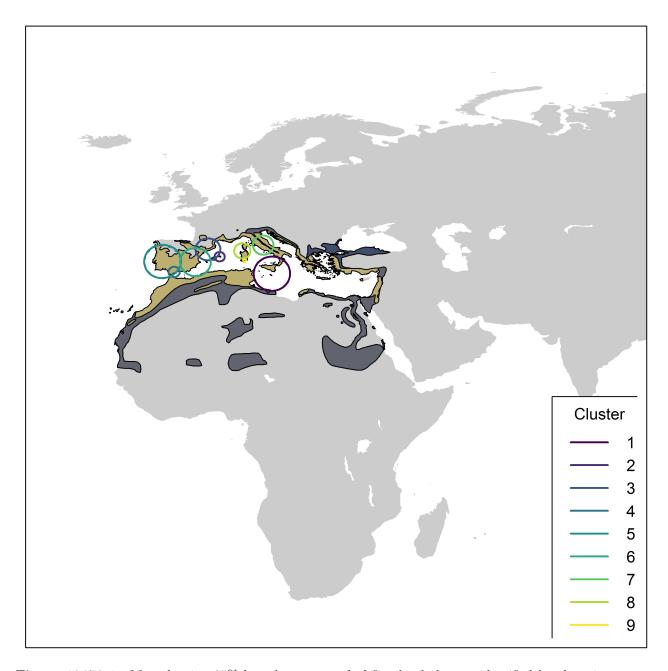


Figure 12670-1. Map showing 95% kernel contours of of first-level clusters identified by the migratory connectivity analysis, if any, or 95% kernel contours of all encounters, in case of no clustering structure. Solid lines indicate the clusters in the breeding range, dotted lines those in the non-breeding range. Different contour colours correspond to different clusters, as reported in legend. The species distribution range is also shown (breeding range: blue; non-breeding range: dark grey; resident range: beige; from BirdLife International, 2019).

1.2 Sensitivity analysis

Results of power analysis and validation. Analyses at the species level were re-run on subsamples of individuals of decreasing size (100 repetitions per subsample size), according to simple random sampling of individuals (Figure 12670-2) and stratified sampling of individuals within the breeding range (Figure 12670-3) and the non breeding range (Figure 12670-4). For stratified sampling, we selected individuals with a

probability inversely proportional to the number of observation in each country. Figures below report the results of the procedure.

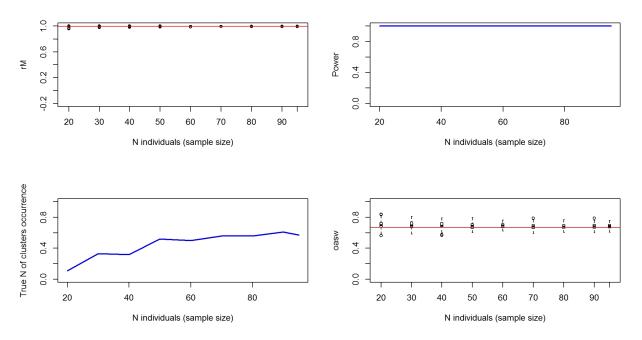


Figure 12670-2. Top left: simulated distribution (boxplots) and observed value (red line) of connectivity. Top right: Simulated power of the analysis (i.e. proportion of times the analyses on the subset of individuals was significant). Bottom left: Proportion of times the analysis provides the observed best number of cluster. Bottom right: simulated distribution (boxplots) and observed value (red line) of clustering intensity.

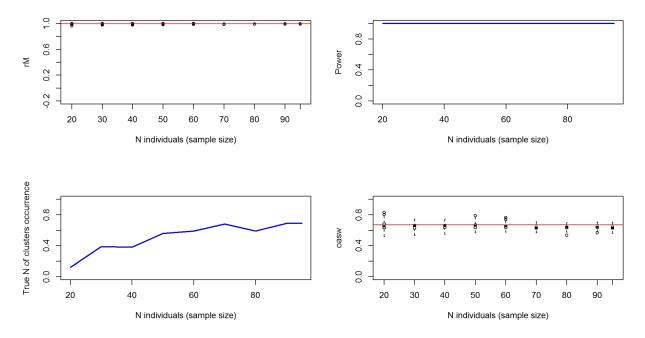


Figure 12670-3. Top left: simulated distribution (boxplots) and observed value (red line) of connectivity. Top right: Simulated power of the analysis. Bottom left: Proportion of times the analysis provides the

observed best number of cluster. Bottom right: simulated distribution (boxplots) and observed value (red line) of clustering intensity.

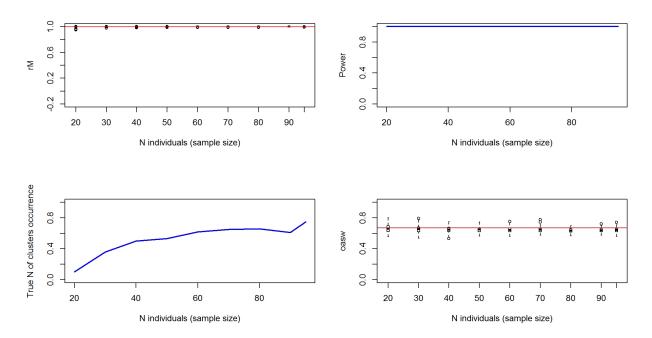


Figure 12670-4. Top left: simulated distribution (boxplots) and observed value (red line) of connectivity. Top right: Simulated power of the analysis. Bottom left: Proportion of times the analysis provides the observed best number of cluster. Bottom right: simulated distribution (boxplots) and observed value (red line) of clustering intensity.

2. Connectivity between pre-defined regions

The species shows high connectivity (MC = 1; MC = 1 when adjusted for absolute abundance) between 4 breeding regions and 4 non breeding regions (Table 12670-2; Figure 12670-6).

Table 12670-2. Transition probabilities between pre-defined regions. Estimated abundance (number of individuals) in each breeding region is also reported.

Breeding region	Abundance	Non breeding region	Transition probability
North Africa	1000	North Africa	1
South-central Europe	2426069	South-central Europe	1
South-east Europe	4534000	South-east Europe	1
South-west Europe	16751433	South-west Europe	1



Figure 12670-6. Map showing pre-defined regions in different colours, with black arrows linking centroids of individual encounters in different regions. Arrow width is proportional to transition probability.

Reference

BirdLife International and Handbook of the Birds of the World (2019). Bird species distribution maps of the world. Version 2019.1. Available at http://datazone.birdlife.org/species/requestdis.