

An example of usage of the niche metric functions (former 'resniche' package)

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1 Introduction

In this document we show how to use the functions described in [De Cáceres et al., 2011] by following an example of dietary preferences in pigeons belonging to two populations (Moià and Barcelona) in Catalonia (north east of Spain). We begin by loading the library and the data:

```
> library(indicspecies)
> data(pigeons)
```

The data consists of three items: the resource use matrix for each pigeon population and the matrix of dissimilarities between pairs of resources (seeds).

```
> ls()

[1] "dfood"          "diet.barcelona" "diet.moia"
```

2 Resemblance between diet resources

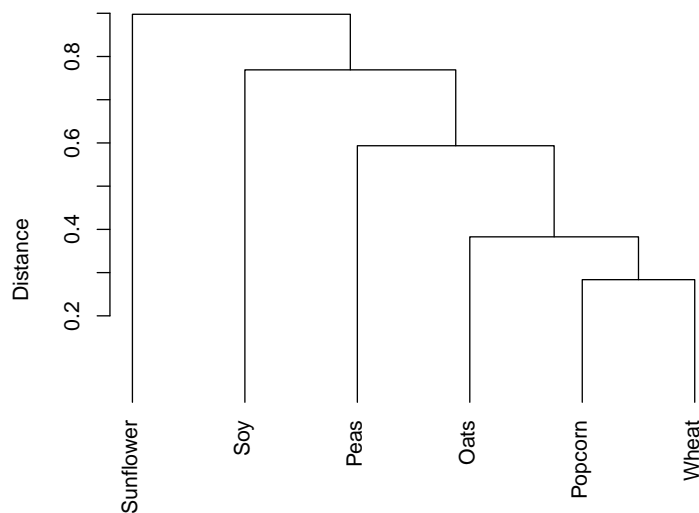
Before starting any resource niche calculation, we can inspect the matrix of dissimilarities between the $r = 6$ resources, **D**:

```
> dfood
```

	Oats	Peas	Popcorn	Soy	Sunflower
Peas	0.7029375				
Popcorn	0.3706196	0.5070959			
Soy	0.7738801	0.6645980	0.8638219		
Sunflower	0.7656489	0.9773428	0.8784905	0.8669801	
Wheat	0.3948828	0.5705974	0.2836482	0.7740283	1.0000000

These can be summarized using a dendrogram. For example:

```
> plot(hclust(dfood, method="average"), h=-1, xlab="",
+       ylab="Distance", main="", sub="", ylim=c(0.2,1))
```



Some seeds are quite similar (e.g. popcorn, wheat or oats), whereas sunflower differs substantially from all other resources.

3 Resource niche analysis at the population level

3.1 Resource use of populations

We begin by showing the resource use data for pigeons of Barcelona and Moia, expressed as proportions (i.e., the vector \mathbf{f} for each population):

```
> diet.pop.barcelona = colSums(diet.barcelona)
> round(diet.pop.barcelona/sum(diet.pop.barcelona), dig=3)
```

Oats	Peas	Popcorn	Soy	Sunflower	Wheat
0.049	0.001	0.028	0.024	0.455	0.442

```
> diet.pop.moia = colSums(diet.moia)
> round(diet.pop.moia/sum(diet.pop.moia), dig=3)
```

Oats	Peas	Popcorn	Soy	Sunflower	Wheat
0.002	0.018	0.091	0.045	0.054	0.791

Whereas pigeons in Moia feed almost exclusively on wheat, those of Barcelona combine wheat and sunflower seeds.

3.2 Niche breadth in populations

We will determine the resource niche breadth of each of the two populations as [De Cáceres et al., 2011]:

$$B_{pop} = (1/2) \sum_{j=1}^r \sum_{k=1}^r f_j f_k d_{jk}$$

We first conduct our calculations without taking into account the resemblance between resources (which is equal to stating that $d_{jk} = 1$ in all cases):

```
> nichevar(P=diet.barcelona, mode="single")
```

	B	LC	UC
Niche	0.2966578	0.2709855	0.3385325

```
> nichevar(P=diet.moia, mode="single")
```

	B	LC	UC
Niche	0.1807663	0.09549493	0.260983

In general, we can say that the niche breadth of the population in Moia is smaller than the niche breadth of the population in Barcelona. If we repeat the same calculations with the matrix of resource resemblance, we realize that the niche breadth of both populations becomes smaller:

```
> popvar.barcelona = nichevar(P=diet.barcelona, D=dfood,
+                               mode="single")
> popvar.barcelona
```

```
          B          LC          UC
Niche 0.2453822 0.2333452 0.2579433
```

```
> popvar.moia = nichevar(P=diet.moia, D=dfood, mode="single")
> popvar.moia
```

```
          B          LC          UC
Niche 0.0853328 0.02260875 0.167566
```

The reason is that the first analysis was assuming that all resources were equally (and maximally) distinct, while the second analysis accounts for the similarity between some resources. Moreover, note that the niche breadth of Moia has decreased more than the niche breadth of Barcelona. This reflects that the resources being used by pigeons of Moia are more similar than the resources used by pigeons of Barcelona.

3.3 Overlap between populations

We can now calculate the niche overlap between the two pigeon populations:

$$O_{12} = \frac{\sum_{j=1}^r \sum_{k=1}^r f_{1j} f_{2k} (1 - d_{jk}^2)}{\sqrt{\sum_{j=1}^r \sum_{k=1}^r f_{1j} f_{1k} (1 - d_{jk}^2) \sum_{j=1}^r \sum_{k=1}^r f_{2j} f_{2k} (1 - d_{jk}^2)}}$$

Using function `nicheoverlap`:

```
> nicheoverlap(P1=diet.barcelona, P2=diet.moia, mode="single")
```

```
          0          LC          UC
Overlap 0.7419319 0.4833565 0.9399538
```

```
> nicheoverlap(P1=diet.barcelona, P2=diet.moia, mode="single", D = dfood)
```

```
          0          LC          UC
Overlap 0.7912472 0.5490802 0.9467254
```

If we include the resemblance between resources, the degree of overlap increases, for the same reason that we obtained smaller niche breadth statistics when resemblances were included.

4 Resource niche analysis at the individual level

In this section, we perform a resource niche analysis at individual level. In particular, we are interested in assessing how much the resource niche of individuals differs from that of their corresponding population. For this, we need to calculate a measure of the degree of individual specialization.

4.1 Resource use of individuals

We begin by showing the resource use data for all 23 pigeons in the sample from the Barcelona population, expressed as proportions (i.e., matrix **F** for Barcelona):

```
> round(sweep(diet.barcelona, 1, FUN="/",
+             rowSums(diet.barcelona)), dig=3)
```

	Oats	Peas	Popcorn	Soy	Sunflower	Wheat
1	0.000	0.000	0.000	0.000	0.020	0.980
2	0.000	0.000	0.000	0.000	1.000	0.000
3	0.000	0.000	0.000	0.000	1.000	0.000
4	0.025	0.000	0.153	0.000	0.000	0.822
5	0.000	0.000	0.000	0.000	0.965	0.035
6	0.000	0.023	0.136	0.000	0.648	0.193
7	0.000	0.000	0.000	0.000	0.795	0.205
8	0.042	0.000	0.000	0.000	0.000	0.958
9	1.000	0.000	0.000	0.000	0.000	0.000
10	0.014	0.000	0.000	0.000	0.014	0.973
11	0.000	0.000	0.000	0.000	0.350	0.650
12	0.000	0.000	0.000	0.000	0.047	0.953
13	0.000	0.000	0.000	0.000	0.437	0.562
14	0.000	0.000	0.000	0.000	0.389	0.611
15	0.000	0.000	0.000	0.549	0.451	0.000
16	0.000	0.000	0.000	0.000	0.913	0.087
17	0.000	0.000	0.113	0.075	0.755	0.057
18	0.000	0.000	0.043	0.000	0.743	0.214
19	0.000	0.000	0.000	0.000	0.667	0.333
20	0.000	0.000	0.000	0.000	0.367	0.633
21	0.000	0.000	0.000	0.000	0.910	0.090
22	0.000	0.000	0.019	0.019	0.786	0.175
23	0.442	0.000	0.000	0.000	0.000	0.558

We see that most individuals in Barcelona feed on either sunflower or wheat, but there are some individuals (like pigeon 9) which prefer oats. Now we display the resource use data for the 19 pigeons representing the population in Moia (i.e., matrix **F** for Moia):

```
> round(sweep(diet.moia, 1, FUN="/",
+             rowSums(diet.moia)), dig=3)
```

	Oats	Peas	Popcorn	Soy	Sunflower	Wheat
1	0.000	0.000	0.000	0.324	0.000	0.676
2	0.000	0.000	0.000	0.012	0.000	0.988
3	0.000	0.000	0.000	0.000	0.000	1.000

4	0.000	0.000	0.000	0.000	0.000	1.000
5	0.000	0.000	0.000	0.000	0.000	1.000
6	0.000	0.000	0.267	0.000	0.000	0.733
7	0.000	0.000	0.319	0.000	0.000	0.681
8	0.027	0.311	0.041	0.000	0.000	0.622
9	0.000	0.000	0.000	0.000	0.000	1.000
10	0.000	0.000	0.148	0.000	0.000	0.852
11	0.000	0.000	0.200	0.000	0.000	0.800
12	0.000	0.000	0.235	0.000	0.000	0.765
13	0.000	0.000	0.000	0.000	0.000	1.000
14	0.000	0.000	0.119	0.339	0.136	0.407
15	0.000	0.000	0.000	0.000	0.000	1.000
16	0.000	0.000	0.000	0.000	0.000	1.000
17	0.000	0.000	0.430	0.000	0.000	0.570
18	0.000	0.000	0.000	0.000	0.583	0.417
19	0.000	0.000	0.000	0.000	1.000	0.000

Many pigeons from Moià feed on wheat seeds almost exclusively, but some of them have broader preferences.

4.2 Measuring the degree of individual specialisation

We begin our resource niche analysis by calculating the niche breadth of each individual in the population, B_i :

$$B_i = (1/2) \sum_{j=1}^r \sum_{k=1}^r f_{ij} f_{ik} d_{jk}$$

We calculate the values for the individuals of both populations:

```
> indvar.barcelona<-nichevar(P=diet.barcelona, D=dfood)
> summary(indvar.barcelona)
```

```

      B
Min.   :0.00000
1st Qu.:0.01755
Median :0.08153
Mean    :0.11168
3rd Qu.:0.19890
Max.    :0.24609
```

```
> indvar.moia<-nichevar(P=diet.moia, D=dfood)
> summary(indvar.moia)
```

```

      B
Min.   :0.00000
```

```

1st Qu.:0.00000
Median :0.01015
Mean   :0.04031
3rd Qu.:0.01861
Max.    :0.24306

```

Most individuals have niche breadths that are smaller than the niche breadth of their corresponding population, although a few individuals in Moià have niche breadths larger than the population value. A niche breadth equal to zero indicates that only one resource is exploited.

We can compare the niche breadth values of the two populations using a non-parametric test.

```
> wilcox.test(indvar.barcelona$B, indvar.moia$B)
```

```
Wilcoxon rank sum test with continuity correction
```

```

data: indvar.barcelona$B and indvar.moia$B
W = 328, p-value = 0.005446
alternative hypothesis: true location shift is not equal to 0

```

The Wilcoxon test confirms that the niche breadth of pigeons in Barcelona is generally higher than that of pigeons in Moià, as we saw at the population level.

In order to calculate the degree of individual specialization, Bolnick et al. [2002] defined WIC/TNW, i.e. the ratio between the within individual component (i.e. average niche width) and the total niche width of the population. Similarly we define the following specialization measure, that takes into account the resemblance between resources:

$$S_{pop} = \frac{\sum_{i=1}^n B_i / n}{B_{pop}}$$

where B_i is the niche breadth of each individual, and B_{pop} is the niche breadth of the population. Note that it is possible that B_i values can be larger than B_{pop} . However, we do not expect the average of B_i values to be larger than B_{pop} . If we calculate S_{pop} for the two populations we have:

```
> Spec.barcelona = mean(indvar.barcelona$B)/popvar.barcelona$B
> Spec.barcelona
```

```
[1] 0.4551234
```

```
> Spec.moia = mean(indvar.moia$B)/popvar.moia$B
> Spec.moia
```

```
[1] 0.4723618
```

Surprisingly, the degree of specialization in Moia seems slightly higher than that in Barcelona. To see whether this holds statistically, we can calculate the degree of specialization of each individual:

$$S_i = \frac{B_i}{B_{pop}}$$

which, in R, is:

```
> Spec.ind.barcelona = indvar.barcelona$B/popvar.barcelona$B
> Spec.ind.moia = indvar.moia$B/popvar.moia$B
```

Finally, we compare this two vectors in a Wilcoxon test:

```
> wilcox.test(Spec.ind.barcelona, Spec.ind.moia)
```

```
Wilcoxon rank sum test with continuity correction
```

```
data: Spec.ind.barcelona and Spec.ind.moia
```

```
W = 277, p-value = 0.1392
```

```
alternative hypothesis: true location shift is not equal to 0
```

Which tells us that those differences in individual specialization are not statistically significant.

4.3 Measuring the degree of overlap between individuals

The idea of this section is to determine how much the niche of each individual overlaps with the niche of other individuals in the population. This can be done by calling function `nicheoverlap` in the ‘pairwise’ mode:

```
> O.barcelona = nicheoverlap(diet.barcelona,D=dfood, mode="pairwise")
> O.moia = nicheoverlap(diet.moia,D=dfood, mode="pairwise")
```

These calls to `nicheoverlap` return a symmetric square matrix with as many rows and columns as individuals in the resource use data frame. Each cell value in the symmetric matrix is the overlap between two individuals of the population. Using these matrices we can derive the average overlap in each population:

```
> mean(O.barcelona[lower.tri(O.barcelona)])
```

```
[1] 0.6726374
```

```
> mean(O.moia[lower.tri(O.moia)])
```

```
[1] 0.8438816
```


We can also calculate the average overlap between each individual and the remaining individuals in its population:

```
> O.barcelona.ind = (rowSums(O.barcelona)-1)/(nrow(O.barcelona)-1)
> summary(O.barcelona.ind)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.5400	0.6028	0.6558	0.6726	0.7595	0.8055

```
> O.moia.ind = (rowSums(O.moia)-1)/(nrow(O.moia)-1)
> summary(O.moia.ind)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.09157	0.89915	0.90649	0.84388	0.90965	0.91134

We subtracted one in the numerator and denominator in order to exclude the target individual from the average (the overlap between a resource niche and itself is always one). Apparently, the individuals in Moia have a larger degree of overlap with individuals of their population than individuals in Barcelona. A non-parametric test seems to confirm this difference:

```
> wilcox.test(O.barcelona.ind, O.moia.ind)
```

Wilcoxon rank sum test with continuity correction

data: O.barcelona.ind and O.moia.ind

W = 37, p-value = 4.549e-06

alternative hypothesis: true location shift is not equal to 0

References

Daniel I. Bolnick, Louie H. Yang, James a. Fordyce, Jeremy M. Davis, and Richard Svanbäck. Measuring individual-level resource specialization. *Ecology*, 83(10):2936–2941, October 2002.

Miquel De Cáceres, Daniel Sol, Oriol Lapiedra, and Pierre Legendre. A framework for estimating niche metrics using the resemblance between qualitative resources. *Oikos*, 120(February):1341–1350, 2011.