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Ranking of critical species to preserve the functionality of mutualistic networks using the k-core decomposition

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Mutualistic communities play an important role in biodiversity preservation. They are modeled as bipartite networks and measurements of centrality and degree help to order species and their relative importance for network robustness. Identifying the most endangered ones or those more prone to trigger cascade extinctions is essential to define conservation policies. In this work, we explain how a classical graph analysis tool, the k-core decomposition, provides new ranking magnitudes that reach outstanding performance for these purposes.

1 Ranking of critical species to preserve the 2 functionality of mutualistic networks using 3 the k-core decomposition

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14 ABSTRACT

15 Mutualistic communities play an important role in biodiversity preservation. They are modeled as
16 bipartite networks and measurements of centrality and degree help to order species and their relative
17 importance for network robustness. Identifying the most endangered ones or those more prone to trigger
18 cascade extinctions is essential to define conservation policies. In this work, we explain how a classical
19 graph analysis tool, the k-core decomposition, provides new ranking magnitudes that reach outstanding
20 performance for these purposes.

21 INTRODUCTION

22 Biotic interaction networks play an essential role in the stability of ecosystems (Tylianakis et al., 2010),
23 as well as in the maintenance of biodiversity (Bascompte et al., 2006). Because community dynamics
24 greatly depend on the way species interact, these networks have been described as the “biodiversity
25 architecture” (Bascompte and Jordano, 2007). Network analysis has become an important approach
26 to provide information on community organization and to predict dynamics and species extinctions in
27 response to ecosystem disturbance (Tylianakis et al., 2010; Thébault and Fontaine, 2010; Traveset and
28 Richardson, 2014). Among other assessments, these studies can point out key species, whose stability
29 would prevent cascading extinctions, and the consequent loss of biodiversity (Sole and Montoya, 2001;
30 Suweis et al., 2013; Dakos et al., 2014; Santamaría et al., 2015). Research on cascading species extinctions
31 as a result of perturbations in biotic interactions has tackled two main issues: the different ways to rank a
32 hypothetical extinction sequence and the robustness and fragility measures (Pocock et al., 2012). There
33 are different strategies both to sort species according to their importance and to measure their influence
34 on extinction. For instance, in early studies on the resilience of food webs Dunne et al. ranked species by
35 degree (i.e., the number of interactions) using three different scenarios of removal: a) from the species
36 with the highest degree to the species with the lowest degree; b) from the lowest to the highest; c) species
37 selected in a random way (Dunne et al., 2002). Memmott et al. worked the same idea to assess the
38 robustness of mutualistic communities, removing active species and measuring the fraction of remaining
39 passive species (Memmott et al., 2004).

40 An observed property of mutualistic interactions is the existence of generalists, highly interconnected,
41 and specialists, with few interactions linked to the generalists, but rarely among them. The nucleus
42 of interactions among generalists seems to be the foundation of resilience. This property has been
43 traditionally identified with nestedness (Bascompte et al., 2003), although there are new approaches to
44 describe it in a more general way as a core-periphery organization (Csermely et al., 2013; Rombach et al.,

45 2014).

46 Identification of key nodes for community preservation is another active field of research. Besides
47 classical measures of centrality, new rankings are available and provide efficient ways to find out them in
48 bipartite networks (Tacchella et al., 2012; Domínguez-García and Muñoz, 2015).

49 In this paper, we aim to explain how the *k-core decomposition*, sheds light on the understanding
50 of robustness in mutualism. The tool classifies the nodes of the network in shells, as in an onion-like
51 structure with the most connected nodes in its center. Taking into account just the very basic topological
52 properties, the decomposition helps to assess in detail the structure of mutualism and enlightens on the
53 processes of species extinction cascades. Derived from the *k-core decomposition* we introduce three
54 new magnitudes, hereafter called *k-magnitudes*, that describe network compactness (*k-radius*), combined
55 quantity and quality of interactions (*k-degree*) and species vulnerability to trigger extinction cascades
56 (*k-risk*). We assess the best criteria for identifying the species for which the networks are most vulnerable
57 to cascade extinctions by comparing *k-degree* and *k-risk* ranking criteria with ranking by well-known
58 indexes and applying them in two network destruction procedures. To conduct the test, we use one of the
59 most complete available data sets (Fortuna et al., 2014).

60 MATERIALS AND METHODS

61 Data

62 We have analyzed the *Web of life* collection (Fortuna et al., 2014), comprised by 89 mutualistic networks,
63 with 59 communities of plants and pollinators and 30 of seed dispersers (<http://www.web-of-life.es/>). There are 57 communities with binary adjacency matrix (i.e., the interaction between the two
64 species is recorded but not its strength), and 32 with weighted matrix, where the strength is accounted for.
65 Network sizes range from 6 to 997 species, the minimum number of links is 6 and the maximum is 2933.
66

67 Decomposition and *k-magnitudes*

68 The idea of core decomposition was first described by Seidman to measure local density and cohesion
69 in social graphs (Seidman, 1983). It has been successfully applied to visualize large systems and
70 networks (Alvarez-Hamelin et al., 2005; Kitsak et al., 2010; Zhang et al., 2010; Barberá et al., 2015).

71 The *k-core* of a network is a maximal connected sub-network of degree greater or equal than *k*. That
72 means that each node is tied to at least *k* other nodes in the same sub-network.

73 A simple algorithm to perform the *k-core decomposition* prunes links of nodes of degree equal or less
74 than *k* (Batagelj and Zaversnik, 2003). The process starts removing links with one of their edges in a
75 node of degree 1. This procedure is recursive and ends when all the remaining nodes have at least two
76 links. The isolated nodes are the *1-shell*. Then it continues with *k* = 2, and so on. After performing the
77 *k-decomposition*, each species belongs to one of the *k-shells* (Fig. 1). The *m-core* includes all nodes of
78 *m-shell*, *m* + 1-shell...

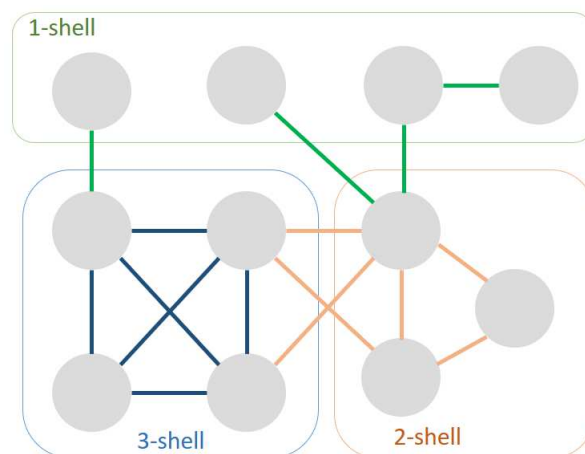


Figure 1. *k-core* decomposition of a fictional network. Green links are pruned during the first iteration, orange during the second and blue during the last one.

79 Mutualistic networks are bipartite, with two guilds of species (plant-pollinator or plant-seed disperser
80 in the studied collection). Links among nodes of the same class are forbidden. We will call these guilds *A*
81 and *B*.

82 Based on the *k*-core decomposition, we define three *k*-magnitudes. In order to quantify the distance
83 from a node to the innermost shell of the partner guild, we define k_{radius} . The k_{radius} of node *m* of guild *A*
84 is the average distance to all species of the innermost shell of guild *B*. We call this set N^B .

$$k_{radius}^A(m) = \frac{1}{|N^B|} \sum_{j \in N^B} dist_{mj} \quad m \in A \quad (1)$$

85 where $dist_{mj}$ is the shortest path from species *m* to each of the *j* species that belong to N^B . The minimum
86 possible k_{radius} value is 1 for one node of the innermost shell directly linked to each one of the innermost
87 shell set of the opposite guild.

88 To obtain a measure of centrality in this *k*-shell based decomposition, we define k_{degree} as

$$k_{degree}^A(m) = \sum_j \frac{a_{mj}}{k_{radius}^B(j)} \quad m \in A, \forall j \in B \quad (2)$$

89 where a_{mj} is the element of the interaction matrix that represents the link, considered as binary. If the
90 network is weighted, a_{mj} will count as 1 for this purpose if there is interaction, 0 otherwise. $k_{degree}(m)$ is
91 a weighted *degree* where each node *i* linked to node *m* adds the inverse of its $k_{radius}(i)$. Generalists score
92 high k_{degree} , whereas specialists, which have only one or two links, with similar k_{radius} , score lower k_{degree} .
93 This magnitude reminds the definition of the *Harary index* (Plavšić et al., 1993) but only considering
94 paths from the nodes tied from *m* to the nodes of the innermost shell.

95 Figure 2 shows how k_{degree} works for one particular network. There are many nodes with the same
96 *degree* value, such as specialists with just one or two links, that from a ranking point of view are equivalent.
97 On the contrary k_{degree} , maps the degree distribution onto a more continuous one, because of the weight
98 of the inverse of k_{radius} . In Fig. 2C the cumulative distributions of both indexes are overlapped over the
99 *degree* scale.

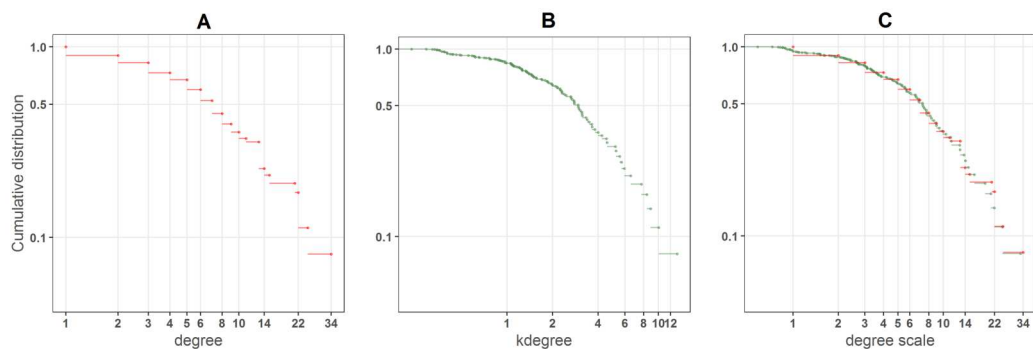


Figure 2. (A)- Degree, (B)- k_{degree} and (C) combined degree and k_{degree} distributions of a big plant pollinator community in Central Los Andes, Chile Arroyo et al. (1982).

100 Finally, we introduce k_{risk} as a way to measure how vulnerable is a network to the loss of a particular
101 species:

$$k_{risk}^A(m) = \sum_i a_{mi} (k_{shell}^A(m) - k_{shell}^B(i)) + \epsilon k_{shell}^A(m) \quad m \in A, \forall i \in B, k_{shell}^B(i) < k_{shell}^A(m) \quad (3)$$

102 The k_{risk} of one species is the sum of the nodes of lower *k*-shells that are tied to it. Each one is
103 weighted by the difference of the *k* indexes. The second element of equation 3 is meant to solve ties
104 among species when they belong to different *k*-shells, and is a very small quantity (in our implementation
105 we use 0.01, two orders of magnitude lesser than the sum).

106 In an intuitive way, if we remove one node strongly connected to others of lower k -shells, these species
 107 are in high risk of being dragged by the primary extinction. On the other hand, the extinction is much less
 108 dangerous for the species of higher k -shells linked to the same node, because they enjoy more redundant
 109 paths towards the network nucleus.

110 Applying the k -magnitudes to a network

111 Fig. 3 is an small seed disperser network with five species of plants, four species of thrushes and
 112 eleven links. We call, by convention, *guild A* the set of plants, and *guild B* the set of birds. The k -core
 113 decomposition was performed with the R *igraph* package (Csardi and Nepusz, 2006) . The maximum
 114 k index is 2. The four bird species belong to *2-shell*; there are three plant species in *1-shell* and two in
 115 *2-shell*. In this example each species of *2-shell* is directly tied to all species of the opposite guild *2-shell*,
 116 but this is not a general rule.

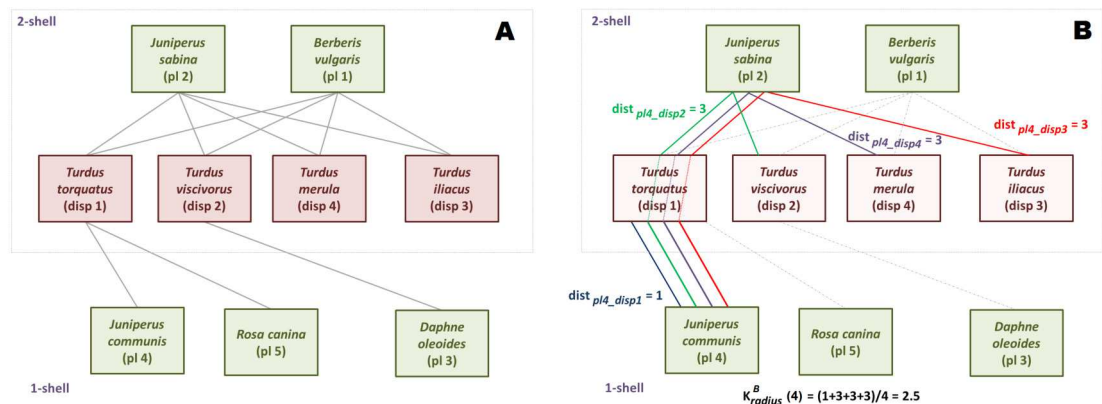


Figure 3. Computation of the k -magnitudes. Seed disperser network in Santa Bárbara, Sierra de Baza (Spain) (Jordano, 1993). A: Decomposed network. B: Computing $k_{radius}^B(4)$.

117 The shortest path from plant species 2 to each of the four bird species of *2-shell* is 1, because of the
 118 direct links. So, $k_{radius}^A(2)$ is 1. The same reasoning is valid for plant species 1. The reader may check that
 119 the k_{radius} of bird species of *2-shell* is 1 as well, measuring their shortest paths to plants species 1 and 2.

120 Computation of this magnitude is simple although a bit more laborious for *1-shell* plant species. We
 121 work plant species 4 as an example. First, we find the shortest paths to each bird species of *2-shell*.
 122 Shortest paths are depicted with different colors. Plant species 4 is tied to seed disperser species 1, so
 123 distance is 1. On the other hand, there is no direct link with bird species 2. Shortest path is *pl4-disp1*-
 124 *pl2-disp2*, and distance is 3. It is easy to check that distances from plant species 4 to bird species 3 and 4
 125 are also 3. Once we have found the four distances, we compute $k_{radius}^B(4)$ as the average of 1, 3, 3 and 3,
 126 that is 2.5.

127 The values of k_{degree} are straightforward to compute. For instance, the k_{degree} of disperser species 1 is:

$$k_{degree}^B(1) = \frac{1}{k_{radius}^A(1)} + \frac{1}{k_{radius}^A(2)} + \frac{1}{k_{radius}^A(4)} + \frac{1}{k_{radius}^A(5)} = 2.8 \quad (4)$$

The last k -magnitude we defined was k_{risk} . We use again the disperser species 1 as example. Links to
 species of the same or upper k -shells are irrelevant to compute k_{risk} , so only bird species 4 and 5 are taken
 into account.

$$k_{risk}^B(1) = k_{shell}^B(1) - k_{shell}^A(4) + k_{shell}^B(1) - k_{shell}^A(5) + \epsilon k_{shell}^B(1) = (2-1) + (2-1) + 0.01 \times 2 = 2.02 \quad (5)$$

128 This magnitude may seem counter-intuitive, because the k_{risk} of a highly connected species like plant
 129 1 is 0.02, almost the same of that of peripheral plant 3. This is because plant 1 has no ties with lower
 130 k -shell animal species. The k_{risk} ranks species to assess resilience, it has not an absolute meaning. It just
 131 tells us that it is more dangerous for the network to remove the disperser 1 than plant 1, and plant 1 than
 132 plant 3.

133 The k -magnitudes of the example network are shown in table 1.

<i>Species</i>	k_{shell}	k_{radius}	k_{degree}	k_{risk}
pl1	2	1	4	0.02
pl2	2	1	4	0.02
pl3	1	2.5	1	0.01
pl4	1	2.5	1	0.01
pl5	1	2.5	1	0.01
disp1	2	1	2.8	2.02
disp2	2	1	2.4	1.02
disp3	2	1	2	0.02
disp4	2	1	2	0.02

Table 1. K -magnitudes of the network of Fig. 3.

134 Extinction procedures

135 We carried out two static extinction procedures. Static assumption implies that there is not rewiring (e.g.,
 136 plants that have lost their pollinators are not pollinated by other insects), despite this kind of network
 137 reorganization is observed in nature (Ramos-Jiliberto et al., 2012; Goldstein and Zych, 2016; Timóteo
 138 et al., 2016). Nodes are ranked once, before the procedure starts.

139 In the first method, one species is removed each step, in decreasing order according to the chosen
 140 index, no matter to which guild it belongs. Four ranking indexes are compared: k_{risk} , k_{degree} , $degree$
 141 and $eigenvector\ centrality$. The k indexes were computed with the R package `kcorebip`; $degree$ and
 142 $eigenvector\ centrality$ with the `degree` and `evcent` functions of the `igraph` package.

143 To estimate the damage caused to the network, the fraction of remaining giant component (i.e., the
 144 highest connected component of a given network) was used. The procedure stops when this ratio is equal
 145 or less than 0.5. To break ties, we ran 100 experiments for each network and index, shuffling species with
 146 the same ranking value. The percentage of removed species needed to get to 0.5 of the remaining giant
 147 component is used to measure the performance of the ranking. The lower the percentage of removed
 148 species, the more efficient the ranking is in destroying the network. The top performer scores the least
 149 average removal percentage. (Fig. 4).

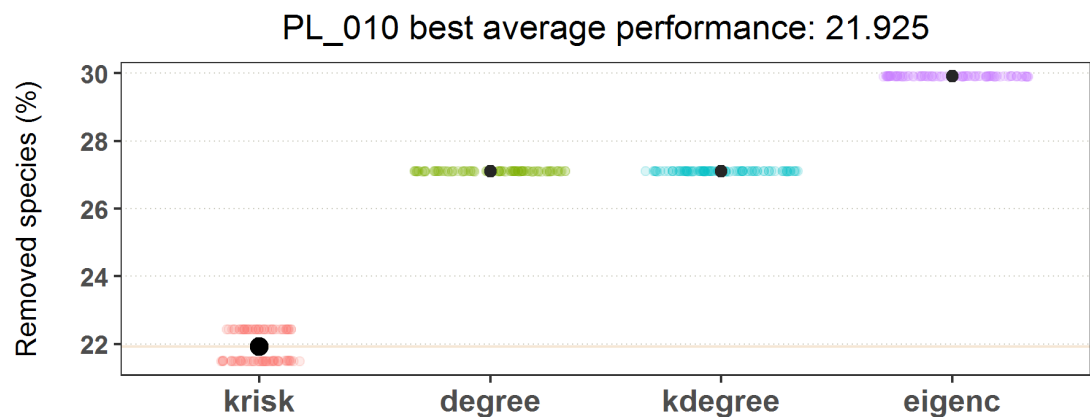


Figure 4. First extinction procedure. Performance of the four ranking indexes for a pollinator community described by Elberling and Olesen in Zackenberg Station (Greenland, unpublished). Individual dots are the results of each experiment while black dots are the average values

150 The second extinction procedure that we followed is more common in the literature. Only animal
 151 species are actively removed (primary extinctions); secondary extinctions happen when nodes become
 152 isolated (Memmott et al., 2004).

153 The fraction of surviving plant species is measured as a function of the removed fraction of animal
 154 species (Fig. 5A,C) and the area under the curve is the value to compare performance. We averaged the

155 results of 100 repetitions.

156 In this case, in addition to the four indexes of the first experiment, we include *MusRank* a non-linear
 157 ranking algorithm for bipartite networks (Tacchella et al., 2012), inspired by *PageRank* (Allesina and
 158 Pascual, 2009). This algorithm is not valid for the first extinction method. Domínguez-García and Muñoz
 159 showed that *MusRank* achieves excellent performance for this extinction procedure (Domínguez-García
 160 and Muñoz, 2015).

161 In the second extinction procedure, we also measured the fraction of remaining giant component (Fig.
 162 5B,D) . Extinction sequences are identical, the only difference is that both magnitudes are measured for
 163 each step.

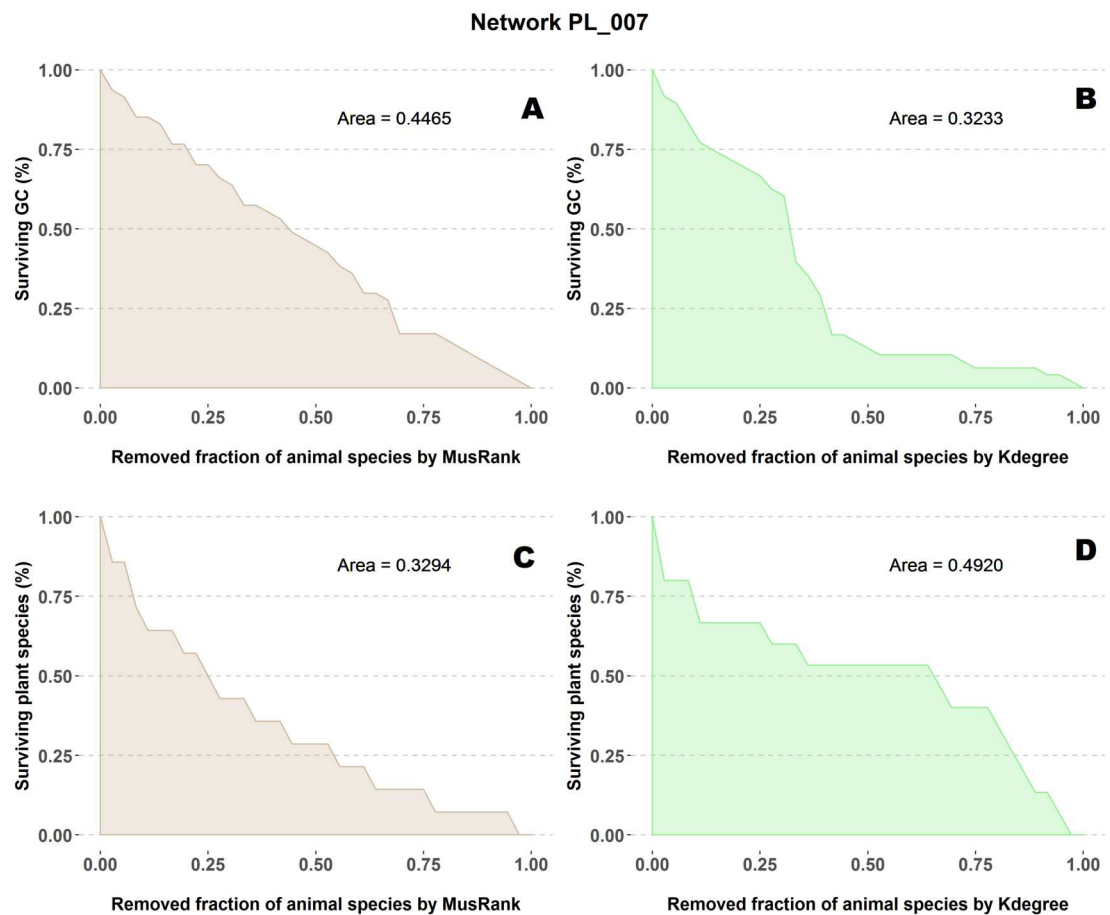


Figure 5. Extinction curves of the second algorithm for a pollination network in Suffolk, UK (Dicks et al., 2002). A,C: Percentage of surviving giant component and percentage of surviving plant species removing animal species ranked by *MusRank*. B,D: Percentage of surviving giant component and percentage of surviving plant species removing animal species ranked by k_{degree} .

164 RESULTS

165 First extinction method

166 k_{risk} was the ranking method with the lowest average species removal percentage to destroy half of
 167 the Giant Component in most of the networks (67 out of 89 networks) (Table 2). Figure 6 shows the
 168 performance comparison of the four ranking criteria. There are some ties, more frequent when networks
 169 are small. Network size is the key factor to explain why the performance range is so wide.

170 As size increases, the removal percentage to break the giant component decreases. When the network
 171 is big, the primary extinction of key nodes triggers an important amount of secondary ones. If the

172 community has 100 or more species, k_{risk} is even a better predictor of the most damaging extinction
 173 sequence and outperforms the other indexes for 28 out of 32 networks.

Table 2. Average number of removed species to destroy half the Giant Component, according to the different indexes: k_{risk} is the top performer for 67 networks, $degree$ for 48, k_{degree} for 39 and $eigenvector$ centrality for 28 networks.

<i>Network</i>	<i>GCsize</i>	k_{risk}	<i>degree</i>	k_{degree}	<i>eigen</i>	<i>Network</i>	<i>GCsize</i>	k_{risk}	<i>degree</i>	k_{degree}	<i>eigen</i>
PL_001	177	21.73	22.13	22	23	PL_046	60	11	11.94	13	14
PL_002	103	14.46	12.51	13	15	PL_047	205	4	4	4	4
PL_003	61	5	5.35	6	6	PL_048	266	10	10	9	12
PL_004	112	3	3	3	3	PL_049	262	11	13	15	16
PL_005	361	25	30.73	36	42	PL_050	49	6	6.36	7	7
PL_006	78	3	3	3	3	PL_051	104	3	3	3	3
PL_007	50	5	4	4	4	PL_052	52	6	6	6	7
PL_008	49	6	7	7	11	PL_053	364	19	22.49	23	34
PL_009	142	7	7.52	8	12	PL_054	414	23	25.23	27	30
PL_010	107	23.46	29	29	32	PL_055	253	16.40	17	19	
PL_011	27	4	5.04	6	6	PL_056	456	22	28.71	33	43
PL_012	84	7	7	7	7	PL_057	997	17	17	20	36
PL_013	65	4	4	4	5	PL_058	111	14	17.55	19	20
PL_014	108	6	5	5	6	PL_059	26	6	5	5	5
PL_015	793	48	56	60	87	SD_001	28	3	3	3	5
PL_016	205	9	9	10	17	SD_002	40	5	5	5	5
PL_017	104	9	10.52	11	11	SD_003	41	4	4	4	4
PL_018	144	18	19	23	24	SD_004	52	4	4	4	4
PL_019	123	14	15.37	16	18	SD_005	34	3	3	3	3
PL_020	109	3	3	3	3	SD_006	34	4	4.31	5	5
PL_021	766	12	12	12	38	SD_007	79	3	3	3	3
PL_022	66	4	4	4	4	SD_008	26	9.83	8.45	8	11
PL_023	90	3	3	3	3	SD_009	25	3	4	4	5
PL_024	22	4	3.55	3	3.0	SD_010	64	8	8	9	13
PL_025	57	6	6	6	10	SD_011	25	6	5.33	5	6
PL_026	150	2	2	2	2	SD_012	64	12.71	12.53	12	14
PL_027	75	8.54	8	9	11	SD_013	55	11	8	19	14
PL_028	180	13	14	16	24	SD_014	33	9	10	10	10
PL_029	167	17	16.97	17	19	SD_015	32	4	4	4	4
PL_030	70	10.23	6.65	7	13	SD_016	85	17	18	20	23
PL_031	91	9.53	7.92	13	17	SD_017	24	7.28	6.62	10	10
PL_032	40	2	2	2	2	SD_018	53	5	5	5	5
PL_033	47	8.65	8	10	12	SD_019	209	13	16.48	20	21
PL_034	151	6	7.54	8	9	SD_020	58	7.65	9.22	10	10
PL_035	97	6	7.54	8	9	SD_021	46	9	10	10	10
PL_036	22	3.68	2	2	2	SD_022	317	39	50.43	53	60
PL_037	50	5	5	5	7	SD_023	23	4	4	4	4
PL_038	50	4	4	4	7	SD_024	19	6	6	7	8
PL_039	68	6	6.68	9	10	SD_025	13	4.38	4.16	4.16	4
PL_040	70	8.56	7	9	10	SD_026	6	2	2	2	2
PL_041	70	10	10	11	12	SD_027	16	3	3	3	3
PL_042	16	2	2	2	2	SD_028	13	3	3	3	3
PL_043	110	12	14	14	19	SD_029	9	2	2	2	2
PL_044	712	21	23	25	49	SD_030	9	2	2.65	2	2
PL_045	41	6	5.45	7	8						

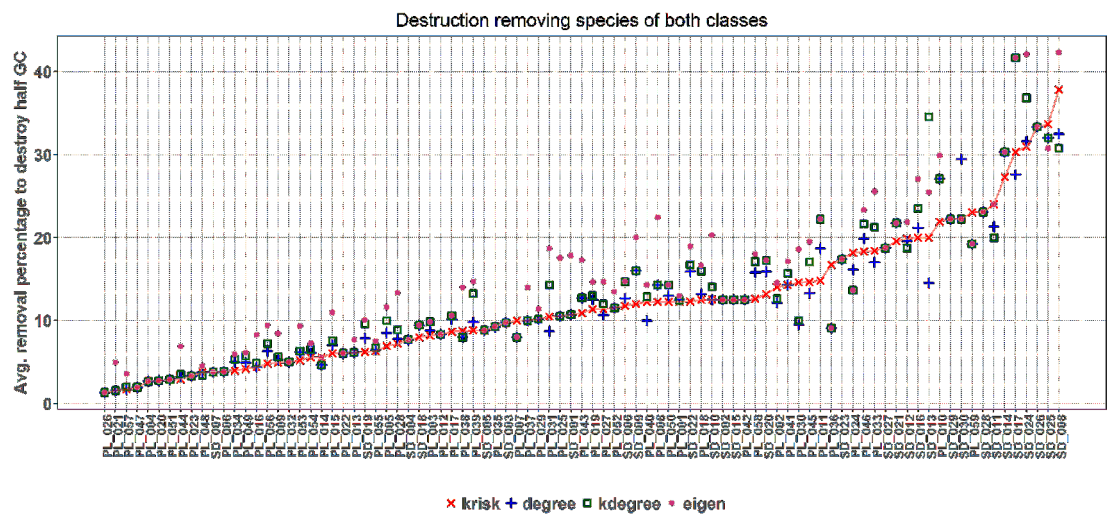


Figure 6. First extinction method. The average percentage of removed species to destroy the Giant Component is depicted for each network and ranking index. Under the X axis, the name of each network as coded in the *web of life* database. The overall top performer is k_{risk} (see Table 2). Species are ordered by the percentage of primary extinctions, ranked by k_{risk} . The red line joins the k_{risk} destruction percentage values as a visual reference to compare them with those of the other indexes.

174 **Second extinction method**

175 *MusRank* ranking method had the lowest area under the extinction curve for 85 of the 89 studied networks
 176 (Figure 7), and in the other 4 the difference is so small that may be just an effect of the averaging procedure.
 177 So, *MusRank* is the optimal ranking index to destroy the network following this algorithm.

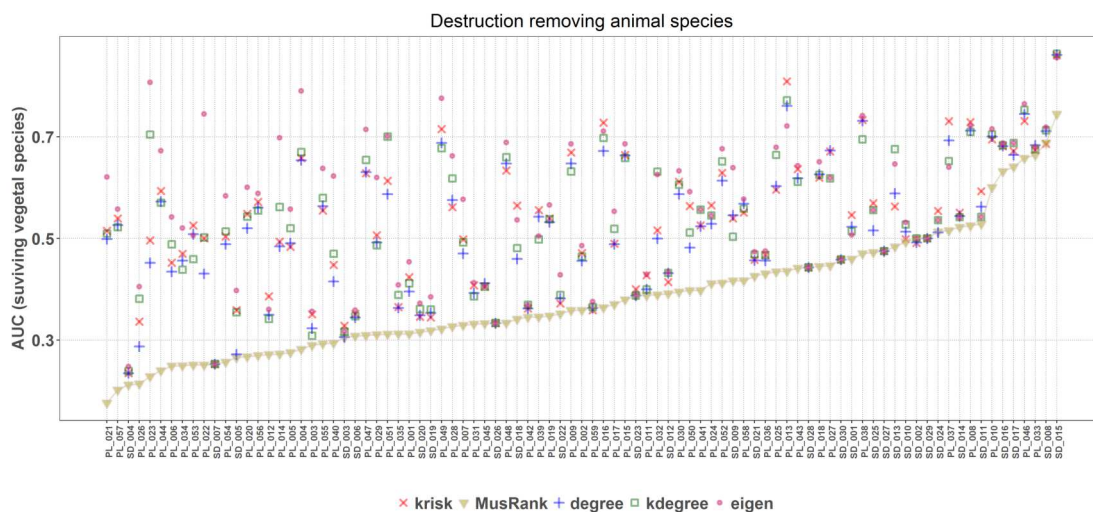


Figure 7. Second extinction procedure, measuring the surviving plant species fraction. The Area Under the extinction Curve is depicted for each network and index. The overall top performer is *MusRank* (see Table 3). The solid line joins the *MusRank* values. Species are ordered by the percentage of primary extinctions, ranked by *MusRank*.

178 On the contrary, when the efficiency of the network destruction was measured through the area under
 179 the curve of the surviving Giant Component fraction the *MusRank* index had the highest values, placing it
 180 as the least efficient ranking method according to this criterion (Figure 8). In this case, k_{degree} is the most
 181 efficient index for 42 out of 89 networks. We must underline that the extinction sequences are the same,
 182 the only difference is the measured output.

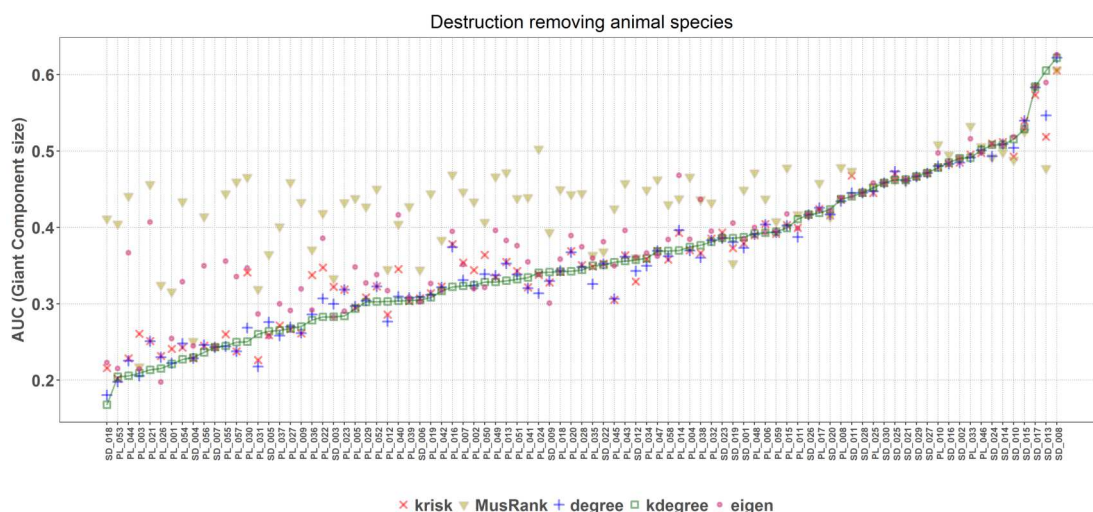


Figure 8. Second extinction procedure, measuring the surviving Giant Component fraction. The Area Under the extinction Curve is depicted for each network and index. The overall top performer is k_{degree} (see Table 4). The solid line joins the k_{degree} values. Species are ordered by the percentage of primary extinctions, ranked by k_{degree} .

183 Fig. 5 is an example of this counterintuitive response. On the upper row (subplots A and B), the

184 difference for both ranking indexes when measuring the giant component. While this magnitude decreases
 185 at a constant pace for *MusRank*, there is a sharp reduction of the component size when one third of animal
 186 species are removed following the k_{degree} ranking. On the lower row (subplots C and D), opposite results
 187 are obtained when accounting for the fraction of surviving plant species.

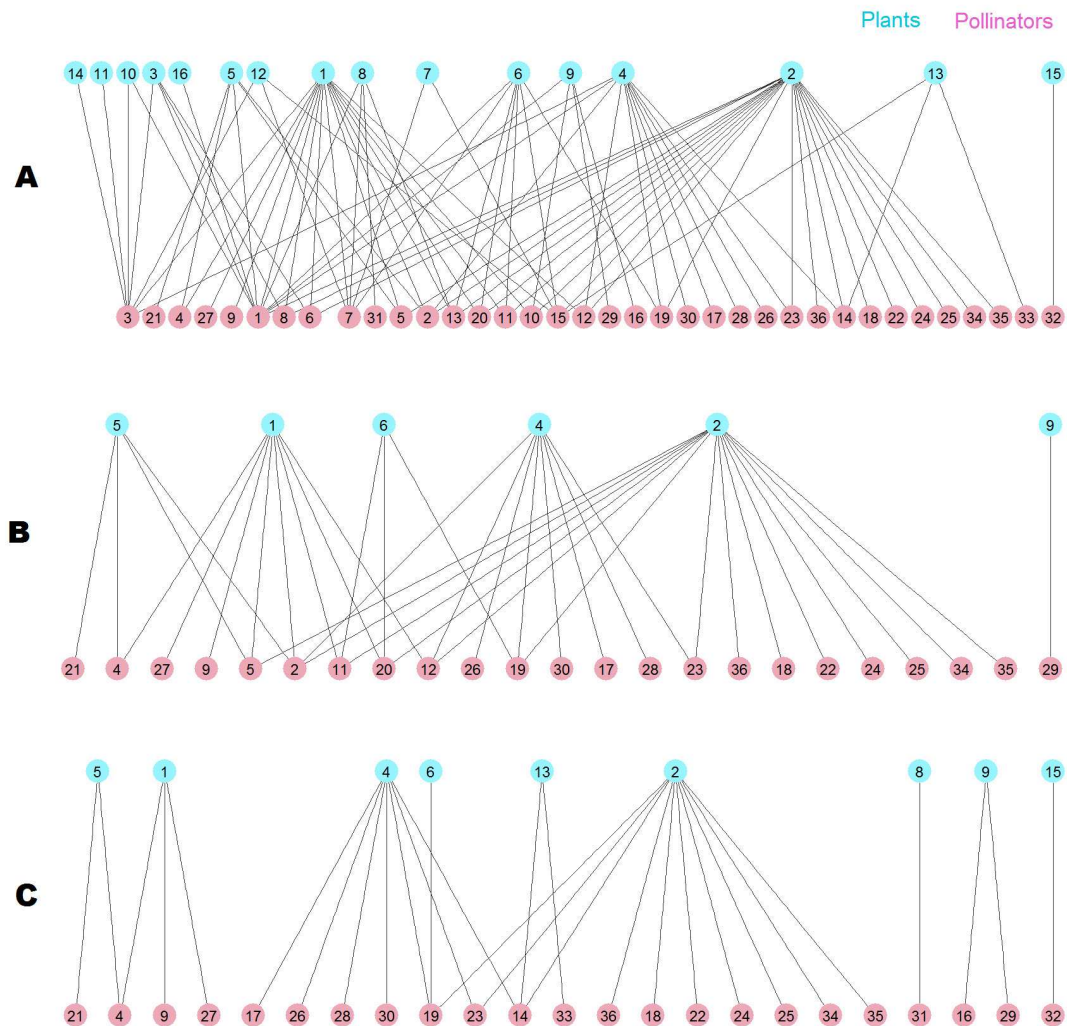


Figure 9. Pollinator network 007 (Dicks et al., 2002). A: Original configuration; B: Structure after the removal of the 13 top *MusRank*-ranked animal species. C: Structure after the removal of the 13 top k_{degree} -ranked animal species

188 The destruction of this pollinator network sheds light on the root cause of the difference. The network
 189 has 36 pollinator and 16 plant species (Fig.9A), 2 of them are outside the giant component. When the 13
 190 top animal species ranked by *MusRank* are removed (pollinators 3, 1, 7, 15, 32, 6, 14, 33, 13, 31, 8, 16, 10),
 191 the community reaches the degraded structure of Fig.9B. The size of the giant component is 27 (54% of
 192 the original), and there are 23 pollinator and 6 plant species.

193 If we remove the 13 top animal species ranked by k_{degree} (pollinators 1, 3, 7, 13, 15, 2, 11, 20, 12, 8, 6, 5, 10)
 194 instead, the community structure is that of Fig.9C. Now, the size of the giant component is 19 (38% of the
 195 original), and there are 23 pollinator and 9 plant species. *MusRank* has killed more plant species, but the
 196 giant component is clearly smaller ranking by k_{degree} .

Table 3. Average area under the extinction curve, when the surviving fraction of plant species is measured. *MusRank* is the top performer for 85 networks, k_{risk} for 9, *degree* for 8, k_{degree} for 7 and *eigenvector centrality* for 7.

<i>Network</i>	<i>MR</i>	k_{risk}	k_{degree}	<i>degree</i>	<i>eigenv</i>	<i>Network</i>	<i>MR</i>	k_{risk}	k_{degree}	<i>degree</i>	<i>eigenv</i>
PL_001	0.3121	0.4236	0.4115	0.3956	0.4535	PL_046	0.6577	0.7308	0.7528	0.7445	0.7642
PL_002	0.3588	0.4710	0.4641	0.4555	0.4856	PL_047	0.3091	0.6283	0.6541	0.6302	0.7143
PL_003	0.2896	0.3505	0.3085	0.3232	0.3563	PL_048	0.3336	0.6334	0.6594	0.6471	0.6887
PL_004	0.2821	0.6579	0.6695	0.6536	0.7900	PL_049	0.3220	0.7150	0.6773	0.6880	0.7757
PL_005	0.2755	0.4836	0.5198	0.4906	0.5573	PL_050	0.3976	0.5635	0.5114	0.4815	0.5918
PL_006	0.2490	0.4520	0.4882	0.4344	0.5420	PL_051	0.3115	0.6132	0.7000	0.5870	0.7008
PL_007	0.3294	0.4982	0.4920	0.4702	0.5769	PL_052	0.4121	0.6289	0.6514	0.6132	0.6761
PL_008	0.5251	0.7287	0.7093	0.7117	0.7213	PL_053	0.2510	0.5253	0.4590	0.5081	0.5058
PL_009	0.3583	0.6689	0.6315	0.6473	0.6857	PL_054	0.2568	0.5035	0.5136	0.4885	0.5835
PL_010	0.6004	0.6948	0.7044	0.7003	0.7150	PL_055	0.2929	0.5548	0.5795	0.5633	0.6378
PL_011	0.3876	0.4272	0.3994	0.3999	0.4290	PL_056	0.2696	0.5717	0.5545	0.5605	0.5885
PL_012	0.2716	0.3860	0.3417	0.3494	0.3604	PL_057	0.2019	0.5392	0.5217	0.5261	0.5576
PL_013	0.4353	0.8088	0.7716	0.7606	0.7212	PL_058	0.4168	0.5507	0.5585	0.5683	0.5773
PL_014	0.2726	0.4935	0.5617	0.4840	0.6980	PL_059	0.3639	0.3587	0.3649	0.3631	0.3757
PL_015	0.3792	0.6637	0.6579	0.6633	0.6859	SD_001	0.4592	0.5459	0.5141	0.5229	0.5068
PL_016	0.3641	0.7275	0.6973	0.6719	0.7108	SD_002	0.5000	0.4911	0.5000	0.4919	0.5000
PL_017	0.3703	0.4895	0.5187	0.4886	0.5532	SD_003	0.3051	0.3281	0.3166	0.3059	0.3166
PL_018	0.4451	0.6194	0.6249	0.6259	0.6504	SD_004	0.2121	0.2351	0.2389	0.2350	0.2476
PL_019	0.3474	0.5391	0.5373	0.5320	0.5658	SD_005	0.2656	0.3587	0.3546	0.2718	0.3974
PL_020	0.2674	0.5480	0.5427	0.5199	0.6005	SD_006	0.3078	0.3515	0.3471	0.3444	0.3588
PL_021	0.1761	0.5147	0.5092	0.4985	0.6207	SD_007	0.2528	0.2528	0.2528	0.2528	0.2528
PL_022	0.2511	0.5003	0.5019	0.4304	0.7444	SD_008	0.6875	0.6861	0.7125	0.7108	0.7188
PL_023	0.2285	0.4959	0.7041	0.4517	0.8068	SD_009	0.4167	0.5395	0.5033	0.5453	0.6389
PL_024	0.4111	0.5650	0.5449	0.5283	0.5444	SD_010	0.4929	0.4986	0.5271	0.5129	0.5314
PL_025	0.4344	0.5956	0.6641	0.6030	0.6792	SD_011	0.5286	0.5927	0.5422	0.5623	0.5422
PL_026	0.2138	0.3362	0.3811	0.2874	0.4052	SD_012	0.3912	0.4140	0.4321	0.4316	0.4332
PL_027	0.4466	0.6711	0.6179	0.6728	0.6185	SD_013	0.4835	0.5629	0.6754	0.5885	0.6462
PL_028	0.3266	0.5613	0.6178	0.5754	0.6621	SD_014	0.5221	0.5504	0.5415	0.5441	0.5404
PL_029	0.3107	0.5061	0.4865	0.4919	0.6198	SD_015	0.7444	0.8602	0.8633	0.8607	0.8556
PL_030	0.3949	0.6113	0.6054	0.5870	0.6329	SD_016	0.6318	0.6812	0.6830	0.6816	0.6872
PL_031	0.3314	0.4078	0.3857	0.3926	0.4128	SD_017	0.6406	0.6719	0.6875	0.6639	0.6875
PL_032	0.3889	0.5157	0.6312	0.4995	0.6255	SD_018	0.3413	0.5645	0.4809	0.4596	0.5361
PL_033	0.6640	0.6732	0.6753	0.6832	0.6799	SD_019	0.3185	0.3444	0.3602	0.3535	0.3849
PL_034	0.2497	0.4697	0.4383	0.4563	0.5204	SD_020	0.3155	0.3459	0.3608	0.3485	0.3722
PL_035	0.3121	0.3648	0.3886	0.3633	0.4085	SD_021	0.4254	0.4579	0.4682	0.4567	0.4737
PL_036	0.4306	0.4677	0.4667	0.4560	0.4750	SD_022	0.3517	0.3720	0.3886	0.3820	0.4284
PL_037	0.5153	0.7303	0.6521	0.6924	0.6400	SD_023	0.3875	0.4000	0.3875	0.3875	0.3875
PL_038	0.4702	0.7328	0.6948	0.7310	0.7411	SD_024	0.5130	0.5542	0.5357	0.5114	0.5357
PL_039	0.3454	0.5556	0.4979	0.5425	0.5046	SD_025	0.4722	0.5692	0.5556	0.5156	0.5556
PL_040	0.2940	0.4478	0.4696	0.4153	0.6227	SD_026	0.3333	0.3333	0.3333	0.3333	0.3333
PL_041	0.3978	0.5253	0.5562	0.5238	0.5566	SD_027	0.4750	0.4750	0.4750	0.4750	0.4750
PL_042	0.3452	0.3621	0.3690	0.3621	0.3690	SD_028	0.4429	0.4429	0.4429	0.4429	0.4429
PL_043	0.4414	0.6364	0.6113	0.6183	0.6430	SD_029	0.5000	0.5000	0.5000	0.5000	0.5000
PL_044	0.2401	0.5932	0.5701	0.5723	0.6721	SD_030	0.4583	0.4583	0.4583	0.4583	0.4583
PL_045	0.3324	0.4052	0.4047	0.4116	0.4063						

Table 4. Average area under the extinction curve, when the surviving fraction of the original giant component is measured. The top performer is k_{degree} for 42 networks, $degree$ for 24, k_{risk} for 21, $eigenvector$ centrality for 18 and $MusRank$ for 16.

Network	MusRank	k_{risk}	k_{degree}	degree	eigenv	Network	MusRank	k_{risk}	k_{degree}	degree	eigenv
PL_001	0.3158	0.2410	0.2212	0.2224	0.2546	PL_046	0.5060	0.4975	0.5013	0.5008	0.4987
PL_002	0.4337	0.3441	0.3244	0.3237	0.3198	PL_047	0.4626	0.3694	0.3686	0.3694	0.3622
PL_003	0.2176	0.2607	0.2093	0.2056	0.2145	PL_048	0.4715	0.3895	0.3920	0.3909	0.3998
PL_004	0.4661	0.3700	0.3744	0.3697	0.3843	PL_049	0.4662	0.3350	0.3286	0.3372	0.3960
PL_005	0.4380	0.2954	0.2935	0.2968	0.3482	PL_050	0.4068	0.3641	0.3285	0.3387	0.3214
PL_006	0.4374	0.4044	0.3929	0.4038	0.3932	PL_051	0.4380	0.3428	0.3322	0.3384	0.3762
PL_007	0.4465	0.3539	0.3233	0.3310	0.3513	PL_052	0.4502	0.3228	0.3029	0.3227	0.3381
PL_008	0.4786	0.4363	0.4370	0.4344	0.4386	PL_053	0.4046	0.2018	0.2045	0.1980	0.2153
PL_009	0.4331	0.2613	0.2703	0.2617	0.3195	PL_054	0.4337	0.2427	0.2273	0.2481	0.3287
PL_010	0.5086	0.4789	0.4783	0.4802	0.4974	PL_055	0.4443	0.2602	0.2449	0.2443	0.3557
PL_011	0.4169	0.3994	0.4108	0.3874	0.3985	PL_056	0.4142	0.2451	0.2364	0.2460	0.3498
PL_012	0.3451	0.2859	0.3030	0.2768	0.3173	PL_057	0.4597	0.2378	0.2498	0.2377	0.3355
PL_013	0.4718	0.3548	0.3302	0.3527	0.3828	PL_058	0.4301	0.3578	0.3694	0.3622	0.3840
PL_014	0.4378	0.3928	0.3697	0.3965	0.4679	PL_059	0.4077	0.3916	0.3944	0.3934	0.3954
PL_015	0.4780	0.4030	0.3991	0.4025	0.4177	SD_001	0.4489	0.3792	0.3871	0.3732	0.3854
PL_016	0.4689	0.3781	0.3221	0.3743	0.3946	SD_002	0.4905	0.4849	0.4905	0.4847	0.4905
PL_017	0.4579	0.4240	0.4190	0.4257	0.4234	SD_003	0.3333	0.3222	0.2829	0.2999	0.2829
PL_018	0.4498	0.3419	0.3421	0.3432	0.3585	SD_004	0.2510	0.2286	0.2298	0.2288	0.2448
PL_019	0.4441	0.3138	0.3083	0.3118	0.3265	SD_005	0.3646	0.2592	0.2640	0.2762	0.2569
PL_020	0.4433	0.3688	0.3426	0.3674	0.3894	SD_006	0.3444	0.3072	0.3049	0.3088	0.3025
PL_021	0.4563	0.2512	0.2136	0.2510	0.4071	SD_007	0.2433	0.2433	0.2433	0.2433	0.2433
PL_022	0.4184	0.3475	0.2828	0.3068	0.3858	SD_008	0.6060	0.6056	0.6220	0.6221	0.6260
PL_023	0.4327	0.3187	0.2840	0.3185	0.2901	SD_009	0.3935	0.3280	0.3415	0.3296	0.3009
PL_024	0.5028	0.3380	0.3406	0.3135	0.3389	SD_010	0.4881	0.4926	0.5153	0.5040	0.5187
PL_025	0.4529	0.4451	0.4521	0.4472	0.4582	SD_011	0.4736	0.4678	0.4405	0.4452	0.4405
PL_026	0.3245	0.2316	0.2153	0.2300	0.1976	SD_012	0.3603	0.3291	0.3575	0.3429	0.3609
PL_027	0.4590	0.2669	0.2676	0.2698	0.2912	SD_013	0.4772	0.5185	0.6053	0.5466	0.5897
PL_028	0.4445	0.3505	0.3444	0.3483	0.3743	SD_014	0.4982	0.5117	0.5081	0.5092	0.5074
PL_029	0.4270	0.3084	0.3024	0.3040	0.3272	SD_015	0.5251	0.5392	0.5288	0.5397	0.5299
PL_030	0.4658	0.3410	0.2504	0.2685	0.3469	SD_016	0.4949	0.4832	0.4849	0.4833	0.4852
PL_031	0.3190	0.2263	0.2603	0.2178	0.2866	SD_017	0.5842	0.5734	0.5842	0.5832	0.5842
PL_032	0.4322	0.3853	0.3810	0.3831	0.3951	SD_018	0.4112	0.2159	0.1678	0.1806	0.2231
PL_033	0.5326	0.4953	0.4910	0.4914	0.5160	SD_019	0.3528	0.3729	0.3861	0.3807	0.4058
PL_034	0.4489	0.3585	0.3596	0.3497	0.3664	SD_020	0.4135	0.4156	0.4236	0.4169	0.4229
PL_035	0.3636	0.3488	0.3498	0.3259	0.3595	SD_021	0.4594	0.4617	0.4627	0.4610	0.4627
PL_036	0.3708	0.3377	0.2785	0.2859	0.2917	SD_022	0.3681	0.3525	0.3508	0.3522	0.3814
PL_037	0.4010	0.2716	0.2652	0.2585	0.2997	SD_023	0.3860	0.3934	0.3860	0.3860	0.3860
PL_038	0.4366	0.3665	0.3767	0.3605	0.4366	SD_024	0.4916	0.5098	0.5079	0.4931	0.5079
PL_039	0.4274	0.3036	0.3042	0.3080	0.3083	SD_025	0.4621	0.4702	0.4621	0.4732	0.4621
PL_040	0.4042	0.3454	0.3037	0.3096	0.4164	SD_026	0.4167	0.4167	0.4167	0.4167	0.4167
PL_041	0.4394	0.3216	0.3344	0.3205	0.3550	SD_027	0.4712	0.4712	0.4712	0.4712	0.4712
PL_042	0.3833	0.3227	0.3167	0.3218	0.3167	SD_028	0.4455	0.4455	0.4455	0.4455	0.4455
PL_043	0.4577	0.3634	0.3558	0.3616	0.3959	SD_029	0.4667	0.4667	0.4667	0.4667	0.4667
PL_044	0.4410	0.2286	0.2058	0.2255	0.3669	SD_030	0.4583	0.4583	0.4583	0.4583	0.4583
PL_045	0.4246	0.3052	0.3544	0.3066	0.3496						

197 DISCUSSION

198 The *k-core decomposition* offers a new topological view of the structure of mutualistic networks. We have
199 defined three new magnitudes to take advantage of their properties. Network compactness is described
200 by k_{radius} , a measure of average proximity to top generalists of the partner guild. Second, k_{degree} maps
201 each node's degree onto a finer grain distribution. It has not only information on the number of neighbors
202 but also on how they are connected to the innermost shell. Finally, k_{risk} is set to identify species whose
203 disappearance poses a greater risk to the entire network.

204 Comparing the *k-magnitudes* based extinction indexes (k_{degree} and k_{risk}) with those routinely used
205 when extinctions take place in both guilds, k_{risk} is the best rank if the goal is to identify the key species to
206 preserve most of the giant component. k_{risk} identifies species linked to a high number of nodes of lower
207 *k-shells*. These species provide vulnerability to the network because their extinction may drag many of
208 the species with lower *k-shells* they are linked to, to extinction as well, as they do not enjoy redundant
209 paths to the innermost shell.

210 Applying the well-known method of removing species of the primary class and measuring the
211 extinctions in secondary class, the most effective extinction sequence, if the goal is to identify the key
212 species to preserve most of the giant component, is k_{degree} . However, if the goal is to identify the key
213 species to preserve the greatest species richness in the second class (e.g., plants in a plant-pollinator
214 mutualistic network), the best criterion is *MusRank* as Fig. 7 makes clear. These results confirm those
215 obtained by Domínguez-García and Muñoz (2015), over a larger network collection (89 in this work vs.
216 67 in the original paper).

217 The most striking result of the second method is how different performance is for a same ranking
218 index, depending on the magnitude we measure. The root cause lies on the definitions of the indexes
219 themselves. *MusRank* is optimal to destroy the plant guild. It identifies the most important active nodes of
220 the bipartite network because of how they are linked to the most vulnerable passive ones. It was designed
221 to excel with this extinction sequence and works with local properties. On the other hand, k_{degree} is an
222 excellent performer to destroy the giant component. It contains information on how nodes are connected
223 to the innermost shell, and ranks higher those nodes strongly tied to that stable nucleus.

224 In summary, in this study, we show that the new *k-core* decomposition derived indexes, k_{risk} and
225 k_{degree} provide a new insight into the structure of mutualistic networks. This insight is particularly useful
226 because these indexes fair much better than other traditionally used ranking indexes, when the aim is to
227 identify the species that are key to preserving the interactions and the functionality of the community.
228 As complex network studies on mutualistic interactions are already being used to suggest conservation
229 policies, it is of utmost importance to have a clear framework of what the conservation practitioners look
230 for when implementing conservation and restoration plans. The static view of considering biodiversity
231 conservation as the mere conservation of a list of species has long been substituted by a new paradigm
232 which looks at conservation from a dynamic viewpoint in which species interactions and the functionality
233 of the ecosystems play a major role (Heywood and Iriondo, 2003).

234 ADDITIONAL INFORMATION AND DECLARATIONS

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240 Competing Interests

241 The authors declare no competing interests.

242 Author Contributions

243 J.G.-A., J.M.P and J.G. developed the mathematical analysis. J.G.-A. wrote the R code and J.M.P. the
244 Python code and they both performed simulations. J.M.I. provided advice in the overall design of the
245 work and ecological interpretation of results. All authors wrote the paper.

246 **Code**

247 The R code for *k*-core decomposition and plotting has been published as a package at [https://www.](https://www.github.com/jgalgarra/kcorebip)
248 [github.com/jgalgarra/kcorebip](https://www.github.com/jgalgarra/kcorebip).
249 The rest of software is available at https://github.com/jgalgarra/kcore_robustness
250 Reproducibility instructions are detailed in the README.md file

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