

Interpretation of graph theory

metrics for species

persistence in a

metapopulation.

The Gulf of Lion

study case.



Costa A., Doglioli A.M., Guizien K., Petrenko A.A.

METAPOPULATION MODEL

(see Guizien et al, 2014)

Species: soft-bottom polychaete

Site: Gulf of Lion

$$N_{t+1} = \min(b_j C_{ij} a_i N_t + s_{jj} \delta_{ji} N_t, N_{max})$$

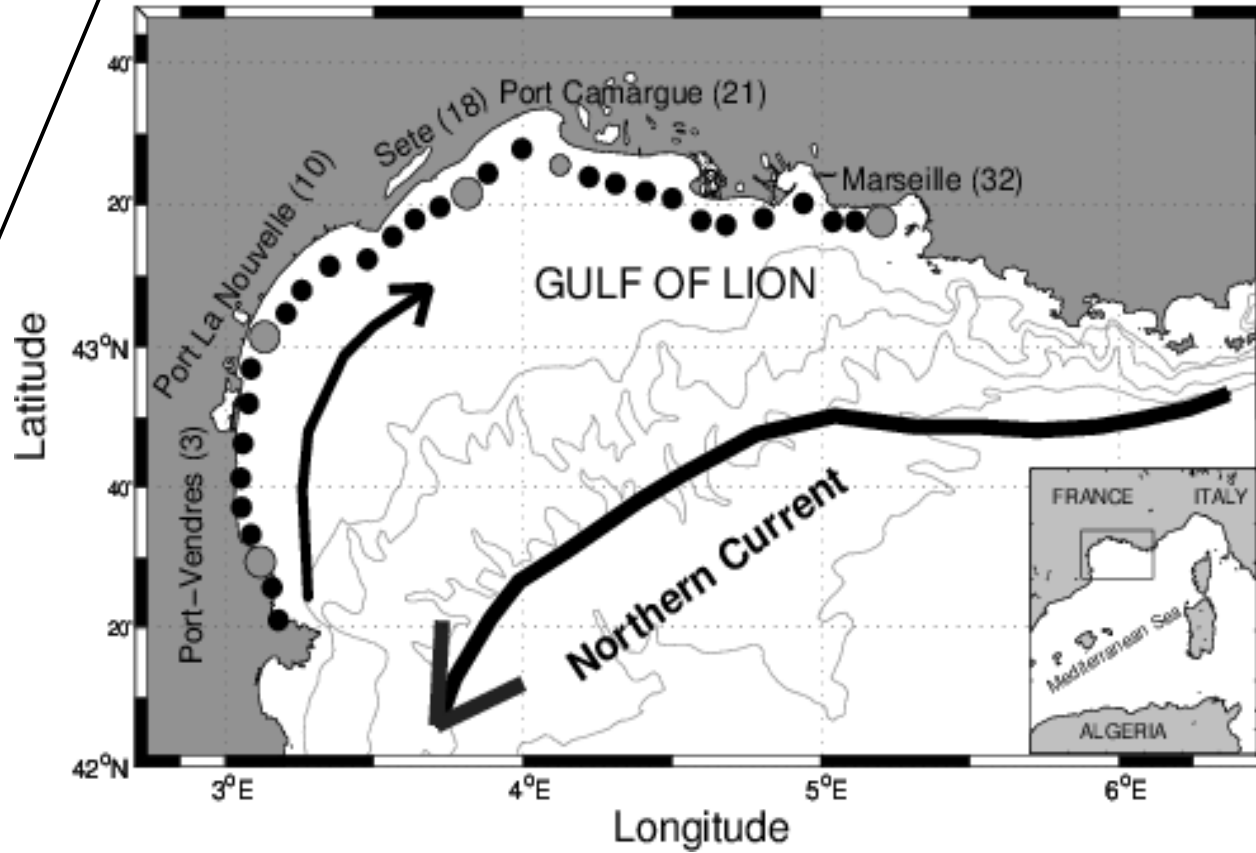


Propagule
production rate

Transfer probability

Recruitment success

Survivorship rate



METAPOPULATION MODEL

(see Guizien et al, 2014)

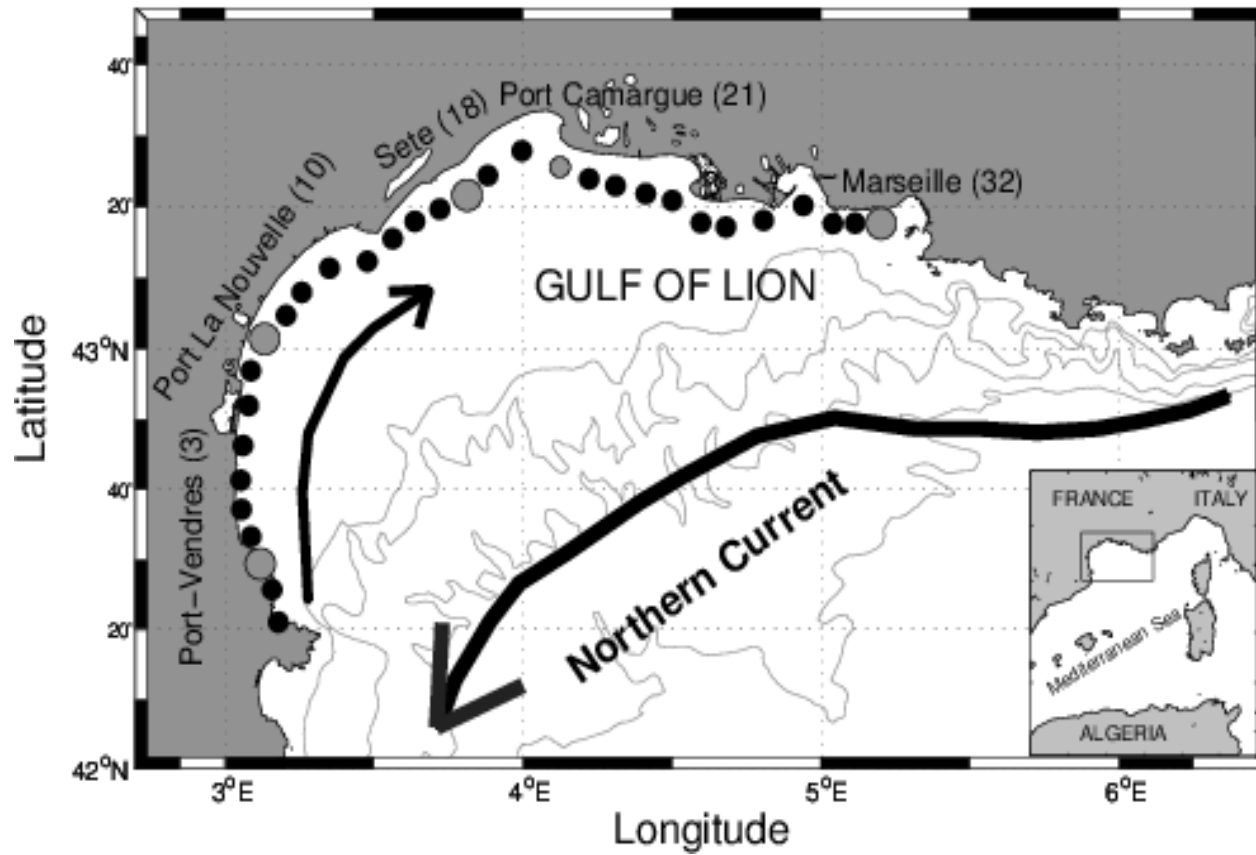
Species: soft-bottom polychaete

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$$N_{t+1} = \min(b_j C_{ij} a_i N_t + s_{jj} \delta_{ji} N_t, N_{max})$$



Four sites analyzed.



METAPOPULATION MODEL

(see Guizien et al, 2014)

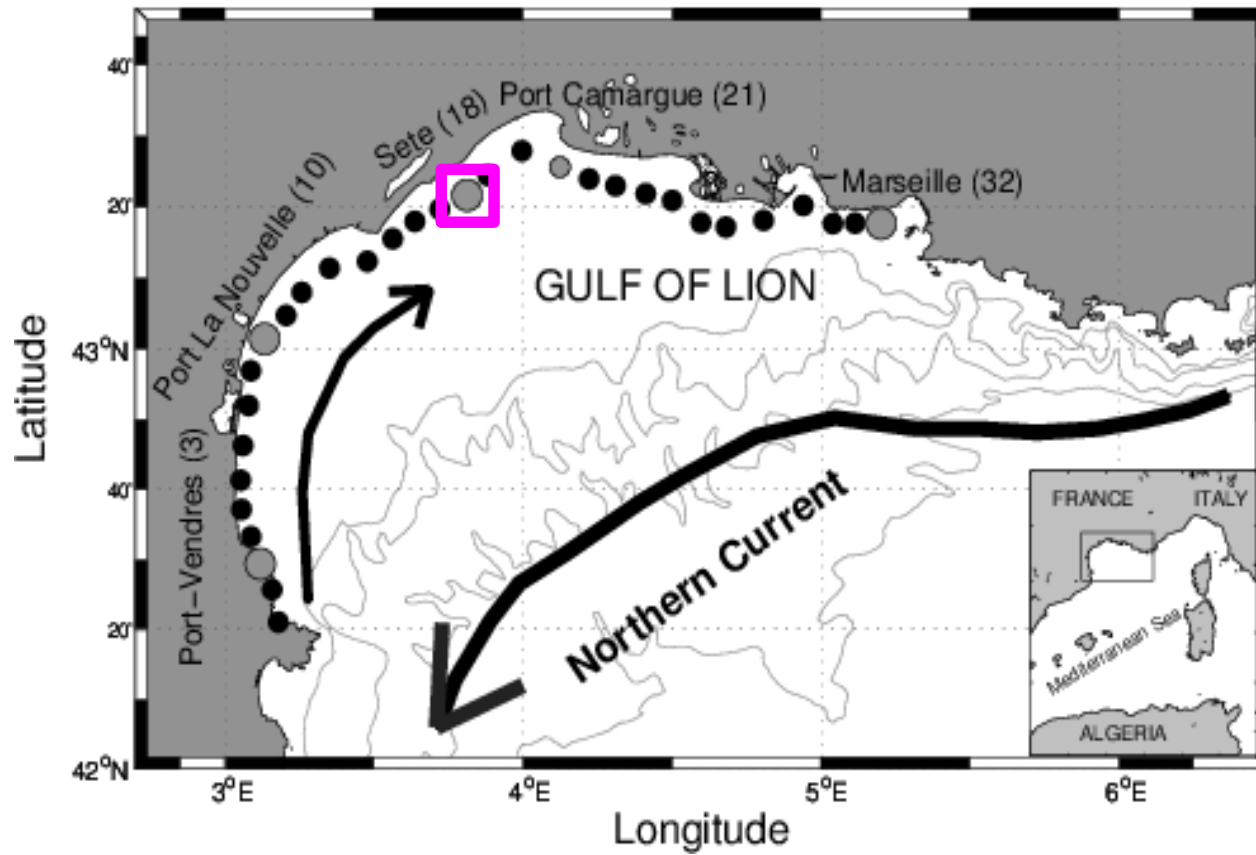
Species: soft-bottom polychaete

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$$N_{t+1} = \min(b_j C_{ij} a_i N_t + s_{jj} \delta_{ji} N_t, N_{max})$$



One essential node for persistence.



METAPOPULATION MODEL

(see Guizien et al, 2014)

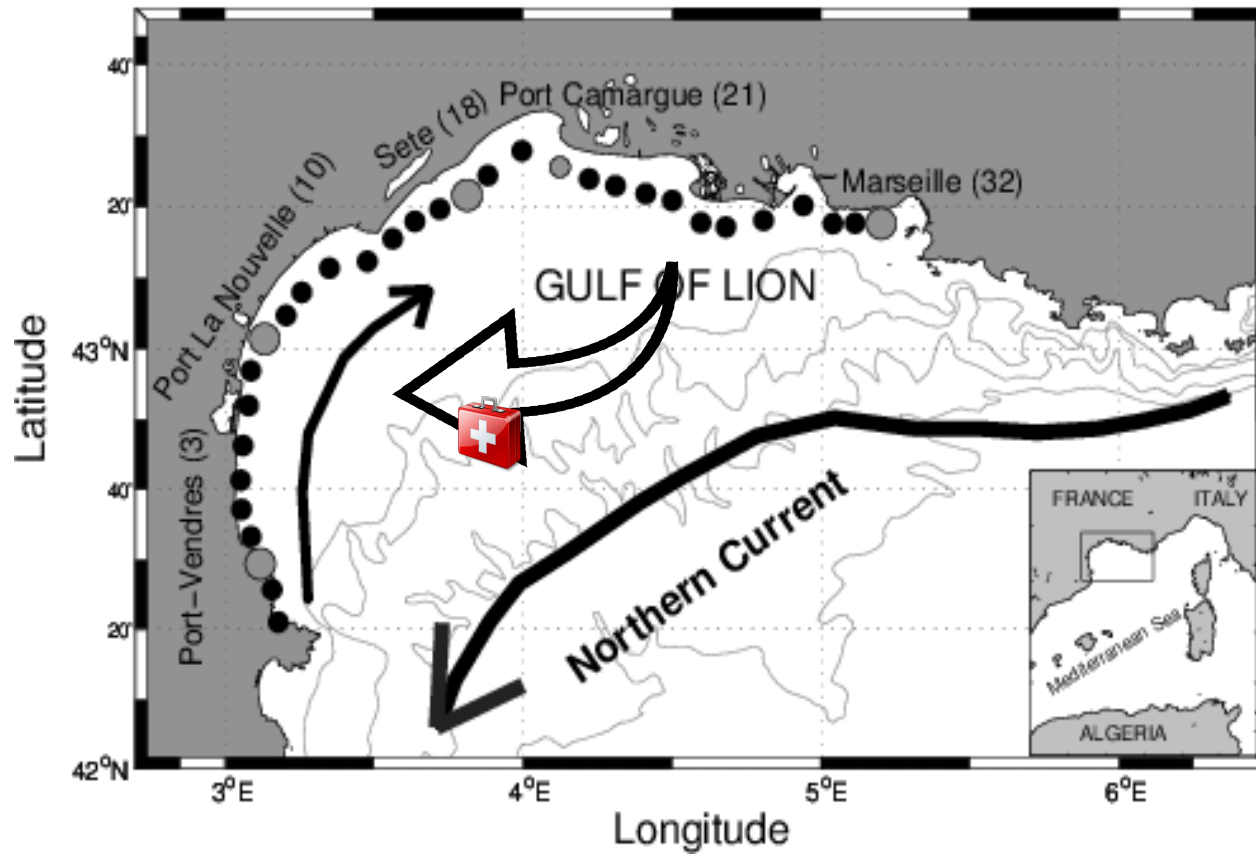
Species: soft-bottom polychaete

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$$N_{t+1} = \min(b_j C_{ij} a_i N_t + s_{jj} \delta_{ji} N_t, N_{max})$$



Presence of a rescue mechanism.



METAPOPULATION MODEL

(see Guizien et al, 2014)

Species: soft-bottom polychaete

Site: Gulf of Lion

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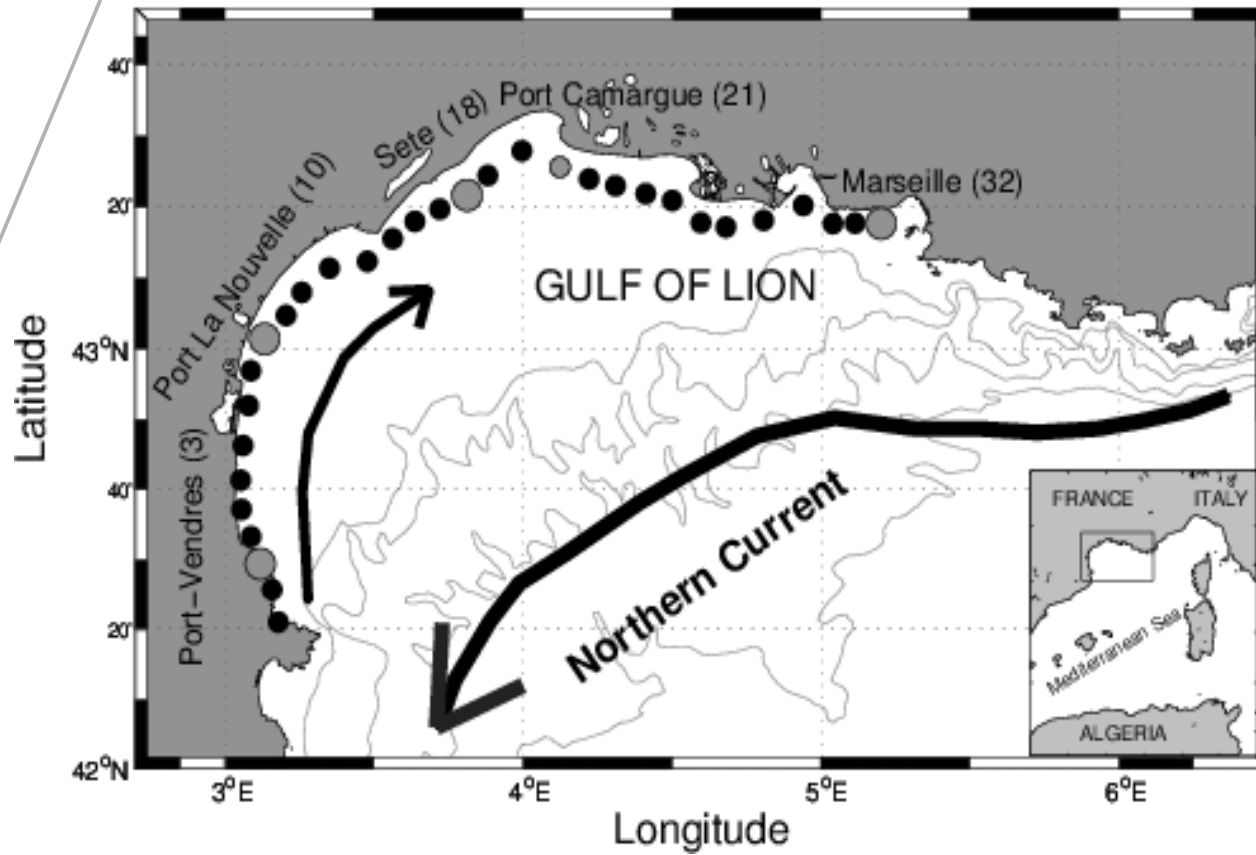


Propagule
production rate

Transfer probability

Recruitment success

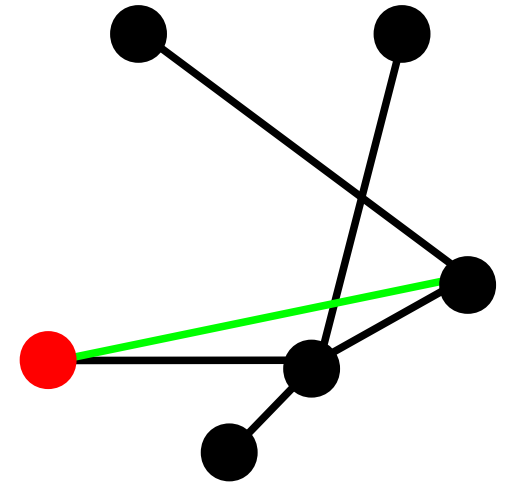
Survivorship rate



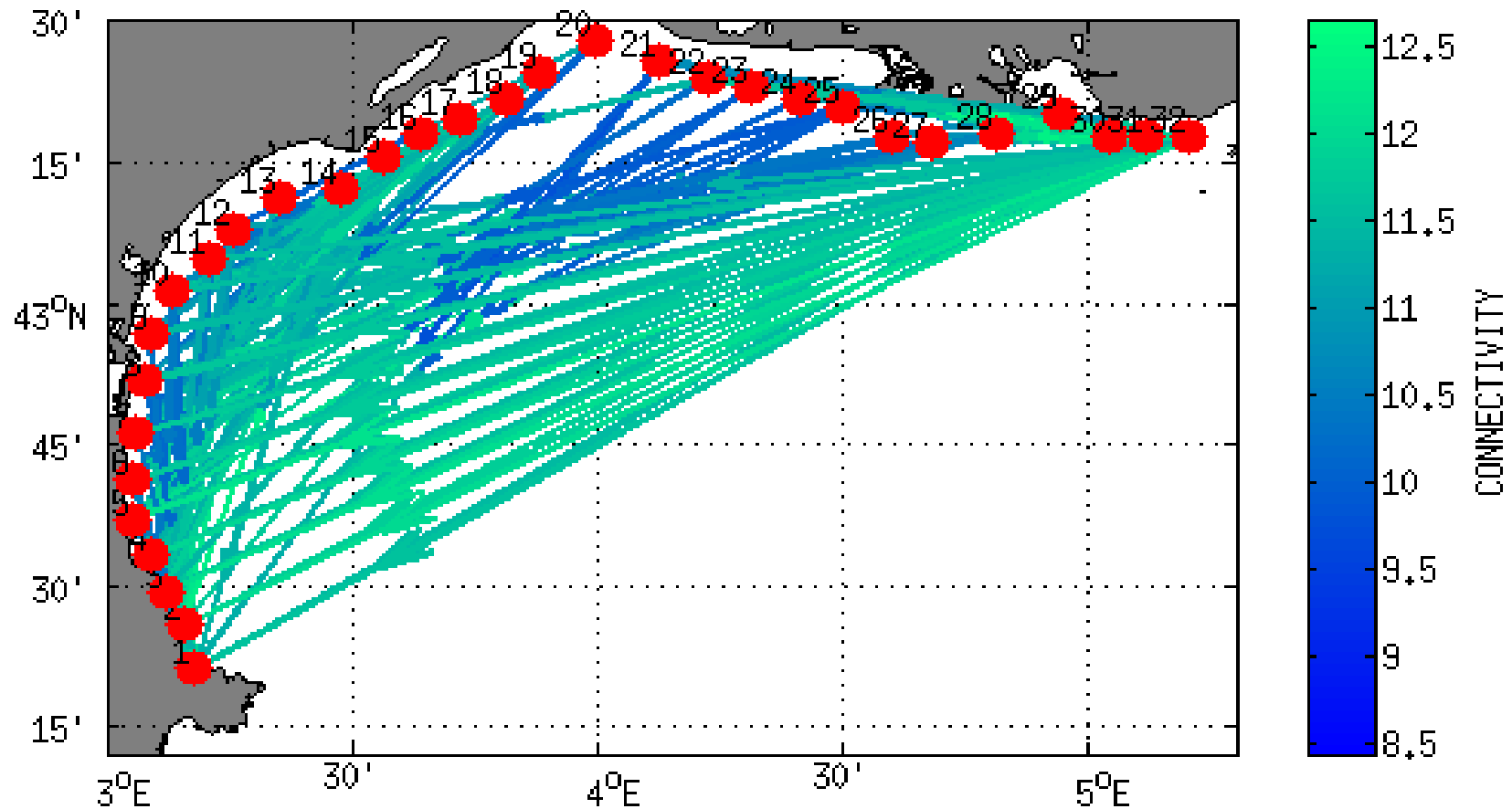
GRAPH THEORY

Graph: representation of pairwise relations between nodes (**sites**).

Relations are represented by links (C_{ij}).



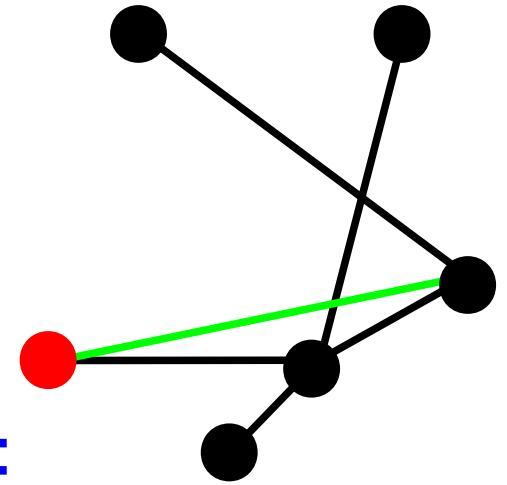
WHAT IT LOOKS LIKE



GRAPH THEORY

Graph: representation of pairwise relations between nodes (**sites**).

Relations are represented by links (C_{ij}).



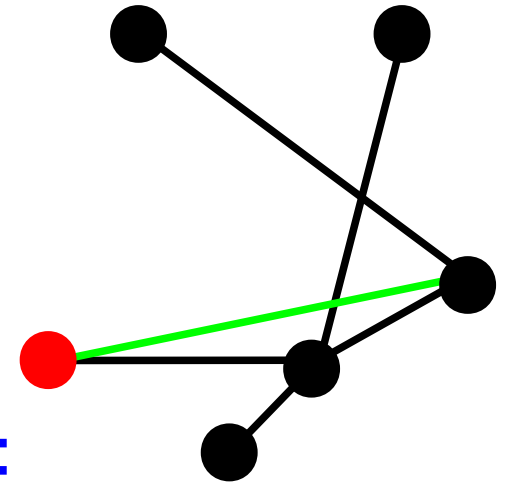
New distance between nodes:

$$d_{i,j} = \log\left(\frac{1}{C_{i,j}}\right)$$

GRAPH THEORY

Graph: representation of pairwise relations between nodes (**sites**).

Relations are represented by links (C_{ij}).



New distance between nodes:

$$d_{i,j} = \log\left(\frac{1}{C_{i,j}}\right)$$

Shortest path
=
most probable

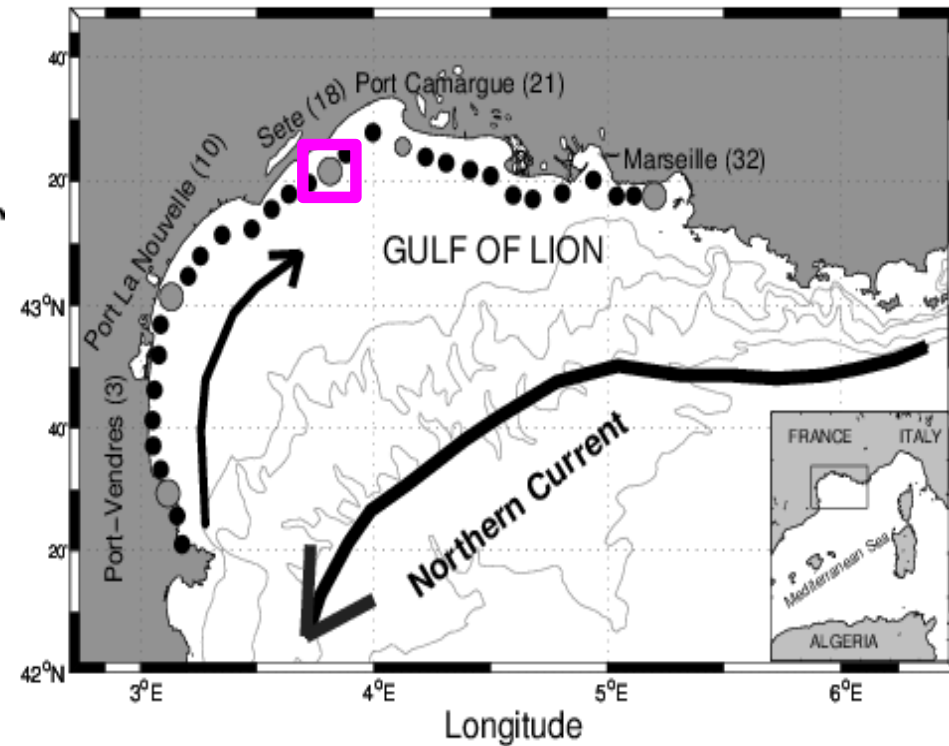
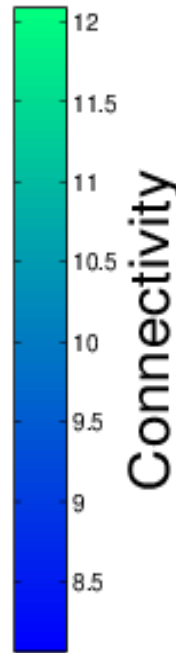
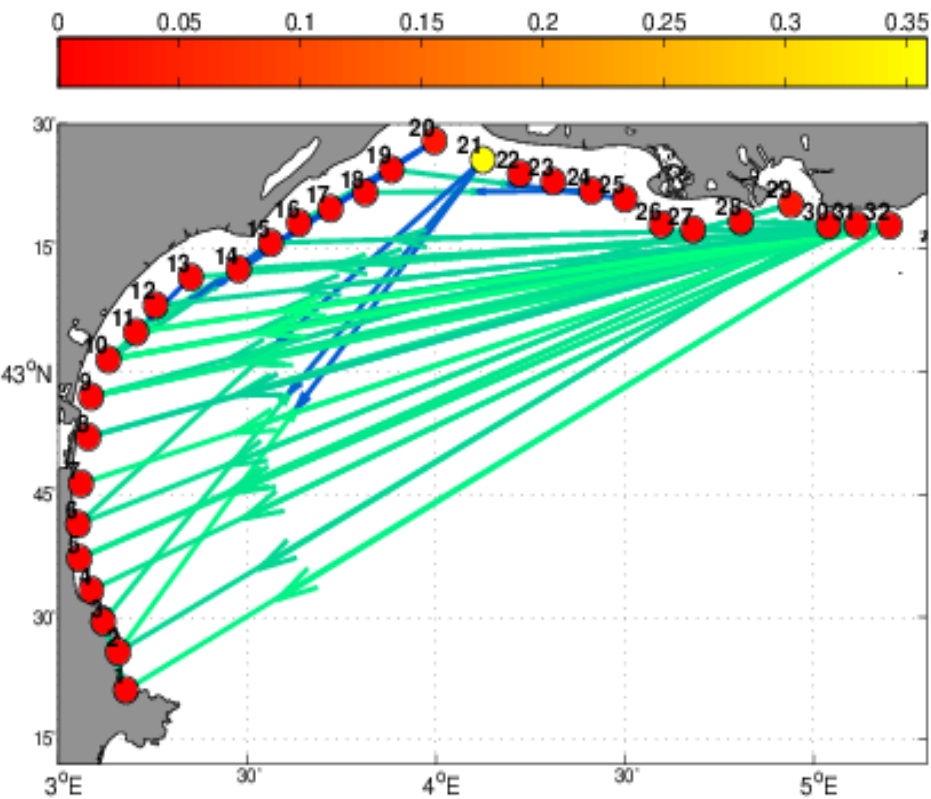
Betweenness: the number of shortest paths that pass through a node

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Gateways → Persistence

Betweenness: the number of shortest paths that pass through a node

Betweenness



Gateways ~~X~~ Persistence

Thus we introduce a new measure.

BRIDGING CENTRALITY

Thus we introduce a new measure.

BRIDGING CENTRALITY

Nodes with:

**High basin-scale
flux of propagule**

**Lying the border
of clusters**

Thus we introduce a new measure.

BRIDGING CENTRALITY

Nodes with:

**High basin-scale
flux of propagule**



High betweenness

**Lying the border
of clusters**



**Topological
factor**

Thus we introduce a new measure.

BRIDGING CENTRALITY

Nodes with:

**High basin-scale
flux of propagule**

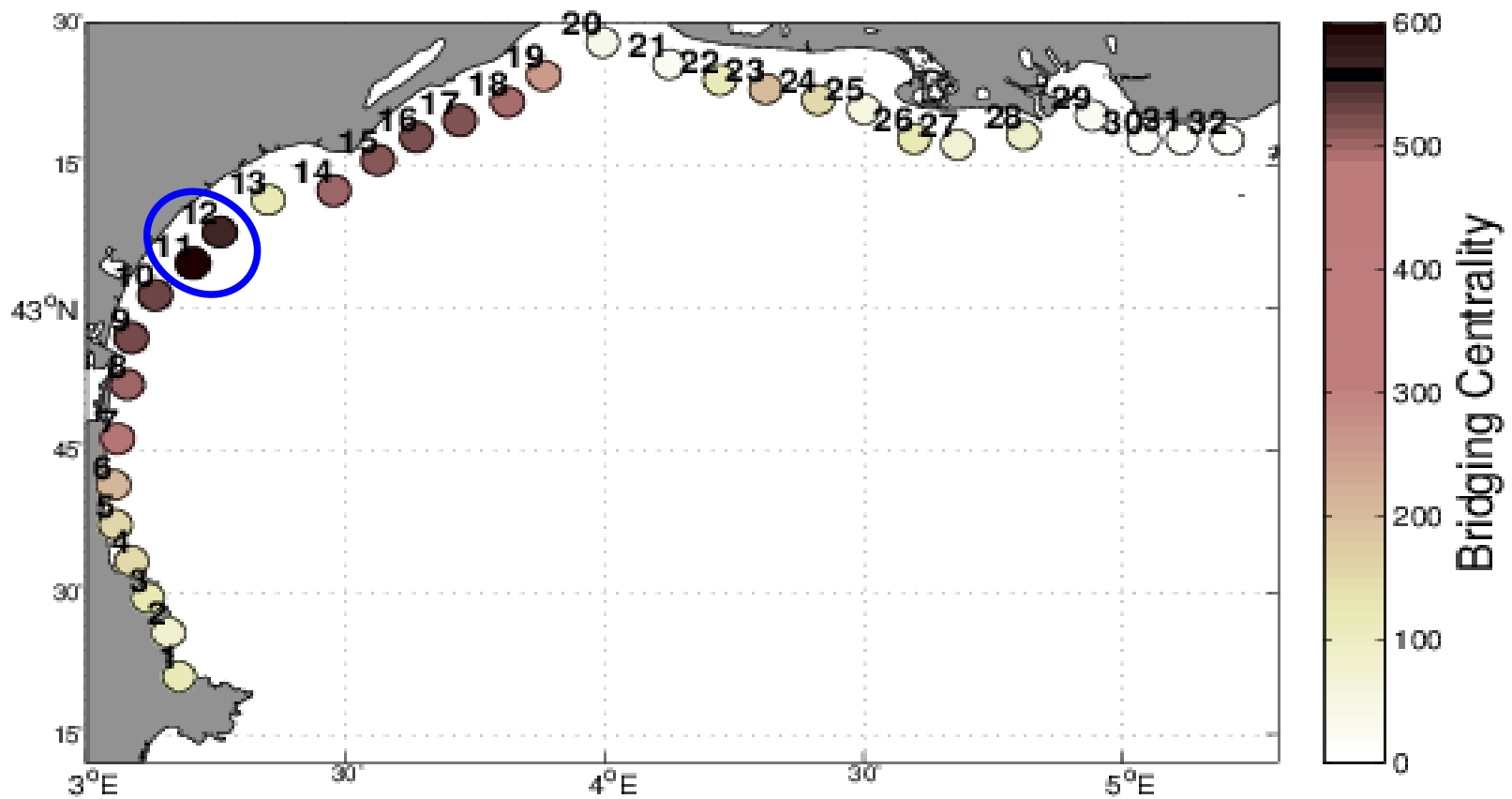


High betweenness

**Lying the border
of clusters**



**Topological
factor**



**High-bridging centrality
nodes ensure
the network's integrity**

Thus we look for clusters.

Thus **we look for clusters.**

Cluster of nodes = Community

Cluster = Statistically surprising
disposition of links

Thus **we look for clusters.**

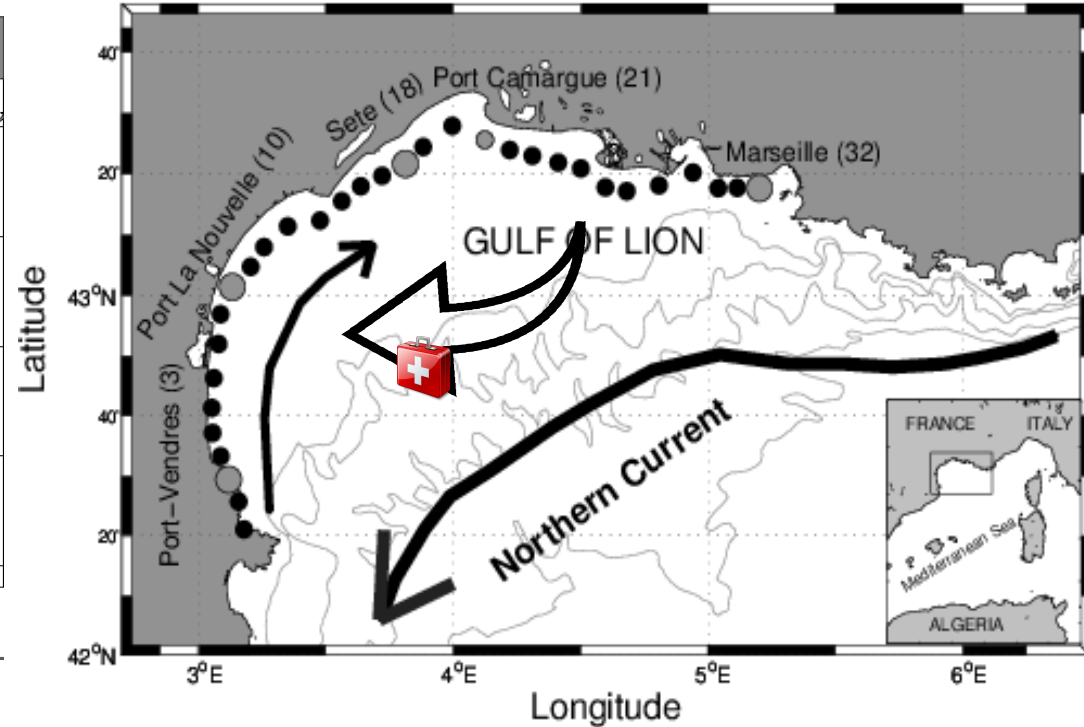
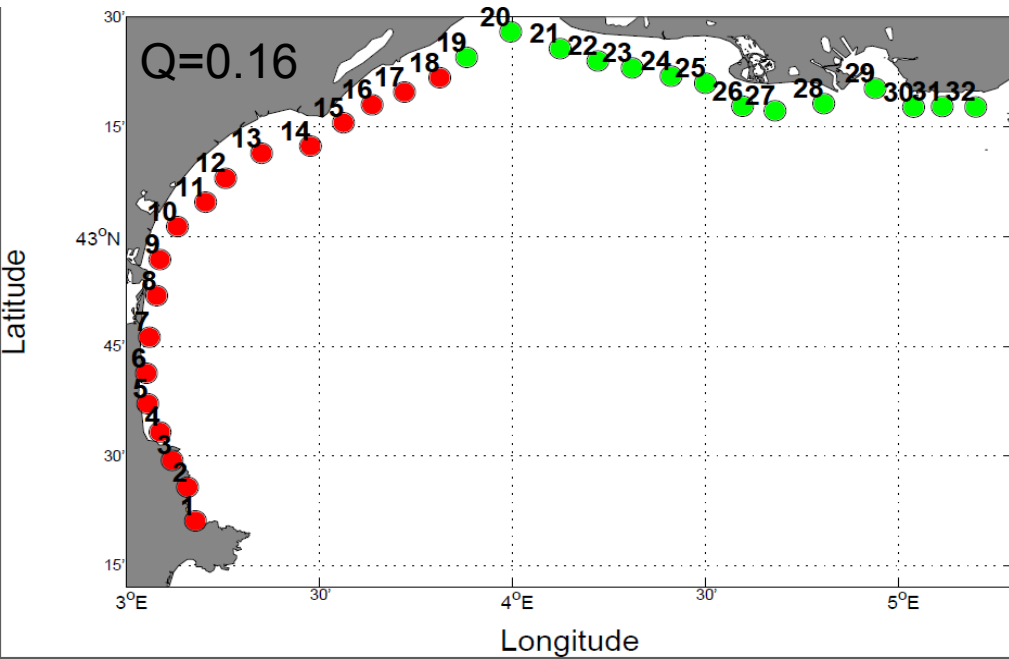
Cluster of nodes = Community

Cluster → Maximization of
modularity Q

Thus we look for clusters.

Cluster of nodes = Community

Cluster → Maximization of modularity Q



**Graph theory
detects
rescue effects.**

CONCLUSIONS

- 1)** We propose a new consistent node-to-node metric.
- 2)** Betweenness is not correlated with persistence.
- 3)** Bridging centrality is a good indicator for persistence.
- 4)** Modularity can identify communities and rescue mechanisms.
- 5)** Methodology independent from demographic parameters.

THANK YOU

FOR YOUR

KIND ATTENTION

BIBLIOGRAPHY

Graph theory:

Costa A., Guizien K., Doglioli A.M., Petrenko A.A., *submitted to Limnology and Oceanography*

Metapopulation model:

Guizien K., Belharet M., Moritz C., Guarini J., *Diversity and Distributions*, 2014.

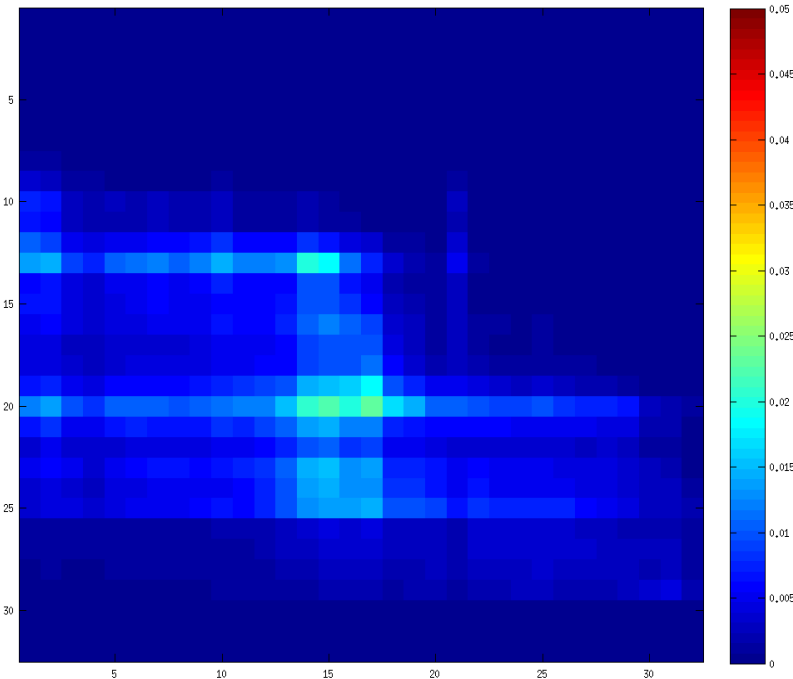
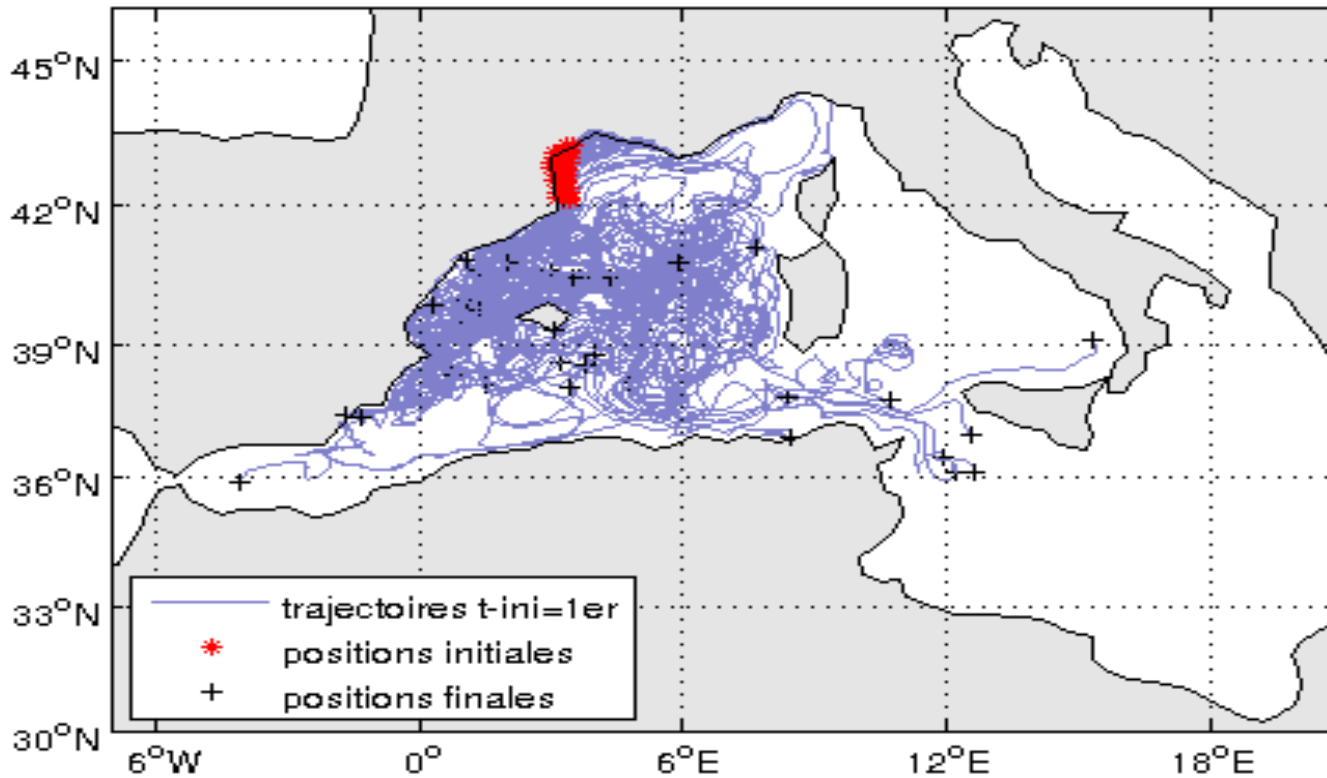
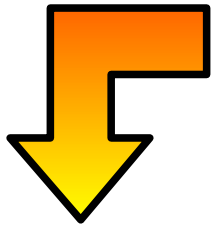
Modularity:

Newman M., Girwan M., *Physical Review E* 69, 2004.

Bridging centrality:

Hwang, W., Taehyong, K., Murali, R., Aidong, Z., *Proceedings KDD '08*, 2008.

EXTRA SLIDES



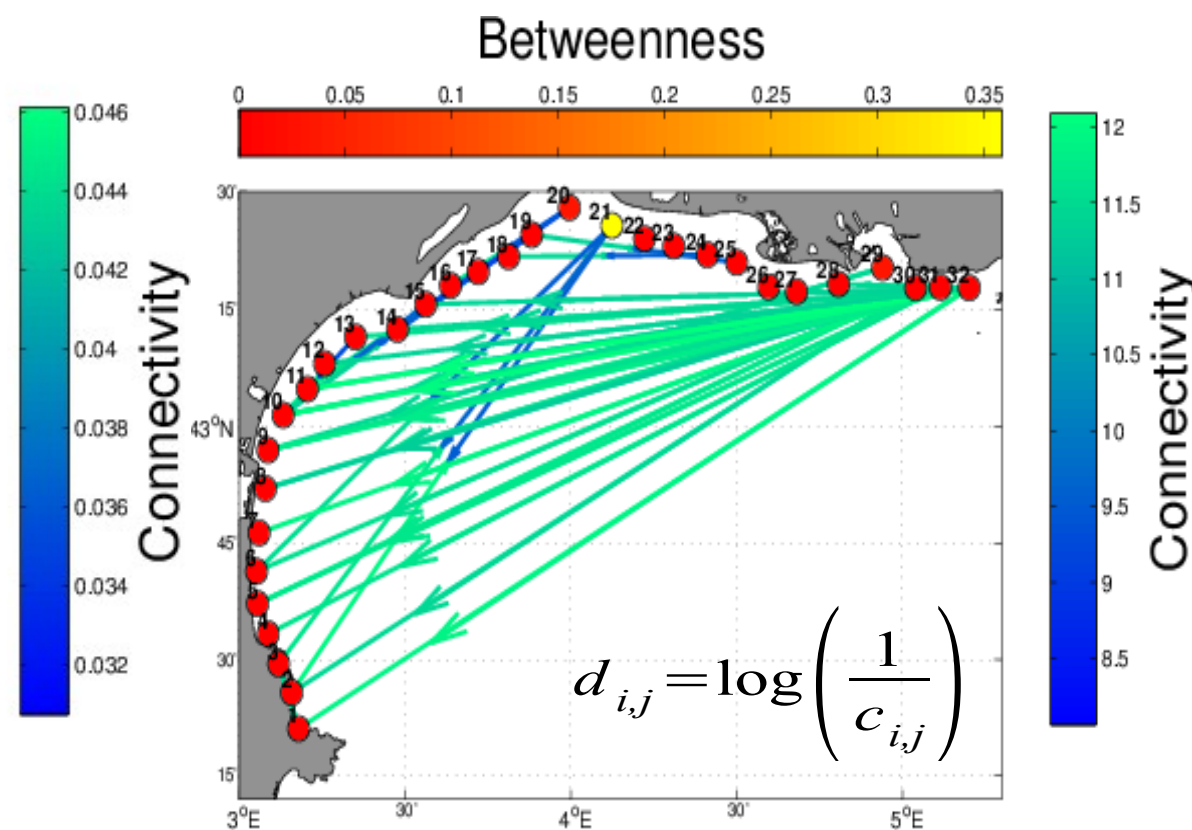
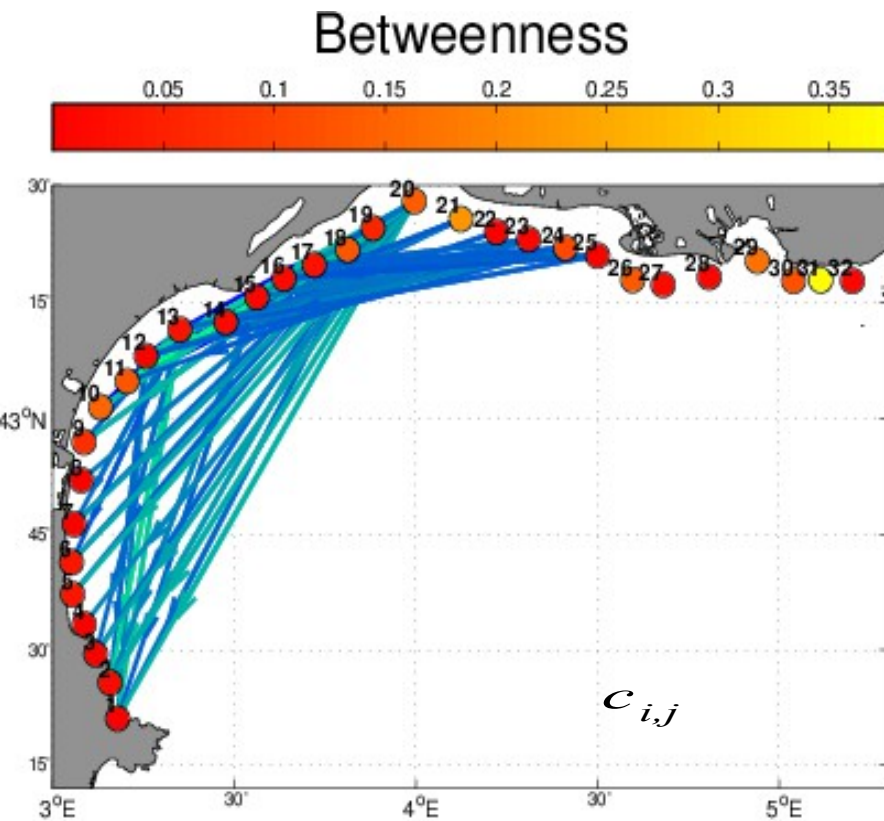
Graph Theory

enables us to
derive more information about
crucial points of the system.

Graph theory for **key site selection**:

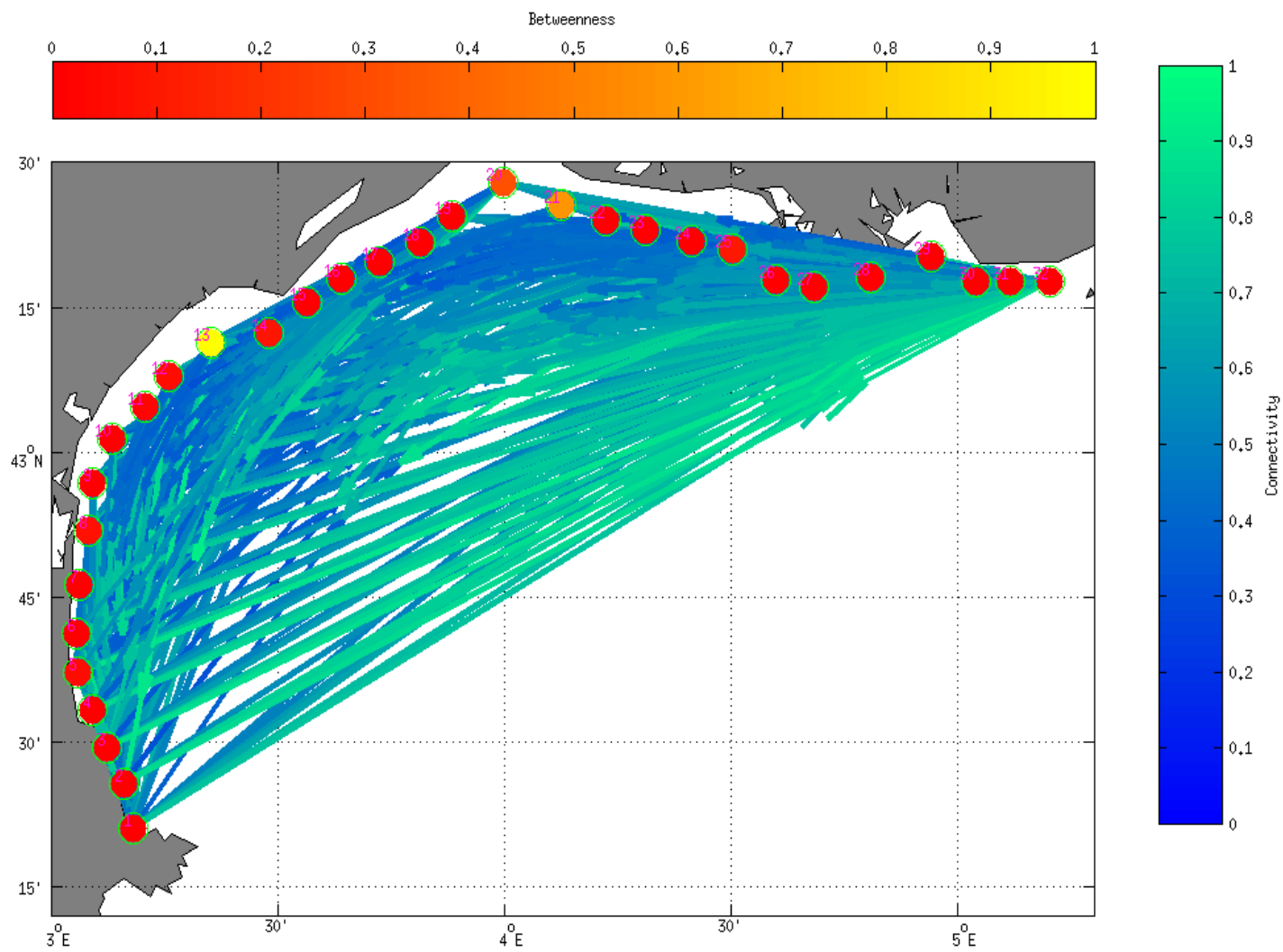
pay great attention on the metrics !!!

Betweenness: identifies gateways (i.e. nodes through which most of the propagules flow, e.g. soft-bottom polychaete in the Gulf of Lion)

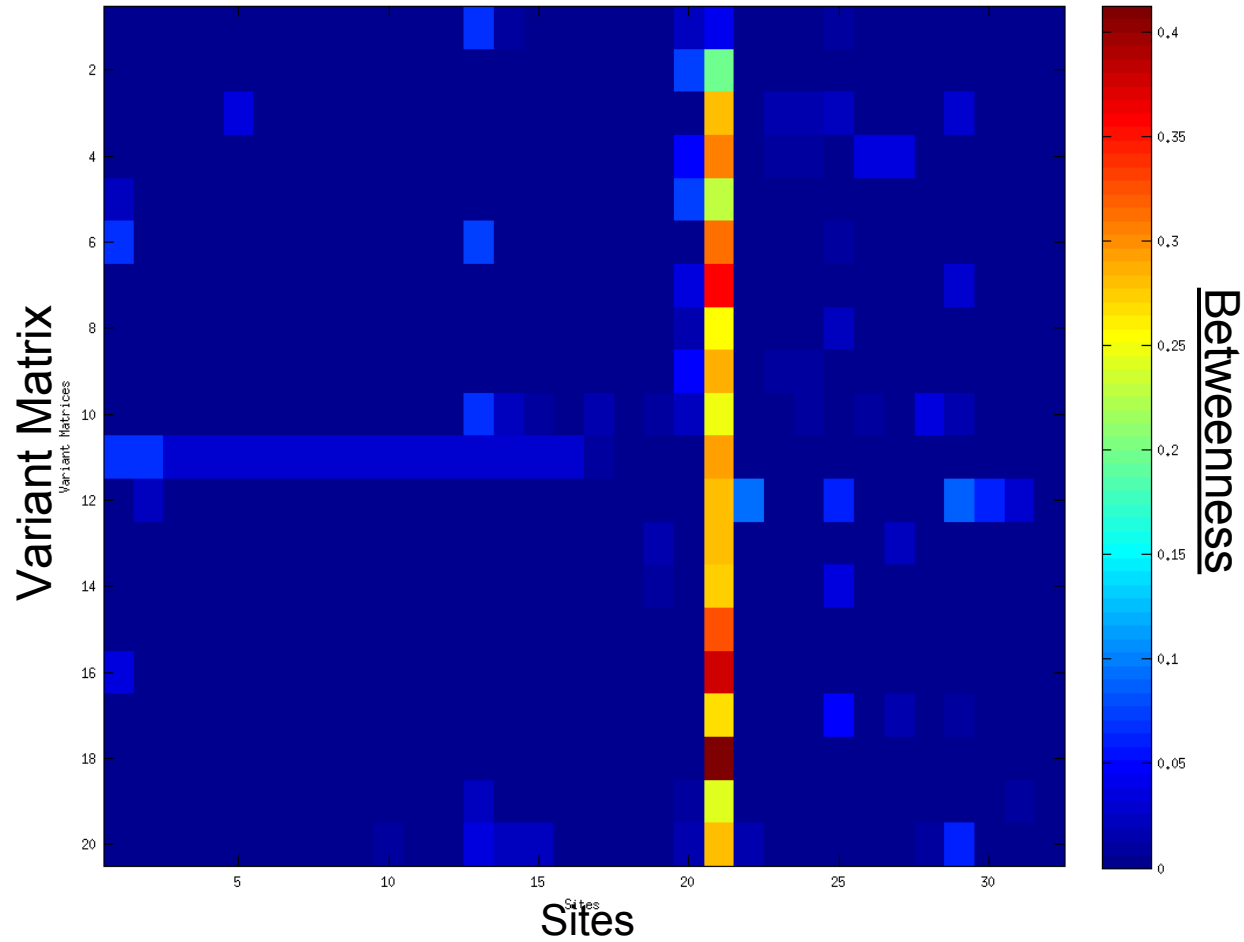


Andrello et al., Low connectivity between Mediterranean Marine Protected Areas: a biophysical modeling approach for the dusky grouper: *Epinephelus Marginatus*, *PlosOne*, 2013

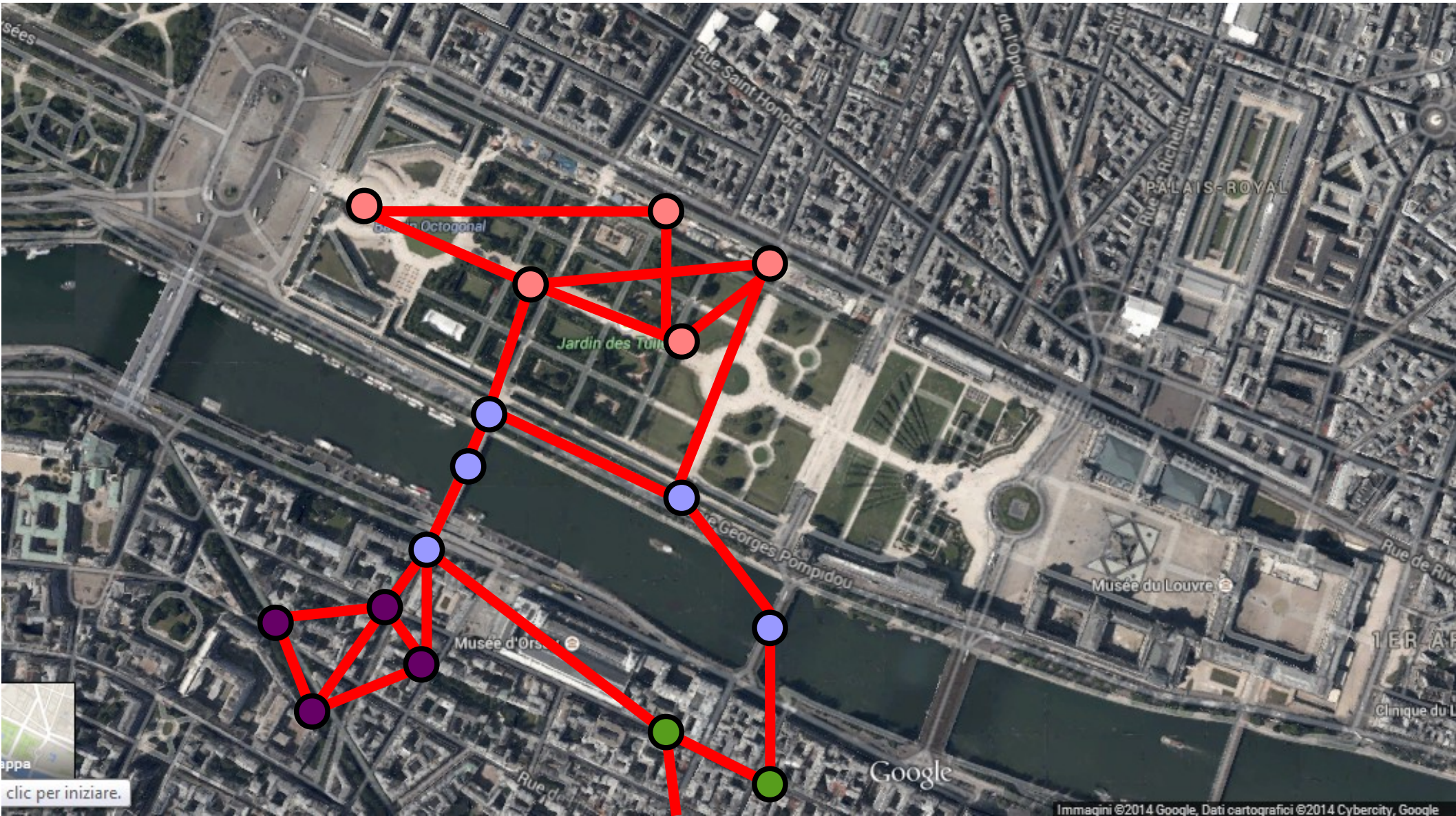
Costa et al., Estimation of connectivity in marine biological networks: Graph Theory versus Metapopulation Models. The Gulf of Lion study case, in preparation



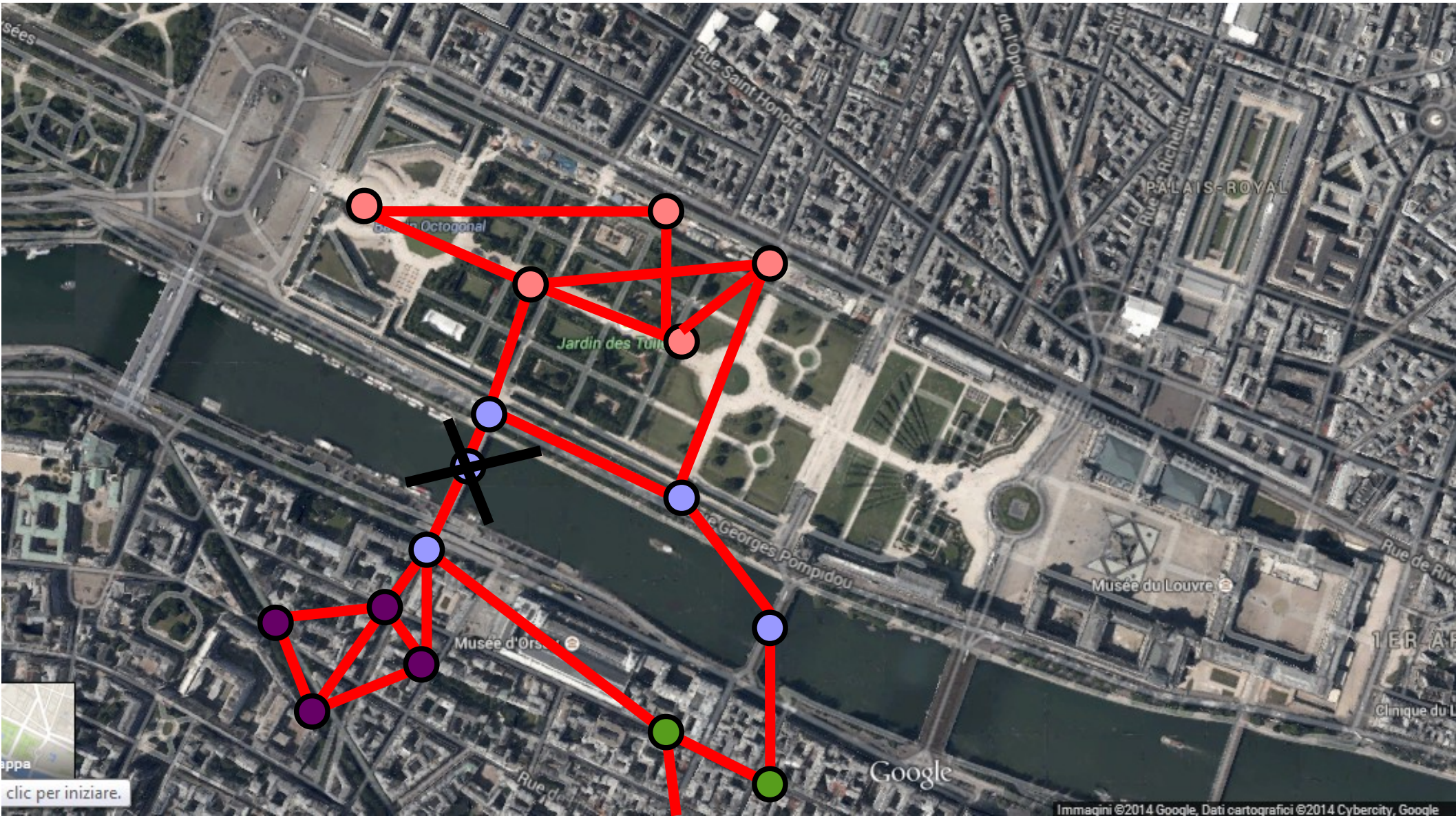
First measure: **betweenness**

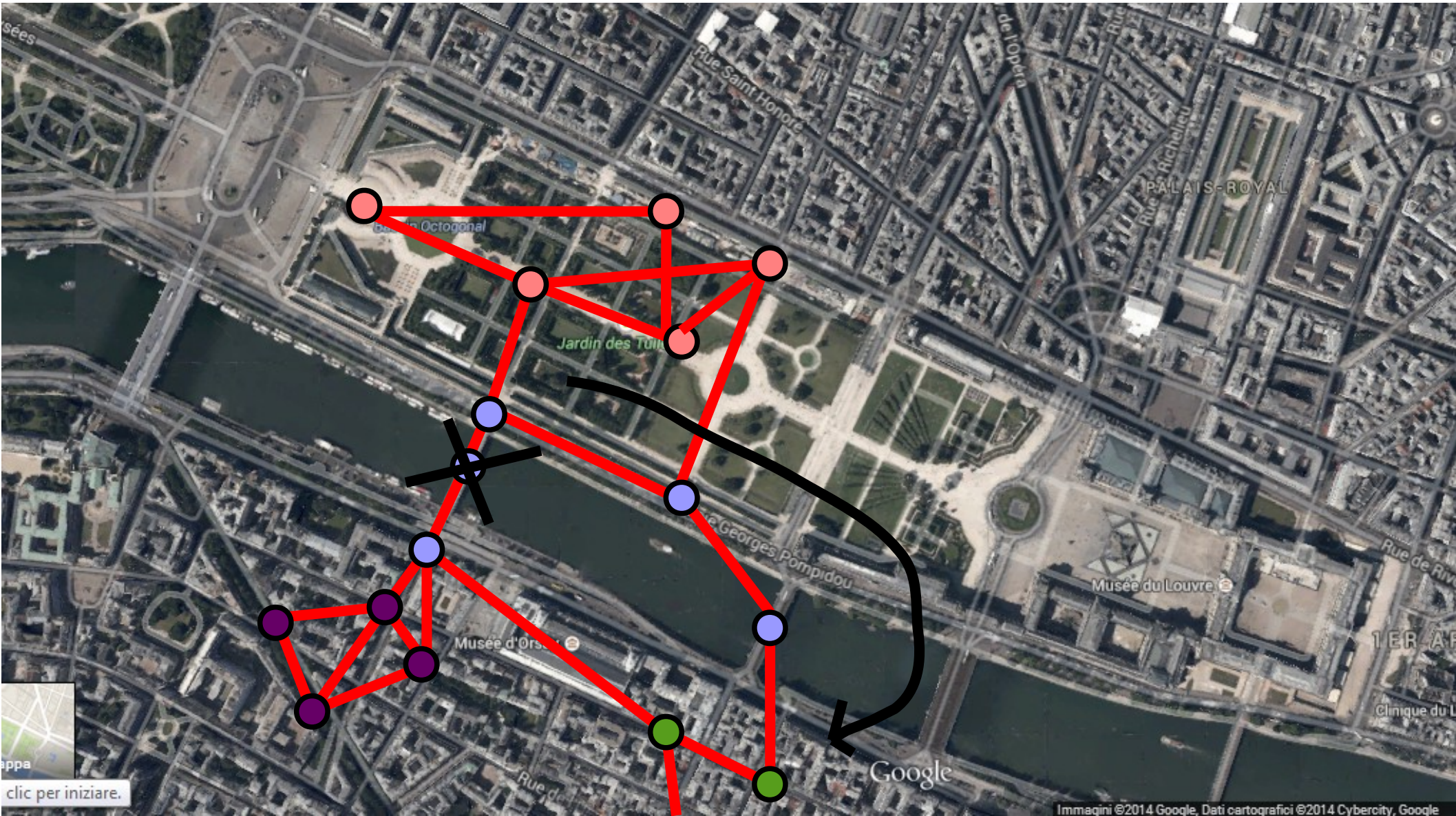


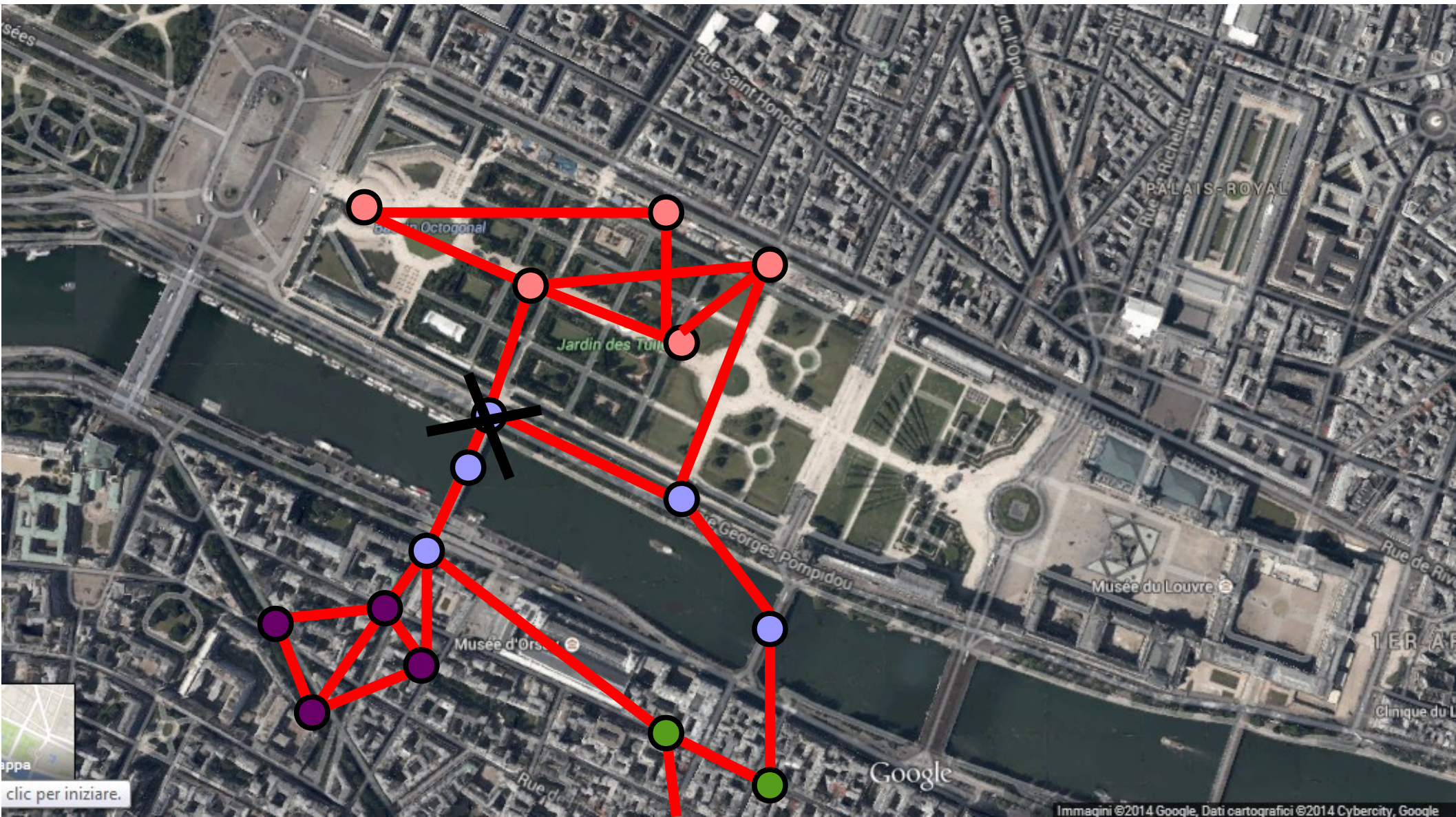
There is **no correspondence between high betweenness and important sites for persistence.**

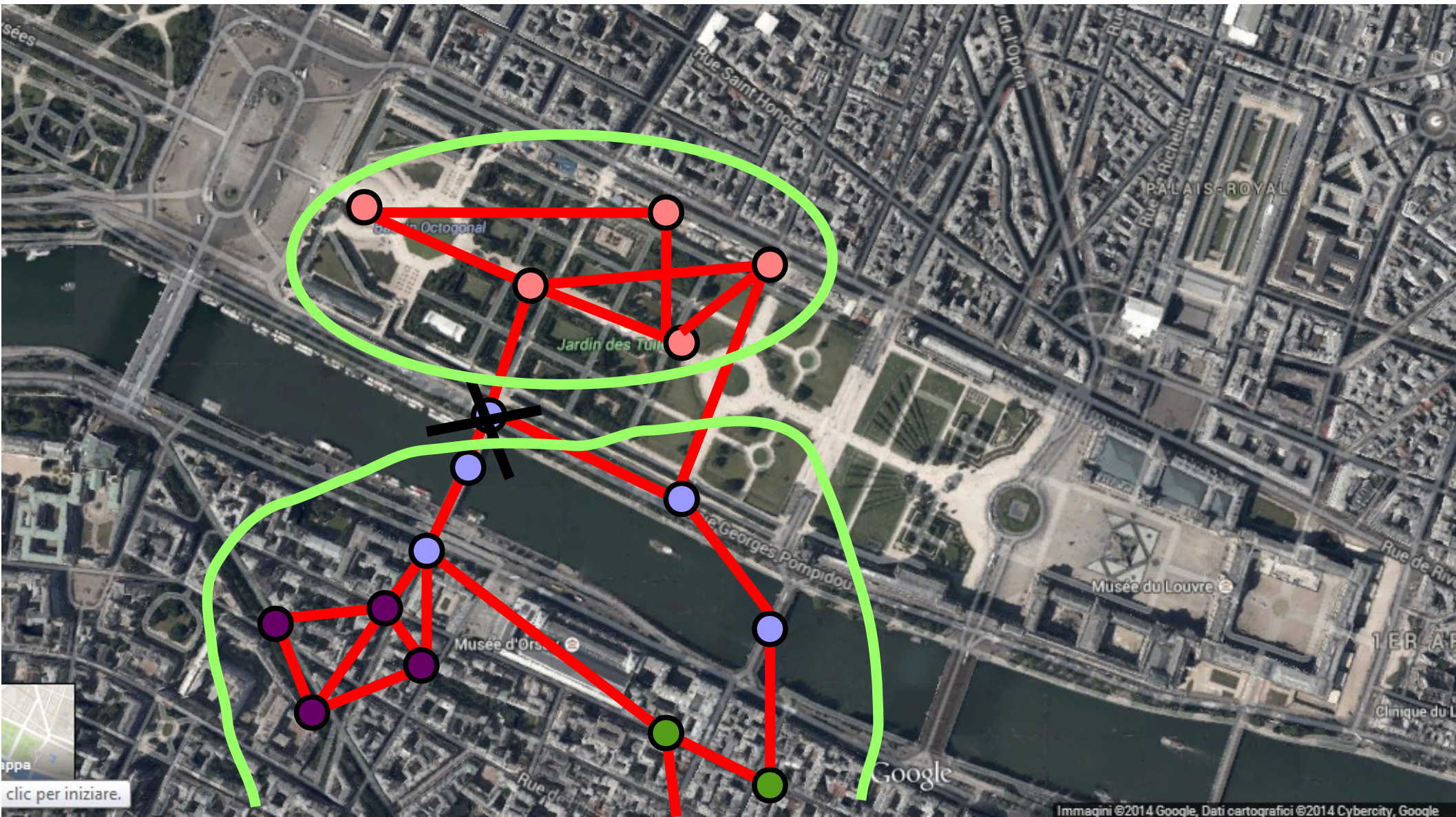


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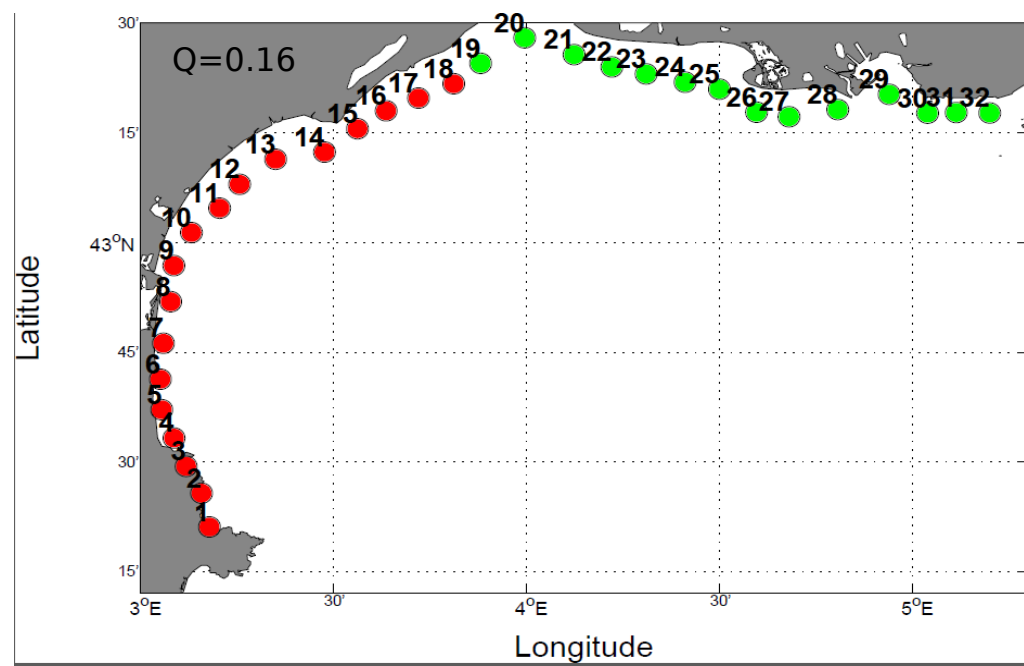






More isolated components!!!

Q is positive

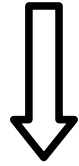


Congruent

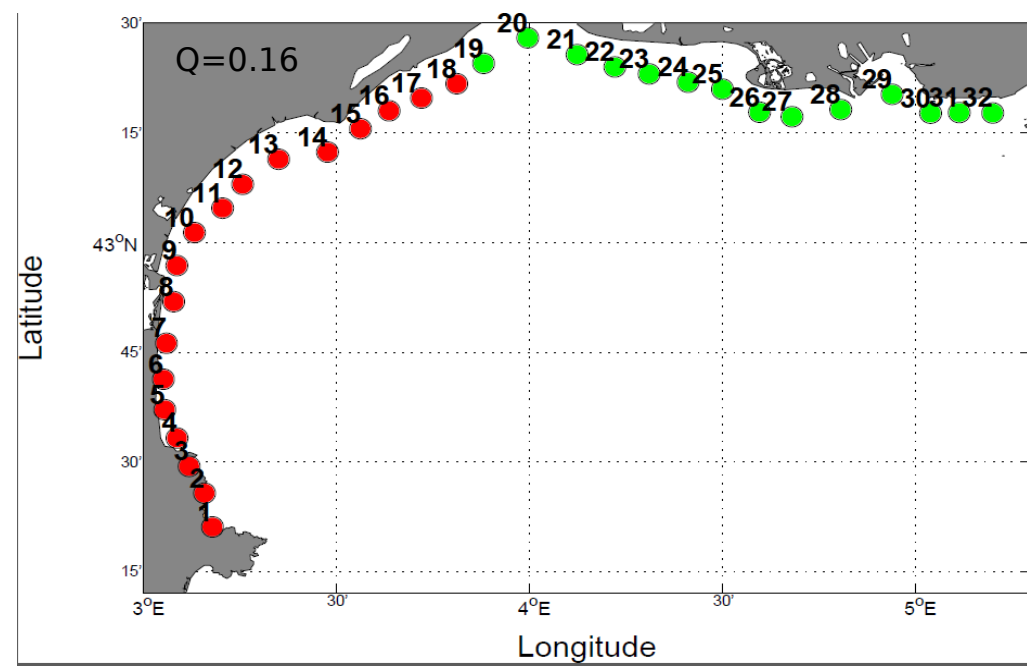
with what emerges from
the metapopulation model analysis.

Second measure: modularity

Q is positive



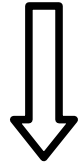
There are distinct communities



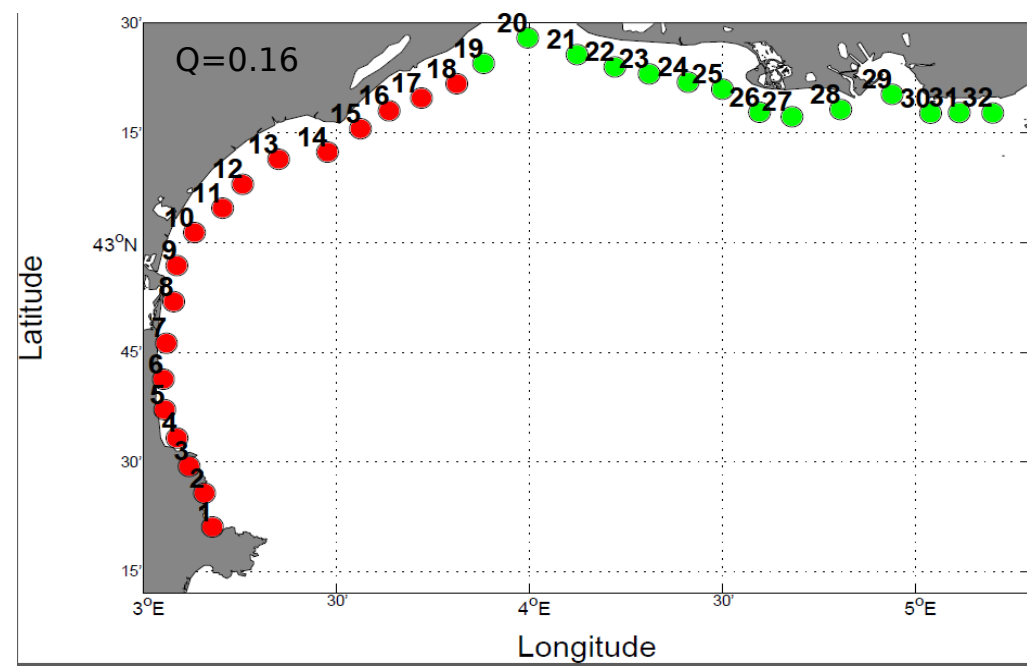
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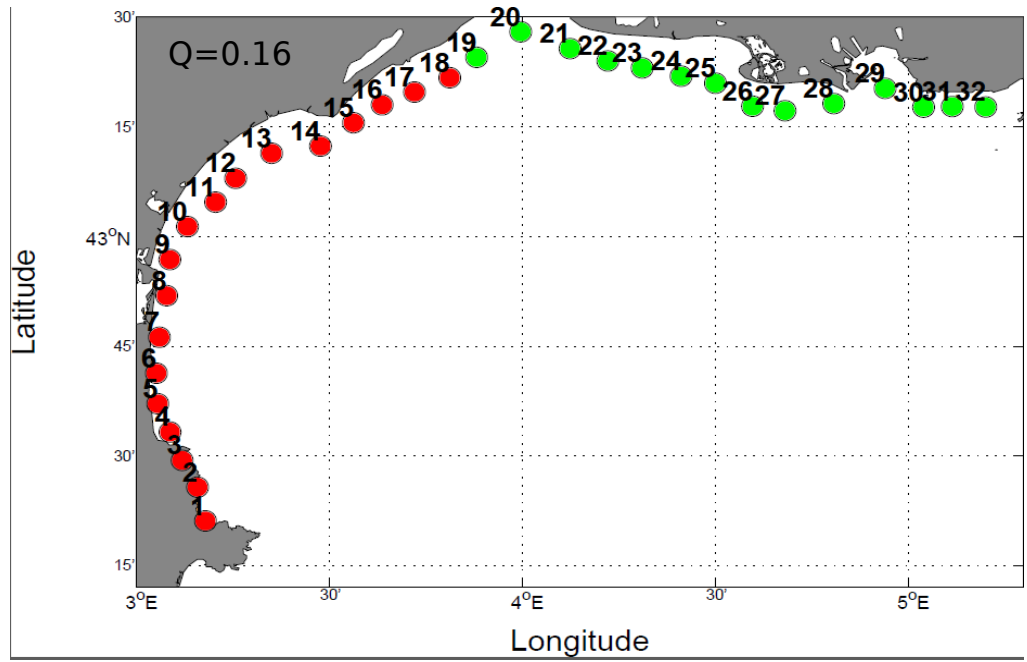
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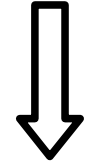
But what about the rescue effect?

Q is positive **but low**

Second measure: modularity

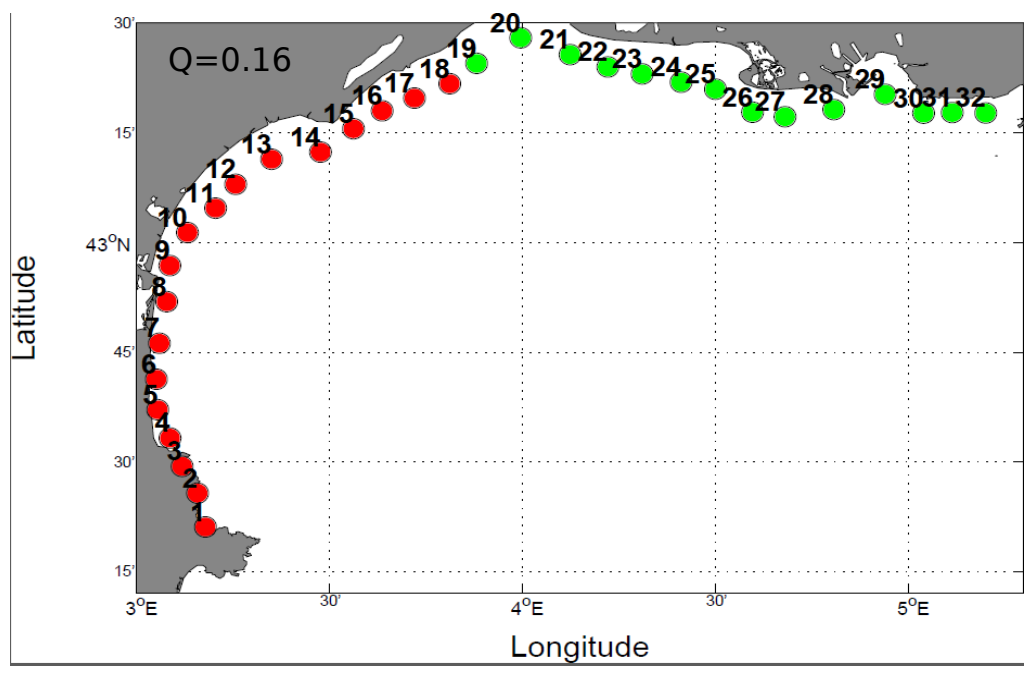


Q is positive **but low**

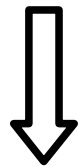


high exchange

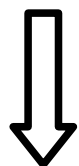
Second measure: modularity



Q is positive **but low**

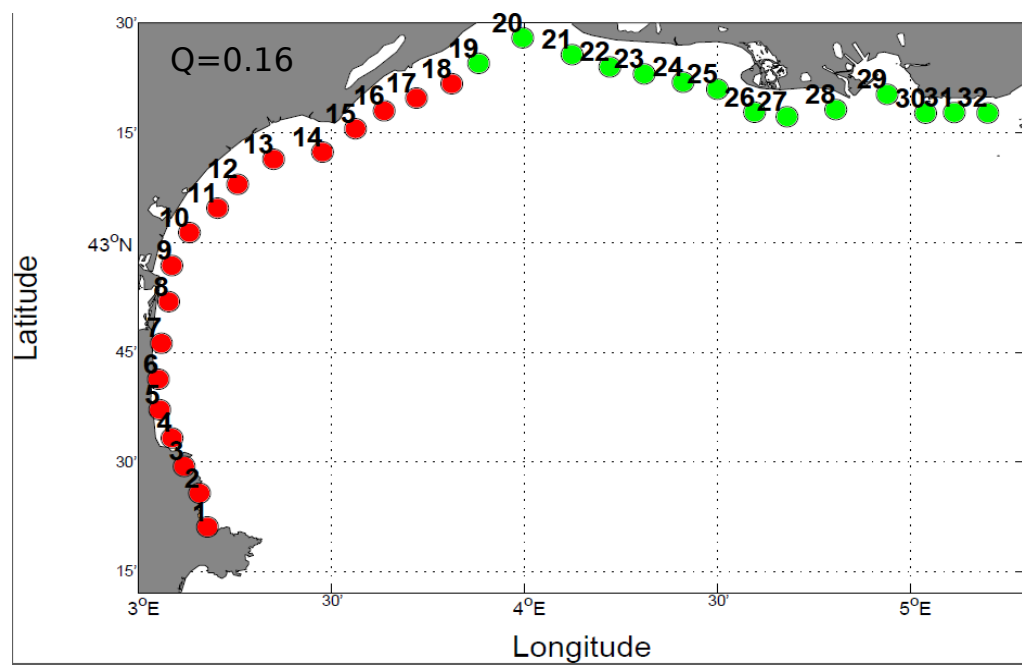


high exchange

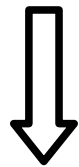


rescue effect is possible

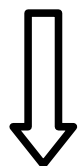
Second measure: modularity



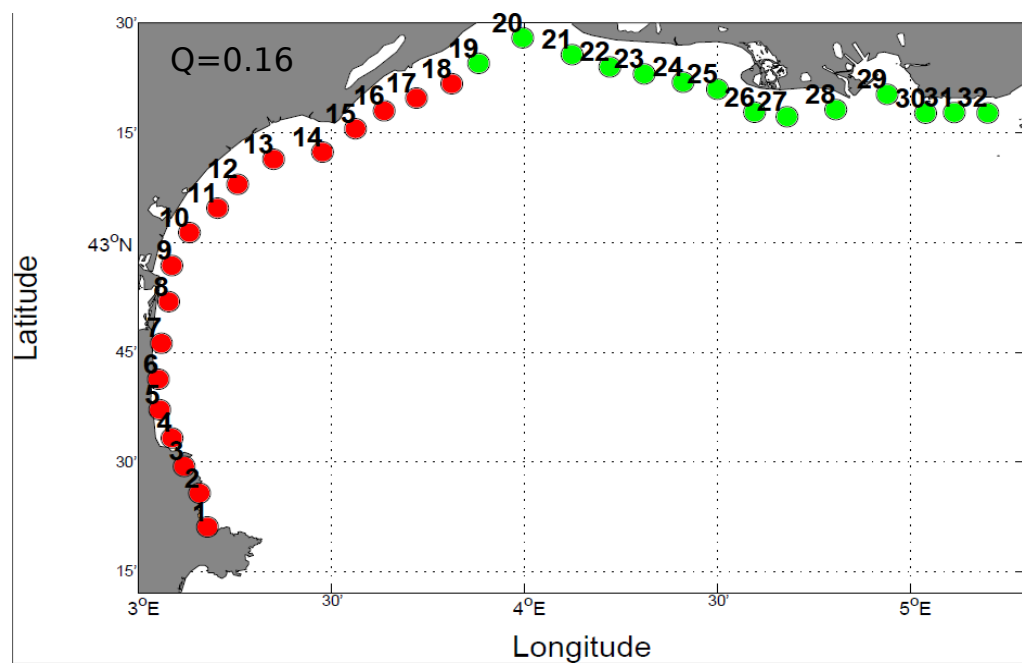
Q is positive **but low**



high exchange



rescue effect is possible



Second measure: modularity

Graph analysis reflects well

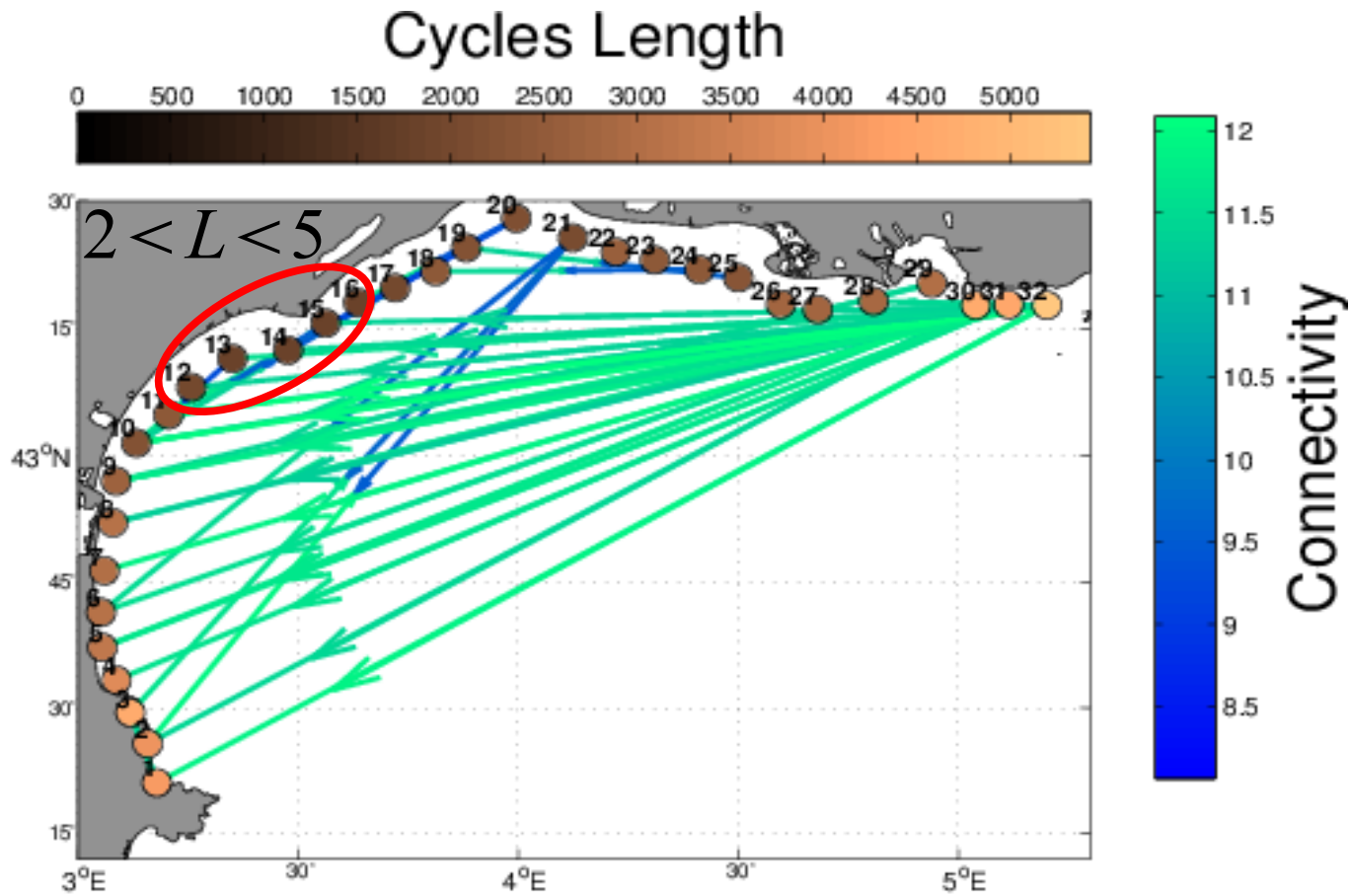
the organization of polychaete population in the gulf
resulting from Guizien et al.(2014)

We search autosufficient sites
using
a simplified metapopulation model
without density dependent factors:

$$N_{t+1} = (b_j C_{ij} a_i) \cdot N_t$$

Hasting (2006) demonstrates that
self sufficient sites have **cycles** starting from it.

The more the cycle is probable,
the more the site is self-sufficient:
it is a source of *larvae*.



Nodes **13-16** composes the majority of the cycles.

Core of well connected nodes in the centre of the GoL.

In the meatapop. model network collapses when nodes around Sète (node 18) are removed.



Clustering coefficient:

$$C(v_i) = E / (V * (V - 1) / 2)$$

V = # of neighbors

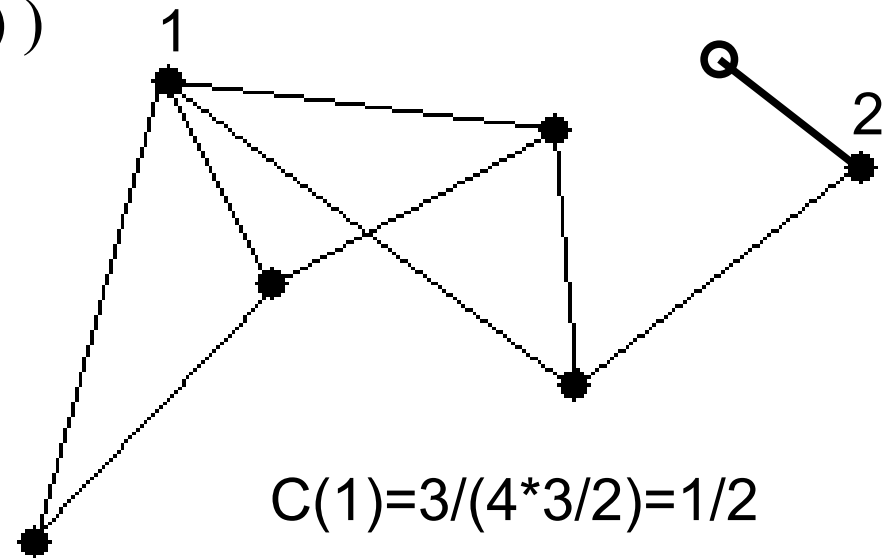
E = # of connections between V

cluster: a set of nodes that are “close” to each other, with respect to a certain measure of distance or similarity

A possible measure is the number of neighbors of a node that are also neighbors between themselves (as Facebook ;-)

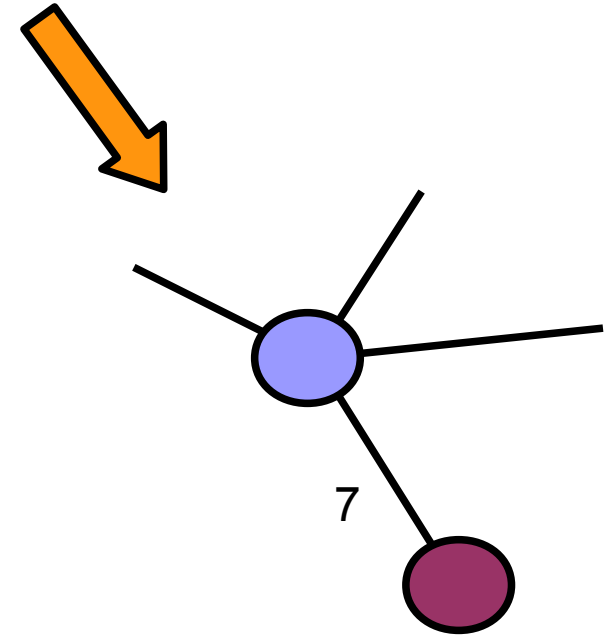
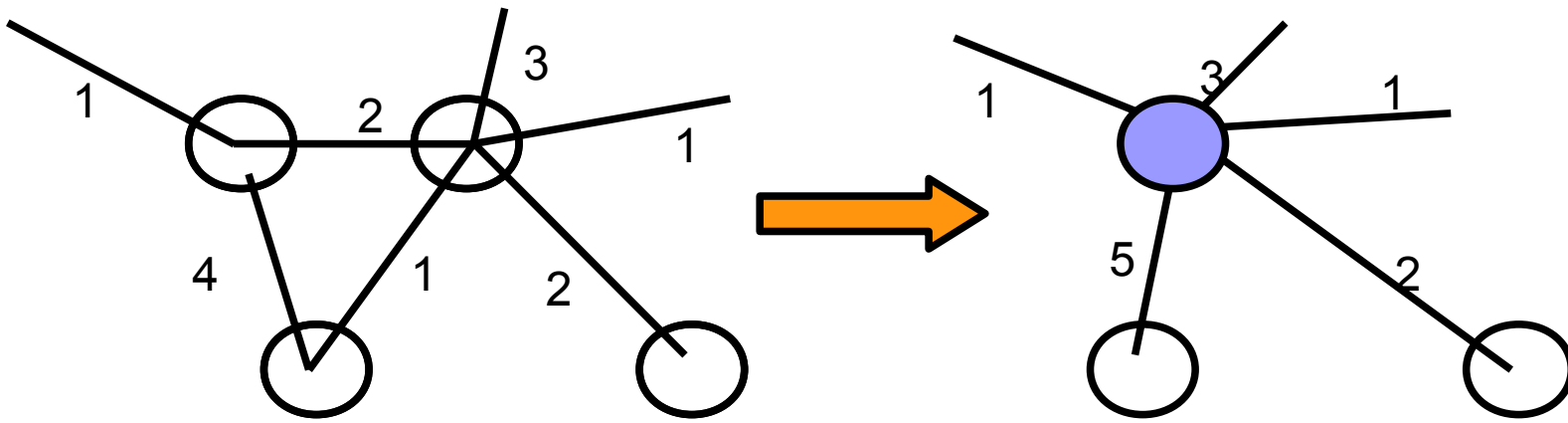


We can identify the more well connected areas of sea, that is where highly connected communities of a particular species .



$$C(1) = 3 / (4 * 3 / 2) = 1/2$$

$$C(2) = 0$$



**Fusion
Algorithm**

Stop: - no more increment of the metric is possible
 - after a certain number of steps

→ **Hierarchy of clusters!** ←

Blondel et al (2008),

“Fast unfolding of communities in large networks”

Increase modularity

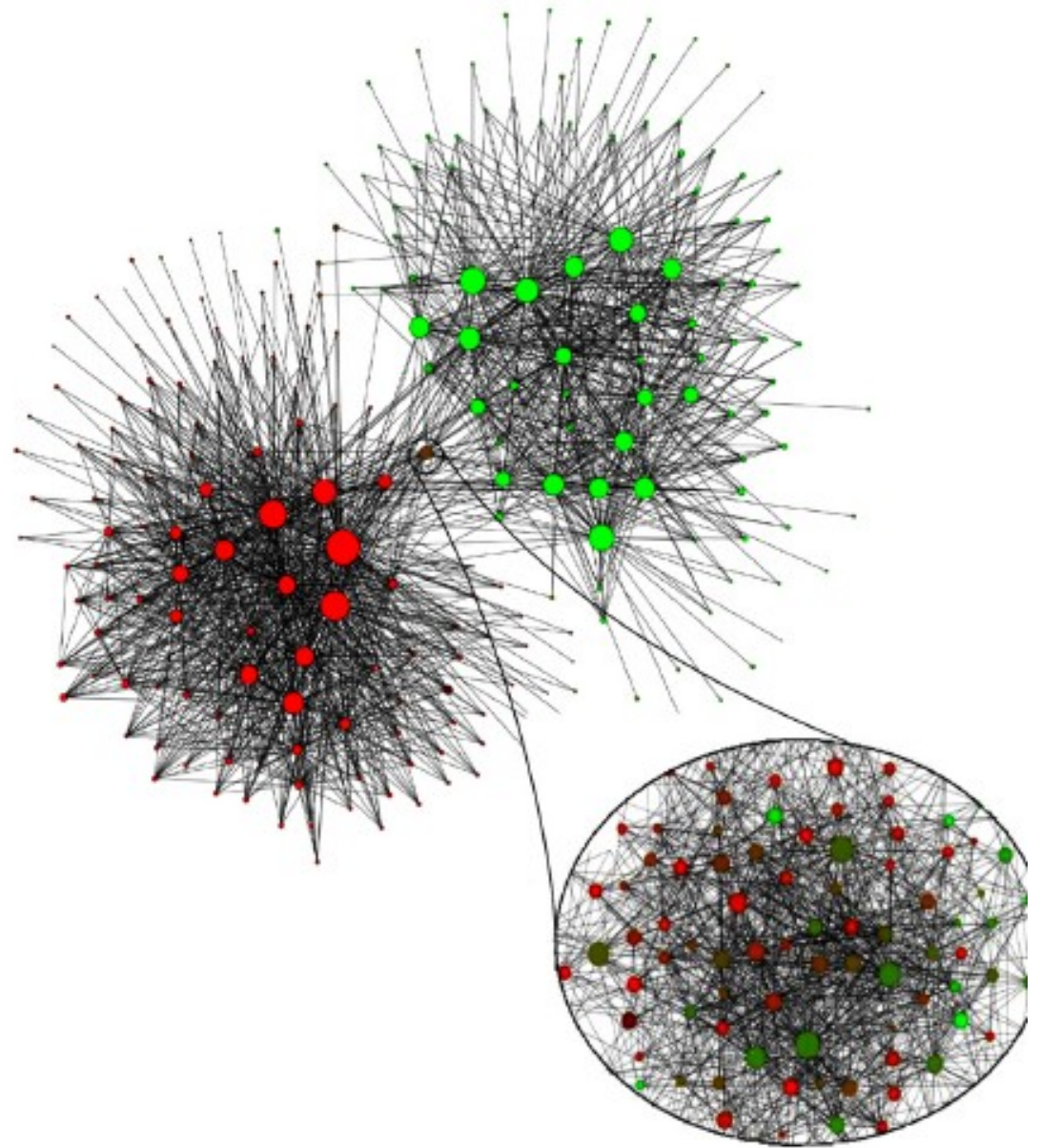


Figure 2. Graphical representation of the network of communities extracted from a Belgian mobile phone network. About 2 million customers are represented in this network. The size of a node is proportional to the number of individuals in the corresponding community and its colour on a red–green scale represents the main language spoken in the community (red for French and green for Dutch). Only the communities composed of more than 100 customers have been plotted. Notice the intermediate community of mixed colours between the two main language clusters. A zoom at higher resolution reveals that it is made of several sub-communities with less apparent language separation.

“Network analysis identifies weak and strong links in a metapopulation system”

A Rozenfeld et al 2008:

→ *Posidonia* genetic analysis and robustness analysis of the network

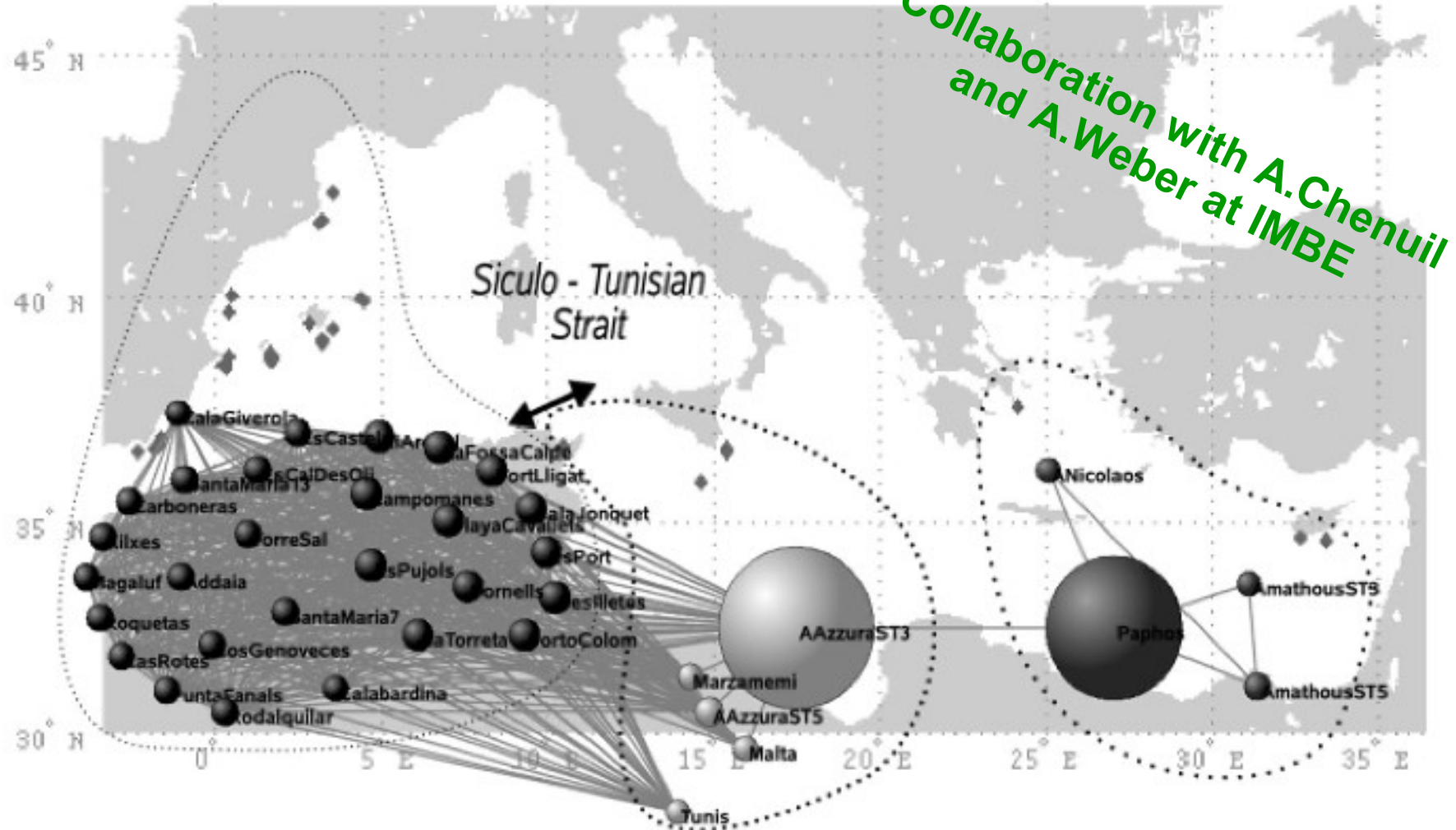


Fig. 1. The network of Mediterranean meadows in which only links with Goldstein distances smaller than the percolation distance $D_p = 91$ (see Fig. 5) are present. Nodes representing populations are roughly arranged according to their geographic origin. The precise geographic locations are indicated as diamonds in the background map. One can identify 2 clusters of meadows, corresponding to the Mediterranean basins (east and west), separated by the Siculo-Tunisian Strait. The size of each node indicates its betweenness centrality (i.e., the proportion of all shortest paths getting through the node).



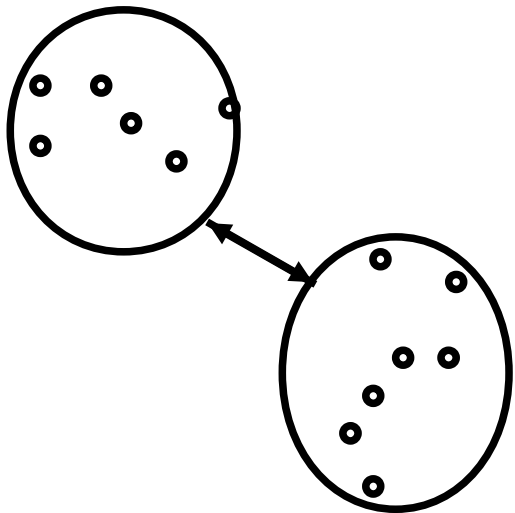
The genetic distance is symmetric,
the current-based one is not.

Which definition of distance we shall use?

I thought to 3 possibilities:

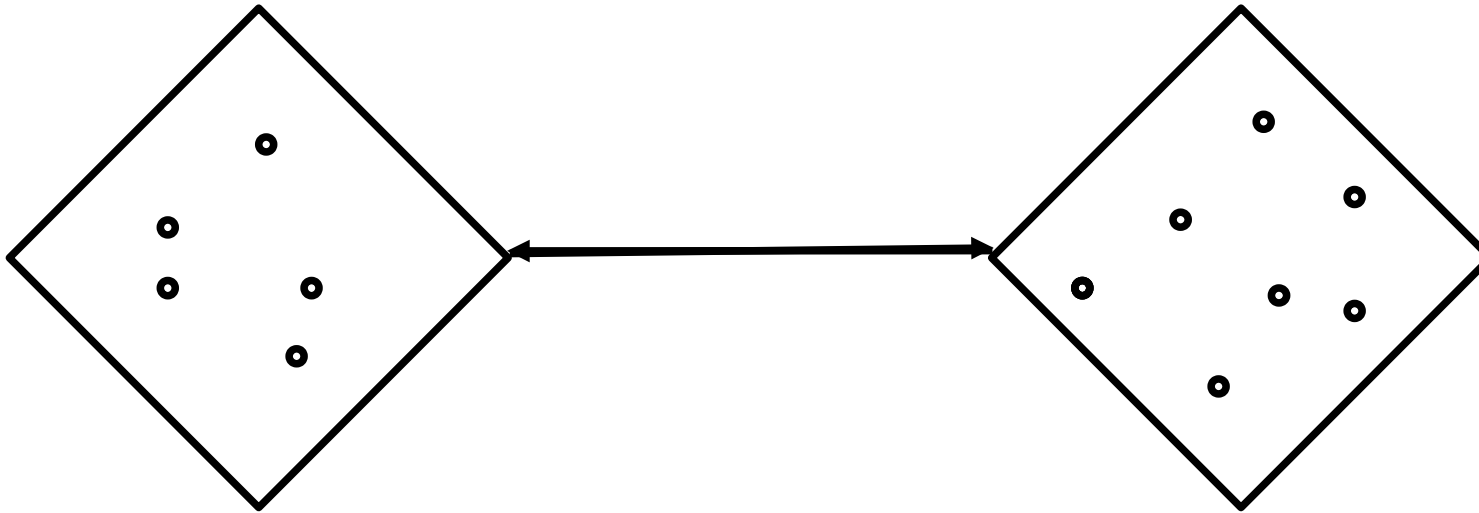
1) a mean genetic variation of the
population respect to a model
and then the difference between
the populations:

$$D_1(1,2) = \left| \frac{\sum_{i \in 1} (g_i - \bar{g})}{I} - \frac{\sum_{j \in 2} (g_j - \bar{g})}{J} \right| = D_1(2,1)$$



2) all vs all:

$$D_2(1,2) = \frac{\sum_{i \in 1} \sum_{j \in 2} (g_i - g_j)}{I \cdot J} = D_2(2,1)$$



(?) g is a vector in the 4-dimensional genetic space

Relatively to case 1: if we have $\text{Div}(G) \gg 0$ we have a genetic differentiation(?)

3) an asymmetric combination of the precedent methods:

$$D_3(1,2) = \frac{D_2(1,2) - \sum_{k, i \in 1 \wedge k \neq i} (g_i - g_k)}{I \cdot J} \neq D_3(2,1)$$

The smaller value points towards the more variable site: it is a source of genetic variability.