



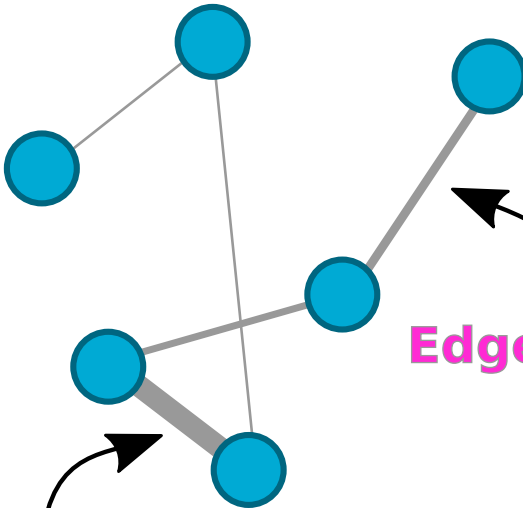
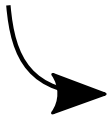
Enhanced extraction of weighted networks backbones

Alessio Cardillo

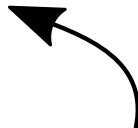
École polytechnique fédérale de Lausanne (EPFL), Lausanne, Switzerland

Wednesday 24 May 2017, Physics of Living Systems Meeting, EPFL, Switzerland

Node



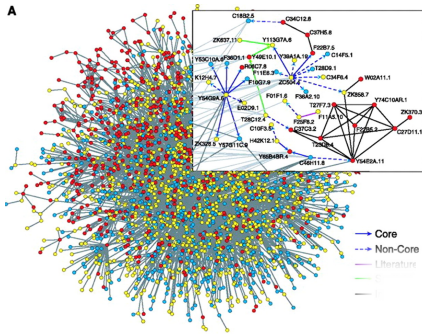
Edge / Link



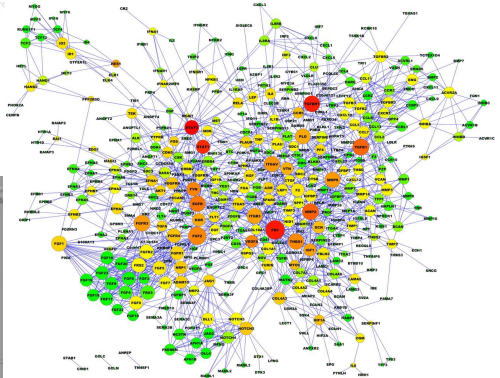
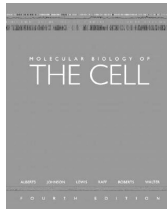
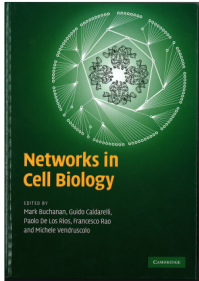
Weight = interaction

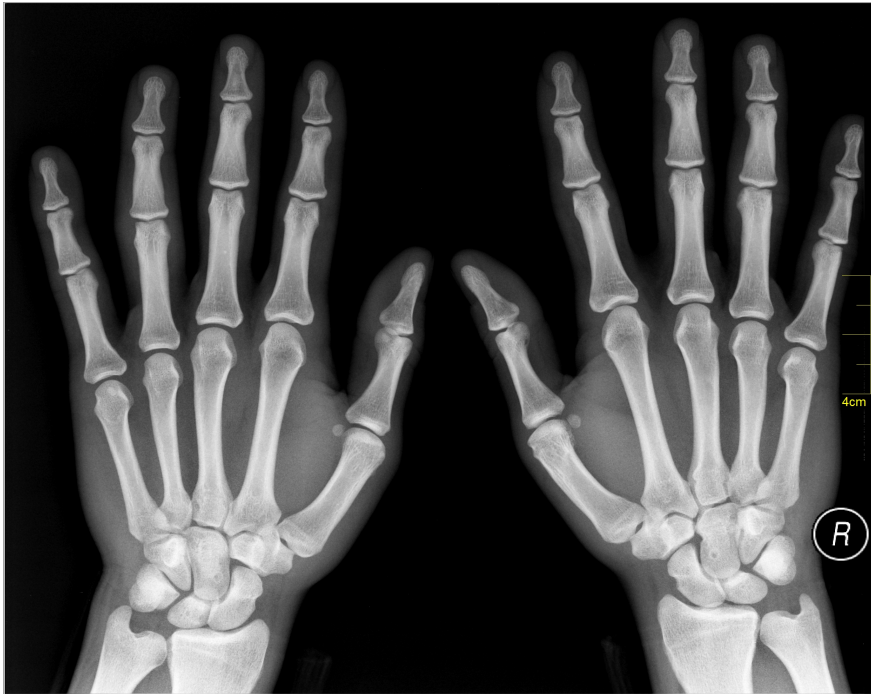


A

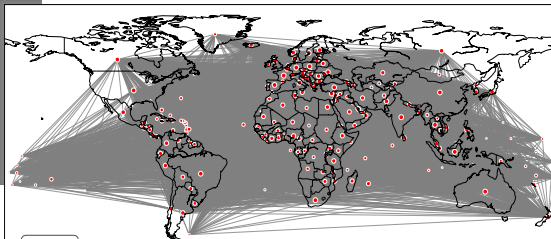
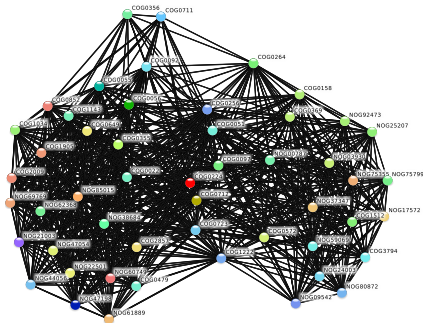
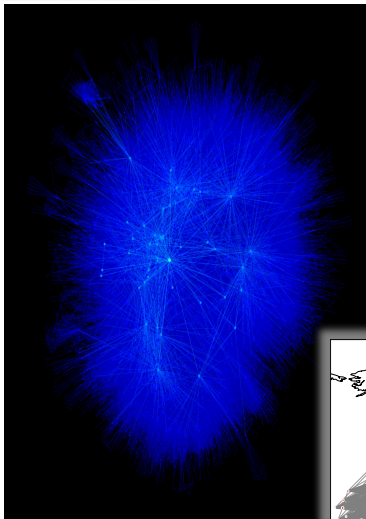


Biology and Networks







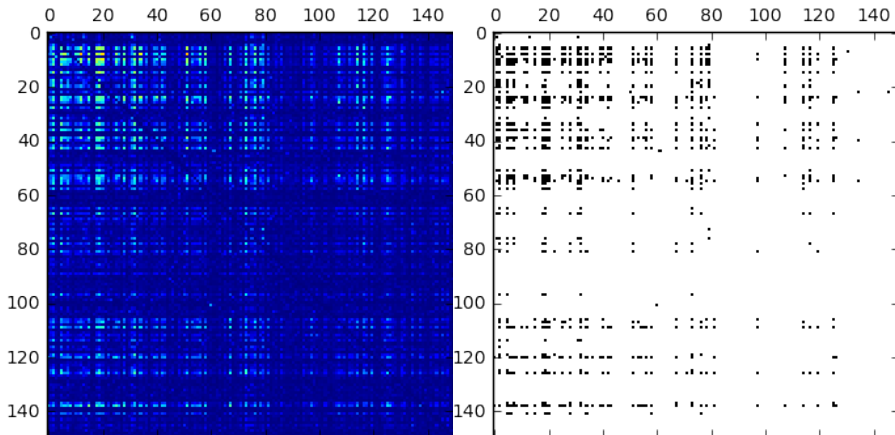


Question:

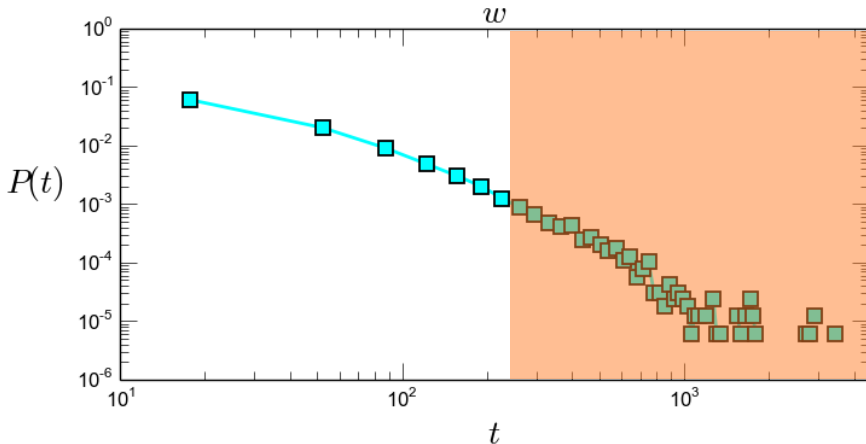
What can we **learn** from a complex system whose **network representation** is **noisy** and/or extremely **dense**?

BRACE YOURSELVES

FILTERING IS COMING



- De Vico Fallani F., et al. *A Topological Criterion for Filtering Information in Complex Brain Networks*. PLoS Comp. Bio. **13** e1005305 (2017).



- Granovetter, M. S. *The Strength of Weak Ties*. Am. Jour. Soc., **78**, 1360 (1973).

Main Features

1. Based on the comparison between the observed network and a null model one.

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2. Null model: maximum-entropy canonical ensemble of networks satisfying given constraints.

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1. Based on the comparison between the observed network and a null model one.
2. Null model: maximum-entropy canonical ensemble of networks satisfying given constraints.
3. Two versions: **local** (focus on links) and **global** (focus on entire network).

• Mastrandrea, R., *et al.* *Enhanced reconstruction of weighted networks from strengths and degrees*. New Jour. Phys., **16**, 043022. (2014).

Results

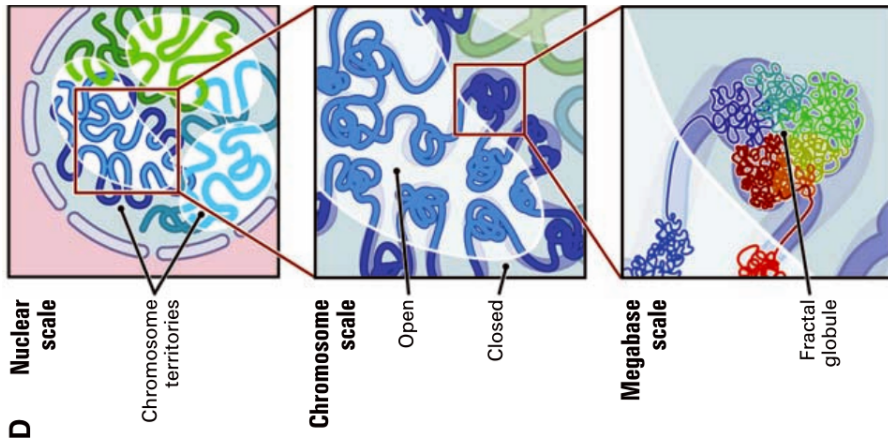


A serious example ...

A long time ago in a galaxy far,
far away....



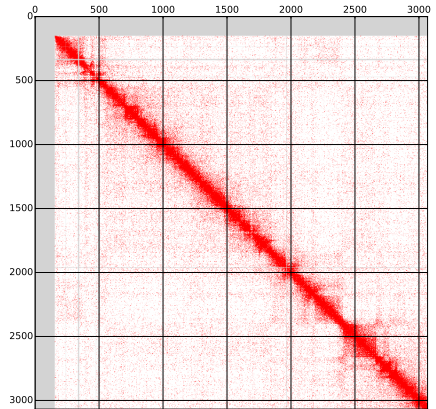




Data

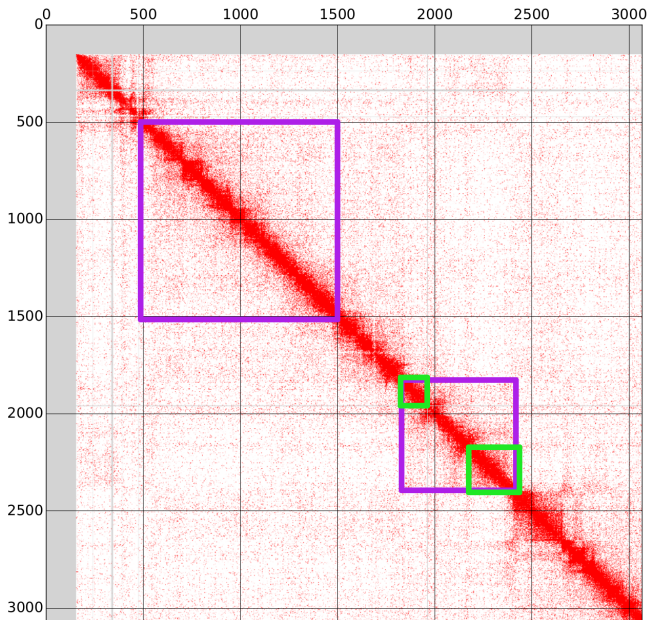
- ▶ Mouse (20 chr)
- ▶ Hi-C / cis-interactions
- ▶ Res. 20 Kbp
- ▶ Network sizes:
 $N \in [3068, 9868]$

chr-19 p-val = 1.0

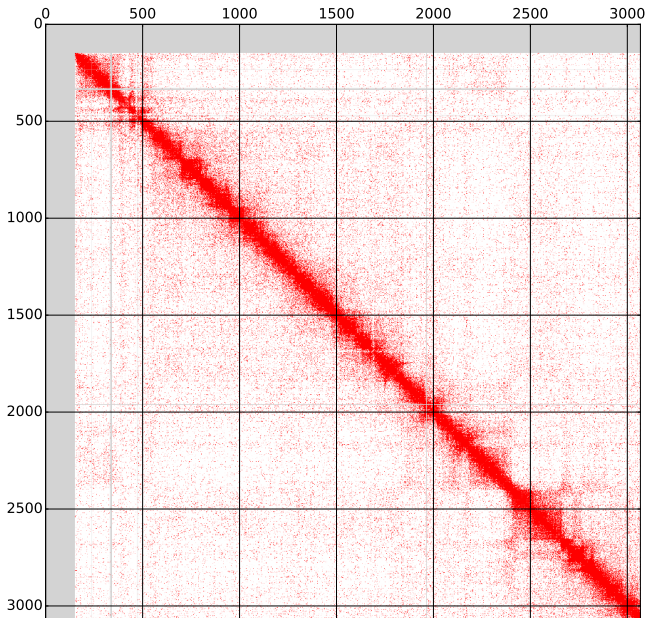


- Imakaev, M. *et al.* Nature Methods, **9**, 999–1003. (2012).
- Giorgetti, L., *et al.* Nature, **535**, 575–579. (2016).

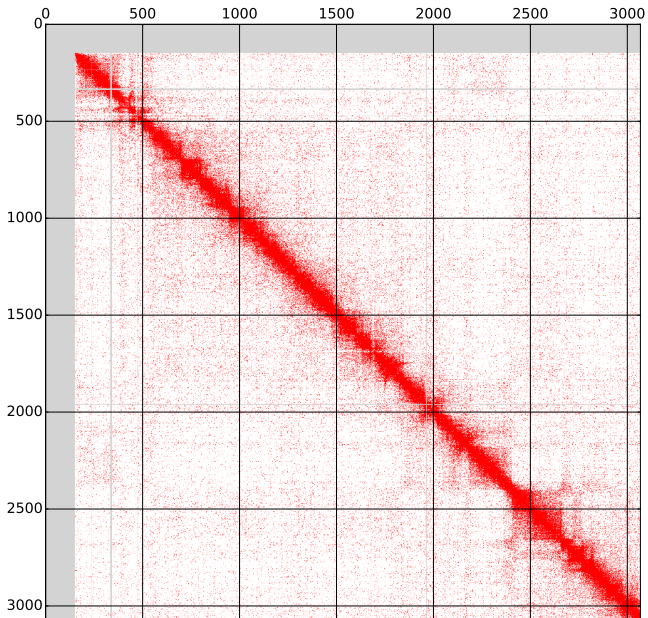
chr-19 p-val = 1.0



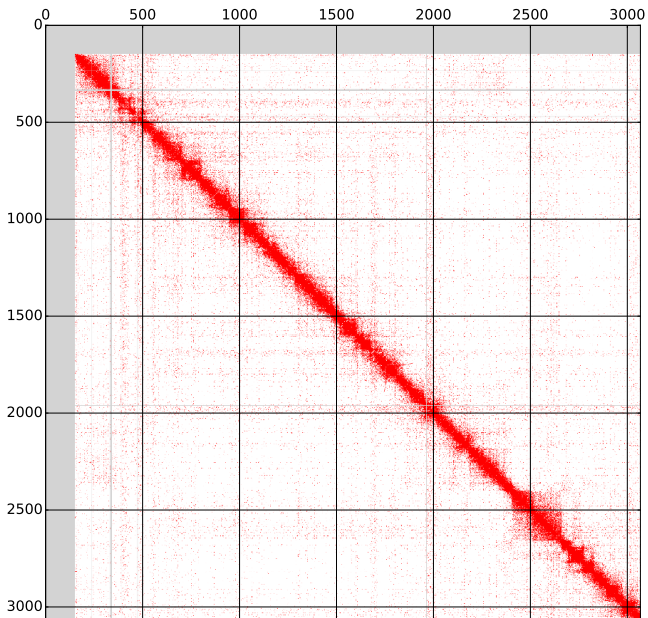
chr-19 p-val = 1.0



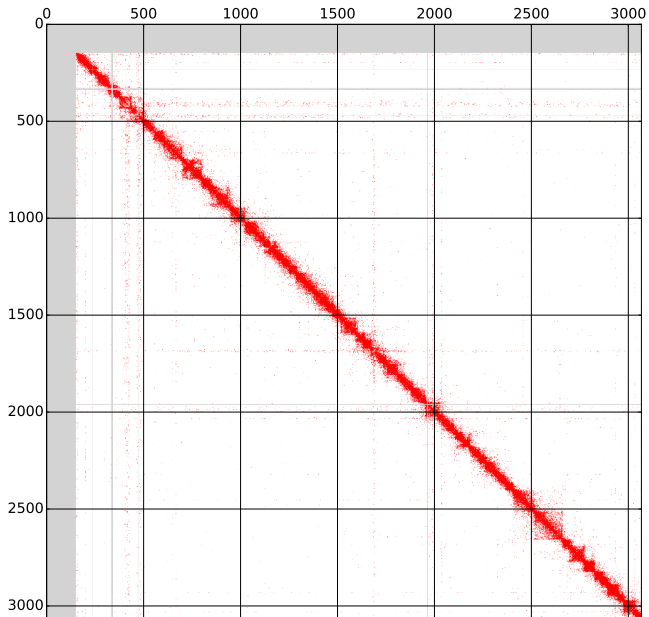
chr-19 p-val = 0.25



chr-19 p-val = 0.1



chr-19 p-val = 0.05

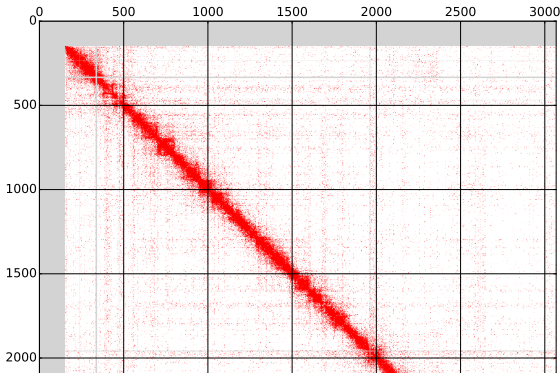


Conclusions

Filtering is becoming a more and more required step to continue using networks in complex systems.



chr-19 p-val = 0.1



ECM filter overcomes limitations of previous approaches and retrieves non trivial features.





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Valerio Gemmetto

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<http://bifi.es/~cardillo/>

Wanna know more?

Stay tuned on the arXiv ...

1. Generate the null model ensemble and compute:

$$q_{ij}(w) \equiv \frac{(x_i x_j)^{\Theta(w_{ij})} (y_i y_j)^{w_{ij}} (1 - y_i y_j)}{1 - y_i y_j + x_i x_j y_i y_j}.$$

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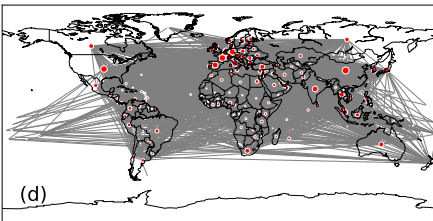
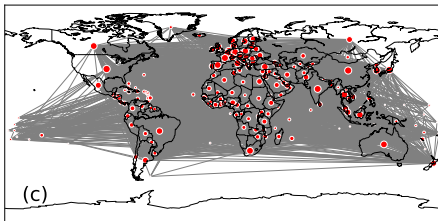
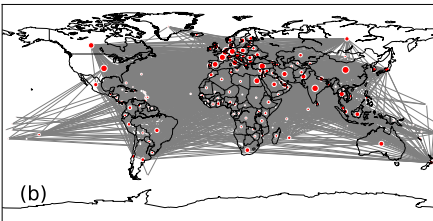
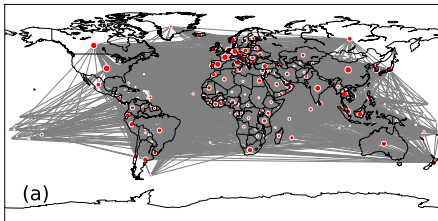
2. Associate for each link a p -value, γ , such that $P(w_{ij} > w_{ij}^*)$.

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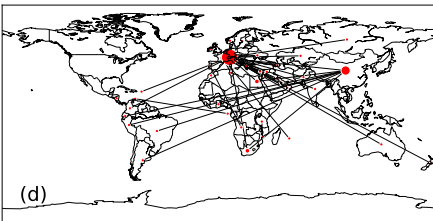
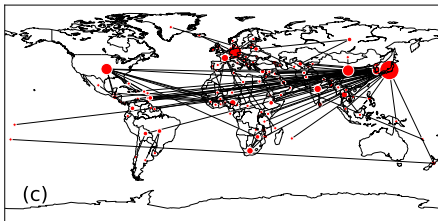
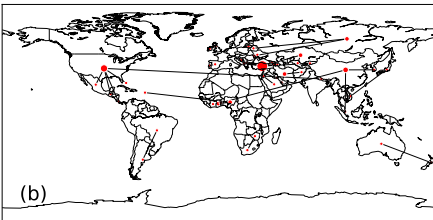
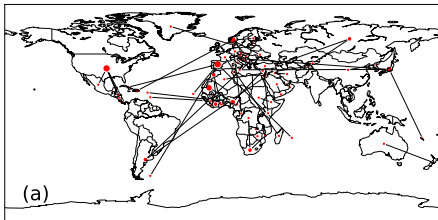
$$q_{ij}(w) \equiv \frac{(x_i x_j)^{\Theta(w_{ij})} (y_i y_j)^{w_{ij}} (1 - y_i y_j)}{1 - y_i y_j + x_i x_j y_i y_j}.$$

2. Associate for each link a p -value, γ , such that $P(w_{ij} > w_{ij}^*)$.
3. Select a threshold $\tilde{\gamma}$ and remove all the links with $\gamma_{ij} > \tilde{\gamma}$.

Trade: single commodities



Trade: single commodities



Trade: single commodities

