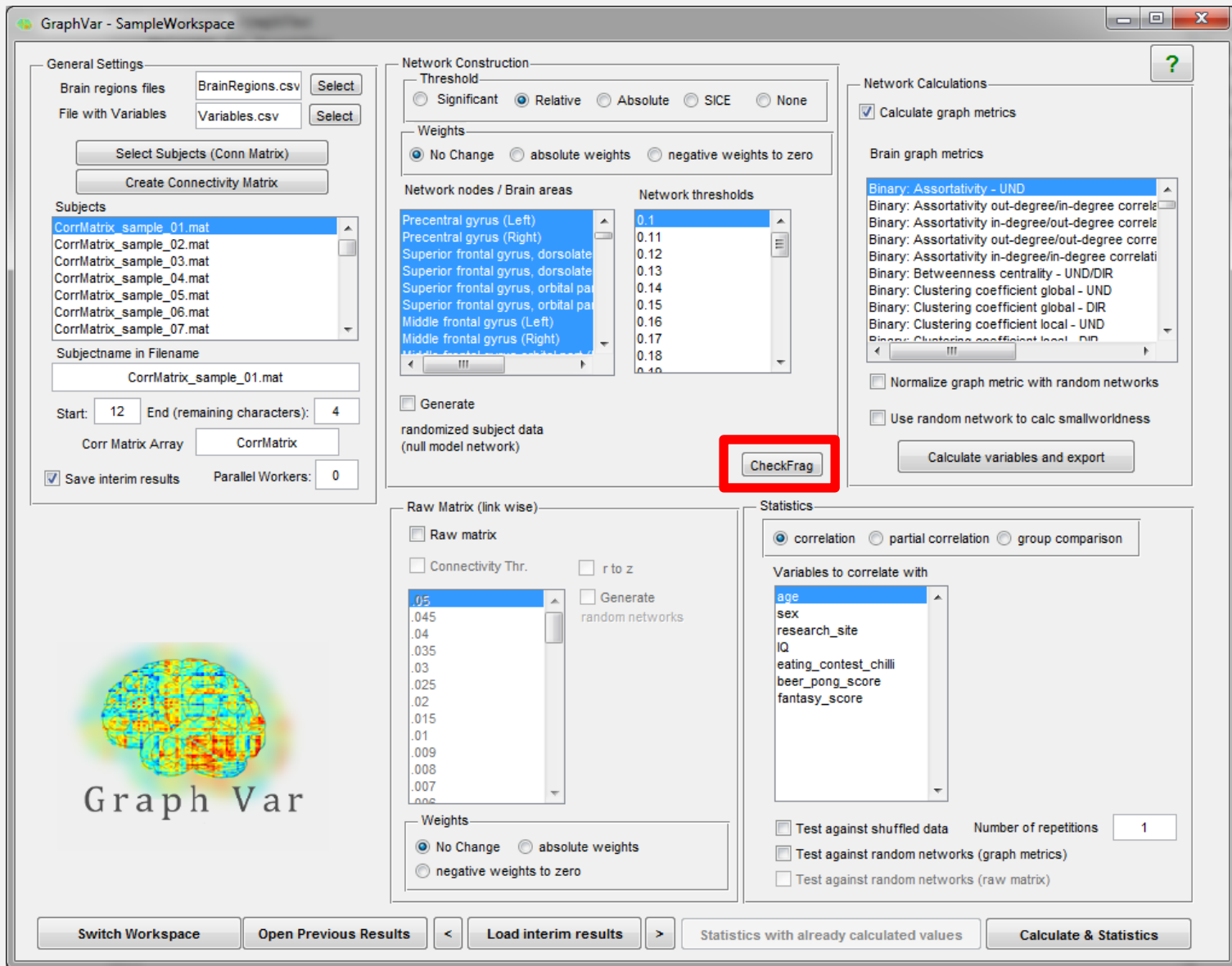
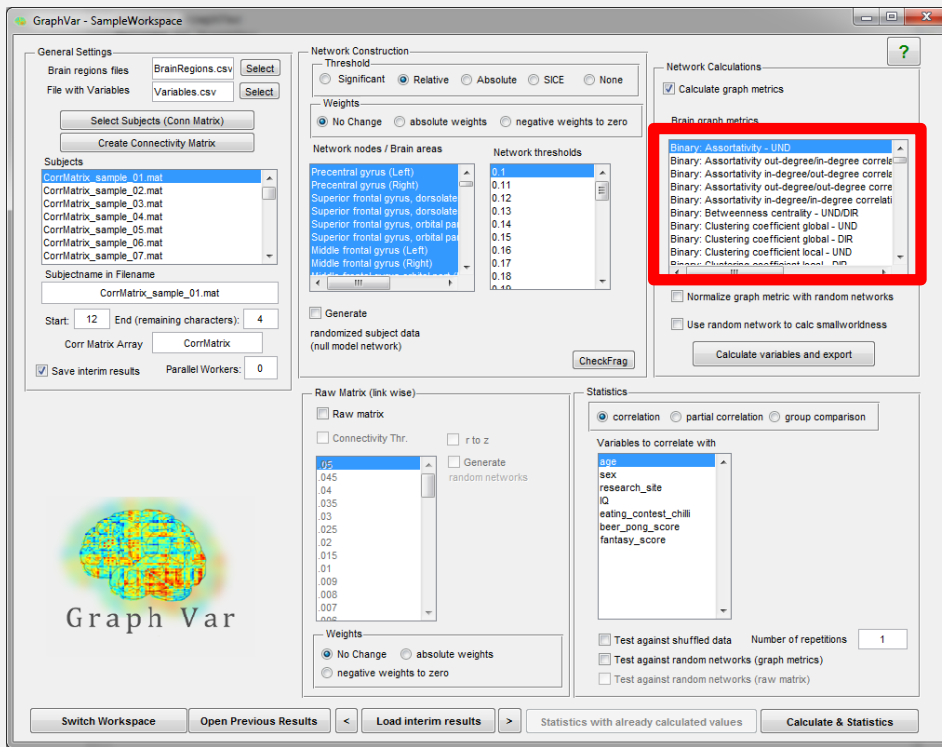


GraphVar beta 0.62 new features

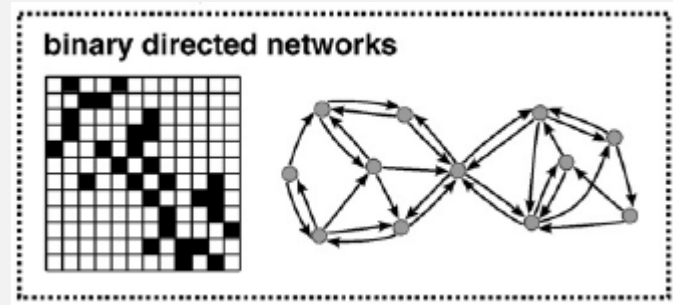


CheckFrag – check network fragmentation

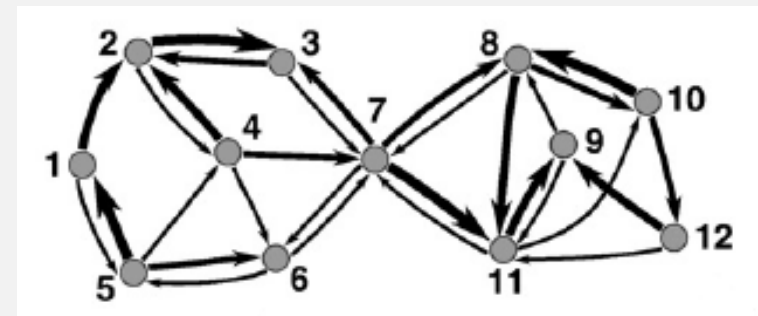
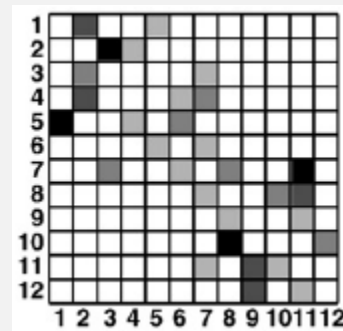




structural datasets: tract tracing  
effective datasets: inference of causality  
from functional data

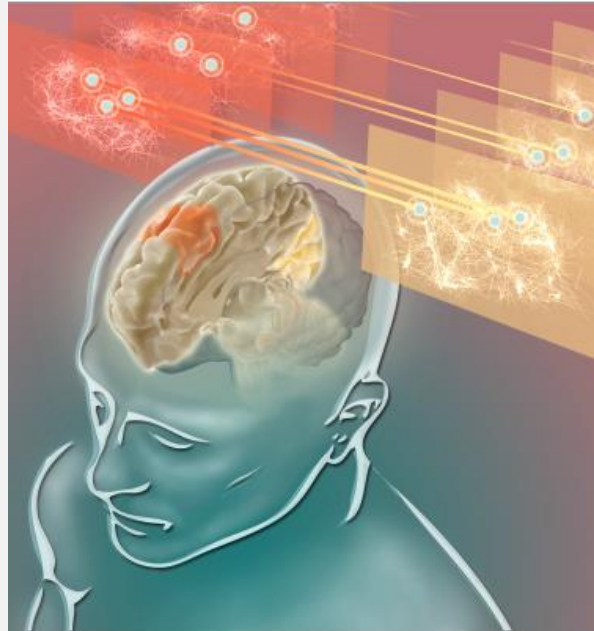


weighted directed networks



Added the directed BCT functions

# Network Community Toolbox



<http://commdetect.weebly.com/>

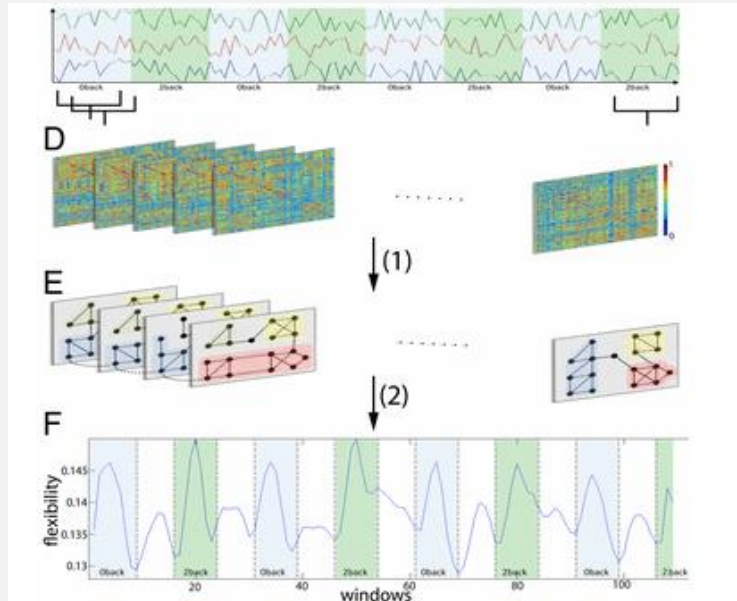
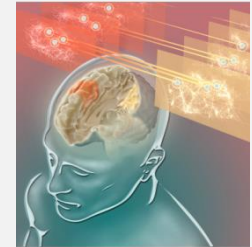
**time-dependent network flexibility**

**time-dependent nodal promiscuity**

Added two new measures from „Network Community Toolbox“

## Dynamic reconfiguration of frontal brain networks during executive cognition in humans

Urs Braun<sup>a,1</sup>, Axel Schäfer<sup>a,1</sup>, Henrik Walter<sup>b,1</sup>, Susanne Erk<sup>b</sup>, Nina Romanczuk-Seiferth<sup>b</sup>, Leila Haddad<sup>a</sup>, Janina I. Schweiger<sup>a</sup>, Oliver Grimm<sup>a</sup>, Andreas Heinz<sup>b</sup>, Heike Tost<sup>a</sup>, Andreas Meyer-Lindenberg<sup>a,1</sup>, and Danielle S. Bassett<sup>c,d,1,2</sup>



### time-dependent network flexibility:

the time-dependent flexibility of a region is defined as the probability that a brain region changed its allegiance to putative functional modules between any two consecutive time windows. Intuitively, flexibility can be thought of as a statistic to quantify the amount of reconfiguration in functional connectivity patterns that a brain region displays over time (Braun et al., 2015).

**GraphVar uses the generalized Louvain algorithm – so please cite:**

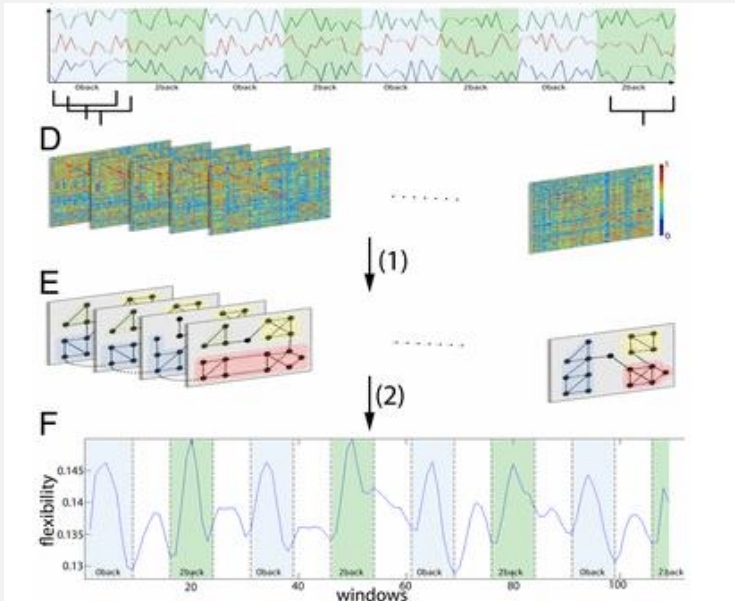
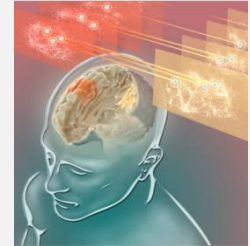
*Inderjit S. Jutla, Lucas G. S. Jeub, and Peter J. Mucha,*

*"A generalized Louvain method for community detection implemented in MATLAB," <http://netwiki.amath.unc.edu/GenLouvain> (2011-2014).*

*Mucha PJ, Richardson T, Macon K, Porter MA, Onnela JP (2010) Community structure in time-dependent, multiscale, and multiplex networks. Science 328(5980):876–878.*

I: „dynamic“ nodal flexibility as in Braun et al. 2015

# Network Community Toolbox



## time-dependent nodal promiscuity:

This function calculates the promiscuity coefficient.

The promiscuity of a temporal or multislice network corresponds to the fraction of all the communities in the network in which a node participates at least once.

**GraphVar uses the generalized Louvain algorithm – so please cite:**

*Inderjit S. Jutla, Lucas G. S. Jeub, and Peter J. Mucha,*

*"A generalized Louvain method for community detection implemented in MATLAB," <http://netwiki.amath.unc.edu/GenLouvain> (2011-2014).*

*Mucha PJ, Richardson T, Macon K, Porter MA, Onnela JP (2010) Community structure in time-dependent, multiscale, and multiplex networks. Science 328(5980):876–878.*

II: „dynamic“ nodal promiscuity“ as <http://commdetect.weebly.com/>

# Network Community Toolbox

<http://commdetect.weebly.com/>

time-dependent network flexibility

time-dependent nodal promiscuity

if normalization with random network is performed NaN's may result  
as these measures can produce zeros as results

(e.g. promiscuity  $P = 0$  if it only participates in one community and  $P=1$  if  
it participates in every community of the network)


... thus check your results in the interim results folder:

multislice\_community\_assignment\_slicesXnodes.mat  
multislice\_community\_assignment\_slicesXnodes-rand.mat

Added two new measures from „Network Community Toolbox“



**Small-World-Propensity:** unbiased assessment of small-world structure in networks of varying densities - developed by Muldoon, Bridgeford and Bassett

 Cornell University  
Library

arXiv.org > q-bio > arXiv:1505.02194

Quantitative Biology > Neurons and Cognition

**Small-World Propensity in Weighted, Real-World Networks**

Sarah Feldt Muldoon, Eric W. Bridgeford, Danielle S. Bassett



(Submitted on 8 May 2015)

Quantitative descriptions of network structure in big data can provide fundamental insights into the function of interconnected complex systems. Small-world structure, commonly diagnosed by high local clustering yet short average path length between any two nodes, directly enables information flow in coupled systems, a key function that can differ across conditions or between groups. However, current techniques to quantify small-world structure are dependent on nuisance variables such as density and agnostic to critical variables such as the strengths of connections between nodes, thereby hampering accurate and comparable assessments of small-world structure in different networks. Here, we address both limitations with a novel metric called the Small-World Propensity (SWP). In its binary instantiation, the SWP provides an unbiased assessment of small-world structure in networks of varying densities. We extend this concept to the case of weighted networks by developing (i) a standardized procedure for generating weighted small-world networks, (ii) a weighted extension of the SWP, and (iii) a stringent and generalizable method for mapping real-world data onto the theoretical model. In applying these techniques to real world brain networks, we uncover the surprising fact that the canonical example of a biological small-world network, the *C. elegans* neuronal network, has strikingly low SWP in comparison to other examined brain networks. These metrics, models, and maps form a coherent toolbox for the assessment of architectural properties in real-world networks and their statistical comparison across conditions.

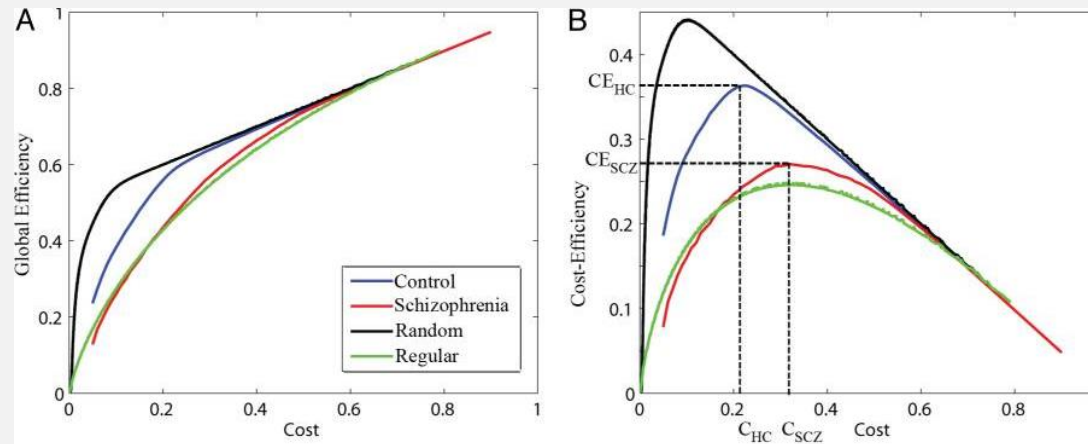
<http://arxiv.org/pdf/1505.02194v1.pdf>

Requires the Bioinformatics Toolbox

Added also Small-World-Propensity

## Cognitive fitness of cost-efficient brain functional networks

[Danielle S. Bassett](#)<sup>a,b,c</sup>, [Edward T. Bullmore](#)<sup>b,d,1</sup>, [Andreas Meyer-Lindenberg](#)<sup>e</sup>, [José A. Apud](#)<sup>a</sup>, [Daniel R. Weinberger](#)<sup>a</sup>, and [Richard Coppola](#)<sup>f</sup>

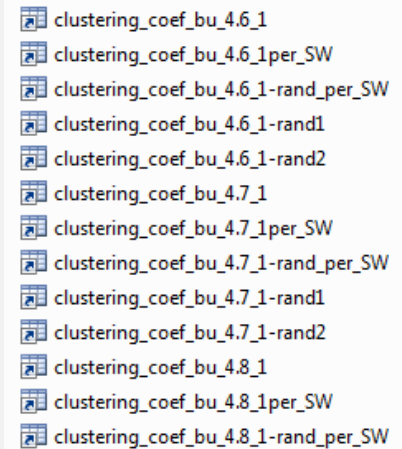


Added also Cost-Efficiency as in Bassett et al., 2009

- GraphVar saves more output files when doing sliding window analyses (not only the dynamic summary measure as before) and will also save all results when only "calculate and Export" for further usage.

Files in your interim results folder are (depending on the computations – here with clustering\_coefficient as example):

- **clustering\_coef\_bu\_4.9\_1.mat**: dynamic summary measure (e.g. variance) of clustering\_coef\_bu across windows for each node on threshold 0.49 for all subjects
- **clustering\_coef\_bu\_4.9\_1per\_SW.mat**: the (normalized) clustering\_coef\_bu for each node in each of the sliding windows on threshold 0.49 for all subjects
- **clustering\_coef\_bu\_4.9\_1-rand1.mat**: dynamic summary measure (e.g. variance) of clustering\_coef\_bu across windows for each node in the first random network on threshold 0.49 for all subjects
- **clustering\_coef\_bu\_4.9\_1-rand\_per\_SW.mat**: the clustering\_coef\_bu for each node in each random network in each of the sliding windows on threshold 0.49 for all subjects (i.e., cell comprised of: subjects x random networks x sliding windows)



- clustering\_coef\_bu\_4.6\_1
- clustering\_coef\_bu\_4.6\_1per\_SW
- clustering\_coef\_bu\_4.6\_1-rand\_per\_SW
- clustering\_coef\_bu\_4.6\_1-rand1
- clustering\_coef\_bu\_4.6\_1-rand2
- clustering\_coef\_bu\_4.7\_1
- clustering\_coef\_bu\_4.7\_1per\_SW
- clustering\_coef\_bu\_4.7\_1-rand\_per\_SW
- clustering\_coef\_bu\_4.7\_1-rand1
- clustering\_coef\_bu\_4.7\_1-rand2
- clustering\_coef\_bu\_4.8\_1
- clustering\_coef\_bu\_4.8\_1per\_SW
- clustering\_coef\_bu\_4.8\_1-rand\_per\_SW

## More output for further usage