

Supplemental material

A benchmark model to assess community structure in evolving networks

C. Granell, R. K. Darst, A. Arenas, S. Fortunato, S. Gómez

	Time window	Jaccard squared error		
		Grow/Shrink	Merge/Split	Mixed
Multislice $\omega = 0.0$	1	0.0720	0.3345	0.0307
	2	0.1365	0.4840	0.0303
	5	0.2336	0.6272	0.0325
Multislice $\omega = 0.5$	1	0.0293	0.3193	0.0276
	2	0.0546	0.4608	0.0282
	5	0.1105	0.6013	0.0303
Multislice $\omega = 2.0$	1	0.0019	0.3326	0.0360
	2	0.0014	0.3605	0.0374
	5	0.0147	0.4488	0.0421

Table S1: Jaccard squared error, for each method tested and each benchmark, considering three different time windows.

	Time window	NMI squared error		
		Grow/Shrink	Merge/Split	Mixed
Multislice $\omega = 0.0$	1	0.0337	0.4932	0.0067
	2	0.0621	0.4806	0.0063
	5	0.1022	0.4855	0.0059
Multislice $\omega = 0.5$	1	0.0143	0.4896	0.0060
	2	0.0262	0.4753	0.0059
	5	0.0479	0.4790	0.0055
Multislice $\omega = 2.0$	1	0.0065	0.4951	0.0094
	2	0.0041	0.4891	0.0094
	5	0.0041	0.4825	0.0100

Table S2: NMI squared error, for each method tested and each benchmark, considering three different time windows.

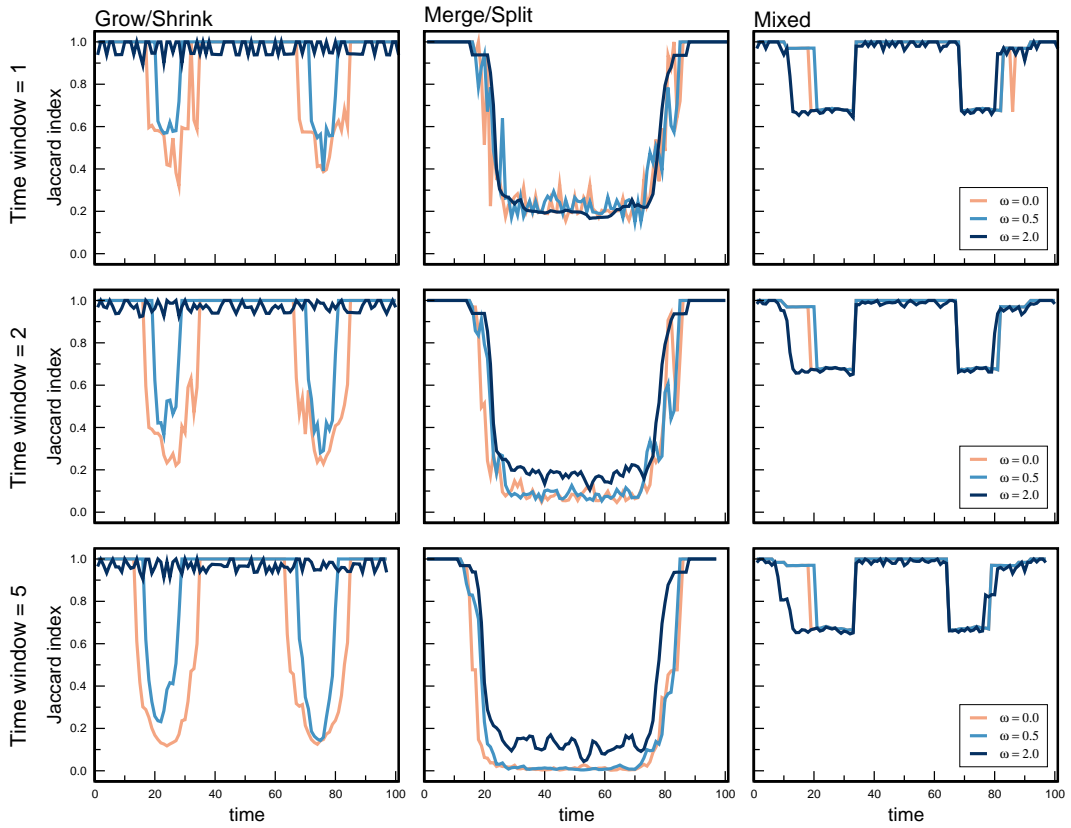


Figure S1: Plots of the Jaccard Index between the planted partition and the results of the multislice algorithm for three different inter-slice couplings and for the three benchmarks proposed. The Jaccard index is computed using the proposed evolving formulation and for three different window sizes: 1, 2 and 5. There is a column for each benchmark, and a row for each time window size.

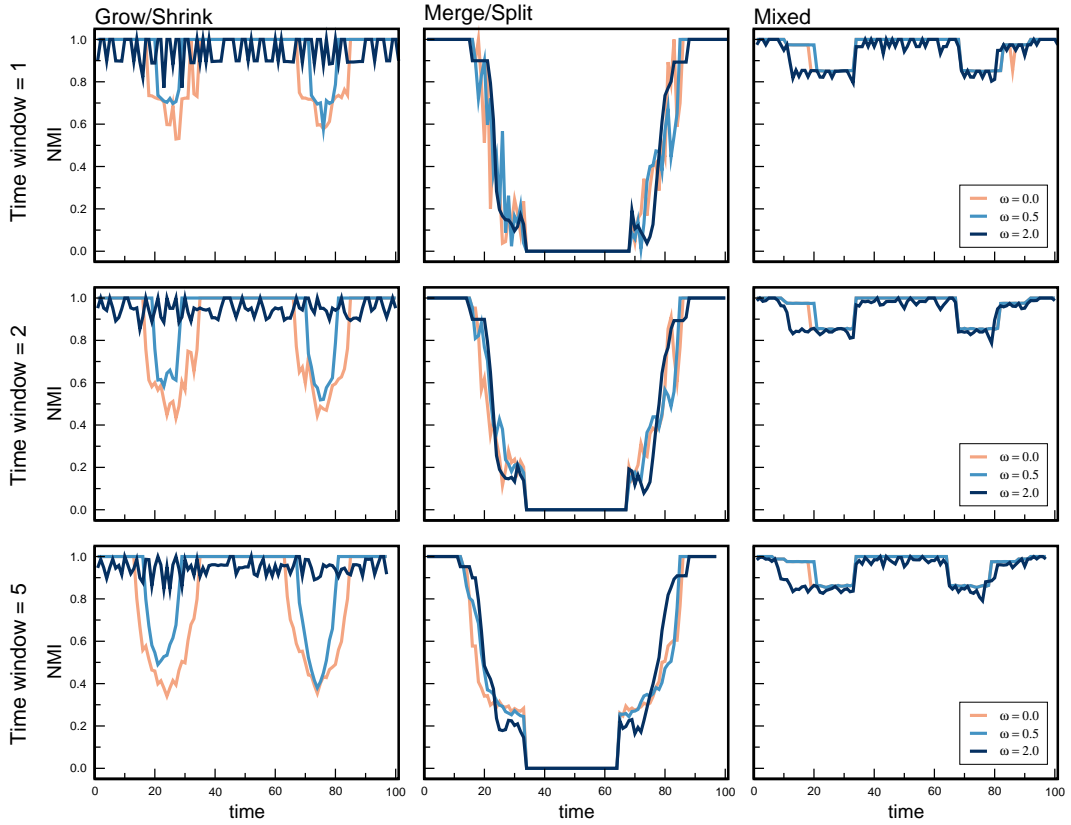


Figure S2: Plots of the Normalized Mutual Information (NMI) between the planted partition and the results of the multislice algorithm for three different inter-slice couplings and for the three benchmarks proposed. The NMI is computed using the proposed evolving formulation and for three different window sizes: 1, 2 and 5. There is a column for each benchmark, and a row for each time window size.