

Motif-driven Dense Subgraph Discovery in Directed and Labeled Networks

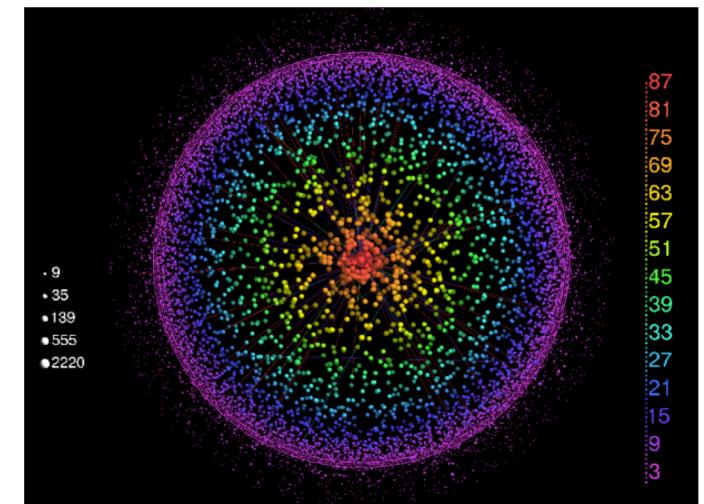
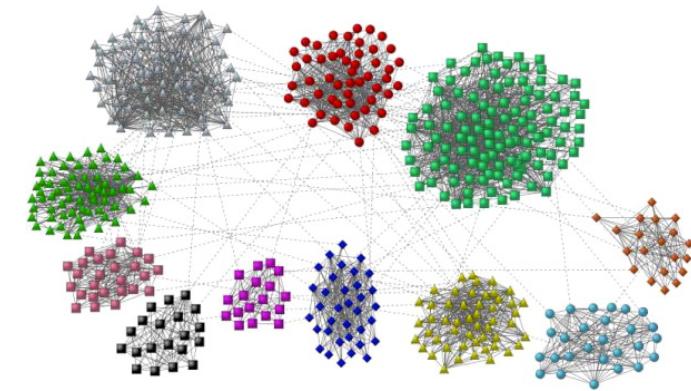
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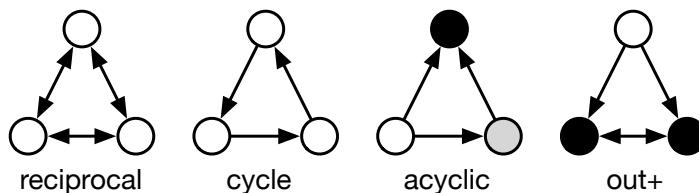
Dense subgraph discovery

- Dense regions are unusual and interesting
 - Anomaly detection, community detection, visualization
- A good proxy for graph clustering
 - Exhibit good cuts [Gleich and C. Seshadhri, 2012]
- Literature is rich for simple, undirected networks
- What about heterogeneous networks?
 - Directed edges
 - Labeled nodes/edges
 - Categorical
 - Numerical
 - How to even define the density?

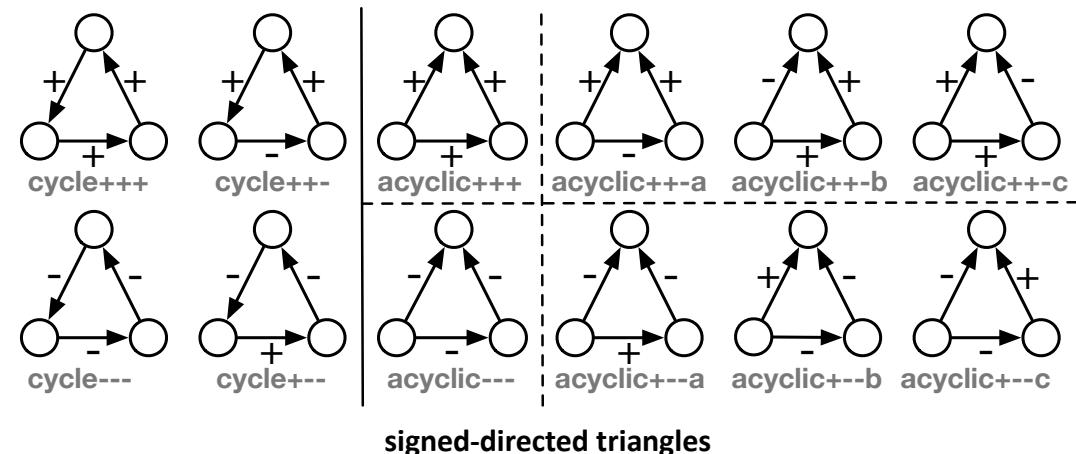


Motifs for help

- Fundamental building blocks in the organization and dynamics of real-world networks
- Captures **higher-order relationships** among multiple nodes
- **Density is the avg. motif degree**
 - Number-of-motifs / number-of-nodes



directed triangles



- Extendible for heterogeneous networks
 - Pros: Customizable; **dense subgraphs w.r.t. motif of interest**
 - Cons: **Spectrum is wide; hard to unify all in a framework**

Idea: Participations of small motifs in larger motifs

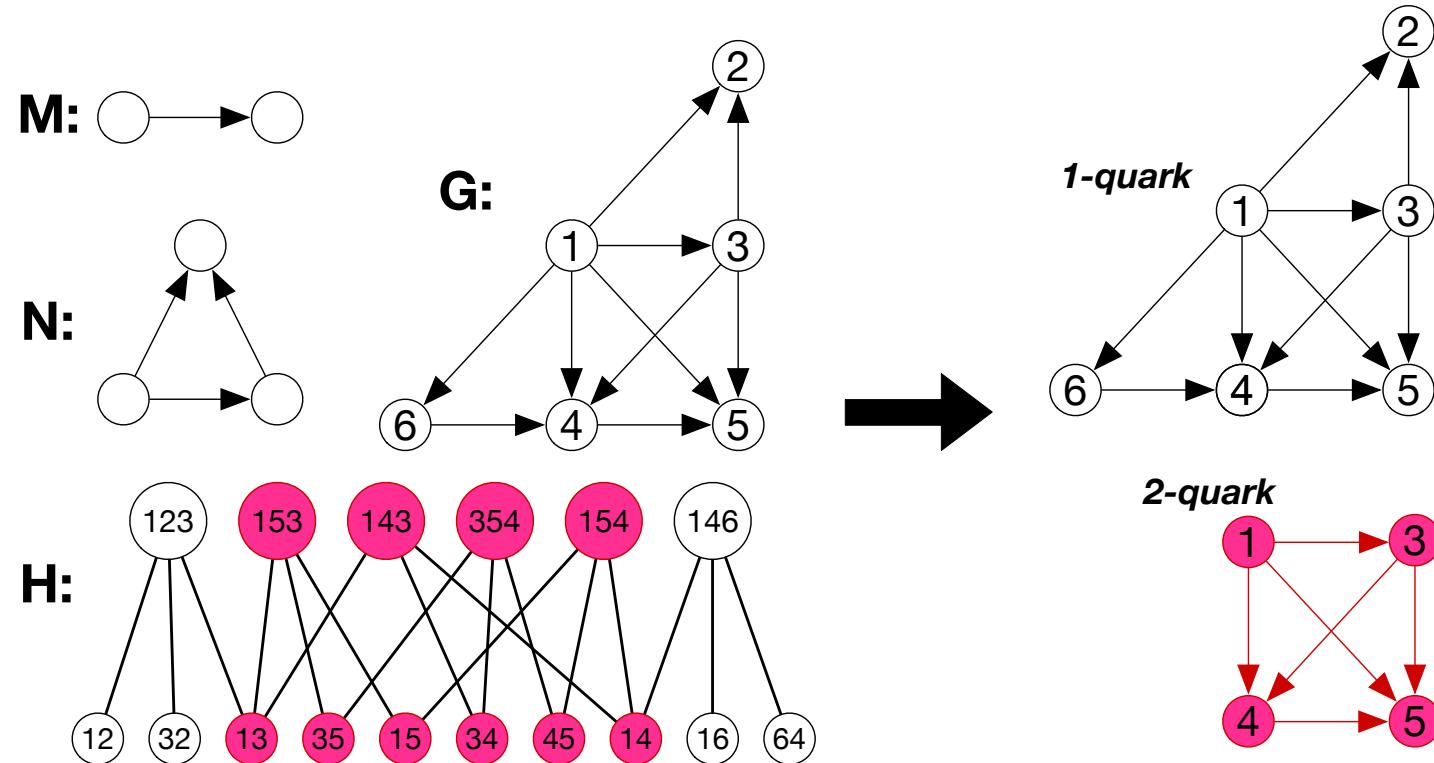
- Given a pair of motifs \mathbf{M} and \mathbf{N} s.t. $\mathbf{M} \subset \mathbf{N}$, find the subgraphs where each \mathbf{M} participates in many \mathbf{N} s
 - Inspired by core and truss decompositions
- \mathbf{M} and \mathbf{N} can have directed edges and categorical labels on nodes/edges
 - No numerical labels – future work
- Motif hypergraph:
 - \mathbf{M} s are the nodes
 - \mathbf{N} s are the hyperedges
 - An \mathbf{M} is connected to an \mathbf{N} iff $\mathbf{M} \subset \mathbf{N}$
- Motif of interest is \mathbf{N}

Quark decomposition

- Given a graph \mathbf{G} and motifs \mathbf{M}, \mathbf{N} ($\mathbf{M} \subset \mathbf{N}$), let \mathbf{H} be motif hypergraph,
 - A \mathbf{k} -quark is a connected and maximal sub-hypergraph where each \mathbf{M} instance participates in at least \mathbf{k} number of \mathbf{N} instances.
 - Quark number of an \mathbf{M} is the largest value of \mathbf{k} s.t. \mathbf{M} belongs to a \mathbf{k} -quark.

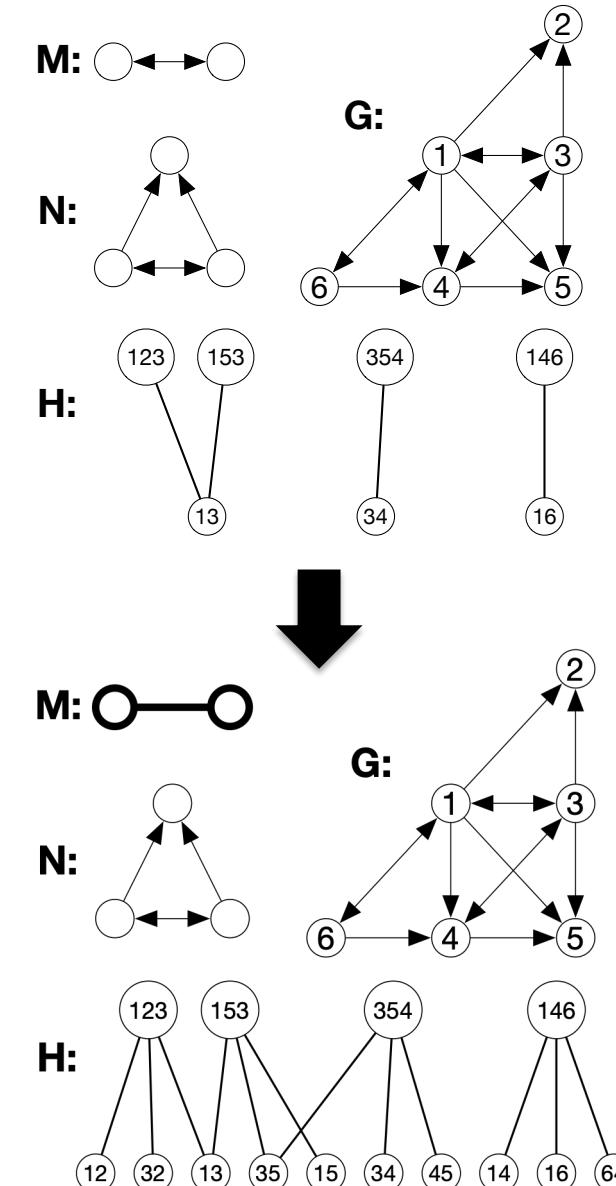
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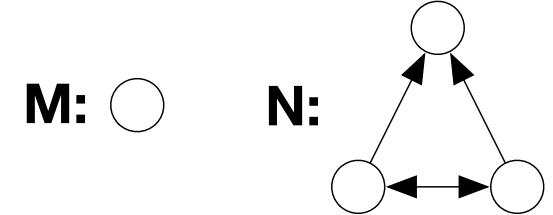
Limitations and practical instantiations

- What if there is only one M in N ?
 - Size of each N in the motif hypergraph becomes one!
 - How to avoid?
- Consider M as vanilla
 - Labelless nodes/edges, directionless edges
- M is better to be an edge (or larger)
 - Overlapping subgraphs!



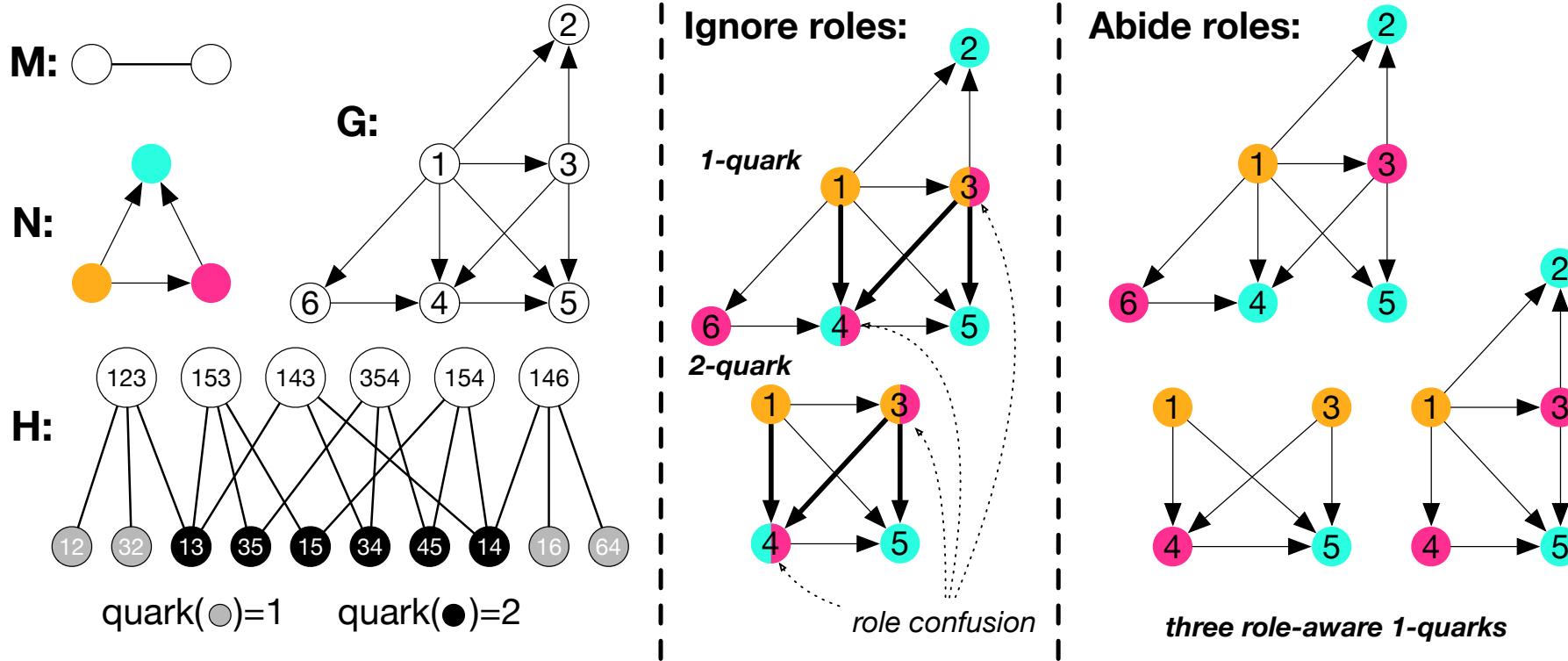
Role confusion problem

- What if \mathbf{M} has different “roles” in \mathbf{N} s it’s part of?
 - Orbits! [Pržulj, 2007]



- How to distinguish the participations where \mathbf{M} is in different orbits?
 - Orbit degrees: Number of \mathbf{N} s that contain \mathbf{M} s.t. \mathbf{M} is in a specific orbit
- Role-aware \mathbf{k} -quark: \mathbf{M} ’s orbit is the same in all the participations.
 - I.e., orbit degree of each \mathbf{M} is at least \mathbf{k}

Role confusion problem



- Role-aware k -quark: \mathbf{M} 's orbit is the same in all the participations.
 - I.e., orbit degree of each \mathbf{M} is at least k

Peeling algorithm works for quark decomposition!

- Both quark and role-aware quark decompositions
- Subgraph and hierarchy construction included
- When M is a node or edge, time complexity is

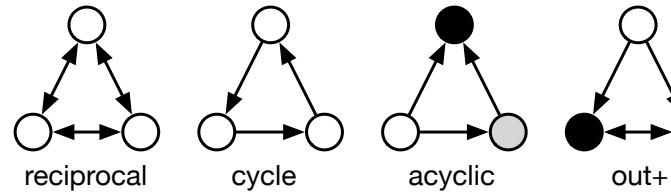
$$O\left(\sum_{v \in V} d(v)^{|V_N|-1}\right)$$



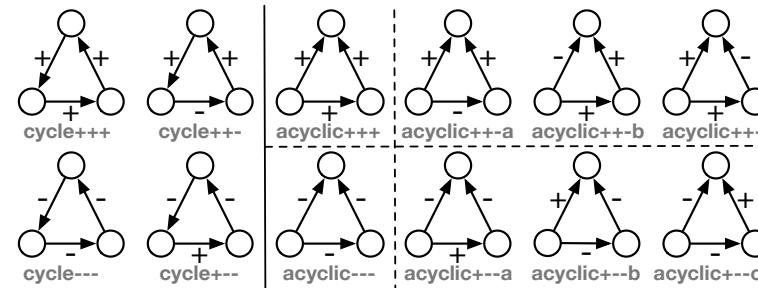
- Existing optimizations for peeling algorithms are applicable
 - Constructing subgraphs during the peeling
 - Parallel, local computations

Experimental evaluation on heterogeneous networks

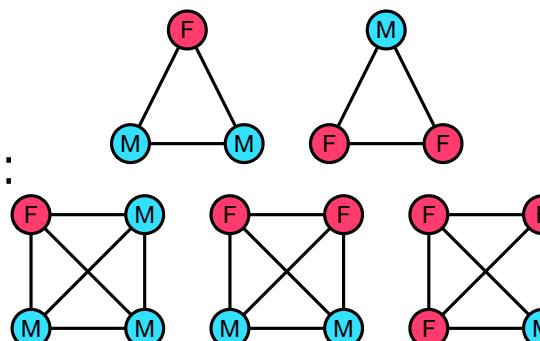
- Directed
 - M is edge
 - N is a triangle:



- Signed-directed
 - M is edge
 - N is a triangle:



- Node-labeled (genders)
 - M is edge or triangle
 - N is triangle or four-clique:

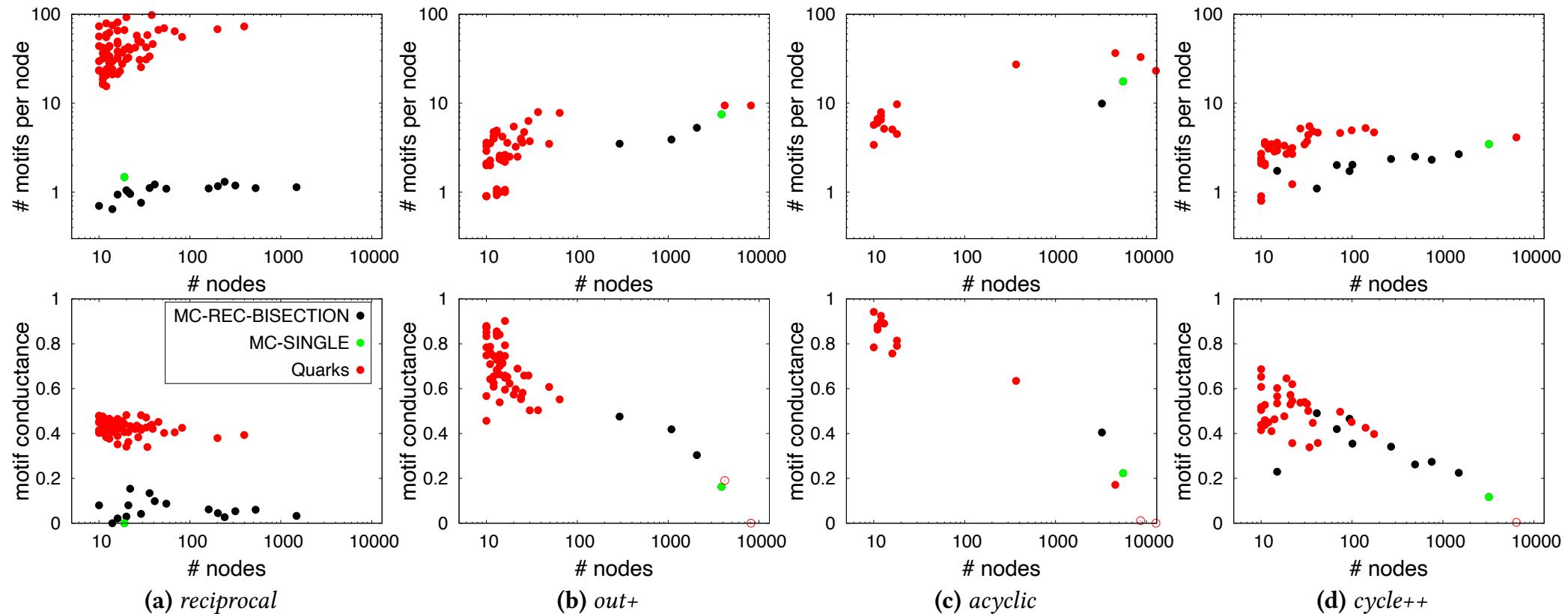


- Baselines:
 - Motif clustering
 - [Benson et al., 2016]
 - Cycle-truss and flow-truss
 - [Takaguchi and Yoshida, 2016]
 - Nucleus decomposition
 - [Sariyuce et al., 2015]

- Metrics
 - Motif conductance
 - Avg. motif degree
 - Edge density
 - For node-labeled

Quark decomposition vs. Motif clustering

- Motif clustering optimizes motif conductance, thus better
- Quark decomposition gives higher avg. motif degrees
- Motif clusters are big due to partitioning, quarks are smaller thanks to bottom-up dec.



(a) reciprocal

(b) out+

(c) acyclic

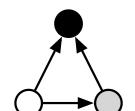
(d) cycle++

Food-web analysis

- Analysis with out^+
- Quarks give consistently better classifications than motif clustering

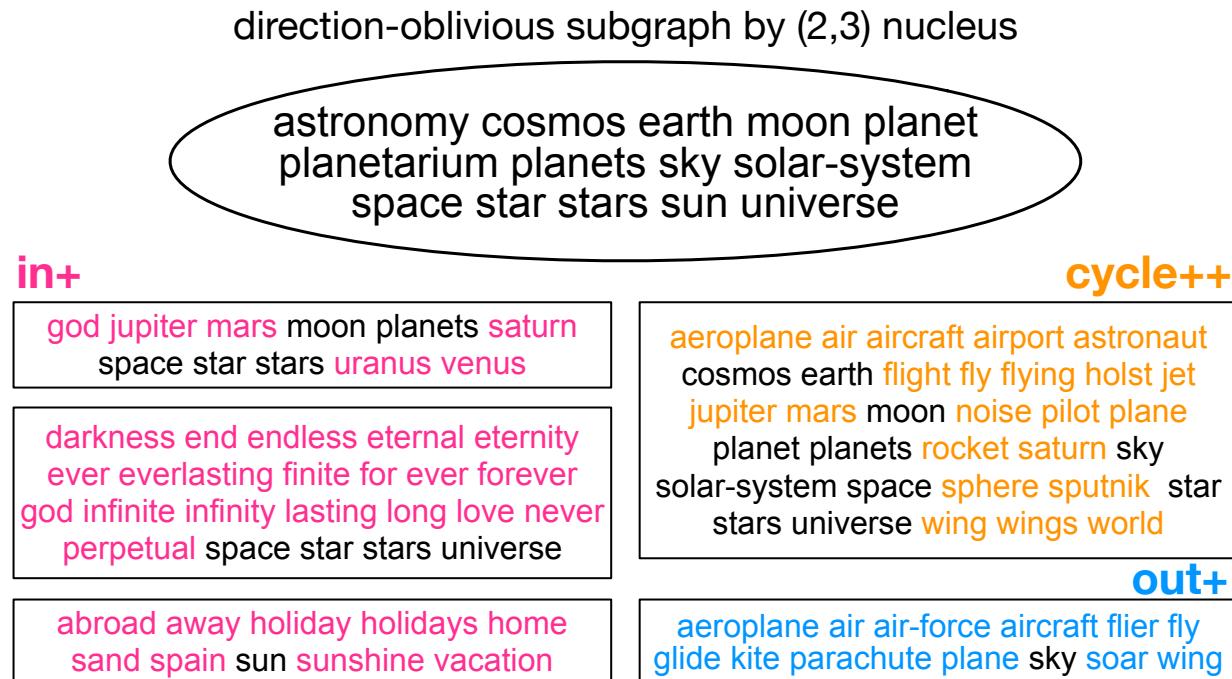
out^+	Metric	Quarks (7 subgraphs)	MC-K-MEANS w/ 4 clusters	MC-K-MEANS w/ 7 clusters
Class 1	ARI	0.3627	0.3005	0.1485
	F1	0.4869	0.4574	0.3794
	NMI	0.5415	0.5040	0.4843
	Purity	0.5968	0.5645	0.5161
Class 2	ARI	0.3816	0.3265	0.1871
	F1	0.5675	0.5380	0.4601
	NMI	0.5206	0.4822	0.4309
	Purity	0.6452	0.6129	0.5645

- Role-aware quark numbers find the preys, predators, and balancers with acyclic
 - Predators: Birds (ducks, herons, greeb)
 - Preys: Clown goby, herbivorous shrimps, zooplankton
 - Balancer: Fishes (anchovy, sardines, mojarra)



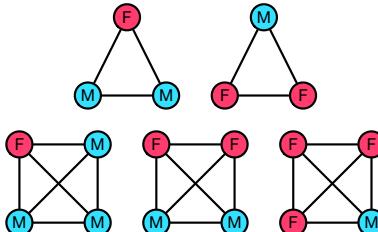
Word-associations

- Diverse subgraphs obtained with different motifs
 - Not possible when directions ignored



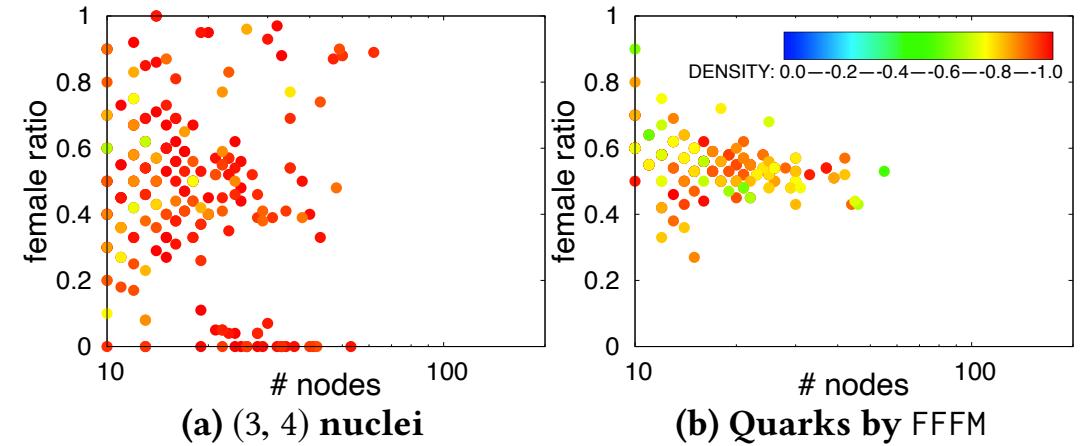
Finding gender-balanced subgraphs

- Facebook100 dataset with genders as node-labels
- How to find gender-balanced dense subgraphs even when the graph is imbalanced?
 - Compared to label-oblivious nucleus dec.
- M is edge, N is triangle
- M is triangle, N is four-clique



	$ V $	$ E $	$\frac{ V_f }{ V }$	edge, triangle		triangle, 4-clique				
	(2,3)n	Quarks	(3,4)n	Quarks	FMM	FFM	FMMM	FFMM	FFFF	
Mich67	3.7K	81.9K	25%	23.0%	45.0%	50.0%	24.5%	40.0%	45.0%	51.6%
Caltech36	769	16.7K	30%	39.4%	46.0%	52.0%	38.5%	43.1%	50.2%	52.8%
Carnegie49	6.6K	250.0K	37%	32.6%	49.0%	52.5%	38.5%	43.5%	49.5%	54.9%
MIT8	6.4K	251.3K	37%	38.8%	48.0%	52.1%	42.0%	44.3%	50.3%	53.9%
Stanford3	11.6K	568.3K	40%	46.8%	48.1%	49.0%	44.1%	45.4%	49.2%	55.4%
Cornell5	18.7K	790.8K	44%	44.3%	47.6%	51.8%	45.6%	43.7%	48.7%	54.9%
Penn94	41.6K	1.4M	44%	49.7%	48.4%	51.4%	52.1%	44.0%	49.8%	55.8%
UPenn7	14.9K	686.5K	44%	37.3%	48.8%	51.1%	46.4%	45.1%	50.4%	55.4%
Average of 18 networks:		40%	42.5%	48.2%	51.5%	44.1%	44.4%	49.7%	54.7%	

Female ratios



Density vs. female ratio for UPenn7

Conclusion & Future Work

- Principled approach for motif-driven dense subgraph discovery in directed and categorical-labeled networks
 - Successfully regularizes the motif degrees to quark numbers
- Role-aware variant considers the orbits and quantifies the roles systematically
- Versatile, efficient, and extendible
 - Code is available with detailed instructions for reproducibility!
- Hierarchy structure had limited success
 - Further analysis of hierarchy w.r.t a given motif
- Extension for networks with numerical node/edge labels
 - While incorporating the ordering

Paper, slides, talk, code: <http://sariyuce.com/WWW21>

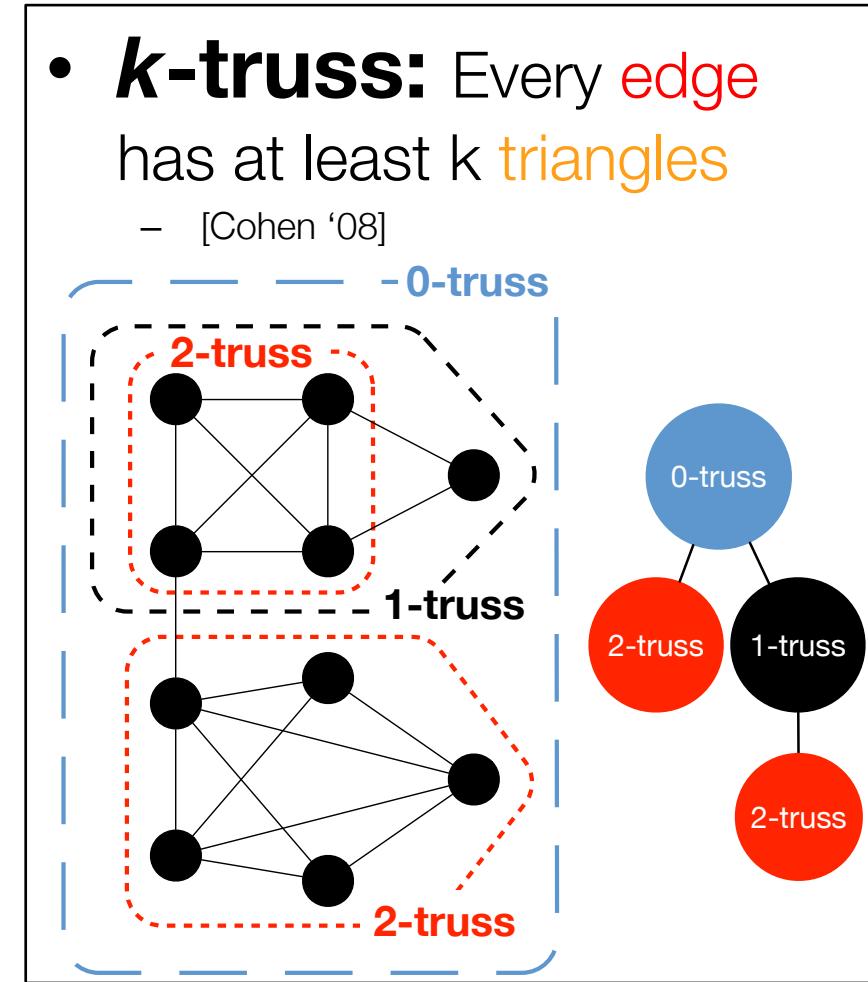
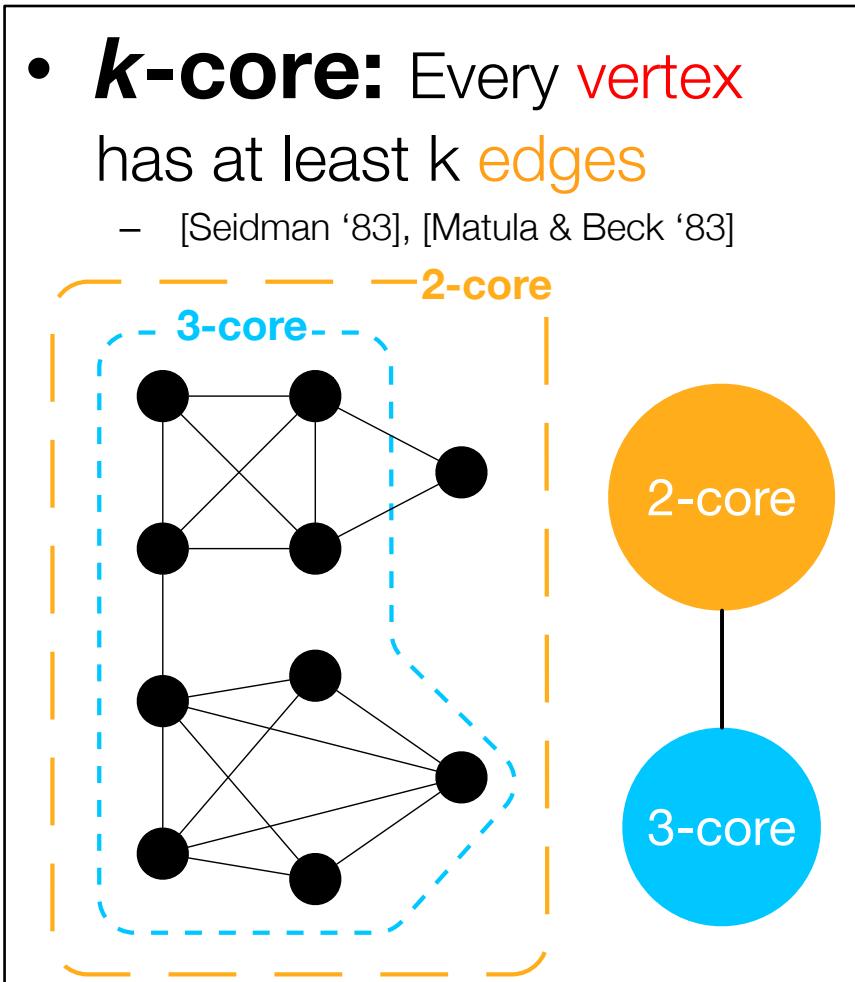
Questions: erdem@buffalo.edu

Thanks!



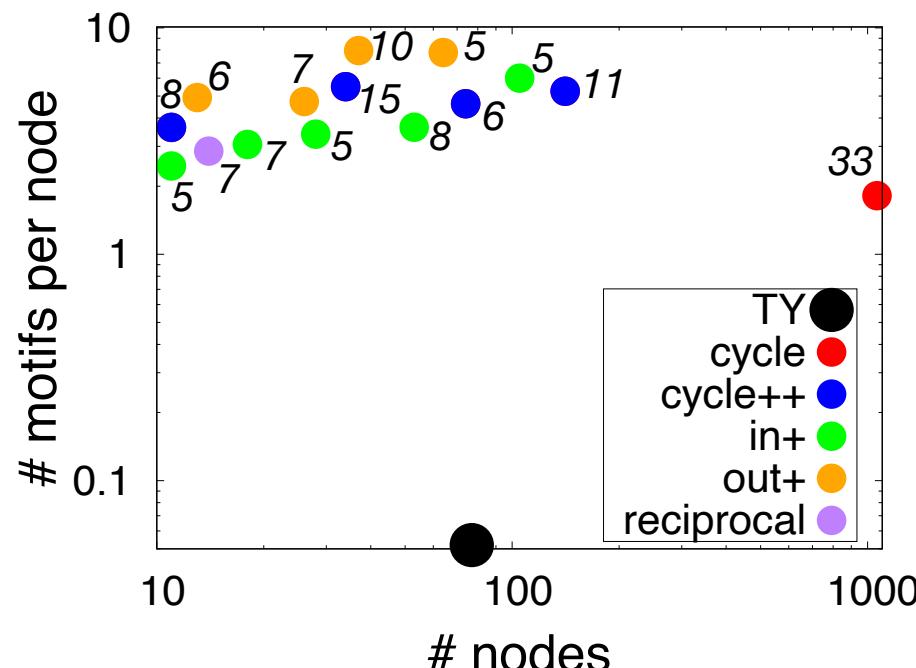
How to model dense subgraphs?

- Two effective models for simple, undirected networks
 - With hierarchical relations

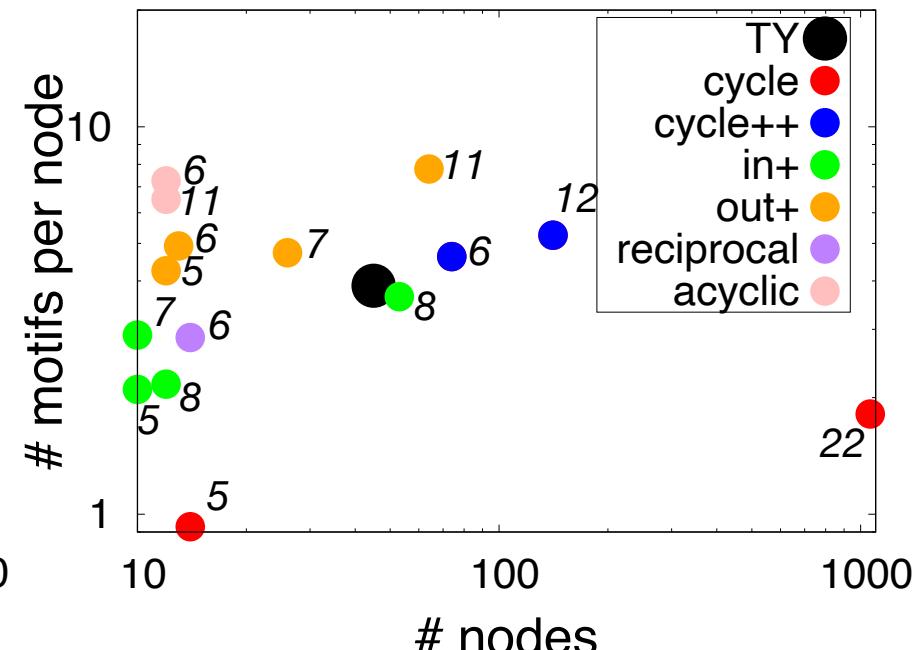


Quarks vs. Cycle- & Flow-truss

- Higher avg. motif degrees with quarks
 - Almost all the nodes in cycle- & flow-trusses are found, in various types
 - Considering each bidirectional edge atomically (instead of two unidirectional edges) highlights the diversity



(a) cycle-truss vs. quarks



(b) flow-truss vs. quarks

Runtime comparison with motif clustering

- Motif clustering with a single optimal cluster
 - Quark decomposition finds all the k -quarks
- Quark decomposition is mostly faster, for all motifs; up to 10x speedups
- Motif clustering is mostly faster for en-Wikipedia and wiki-Talk
 - Spectral clustering is heavy, cost increases when multiple clusters found

	<i>cycle</i>		acyclic		<i>out+</i>		<i>in+</i>		<i>cycle+</i>		<i>cycle++</i>	
	Q	M	Q	M	Q	M	Q	M	Q	M	Q	M
web-ND	0.34	3.31	4.26	16.8	0.62	6.3	2.11	8.54	0.53	10.01	0.78	9.86
amzn	0.74	3.54	3.29	79	2.25	132	1.92	105	1.18	5.29	3.23	107
wiki	28.9	14.0	112	18.2	10.9	16.4	21.1	17.7	20.5	20.2	47.8	16.8
soc-p	23.6	79	66.9	99	37.0	119	34.2	139	48.9	129	98.1	128
live-j	37.4	200	180	943	118	1135	126	1438	112	828	289	2248
en-w	900	501	7746	864	1511	799	1709	677	398	724	2223	677