

Plan for today:

- Ashvin presents: Study on centrality measures in social networks: a survey
- Talkin' proj
- Quick review (triadic and strong triadic closure)
- Vertex centrality
- Diffusion and epidemiology
- CODE MODE (SIR model+centrality)

Talkin' proj

- proposal presentation in a couple weeks
- order will be random
- grad students: also must give lit review as part of your proposal

IMPORTANT → must use or consider graph data and use some kind of "graph mining" technique

(Note: theory studies prob. OK, but

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Topic examples:

- * studying real-world network properties, especially with novel data
- * prediction → link prediction, metadata labels, edge direction
- * Classification → determining properties of verts/edges
- * clustering → classification method, finding communities
- * diffusion / simulation → how information "flows" or propagates through network topology
- * data creation → creating a novel graph dataset
- * visualization → enabling the visual study of graph data

visual study of graph data
and data + above GM stuff

* etc. → ask me

Centrality → vertex centrality

→ a measure of "importance"
or "influence" for a vertex
in a network

Note: usually defined topologically
locally or globally

who has high centrality:

Social networks: popular or
important people

Road/infrastructure: important
intersections or junctions

Epidemiology: someone with
a lot of contact with
others OR limited contact

others OK limited contact
but with important people

Types of centrality

Degree centrality:

Defined as the degree or # of
connections for a vertex

$$\rightarrow d(v) \text{ or } |N(v)|$$

For directed graphs:

$d^+(v)$, $d^-(v)$, or both
out degree in degree

Pros:

Easy to calculate

Gives good insight into local
spread of a diffusive process

Overall useful in various contexts

Cons:

Easy to falsify (spam, bot nets)

Local measurement $\rightarrow$ not useful

Local measurement $\rightarrow$ not useful
for diffusion over entire graph

Closeness centrality

Average shortest paths lengths

$\hookrightarrow$ How "close" a vertex is to
all other vertices

Diffusion: how quickly is the rest
of the network being impacted

Pros: More useful global measure
for diffusive processes

Captures a lot of the same
usefulness as degree centrality

Cons: Computationally expensive
to compute for *all* vertices
(quick for a small subset)

Can be a skewed measure
if attached to a hub

Betweenness Centrality

↳ ratio of shortest u, v -paths that contain some vertex w over all u, v -paths

→ The amount of diffusion that travels through a vertex

Pros: Global measure, very useful for understanding diffusive processes

Cons: Also pretty expensive (can approximate if needed)

Eigenvector centrality

Note: more to come Monday (Pagerank)

→ Basically: a vertex is important if its neighbors are

Consider adjacency matrix A

Solve for x : $Ax = \lambda x$

↑ ↑
eigenvector eigenvalue

Pros: Easier to calculate (power iteration)

Useful in ways that pure path-based centrality measures are not

Cons: Doesn't exactly/nicely correlate with diffusions

Difficult to intuitively infer

Directed graphs

Note: many of the above measures can apply directly to digraphs

↳ consider directed paths, degrees, or adjacency matrix

One specific measure:

Harmonic centrality

Harmonic centrality

↳ Reachability: How many can I reach? How many can reach me?
And how quickly for both?

Pros: easy to calculate
captures similar information
to degree/closeness centrality

Cons: doesn't differentiate many
vertices

Diffusive processes

Previously: we focused on strength
or importance of edges
(explicitly defined)

Now: we'll consider how vertex
centrality impact diffusion

HOW: Simulate an SIR model
and infect/remove vertices
based on centrality

Epidemiology

↳ defined as the study of
how diseases spread

For GM: How a disease spreads on
a network topology

Disease spreading: diffusive process

↳ we'll create an explicit model,
run it on a network, and
observe average results

Our model: SIR



susceptible

↓
not infected
but can be

infected

↓
can infect
others

removed

↓
can't be infected
can't infect others

Our approach:

- Initialize some populations as I, R
- Rest are in S
- p = probability of transmission per I, S interaction
- Iterate until $I = \emptyset$

Assumptions:

- p is fixed
- interactions are fixed
- duration of infection is fixed

What we're looking for:

How do our centrality measures correlate with diffusion dynamics

↳ Given initial I, R sets, what total # of people infected and

total # of people infected and
how long does the epidemic last?