Epidemics in Communication Networks *

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Small World Networks

- significant *clustering* or *locality* (neighbouring nodes have many common neighbours)
- small diameter or average distance between nodes

Examples

- Erdös numbers
- world wide web
- spread of contagious diseases
- neural network of flatworm
- electric power grids

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Probabilistic Model (Watts-Strogatz)

Consider each edge of a network and move one of its endpoints with probability p.

- $p = 0 \Rightarrow$ structured graphs
- $p = 1 \Rightarrow$ random graphs
- ▶ $p \approx .01 \Rightarrow$ small world networks

distance 20% of original graph clustering 95% of original graph

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Initial Graph for Probabilistic Model



Notation

- network is G = (V, E) of order n = |V|
- maximum degree is Δ
- diameter is D_G or D
- clustering is a measure of the connectedness of a graph (definition later)

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Circulant Graphs

- ► $C_{n,\Delta}$, Δ even, is the circulant graph $C(n; 1, 2, \cdots, \frac{\Delta}{2})$
- n nodes labelled with integers modulo n
- Δ links per node, each node *i* adjacent to $i \pm 1, i \pm 2, \cdots, i \pm \frac{\Delta}{2} \pmod{n}$.

•
$$C_{n,\Delta}$$
 has diameter $D_{C_{n,\Delta}} = \left\lceil \frac{n-1}{\Delta} \right\rceil$.

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Example



Circulant graph $C_{n,\Delta}$ with n = 24 and $\Delta = 6$

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Deterministic Model



Choose *h* equally spaced nodes of $C_{n,\Delta}$ to be *hubs*. Interconnect the hubs using a second graph *H* to obtain $C_{n,\Delta,h}$. Subgraph of circulant between two hubs is a *segment*.

Diameter of Deterministic Model

Theorem The diameter of $C_{n,\Delta}$ can be reduced to D using a hub graph H with approximately $h = \frac{2n}{\Delta(D-D_H)}$ nodes.

Theorem When a graph H with h nodes and diameter D_H is used to interconnect h hubs of $C_{n,\Delta}$, the diameter is approximately $\frac{2n}{\Delta h} + D_H$. ($\frac{n}{\Delta h}$ is the maximum distance from any node of $C_{n,\Delta}$ to the nearest hub.)

Examples Consider $C_{n,\Delta}$ with n = 1000, $\Delta = 10$, $D_{C_{n,\Delta}} = 100$.

- ► To reduce the diameter to 34 requires 7 hubs (first theorem).
- With h = 4 hubs, the diameter is $50 + D_H$ (second theorem).

Hub Graphs

- ► K_h (complete graph): diameter 1 but increases degree of hubs by h - 1
- K_{1,h-1} (star graph): diameter 2, adds minimum number of edges, one hub has high degree
- C(n; a, b) (double-step graph): circulant graph with each node i adjacent to i ± a, i ± b (mod n), diameter is
 D = [^{-1+√2n-1}/₂] when a = D and b = D + 1, increases degree of hubs by 4

Examples of Hub Graphs



Complete, star, and double-step hub graphs with h = 13

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Examples of Diameter with Hub Graphs

Consider $C_{n,\Delta}$ with n = 10,000, $\Delta = 10$, and $D_{C_{n,\Delta}} = 1000$. Using h = 50 hubs:

- K_{50} diameter 41, degree of hubs is 59
- ▶ *C*(50; 5, 6) − diameter 45, degree of hubs is 14

50 hubs is 0.5% of *n* but diameter decreases by factor of nearly 25 20 hubs are needed to decrease diameter by factor of 10

Clustering

For each node *i* of a graph *G*, let n_i be the number of neighbours of *i*. Let C_i be the fraction of the $\frac{n_i(n_i-1)}{2}$ possible edges among the neighbours of *i* that are present in *G*. The *clustering parameter* of *G*, C_G , is the average over all nodes *i* of C_i .

Theorem The clustering parameter of $C_{n,\Delta}$ is $C_{C_{n,\Delta}} = \frac{3(\Delta-2)}{4(\Delta-1)} \approx \frac{3}{4}.$

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Comparison of Probabilistic and Deterministic Models



Clustering and diameter for graphs based on $C_{1000,10}$

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Epidemics

- Flu epidemic spreads to many people simultaneously but individuals remain contagious for a short time
- Aids epidemic spreads to one person at a time but individuals remain contagious for a long time

How can we model the infectiousness of a disease and how long an individual is contagious?

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Broadcasting

Infectious diseases spread in a broadcast-like pattern.

- at time t = 0, an individual (originator) starts to spread the disease
- an individual (node) remains active (contagious) for A time units (rounds)
- ▶ an active node can infect $k \leq \Delta$ neighbours during each round
- ▶ an active node becomes permanently *inactive* after A rounds

Special Cases

- $A = \infty$ is normal broadcasting with a *k*-port model.
- A = 1 has been studied with probabilistic models.
- We will study A = 1 with a deterministic model.

Definitions

- $\mathcal{T}_{k,\mathcal{A}}(u)$: broadcast time for originating node u when
 - ► a node can infect k ≤ ∆ of its neighbours during each active time step
 - a node remains *active* for A rounds after infection

The broadcast time of graph G is $\mathcal{T}_{k,\mathcal{A}}(G) = \max\{\mathcal{T}_{k,\mathcal{A}}(u)|u \in V(G)\}.$ Simplify to $\mathcal{T}_k(u)$ and $\mathcal{T}_k(G)$ since only A = 1 is considered.

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Broadcast Time in $C_{n,\Delta}$

Lower Bound Diameter of $C_{n,\Delta}$, Δ even, is $\left\lceil \frac{n-1}{\Delta} \right\rceil$.

Theorem For $n \ge 2\Delta(\lceil \log_k \rceil - 1) + 2$

•
$$T_k(C_{n,\Delta}) = n-1$$
 if $k = 1$

•
$$\mathcal{T}_k(\mathcal{C}_{n,\Delta}) = \left\lceil \frac{n-1}{\Delta} \right\rceil$$
 if $2 \le k \le \Delta$

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Sections of a Circulant



Proof Outline

- ▶ k = 1: one node active during each round $\implies n 1$ rounds
- k = Δ: spreading pattern is flooding, one section on each side of the originator per round
- $\frac{\Delta}{2} < k < \Delta$:
 - Round 1: originator infects farthest $\frac{k}{2}$ neighbours on each side
 - Round 2: infect remaining nodes in section on each side of originator and more than half of nodes in next section on each side
 - continue until round $\left\lceil \frac{n-1}{\Delta} \right\rceil$
- 2 ≤ k ≤ Δ/2: infect nodes in section on each side of originator in ⌈log_k Δ⌉ rounds and then flood for ⌈n-1/Δ⌉ − 1 more rounds to infect remaining sections giving T_k(C_{n,Δ}) ≤ ⌈log_k Δ⌉ + ⌈n-1/Δ⌉ − 1

Example for $\frac{\Delta}{2} < k < \Delta$



Optimal broadcast in $C_{n,\Delta}$ for n = 24, $\Delta = 6$, k = 4

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Example for $2 \le k \le \frac{\Delta}{2}$



Broadcast in $C_{n,\Delta}$ for n = 24, $\Delta = 6$, k = 2, $\mathcal{A} = 1$,

Proof Outline for $2 \le k < \Delta$

- Round 1: originator infects farthest ^k/₂ neighbours in section on each side
- Round 2: infect the farthest $\frac{k^2}{2}$ nodes in next section on each side
- Continue to round $\alpha 1$ where $\alpha = \lceil \log_k \Delta \rceil$
- ► Round \(\alpha\): infect all nodes in section \(\alpha\) and some nodes in section \(\alpha\) 2
- ► Round α + 1: infect remaining nodes in section α − 1, all nodes in section α + 1, and some nodes in section α − 3
- ► Continue flooding in direction away from originator and filling in toward originator until round $2\alpha 1$

Segments of $C_{n,\Delta,h}$



Broadcasting in $C_{n,\Delta,h}$

Theorem $T_k(C_{n,\Delta,h}) \approx \frac{2n}{h\Delta} + D_H$ where

- H = C(h; a, b) (double-step graph)
- ► $3 \le k \le \frac{\Delta}{2}$
- $\Delta \geq$ 4, Δ even

Proof Outline

- ► First phase: originator sends message to nearest hub using chords of length ^A/₂ (approximately ⁿ/_h rounds worst case)
- ► Second phase: the informed hub broadcasts in the hub graph $H(D_H = \left\lceil \frac{-1 + \sqrt{2h-1}}{2} \right\rceil$ rounds)
- Third phase: each hub broadcasts to the two segments on either side of it (approximately ⁿ/_{bΔ} rounds)

Theorem $T_k(C_{n,\Delta,h}) \approx \frac{3n}{h\Delta} + D_H$, k = 2

Broadcast in Tile



Broadcast in a tile with k = 2

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Broadcast in Modified Tile



Modified broadcast in a tile with k = 2

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Stopping the Infection

Theorem If no communications are lost, then all nodes of $C_{n,\Delta}$ will be infected for any $2 \le k \le \Delta$.

Proof Outline

- ► A/2 < k ≤ Δ: the infection spreads at the maximum possible rate (flooding) and it is impossible to leave uninfected gaps.</p>
- ► $2 \le k \le \frac{\Delta}{2}$:
 - permanently uninfected nodes must be permanently unreachable from all active nodes
 - try to construct two blocks of inactive (previously infected) nodes with isolated uninfected nodes between the blocks
 - ▶ blocks must contain at least A/2 nodes to prevent an active node from infecting an isolated node
 - two possible strategies (both fail)

Protection near Originator



Attempt to isolate uninfected nodes close to originator

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Protection far from Originator



Attempts to isolate uninfected nodes far from originator

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