Systems Biology: A Personal View V. Networks: Models I

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Theoretical understanding of networks

- Regular lattice or grid (Physics)
 - average path length $\sim N$ (no. of nodes)
 - clustering high
 - delta function distribution of degree (links/node)
- •Random networks (*Graph theory*) Also known as Erdos-Renyi networks
 - average path length $\sim \log N$
 - clustering low
 - Poisson distribution of degree

Empirical networks are not random – many have certain structural patterns





Example: *small-world* networks



Increasing Randomness

p: fraction of random, long-range connections

Watts and Strogatz (1998): Many biological, technological and social networks have connection topologies that lie between the two extremes of completely regular and completely random.

"It's a small world": The Milgram Experiment



The chains progress from the starting position (Omaha) to the target area (Boston) with each remove. Diagram shows the number of miles from the larget area, with the distance of each remove averaged over completed and uncompleted chains. Stanley Milgram (1933-1984), US social psychologist

Arbitrarily selected individuals in Nebraska were asked to generate acquaintance chains (knowing on first name basis) connecting them to a target individual in Boston

In one experiment, 64 of the 296 chains initiated eventually reached the target – the mean number of intermediaries between source and target being slightly larger than 5

\Rightarrow Six degrees of separation



"Small world": Local properties of regular networks but global properties of random networks



yet have small characteristic path lengths (as in random networks).

Epidemics on "Small world"

Dynamical process:

- Time *t* = 0: single infected individual present.
- Each infected agent can infect any of its neighbours with probability r.
- Infected individuals removed (by immunity or death) after unit period of sickness.

Key Results:

- Critical infectiousness r_{half}, at which the disease infects half the population, decreases with p
- Time required for a maximally infectious disease (r = 1) to spread throughout the entire population T(p) has same form as characteristic path length L(p)
- \Rightarrow rewiring only a few links in the original lattice causes global infection to occur almost as fast as in random network

Implication:

"Control the truck-drivers"



Do small-world networks occur in real life?

| | # nodes | Avg degre | e Avgp | Avg path length | | Clustering coefficien | |
|--------------------------|-----------|--------------------|----------|-----------------|----------|-----------------------|--|
| Network | Size | $\langle k angle$ | l | l rand | С | Crand | |
| WWW, site level, undir. | 153 127 | 35.21 | 3.1 | 3.35 | 0.1078 | 0.00023 | |
| Internet, domain level | 3015-6209 | 3.52-4.11 | 3.7-3.76 | 6.36-6.18 | 0.18-0.3 | 0.001 | |
| Movie actors | 225 226 | 61 | 3.65 | 2.99 | 0.79 | 0.00027 | |
| LANL co-authorship | 52 909 | 9.7 | 5.9 | 4.79 | 0.43 | 1.8×10^{-4} | |
| MEDLINE co-authorship | 1 520 251 | 18.1 | 4.6 | 4.91 | 0.066 | 1.1×10^{-5} | |
| SPIRES co-authorship | 56 627 | 173 | 4.0 | 2.12 | 0.726 | 0.003 | |
| NCSTRL co-authorship | 11 994 | 3.59 | 9.7 | 7.34 | 0.496 | 3×10^{-4} | |
| Math. co-authorship | 70 975 | 3.9 | 9.5 | 8.2 | 0.59 | 5.4×10^{-5} | |
| Neurosci. co-authorship | 209 293 | 11.5 | 6 | 5.01 | 0.76 | 5.5×10^{-5} | |
| E. coli, substrate graph | 282 | 7.35 | 2.9 | 3.04 | 0.32 | 0.026 | |
| E. coli, reaction graph | 315 | 28.3 | 2.62 | 1.98 | 0.59 | 0.09 | |
| Ythan estuary food web | 134 | 8.7 | 2.43 | 2.26 | 0.22 | 0.06 | |
| Silwood Park food web | 154 | 4.75 | 3.40 | 3.23 | 0.15 | 0.03 | |
| Words, co-occurrence | 460.902 | 70.13 | 2.67 | 3.03 | 0.437 | 0.0001 | |
| Words, synonyms | 22 311 | 13.48 | 4.5 | 3.84 | 0.7 | 0.0006 | |
| Power grid | 4941 | 2.67 | 18.7 | 12.4 | 0.08 | 0.005 | |
| C. Elegans | 282 | 14 | 2.65 | 2.25 | 0.28 | 0.05 | |

Albert & Barabasi, 2003