





SMR.1656 - 32

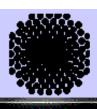
School and Workshop on Structure and Function of Complex Networks

16 - 28 May 2005

Model of Mobile Agents for Contact Networks

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These are preliminary lecture notes, intended only for distribution to participants



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Model of mobile agents for contact networks

Marta C. González

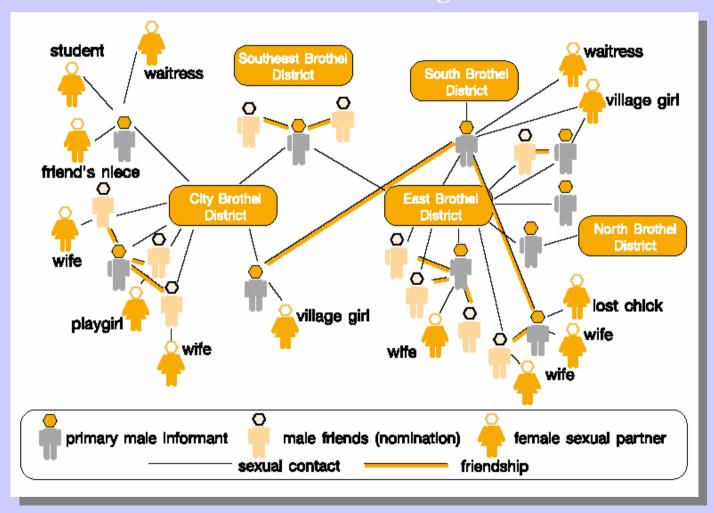
Pedro G. Lind

Hans J. Herrmann

Social Networks



Social and Sexual Network of Young Northern Thai Men

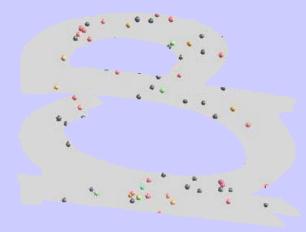


Source: Catherine C. Bond, "Building Connections".

PATH: Program for Appropriate Technology in Health

Me Model





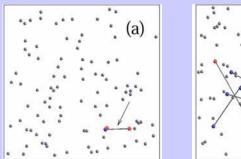
RGB scale for number of partners

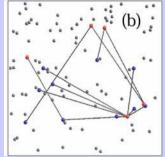
Parameters of the ED simulation

$$r=1$$
 $\rho = N/L^2 << 1$

- Non-connected
- Initial Conditions

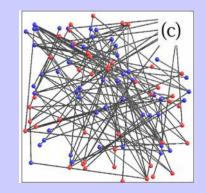
$$\begin{vmatrix} \overrightarrow{v_0} \end{vmatrix} = 1$$
 $N_{ntw} = 1$





Collision rule for "sexual interactions"

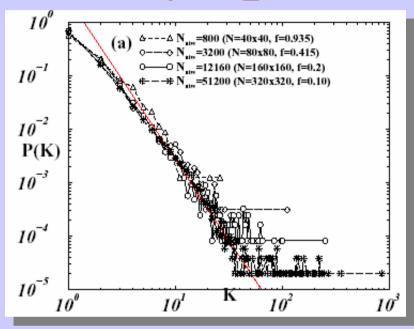
$$\overrightarrow{v}(k_i) = \left(k_i^{\alpha} + \left|\overrightarrow{v}_0(i)\right|\right) \overrightarrow{\omega}$$



- (ii) Between two connected agents with probability (1-5)
 - 5 is called "Selectivity"
 - S=1, only type (i) collisions are allowed

Mocity update rule

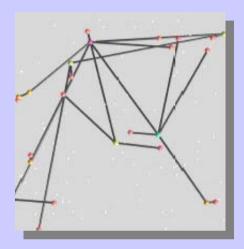




$$v_1 = \underline{r}u_1 + \underline{q}u_2$$

$$p = -(1+f)$$

$$q = f$$
 $f \ll 1$

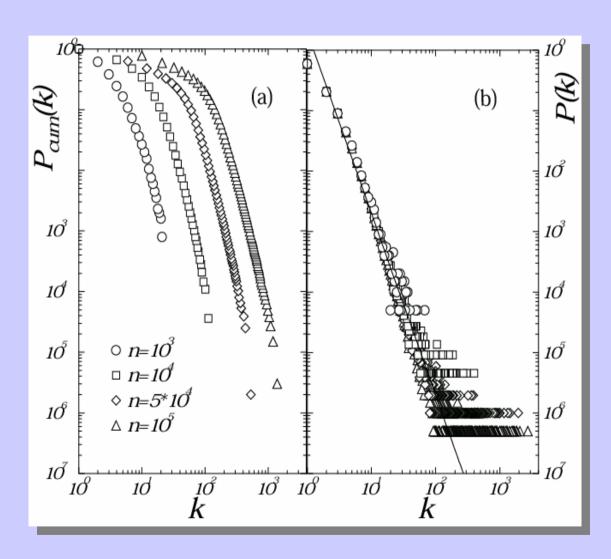


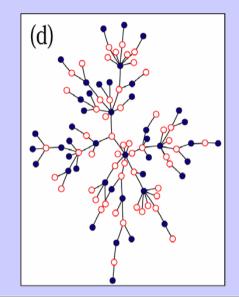
$$\overrightarrow{v}(k_i) = \left(k_i^{\alpha} + \left|\overrightarrow{v}_0(i)\right|\right) \overrightarrow{\omega}$$

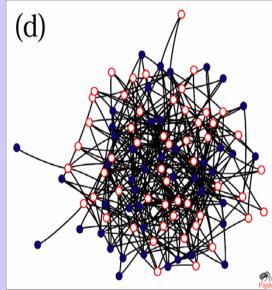
$$\vec{u}_1$$
 \vec{u}_2



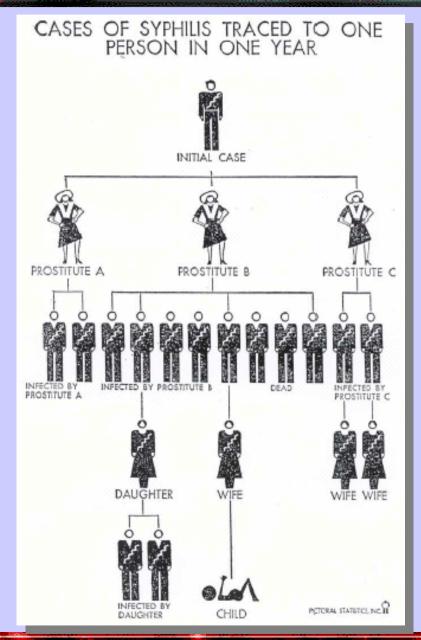


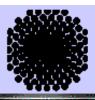






Saxually Transmitted Diseases P

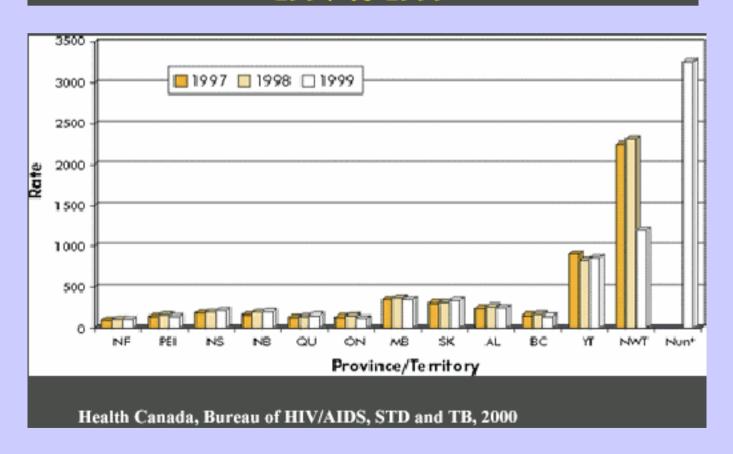




Sexually



Reported Female Genital Chlamydia Rates per 100,000 in Canada by Province/Territory, 1997 to 1999





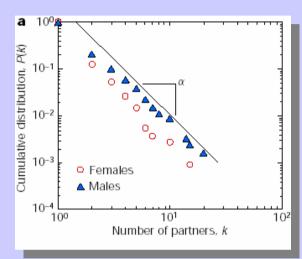


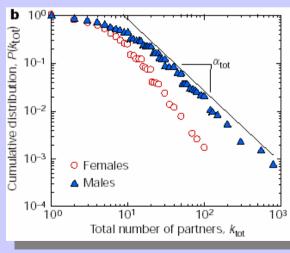
"The Web of Human Sexual Contacts"

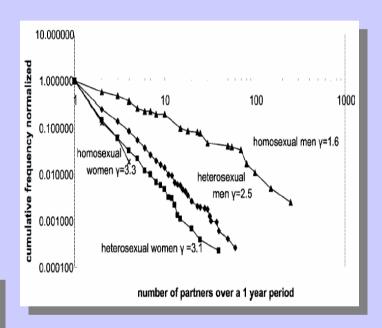
F. Liljeros, C. R. Edling, L. A. N. Amaral, H. E Stanley, and Y. Aberg, Nature 411, 907-908 (2001).



Painting by Idahlia Stanley







A.SCHNEEBERGER, et. al. *Sex. Trans. Dis.*, Vol. 31, No. 6, p.380–387 (2004)

Contact Tracing



Patterns of Chlamydia and Gonorrhea Infection in Sexual Networks in Manitoba, Canada

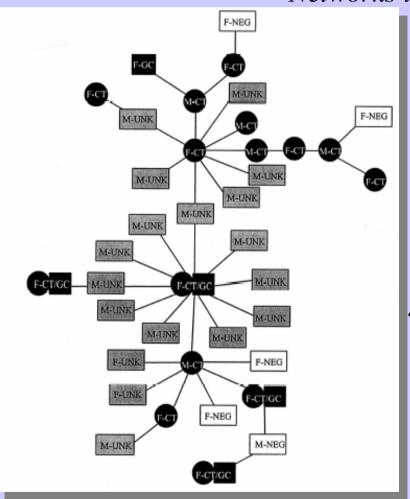


TABLE 1.	Summary	of the N	Number (of Comp	onents	of	Size n
Identified \	Within the	Manitob:	a Sexual	Notwor	k*		

No. of individuals	No. of components	No. of individuals	No. of components	
2	911	12	1	
3	366	13	2	
4	116	14	6	
5	39	16	1	
6	20	17	1	
7	15	18	1	
8	8	19	1	
9	5	39	1	
10	2	41	1	
11	5	82	1	

"Number of cases and contacts in database was 4544; 1503 components with 2 or more people were identified.

4544 Persons (Males:Females 1.27) 1503 components (clusters)

Legend

F: females M:males

■ Positive results

Negative results

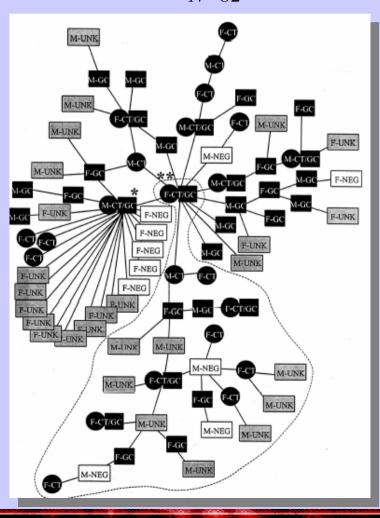
Unknown results

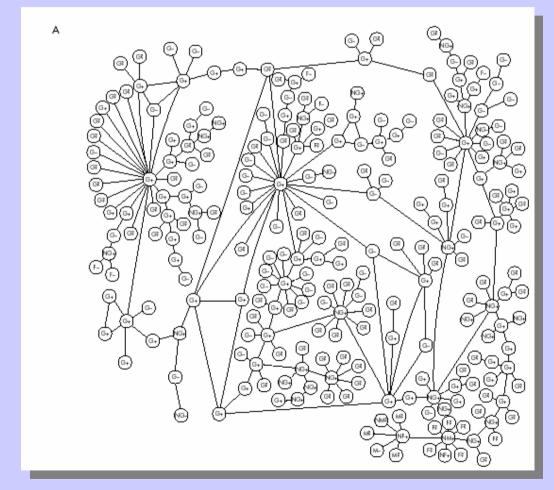
J. L.WYLIE and A.J JOLLY, Sex. Trans. Dis., 28, 14-24 (2001)

Empirical Data



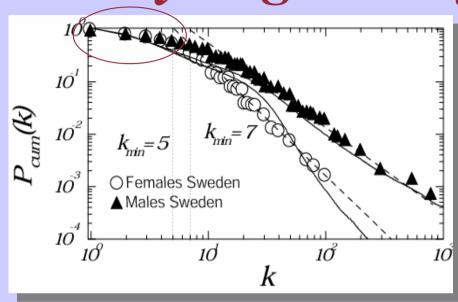
Largest Connected Component Chlamydia and Gonorrhea Infection in Sexual Networks in Manitoba, Canada N=82 Largest connected Component in the early epidemic phase of HIV transmission in Colorado Springs, USA N=250

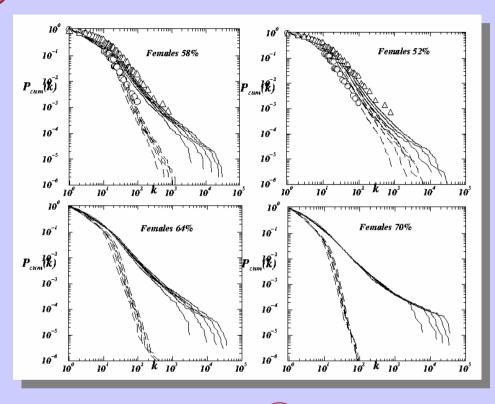




Analyzing Surveys







Discrepancy in empirical Data

$$\gamma_{mal} = 2.6 \pm 0.3$$

$$\gamma_{fem} = 3.1 \pm 0.3$$

$$\vec{v}(k_i) = \left(k_i^{\alpha} + \begin{vmatrix} \vec{v}_0(i) \end{vmatrix}\right) \vec{\omega}$$

$$\left[\frac{\gamma_{fem} - 1}{\gamma_{fem} - 2}\right] K_{\min}^{fem} = \left[\frac{\gamma_{mal} - 1}{\gamma_{mal} - 2}\right] K_{\min}^{mal}$$

$$\alpha = 1.2$$
 $n = 0.2N$

alyzing Contact Tracing



Networks

a20

Q15

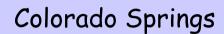
Q10C(K) aa5 $\alpha \infty$

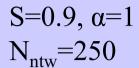
K

10

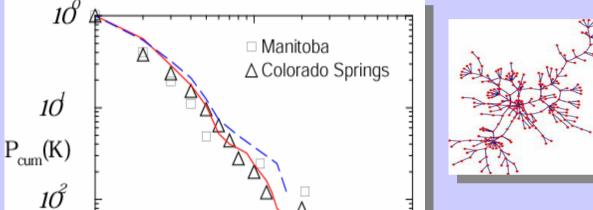
10³

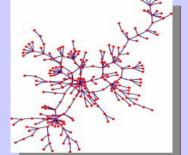
10⁴

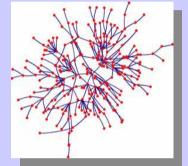


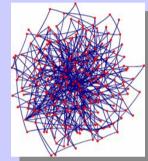




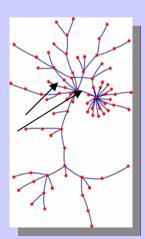








Manitoba

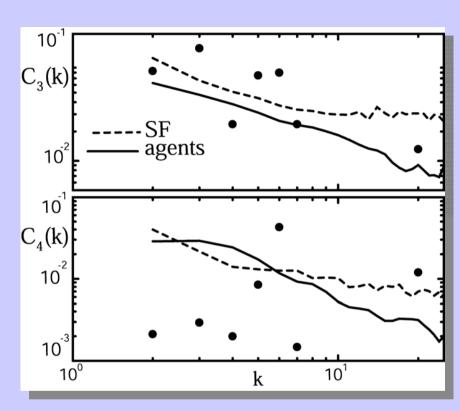


S=0.98,
$$\alpha$$
=1 N=81



Counting Cycles





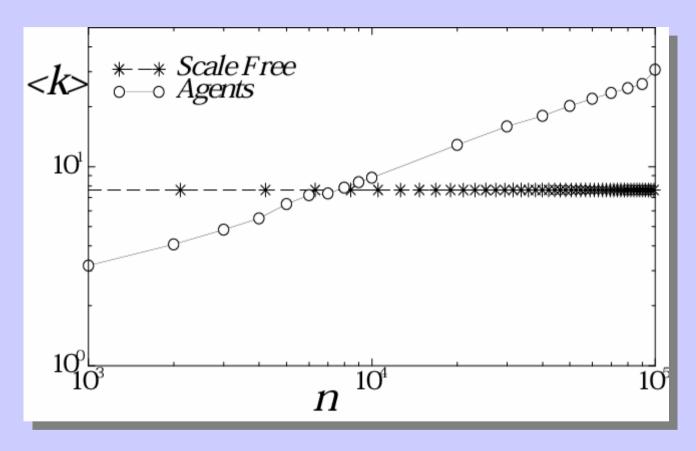
 $C_4(k)$ $C_4(k)$ C

	N	L	T	Q	$\langle C_3 \rangle$	$\langle C_4 \rangle$
Heterosexual	82	84	0	2	0	0.00486
(Fig. 2a)						
Homosexual	250	266	11	6	0.02980	0.00192
(Fig. 2b)						
Heterosexual	82	83.63	0	1.45	0	0.01273
(Agent Model)						
Homosexual	250	287.03	8.23	10.52	0.02302	0.01224
(Agent Model)						
Heterosexual	82	162	0	159.72	0	0.12859
(Scale-free)						
Homosexual	250	498	45.28	256.79	0.08170	0.02787
(Scale-free)						

P.G Lind, MCG, H.J. Herrmann cond-mat/0504241







In order to observe the behavior of the system into a quasi-stationary state we introduce birth and death of agents

Orași-stationary state



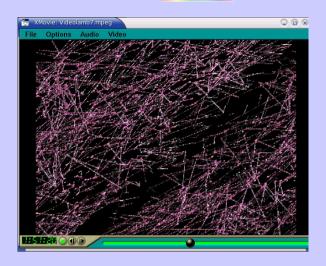
Labeling of Cluster sizes

isolated O

RGB scale for cluster size







 $T_1 = 18.75$

 $T_1 = 30.75$

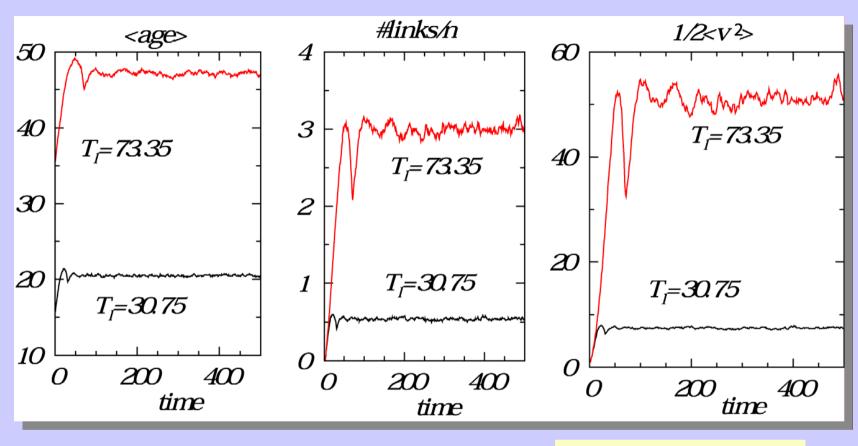
T'=73.35

Rules for generating the network

- Each time one agent collides with other a bond between the two is created.
- Each agent has an "age" distributed uniformly between 0 and a time of life (T_I). The "age" of each agent is incremented each time step until It reaches T_I when it dies (it is cut from the network). When one agent dies other enters into the system with random age.

Omasi-stationary state





Characteristic times

$$\tau_{\mathit{MFP}} = \frac{1}{\rho 2r_{0}} \sqrt{\frac{m}{2\pi < v >}} \qquad \tau_{\mathit{eff}} = T_{l} - < age > \qquad \lambda(T_{l}) \equiv \frac{T_{l} - < age >}{\tau_{\mathit{MFP}}}$$

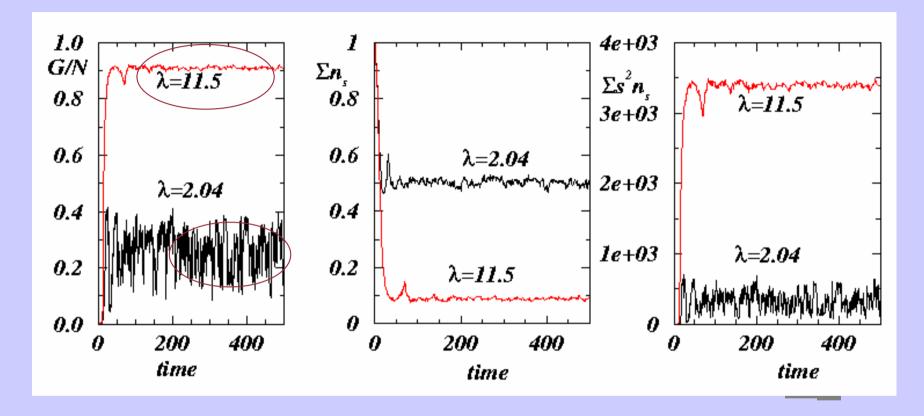
$$\tau_{eff} = T_l - \langle age \rangle$$

Control parameter

$$\lambda(T_l) \equiv \frac{T_l - \langle age \rangle}{\tau_{MFP}}$$

Chister size distribution

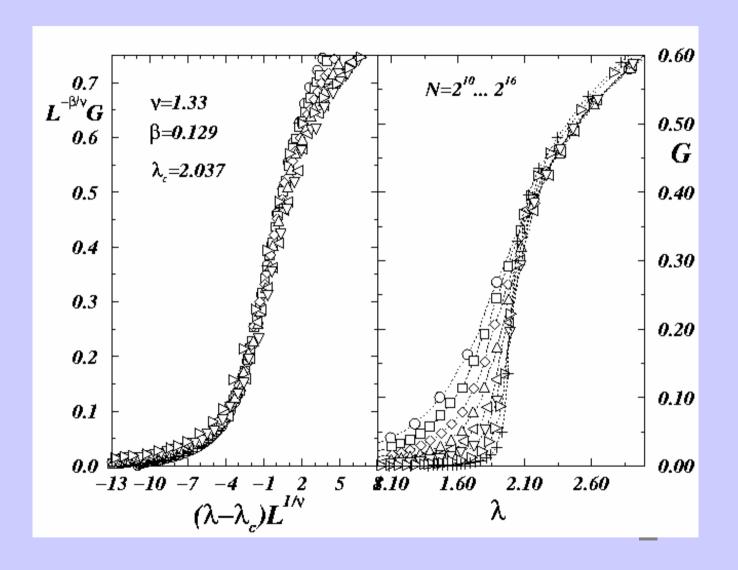




- ullet Number of clusters of size s: N_s
- $lue{}$ Total number of agents: N

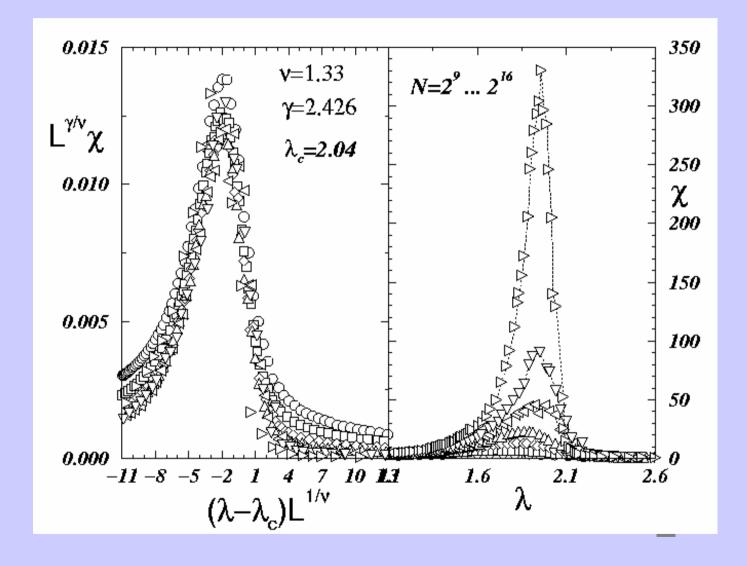
- ullet Fraction of agents at the largest cluster: G
- **Q**Clusters of size distribution: $n_s = \frac{N_s}{N}$

Saling of the largest cluster CP



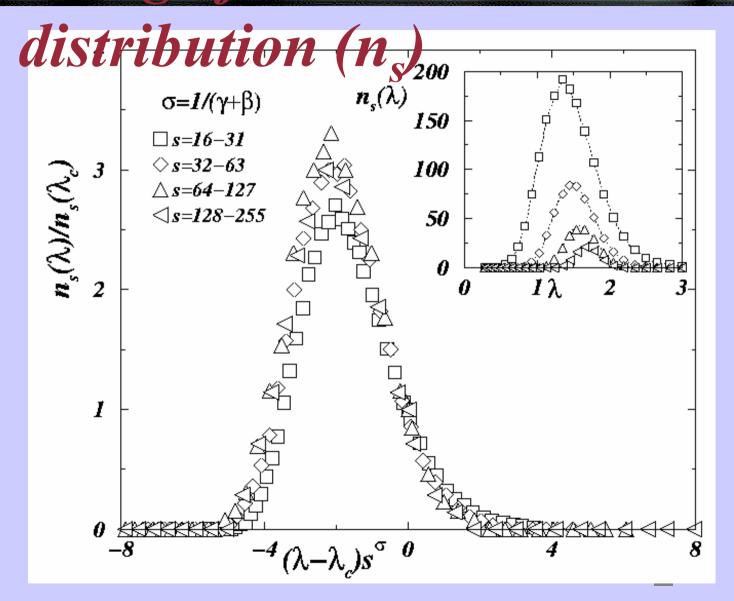
Scaling of the second moment of

ns



Scaling of the cluster size







Critical Exponents



	Mean Filed	SW*	Moving Agents	Percolation 2D
ν	0.5	0.59 ± 0.05	1.3 ± 0.1	1.33333
γ	1	1.3 ± 0.1	2.4 ± 0.1	2.388888
β	1		0.13 ± 0.01	0.138888
σ	0.5		0.40 ± 0.01	0.3956

$$\sigma=1/(\beta+\gamma)$$

$$Dv=2\beta+\gamma$$

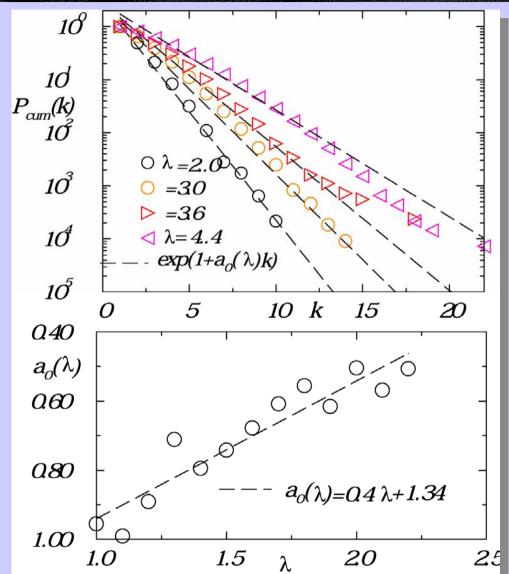
(D=2.01)

(*) Newman and Watts, PRE, Vol. 60, 7332, (SW, $k=1 \phi=0.1$)



Characteristics of the



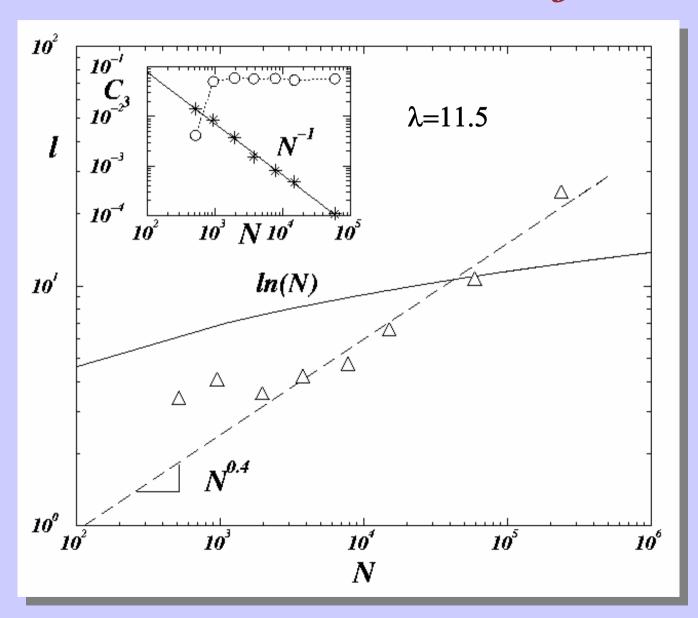


★The degree distribution is an exponential characterized by λ



Characteristics of the



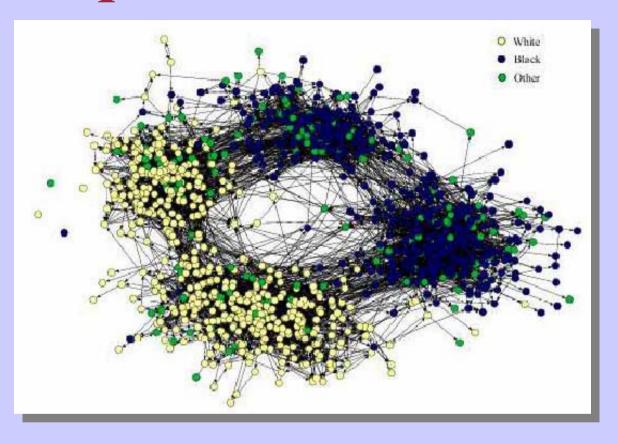


★C₃ vs. λ
and *l* vs. λ
remain to be
studied



Comparison with





Add Health data base. Picture by James Moody

Friendship networks from 84 US schools.

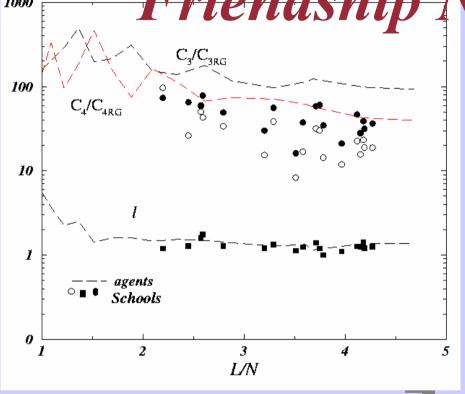
Sizes of the networks (50-2000)

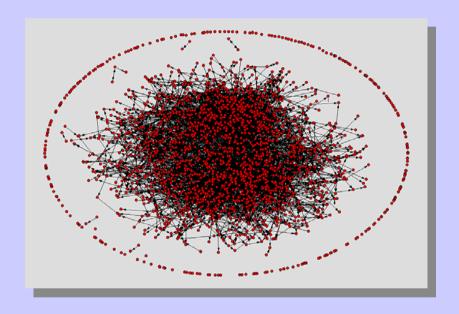


Comparison with









n=2250

Giant=1954

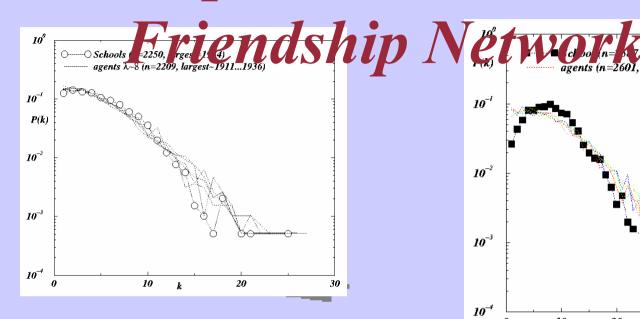
L/N=4.9

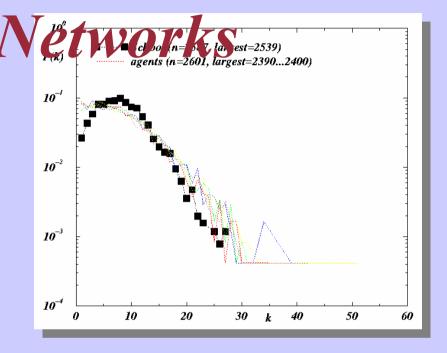
Here e analyze the Giant cluster only

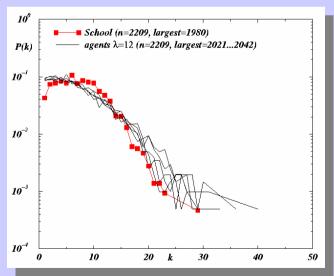


Comparison with







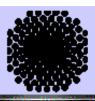


A detailed study of the topological characteristics of the empirical networks is in preparation.

Conclusions



- ✓ We present a model of contact network based on a system of mobile agents.
- ✓ Our model suggests that the growth mechanism of connected components of sexual networks is not pure scale-free, due to interaction among internal agents. This implies an increase on the mean number of partners with time. This differences should affect the results on spreading of infections.
- ✓ In a quasistationary version of the model the resulting contact network is exponential and can be used to model empirical data of friendship networks.











































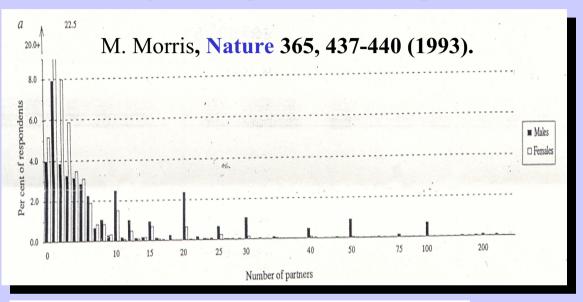








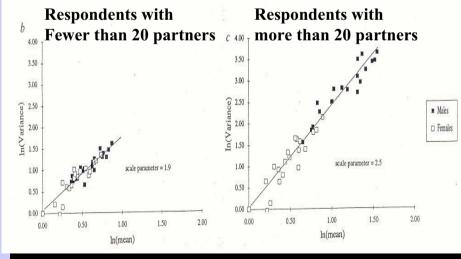
"Telling tails explain the discrepancy in sexual partners reports"



More than 60% of all contacts reported by men are unaccounted for!

Sample bias?

- 160% men with women under 18
- ♠ Each man average of 15 CSW
- ↑ 0.5-1.0% of Female population CSW Reporting bias?
- All men over reporting 65%
- All women under reporting by 200%



Important Result: The explanations for the male:famale discrepancy should be focused on the upper tail of the contact distribution