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Traceroute-like exploration of unknown networks: a statistical analysis

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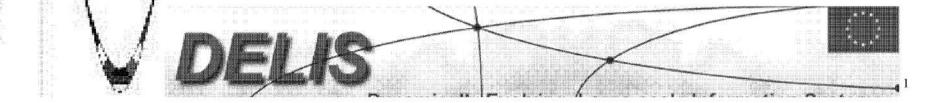
These are preliminary lecture notes, intended only for distribution to participants

Traceroute-like exploration of unknown networks: a statistical analysis

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cond-mat/0406404 to appear in LNCS; cs.NI/0412007 to appear in TCS; Phys. Rev E 71 (2005)





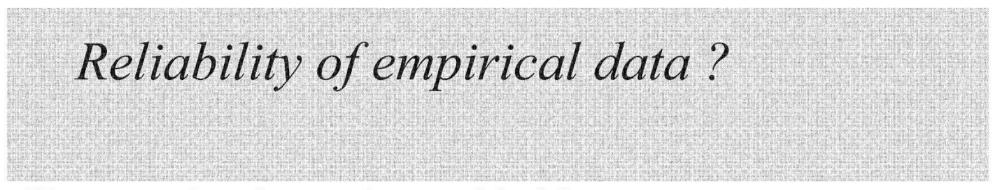
- Context: sampling of complex networks
- Model for traceroute-like sampling
- Theoretical approach
- Numerical results
- Conclusions

Main characteristics of complex networks

- Small-world networks
- Heterogeneous networks: broad degree distributions
- Dynamical evolution, self-organisation

...In contrast with usual random graphs

....development of new paradigms (evolving networks, etc...)



Heterogeneity of networks: empirical fact

- social networks: various samplings/networks with similar results
- transportation network: reliable data
- biological networks: incomplete samplings
 - Internet. various mapping processes



Statistical analysis of the sampling process

Internet representation

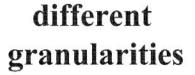
•Multi-probe reconstruction (router-level)

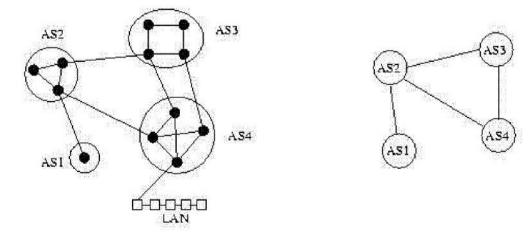
•Use of BGP tables for the Autonomous System level (domains)

Many projects (CAIDA, NLANR, RIPE, IPM, PingER...)

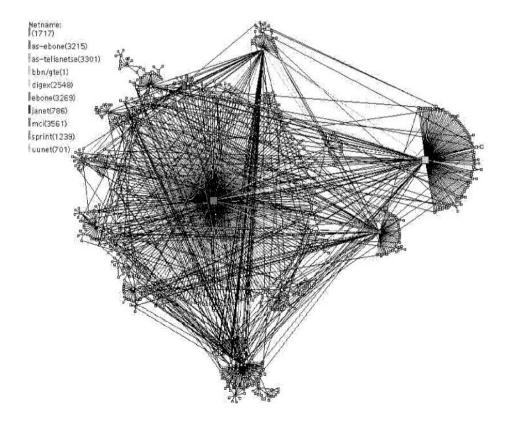


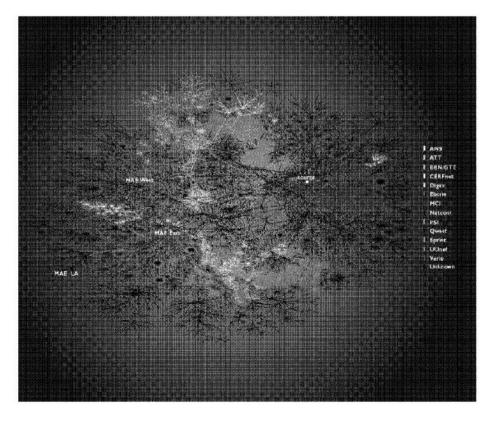
Graph representation





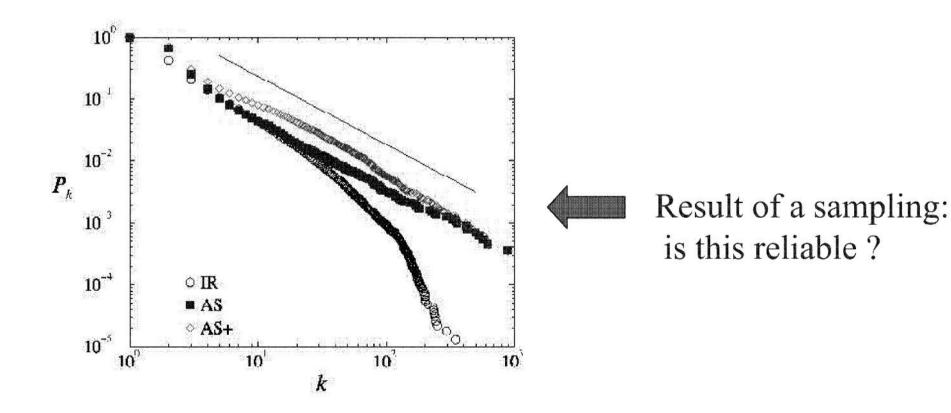
Large-scale visualizations

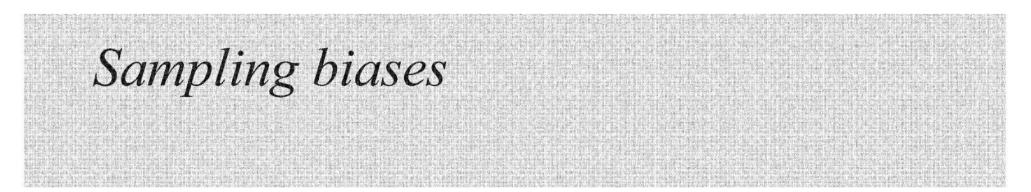




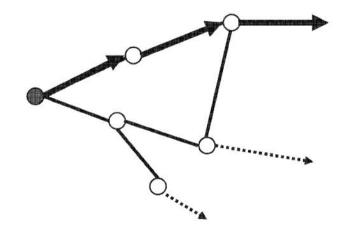
Topological analysis

Broad connectivity distributions: obtained from mapping projects



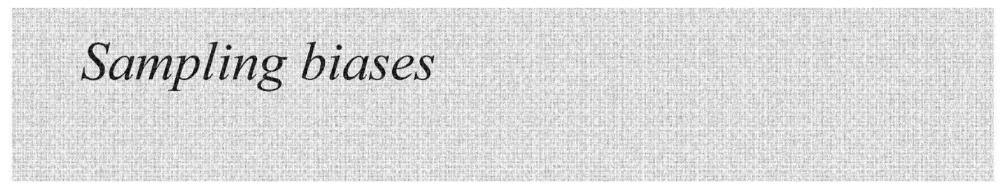


Internet mapping: traceroute



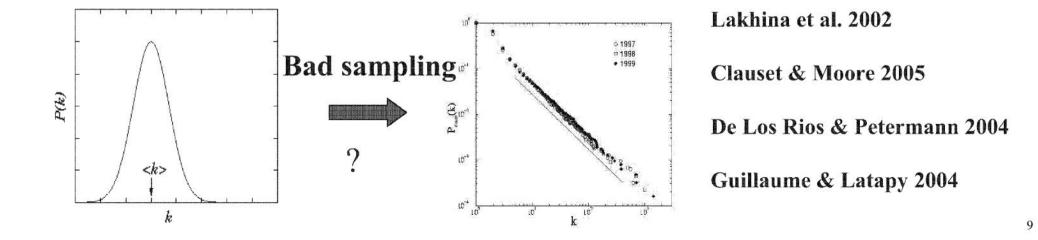
=> spanning tree

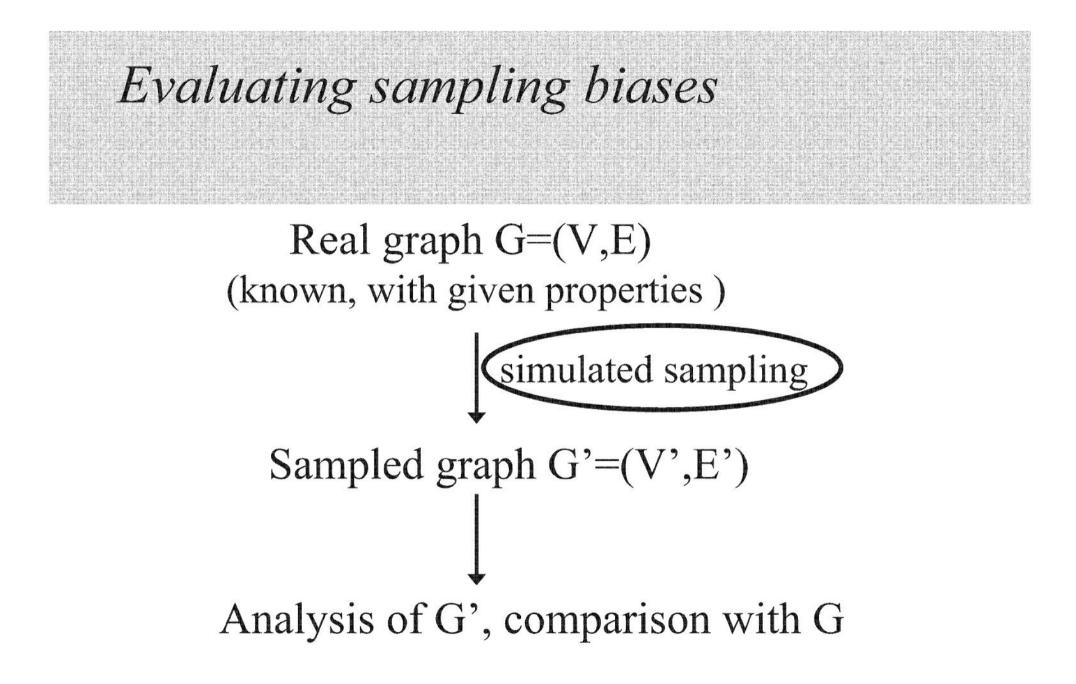
Sampling is incomplete Lateral connectivity is missed (edges are underestimated) Finite size sample



- Vertices and edges best sampled in the proximity of sources
- Bad estimation of some topological properties

Statistical properties of the sampled graph may sharply differ from the original one



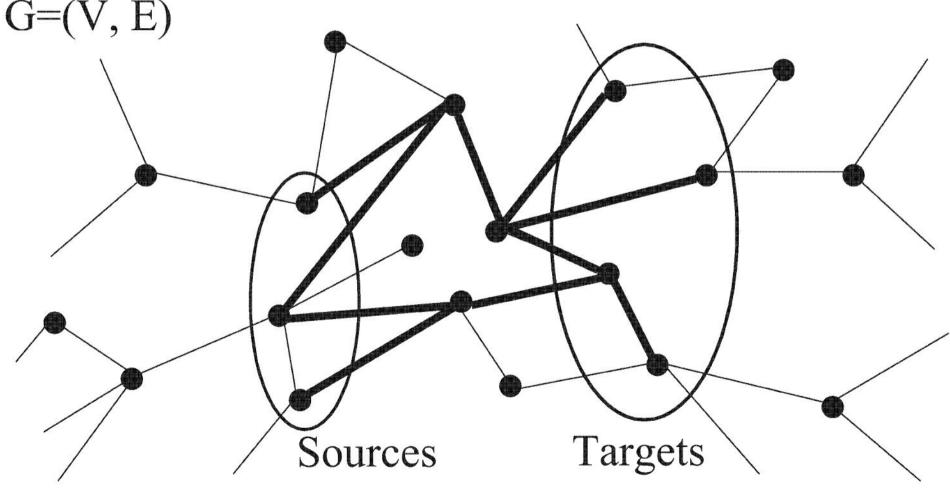


What we have done

- I. model for traceroute sampling
- II. analytical analysis with approximations
 => link between topological properties of the sampled network and the sampling biases
- III. numerical analysis on various networks with different topologies

Model for traceroute

First approximation: union of shortest paths

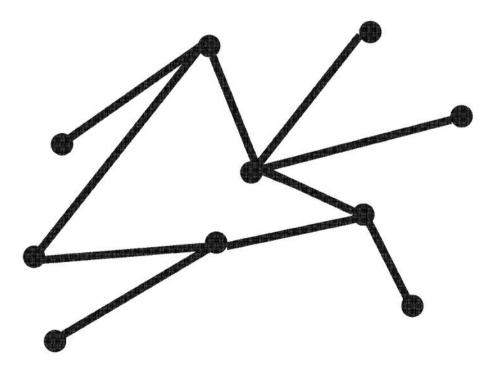


NB: Unique Shortest Path

Model for traceroute

First approximation: union of shortest paths

G'=(V', E')



Very simple model, but: allows for some analytical and numerical understanding

G = (V, E): sparse undirected graph with a set of

- $N_{S} \text{ sources } S = \{i_{1}, i_{2}, ..., i_{N_{S}}\}$
- $N_T \text{ targets } T = \{j_1, j_2, ..., j_{N_T}\}$

randomly placed.

The <u>sampled graph</u> G'=(V',E') is obtained by considering the union of all the traceroute-like paths connecting source-target pairs.

PARAMETERS:
$$\rho_s = \frac{N_s}{N}$$
, $\rho_T = \frac{N_T}{N}$, $\varepsilon = \frac{N_s N_T}{N}$ (probing effort)
Usually N_S=O(1), ρ_T =O(1)

For each set $\Omega = \{ S, \tau \}$, the indicator function that a given <u>edge</u> (*i*, *j*) belongs to the sampled graph is

$$\pi_{ij} = 1 - \prod_{l \neq m} \left(1 - \sum_{s=1}^{N_s} \delta_{li_s} \sum_{t=1}^{N_T} \delta_{mj_t} \sigma_{ij}^{(l,m)} \right)$$

with

$$\sigma_{ij}^{(l,m)} = \begin{cases} 1 & \text{if } (i,j) \in path \text{ between } l,m \\ 0 & \text{otherwise,} \end{cases}$$

Mean-field statistical analysis

Averaging over all the possible realizations of the set $\Omega = \{ S, T \}$,

$$\left\langle \pi_{ij} \right\rangle = 1 - \left\langle \prod_{l \neq m} \left[1 - \sum_{s=1}^{N_s} \delta_{li_s} \sum_{t=1}^{N_T} \delta_{mj_t} \sigma_{ij}^{(l,m)} \right] \right\rangle \approx 1 - \prod_{l \neq m} \left(1 - \rho_s \rho_T \left\langle \sigma_{ij}^{(l,m)} \right\rangle \right)$$

WE HAVE NEGLECTED CORRELATIONS !!

$$< \pi_{ij} > \simeq 1 - \exp(-\rho_{S} \rho_{T} b_{ij})$$
• usually $\rho_{S} \rho_{T} << 1$
• $b_{ij} = \sum_{s \neq i \neq j \neq i \in V} \left[\frac{\sigma_{ij}^{(s,i)}}{\sigma^{(s,i)}} \right]_{USP} = \sum_{l \neq m \neq i \neq j \in V} \langle \sigma_{ij}^{(l,m)} \rangle$

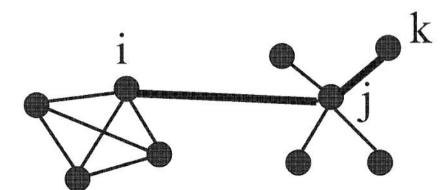
Betweenness centrality b

for each pair of nodes (1,m) in the graph, there are

 $\nu^{\rm lm}$ shortest paths between l and m

 v_{ij}^{lm} shortest paths going through ij

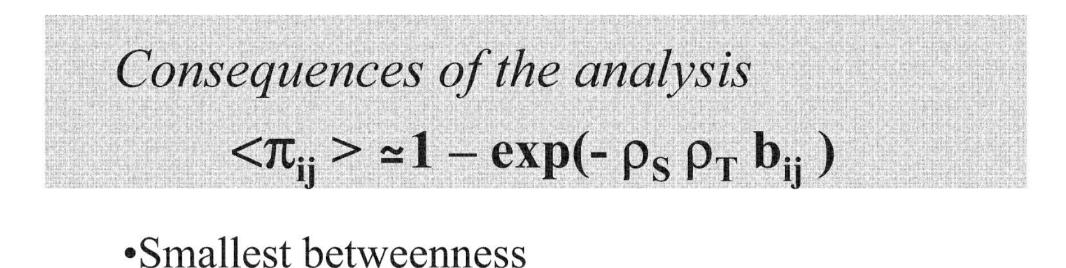
 b_{ij} is the sum of v_{ij}^{lm} / v^{lm} over all pairs (l,m)

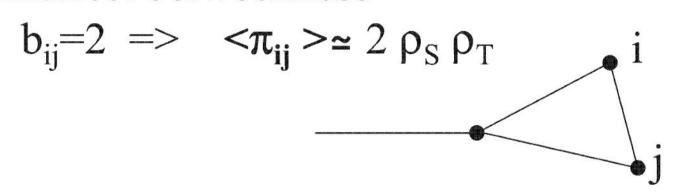


ij: large betweenness

jk: small betweenness

Similar concept: node betweenness b Also: flow of information if each individual sends a message to all other individuals





(i.e. i and j have to be source and target to discover i-j)

•Largest betweenness $b_{ij}=O(N^2) \implies <\pi_{ij} \ge 1$ $\langle \pi_{ij} \rangle \simeq 1 - \exp(-\epsilon b_{ij}/N)$ $\langle \pi_i \rangle \simeq 1 - (1 - \rho_T) \exp(-\epsilon b_i/N) \text{ (discovery probability)}$ $N^*_k / N_k \simeq 1 - \exp(-\epsilon b(k)/N) \text{ (discovery frequency)}$ $\langle k^* \rangle / k \simeq \epsilon (1 + b(k)/N) / k \text{ (discovered connectivity)}$

Results for the vertices

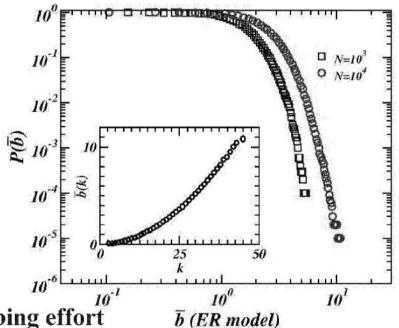
• discovery probability strongly related with the *centrality*;

Summary: • vertex discovery favored by a *finite density* of targets ;

• accuracy increased by increasing *probing effort* ε .

1. Homogeneous graphs:(ex: ER random graphs)

- peaked distributions of k and b
- narrow range of betweenness



=> Good sampling expected only for high probing effort

$$\varepsilon >> \max\left[\overline{b}^{-1}, \overline{b}_{e}^{-1}\right]$$

- 1. Homogeneous graphs
- 2. Heavy-tailed graphs

(ex: Scale-free BA model)

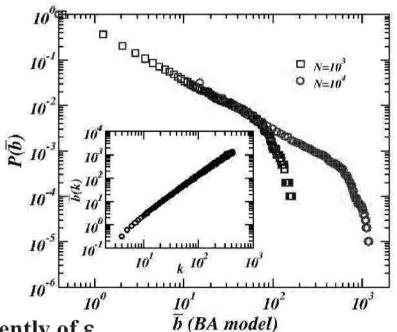
• broad distributions of k and b

 $P(k) \sim k^{-3}$

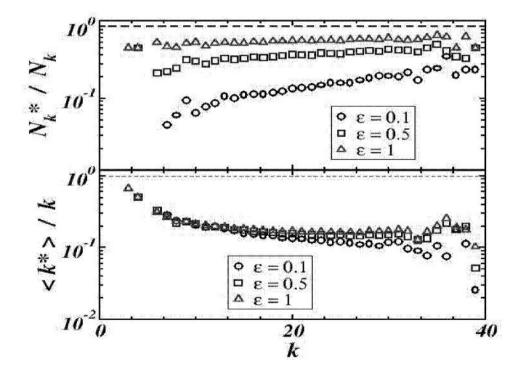
• large range of available values



 $k \gg \varepsilon^{-1/\beta}$ (b(k) ~k^{\beta})

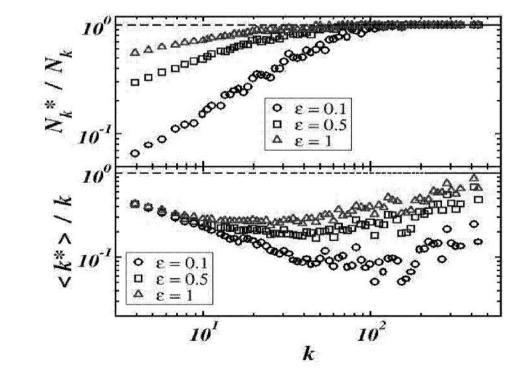


Homogeneous graphs

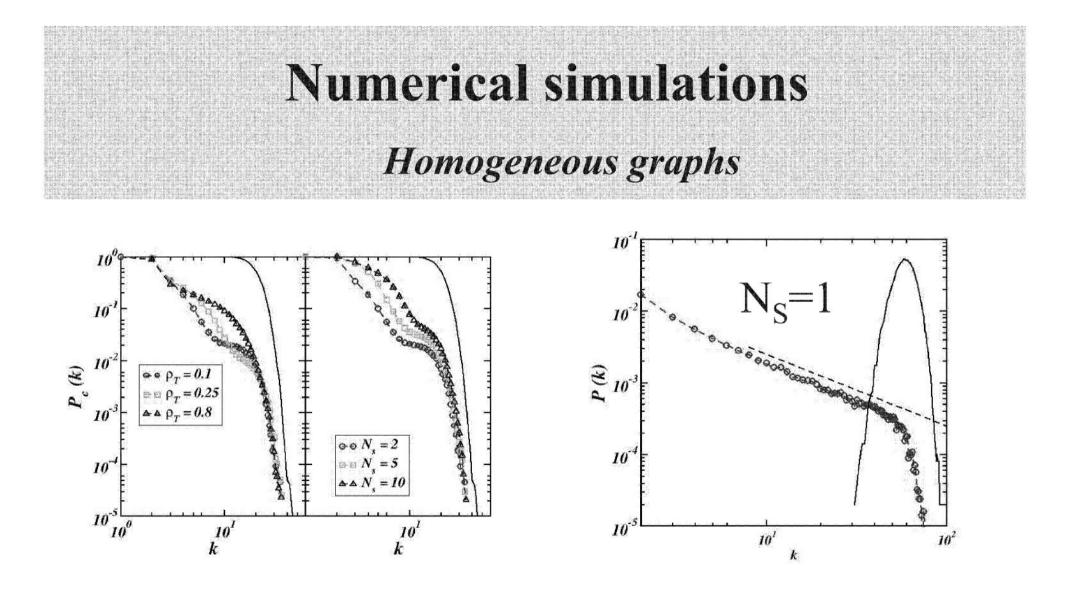


Homogeneously pretty badly sampled

Scale-free graphs



Hubs are well discovered

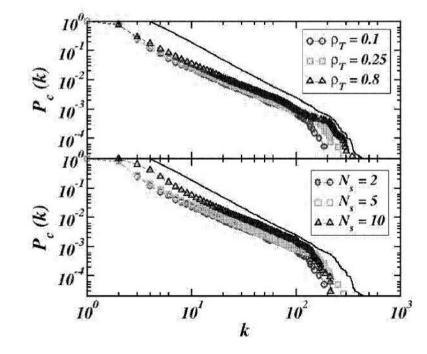


• heavy-tailed $P^*(k)$ only for $N_S = 1$ (cf Clauset and Moore 2005)

• cut-off around <k> => large, unrealistic <k> needed

• bad sampling of P(k)

Scale-free graphs



- good sampling, especially of the heavy-tail;
- almost independent of N_s ;
- slight bending for low degree (less central) nodes => bad evaluation of the exponents. 25



- > Analytical approach to a traceroute-like sampling process
- > Link with the topological properties, in particular the betweenness
- ➤ Usual random graphs more "difficult" to sample than heavy-tails
- ➤ Heavy-tails well sampled
- Bias yielding a sampled scale-free network from a homogeneous network: only in few cases <k> has to be unrealistically large



<u>Heavy tails properties are a</u> <u>genuine feature of the Internet</u>

however

Quantitative analysis might be strongly biased (wrong exponents...)

Perspectives

- •Optimized strategies:
 - -separate influence of ρ_{T}, ρ_{S}
 - •location of sources, targets cf also Guillaume and Latapy 2004
 - investigation of other networks
- •Results on redundancy issues
- •Estimation of the real size of a network from a sampling ?
- Massive deployment

www.tracerouteathome.net ; www.netdimes.org

•The internet is a weighted networks

bandwidth, traffic, efficiency, routers capacity

and...

•Data are scarse and on limited scale

Interaction among topology and traffic