## **Protein and Gene Networks Inference**

**1. New Science:** What are the underlying principles (static and dynamic) of biological networks ?



Dynamical attractors

Scale free static networks



### **Pragmatic problem: search space size**

Random networks (100 nodes) 10<sup>3010</sup> Scale-free networks

Networks with similar dynamics ~10<sup>8</sup> Non-chaotic networks



Jean-Loup Faulon, GTL Modeling & Simulation Workshop, July 23, 2003

# **Protein and Gene Networks Inference**

### 2. Barriers:

- Reaction rates (experimental)
- Static and dynamic network characterization tools (algo & math)
- Data format standard (software & hardware)

2-Hybrid systems, phage display, MS, gene microarray, protein chips, bioinformatics

- Inference algorithm with sensitivity analysis (algo)



Number of data points required to infer unique parsimonious Boolean networks from microarray data and number of clusters with similar dynamics vs. number of networks

#### 4. Resources:

- Database (hardware & software)
- Manpower

