# Feed-forward loops: Linking Function, Plasticity, Evolvability and Abundance

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Bled, February 2011



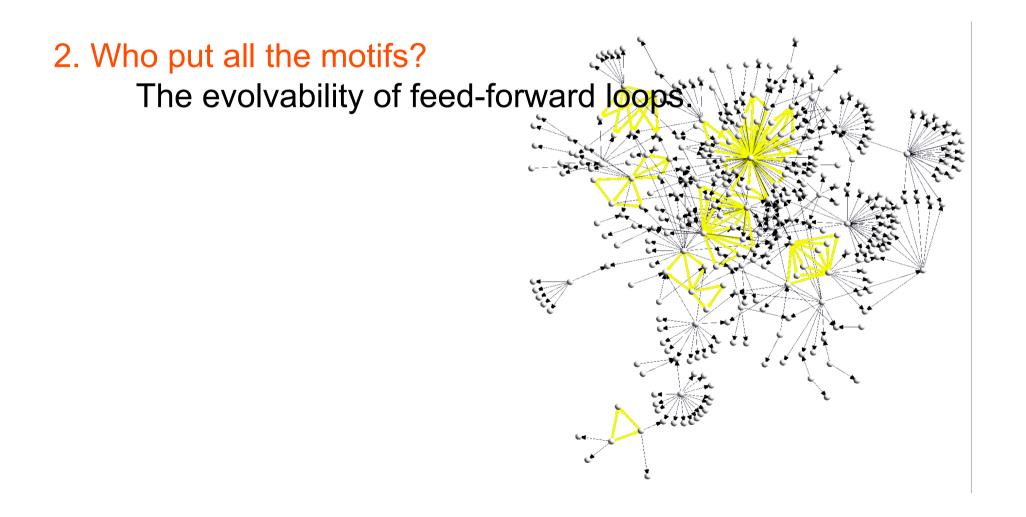




## Outline

1.Why networks?

Systems biology and interaction webs.



# **Early Pioneers**





Alan Hodkin

Andrew Huxley

Modelization of the nerves action potential (giant squid neuron) 1952 Noble Prize in 1963

(Biological) pattern formation as result of simple physical constraints (reaction-diffusion).



Denis Noble

Modelization of a working heart in 1960 and development of the virtual heart using supercomputers.



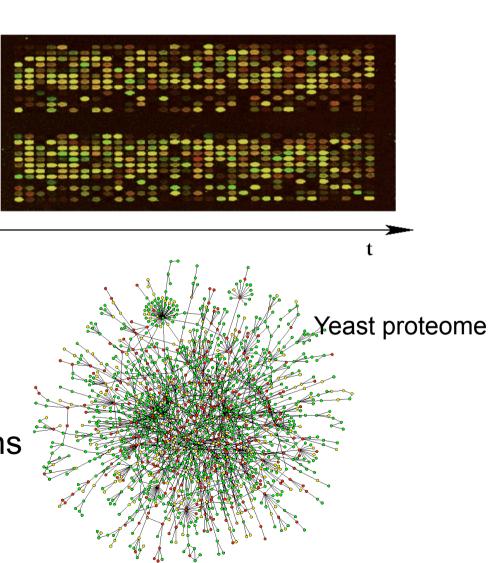
Alan Turing

х

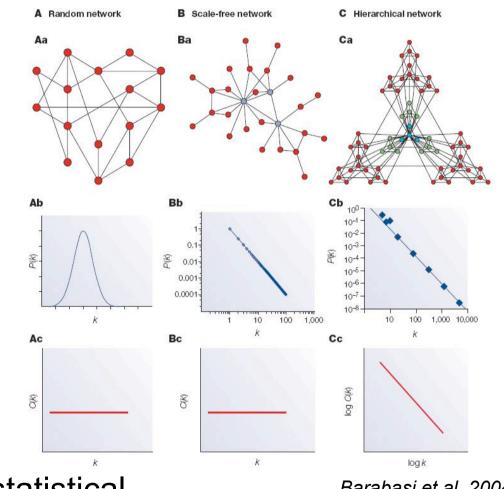
interactions in space and time



networks as representation for maximal physical interactions



metabolic protein interaction transcriptional ecological functional: apoptosis cancer signaling

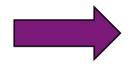




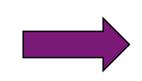
graph theory and statistical measures (eg. k, p(k),  $\langle I \rangle$ , C(k))

Barabasi et al. 2004

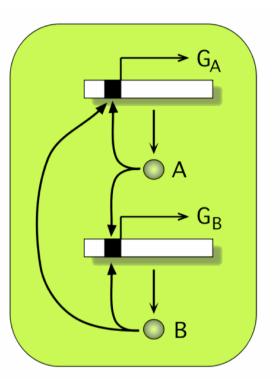
.. subsections of large NWs

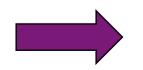


analytically calculable in their entirety for their *local* qualities

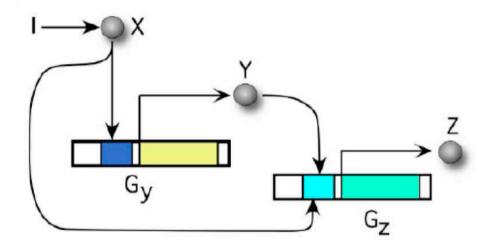


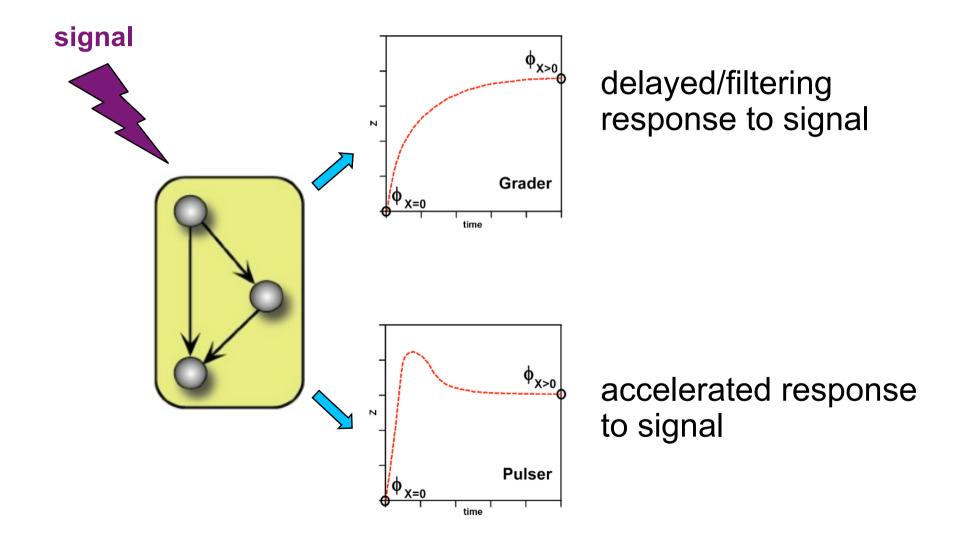
supersection crucially important for understanding their plasticity

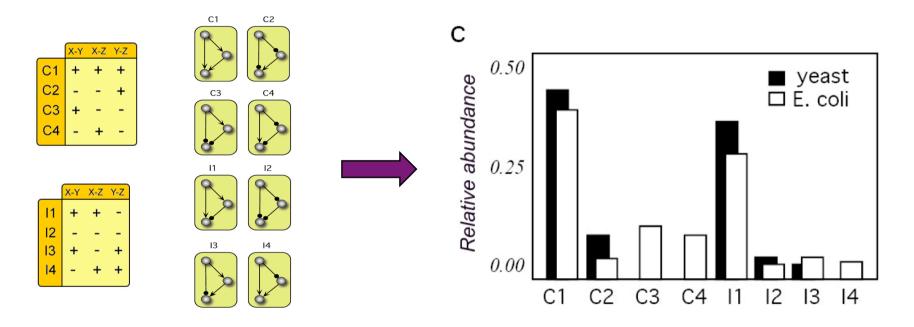




questions on the emergence of the status quo



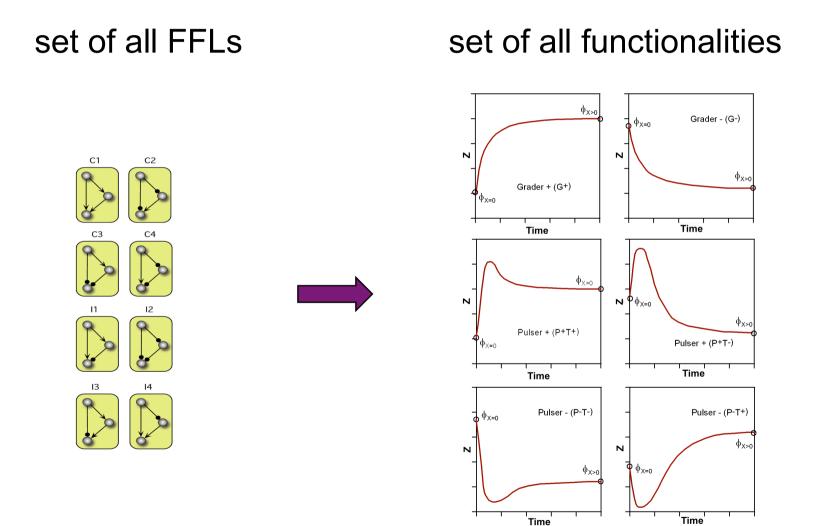




Mangan et al. 2006

The pattern of abundance cannot be explained in a satisfactory manner by functionalist (single topology -> single function) nor by neutralist (by-product of growth rules) arguments.

#### FFLs and their Trajectory



# FFL model

$$\dot{Y} = \gamma_{Y} \left( \frac{1 + \alpha^{X} \omega_{Y}^{X} x^{n}}{1 + \omega_{Y}^{X} x^{n}} \right) - d_{Y} Y$$

$$\dot{Z} = \gamma_{Z} \left( \frac{1 + \beta^{X} \omega_{Z}^{X} x^{n} + \beta^{Y} \omega_{Z}^{Y} y^{m} + \beta^{XY} \omega_{Z}^{XY} x^{n} y^{m}}{1 + \omega_{Z}^{X} x^{n} + \omega_{Z}^{Y} y^{m} + \omega_{Z}^{XY} x^{n} y^{m}} \right) - d_{Z} Z$$

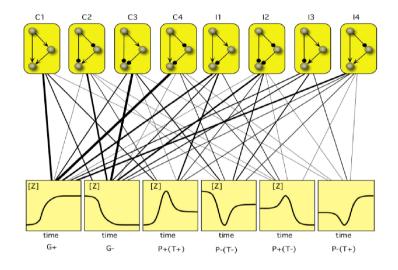


Which functionality can be expected?

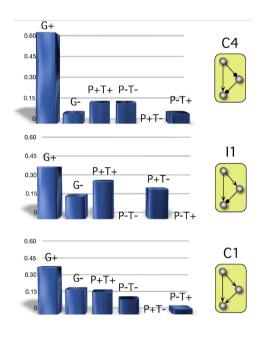


- Intrinsic properties of the system allow to deduce the probability for a given function, independent from the numeric values of the parameters.
  - Key parameter define the characteristic shape of the nullcline (BR)
  - Nullclines' shape confines the trajectory
  - A separatrix defines the limit of two distinct qualit. behavior of the trajectory

#### **Probability of Function**



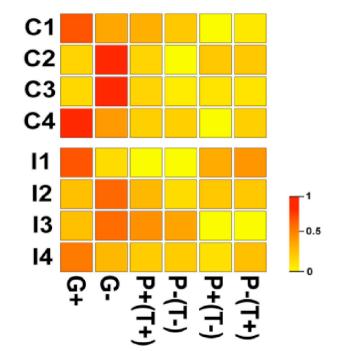
$$P_{ij} = \Omega_k \sum_{i=1}^{T_{jk}} p_{ik} \Psi_{ijk}$$



- $p_{ik}$  prob for a certain set of parameters to implement seq. *i* for motif *k*
- $P_{ij}$  prob for motif *k* to implement functionality *j* described by seq. *i*
- $\Psi_{iik}$  number of equal dynamical outcomes j
- $T_{ik}$  total number of backbone seq. *i* implementing dynamic *j*
- $\Omega_k$  normalization constant

Macía, Widder and Solé 2009

## Plasticity of FFLs



Quantitative description of the FFLs' underlying plasticity independent of parameters.

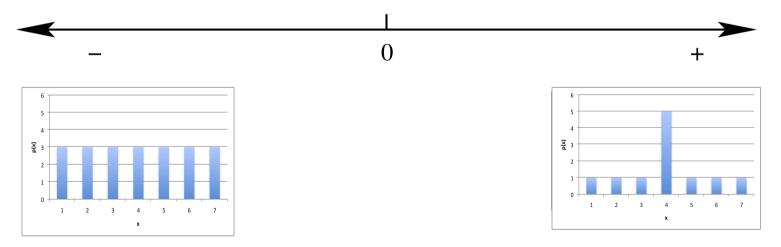
# **Kurtosis**

$$K = \frac{\mu_4}{\sigma^2} - K_0$$

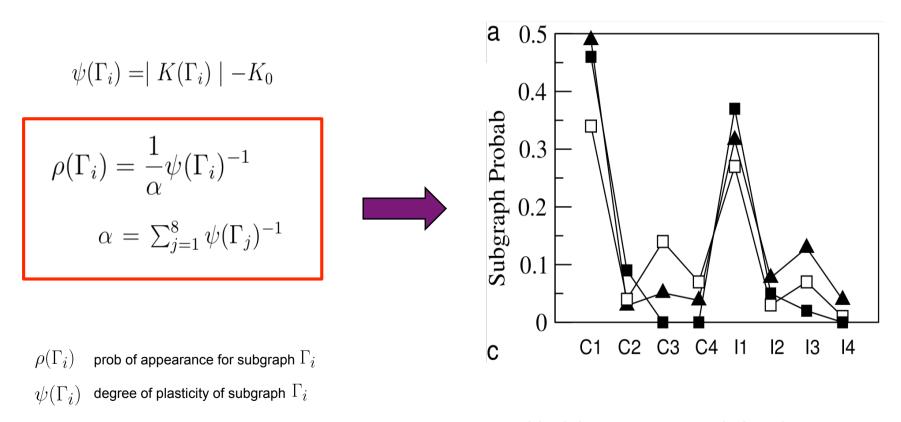


- $\mu_4$  4th moment around the mean
- $^{\sigma}$  standard deviation
- $K_0$  reference value (normal dist. 3)



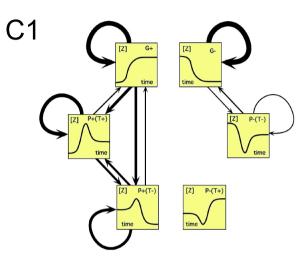


.. the peakedness of a probability distribution



black boxes.....natural abundance yeast white boxes.....natural abundance E.coli black triangles...predicted probabilities

# Linking plasticity and evolvability

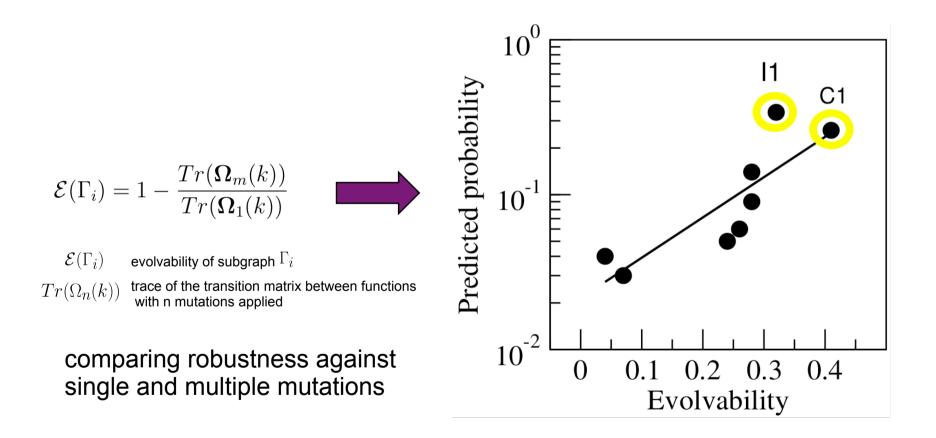


transition probabilities between different functions for a given subgraph upon mutation (of parameters)

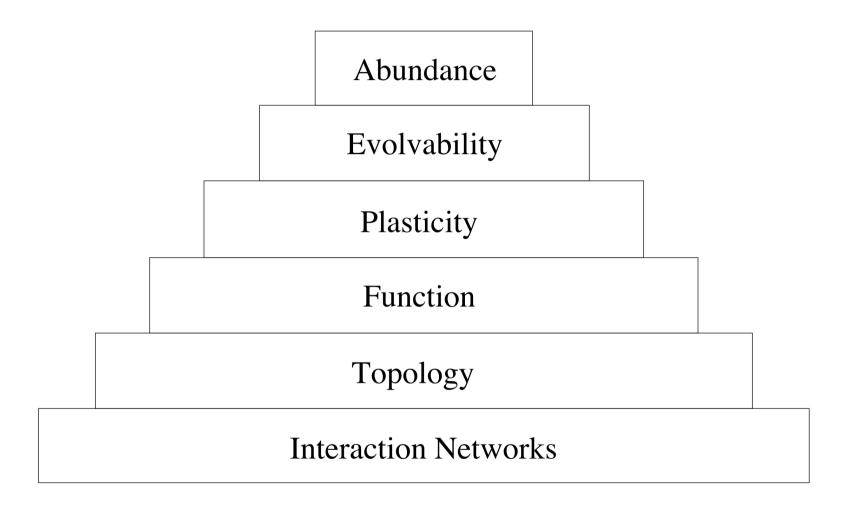
Table 1: Transition probabilities for single mutations, C1

	$G^+$	$G^-$	$P^+T^+$	$P^{-}T^{-}$	$P^+T^-$	$P^{-}T^{+}$
$G^+$	0.251	0	0.027	0	0	0
$G^-$	0	0.329	0	0	0.008	0
$P^+T^+$	0	0	0.180	0	0	0
$P^{-}T^{-}$	0	0	0	0.157	0	0
$P^+T^-$	0	0.016	0	0	0.031	0
$P^{-}T^{+}$	0	0	0	0	0	0

robustness against mutation



## Conclusions





#### Thanks!





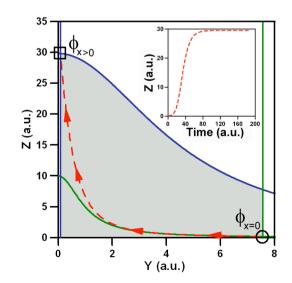


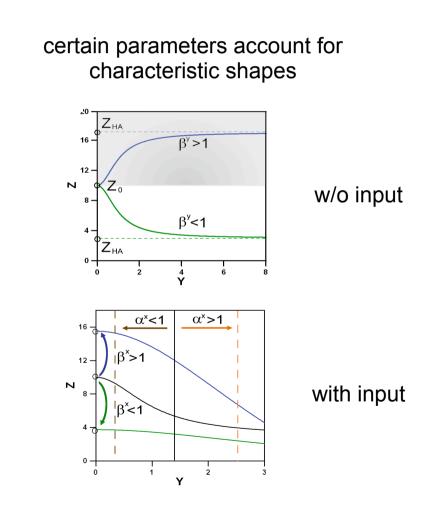
Barcelona Biomedical Research Park

$$\left(Y\right)_{\dot{Y}=0} = \frac{\gamma Y}{dY} \left(\frac{1 + \alpha X \omega_Y^X X^n}{1 + \omega_Y X^n}\right)$$

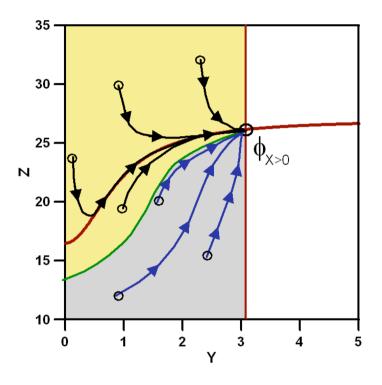
$$(Z)_{\dot{Z}=0} = \frac{\gamma Z}{dZ} \left( \frac{1 + \beta^X \omega_z^X X^n + \beta^Y \omega_Z^Y Y^m + \beta^{XY} \omega_Z^{XY} X^n Y^m}{1 + \omega_z^X X^n + \omega_Z^Y Y^m + \omega_Z^{XY} X^n Y^m} \right)$$

change of the nullcline's shape upon input confine the trajectory





Starting from different ICs in phase space, two distinct qualitative behaviours are encountered: joining of the nullcline before the FP or joining at the FP. The regions are separated by Z(Y).



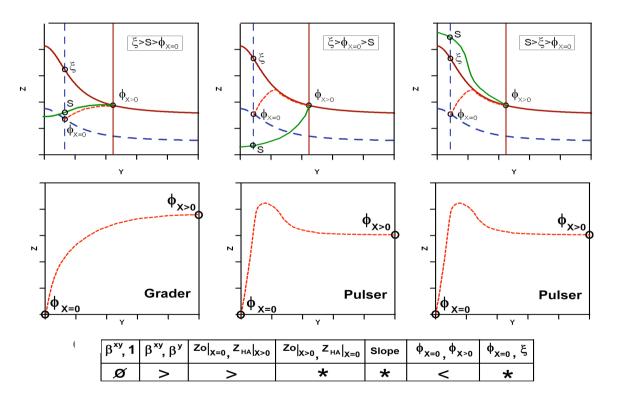
Using condition 
$$\frac{Y_f - Y}{Z_f - Z} = \frac{\dot{Y}}{\dot{Z}}$$

we find the analytical expression of the separatrix

$$Z(Y) = \left(\frac{1}{1 - \frac{d_Y}{d_z}}\right) \left[\frac{\gamma_Z}{d_z} \left(\frac{1 + \beta^X \omega_z^X X^n + \beta^Y \omega_Z^Y Y^m + \beta^{XY} \omega_Z^X X^n Y^m}{1 + \omega_z^X X^n + \omega_Z^Y Y^m + \omega_Z^{XY} X^n Y^m}\right) - \frac{d_Y}{d_Z} Z_f\right]$$

The relative position of 3 points at the crossing with  $Yo|_{x=0}$  determines the functionality of the trajectory for a given nullcline:

Separatrix,  $Zo|_{x=0}$  and  $Zo|_{x>0}$ 



The parametric backbone sequence determines the shape of the nullcline.