### Aspects of analysis

Focus on size of attractors and basins of attraction, number of attractors, stability **Impact of** K (averaged)

- K = N:  $\triangleright$  network is complete graph, successor states chosen at random
  - ▷ attractors can be very large (exponential in *N*), number of attractors of order *N*, maximal sensitivity to intial conditions (chaotic)
- K = 1: > structure decomposes into simple cycles with tails, behavior determined by constant functions and cycles (product of subsystems)
  - ▷ attractor number exponential in *N*, attractor size of order *N*, limited effect of perturbations (ordered)
- K = 2:  $\triangleright$  variability in structure, canalyzing functions
  - ▷ attractor number and size increases superpolynomial with *N*, good stability properties (critical)

#### Understanding RBNs yields insights into more complex real world networks

### Reasons for order

Frozen core: set of "dynamically inactive" elements dividing the system into functionally isolated modules

- fixed value for large part of the dynamical behavior
- decision taking subnetworks
- combinatorial dynamics (attractors)
  - modularity allows for short cycle lengths (K = 1)
  - ► "walls of constancy" absorb perturbations (K = 1), restrict attractors reachable by mutations/noise
  - moderate number of attractors w.r.t. N due to big number of frozen elements

# **Observation:** large connected frozen structure percolates spontaneously if K = 2

### Canalyzing functions

**Definition**  $f : \{0,1\}^n \to \{0,1\}$  is called *canalyzing (in i)*, if there is  $i \in \{1,...,n\}$  and  $c,c' \in \{0,1\}$  such that f(x) = c' for all  $x \in \{0,1\}^n$  with  $x_i = c$ .

The input value c is called *canalyzing value*, and the output c' is called *canalyzed value*.

- elements regulated by canalyzing functions are insensitive to other regulator inputs in the presence of canalyzing value
- forcing structures can percolate through the network if canalyzing functions of several elements interlock:

 $A \rightarrow B$  with A, B regulated by canalyzing functions, and canalyzed value of A is canalyzing value of B (forcing connection)

- concepts of frozen core and canalyzing functions can be generalized for multi-valued functions.
- many biological functions are canalyzing

### Structural modularity

Consider interaction graph G(f) = (V, E) representing the network structure

- find modules based on graph theoretical criteria, shared expression pattern ...
- ▷ include statistical significance [U. Alon et al., Science, 2002]
  - recurring patterns of interconnections with significantly higher occurrence than in random networks (motifs)
  - sets of motifs of distinct networks may differ considerably
  - ► similar motifs in networks with similar function ⇒ homologies, classification, comparison



### Significance of motifs

- evaluating structural significance
  - pairwise disconnectivity index: importance of subgraph *M* for sustaining network connectivity [B. Goemann et al., 2009]:

compare number of connected ordered pairs in G with number in  $G \setminus M$ 

check all (sets of) subgraphs of given size (motifs, anti-motifs)

- no correlation between abundance of pattern and topological significance
- topological role of substructures mainly determined by location/embedding
- evaluating functional significance
  - analyzing behavior of isolated motifs
    - biological meaningful behavior (differentiation, oscillation, pulse, coordinated expression,...)
    - \* not necessarily retrievable in complex system

### **Compositional dynamics**

#### Consider disjoint modules

 $G^1, \ldots, G^k$  interaction graphs associated with discrete functions  $f^1, \ldots, f^k$  and state space  $X^i$ ,  $G := \bigcup_{i=1}^k G^k$ , *f* derived from  $f^1, \ldots, f^k$ 

- ► all attractors of (G, f) can be derived from products of attractors of (G<sup>i</sup>, f<sup>i</sup>), i ∈ {1,...,k}
- asynchronous update
  - # attractors of (G, f) is the product of # attractors of  $(G^{i}, f^{i})$
  - cardinality of compositional attractor A is the product of the cardinalities of the component attractors
- synchronous update
  - # attractors of (G, f) depends on # of attractors of (G<sup>i</sup>, f<sup>i</sup>) as well as least common multiple and greatest common divisor of the component attractor lengths
  - maximal length of a compositional attractor is the maximum of the least common multiple of attractor lengths of component attractors

### Non-isolated modules

#### Interacting modules

consider network (*G*, *f*) with associated state space  $X = X_1 \times \cdots \times X_n$ 

- module attractors generally not preserved
- dynamics of the isolated module is not necessarily imprinted on network dynamics

#### **Recover isolation**

- ▶ find partial fixed points, i. e. find  $I \subset \{1, ..., n\}$  and  $x_i \in X_i$  s. t.  $f_i(y) = x_i$  for  $i \in I \subset \{1, ..., n\}$  and all  $y \in X$  with  $y_i = x_i$  for all  $i \in I$  (frozen core)
- consider part of state space  $X[I, x_i] := \{y \in X \mid \forall i \in I : y_i = x_i\}$
- Consider graph components Z<sup>1</sup>,..., Z<sup>k</sup> of the subgraph of G derived from the vertices *j* ∉ *I*, and associated functions f<sup>j</sup> derived by projection of f|<sub>X[I,x<sub>i</sub>]</sub>
- $\Rightarrow$  all attractors in  $X[I, x_i]$  can be derived from attractors of  $(Z_j, f^j)$

- easy calculation of partial fixed points for networks with input layer
  - fix input values
  - constraint percolation through



#### **Complexity reduction**

input (0,0,0)

 state space sizes: 82944/2592

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#### **Complexity reduction**

input (0,0,0)

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- easy calculation of partial fixed points for networks with input layer
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#### **Complexity reduction**

input (0,0,0)

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[L. Mendoza, Biosystems 84(2), 2006]

## Complexity reduction

### input (0,0,0)

state space sizes:
 82944/2592

### input (1,0,0)

- state space sizes: 82944/4
- additional information,
  e.g. extent of crosstalk, linking structural and dynamical aspects

### Transfer of characteristics

- isolated modules may imprint qualitative characteristics on the network
- isolation in parts of state space may be sufficient
- example: positive and negative circuits
  - isolated circuits show distinct behavior:
    \* positive circuits induce multiple attractors
    \* negative circuits induce cyclic attractors (cardinality >1)
    note differences for synchronous and asynchronous update
  - check characteristics for complex networks containing circuits
    characteristics are conserved for circuits in "conditional" isolation (compositional properties)

#### **Exploiting modularity**

- reduce network complexity
- understand design principles