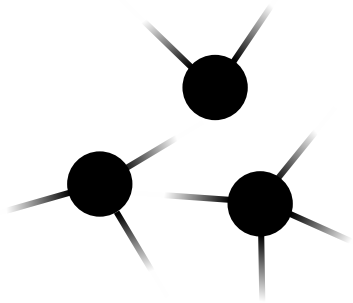




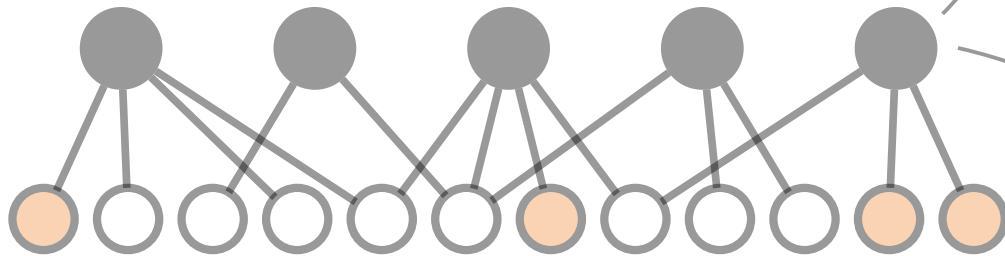
p_n

Clique size
distribution

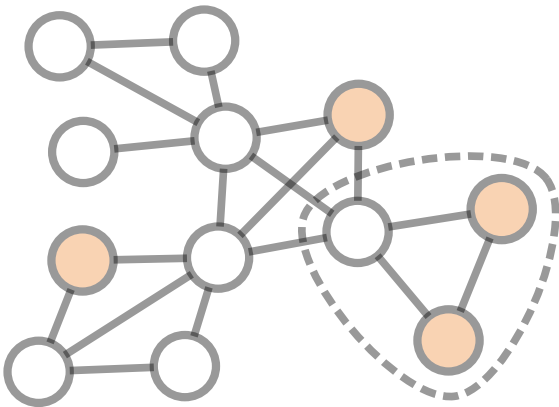


Cliques

Nodes

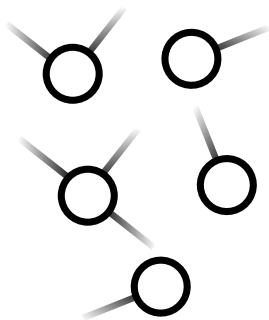


Projection
on nodes

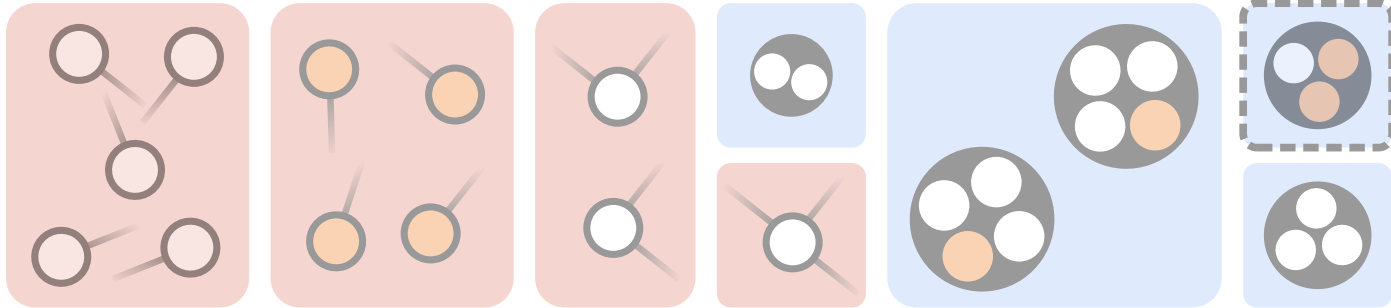


g_m

Membership
distribution

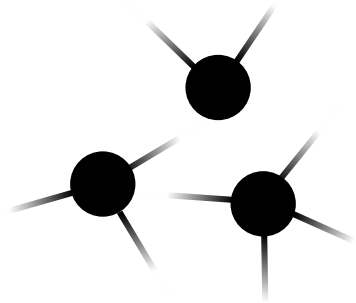


Compartmental formalism



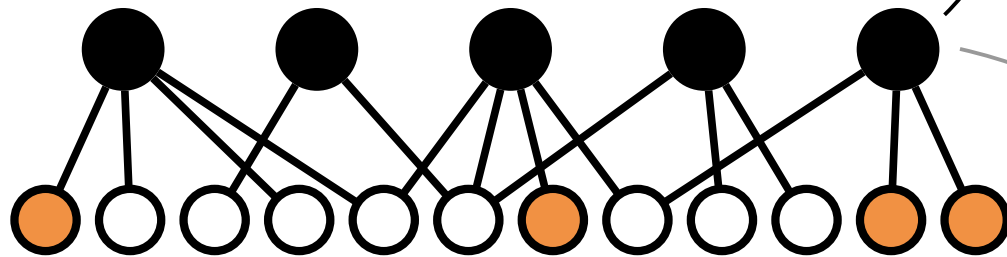
p_n

Clique size
distribution

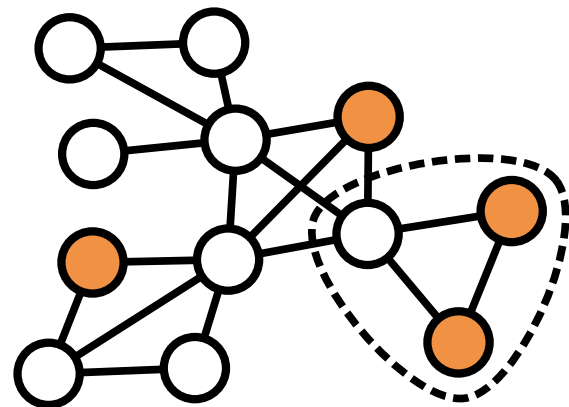


Cliques

Nodes

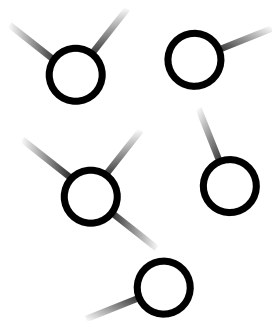


Projection
on nodes

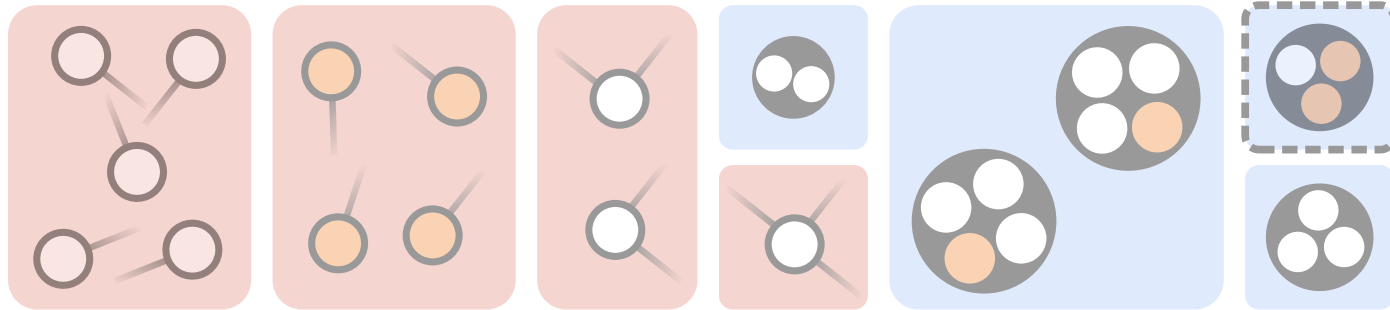


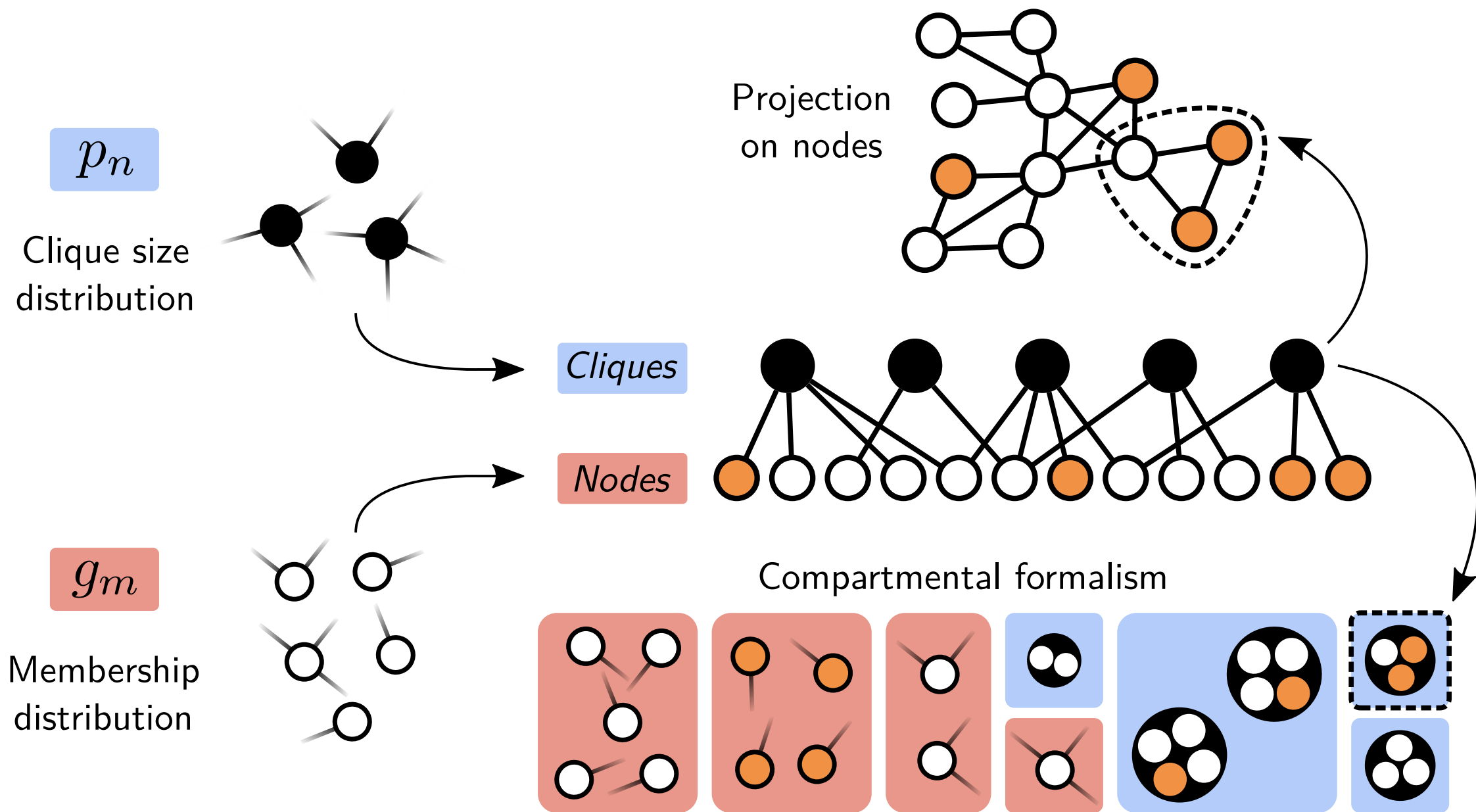
g_m

Membership
distribution



Compartmental formalism





M.E.J. Newman, *Properties of highly clustered networks*, Phys. Rev. E 68, 026121 (2003)

- μ : recovery rate
- β : *weight* associated to transmission
- $\Theta_{n,i,\beta}$: transmission rate in a group of size n , with i infected nodes, and weight β

Dynamics at the groups level

$$\begin{aligned}\frac{dG_{n,i}}{dt} = & \mu(i+1)G_{n,i+1} - \mu i G_{n,i} \\ & + (n-i+1)\Theta_{n,i-1,\beta}G_{n,i-1} - (n-i)\Theta_{n,i,\beta}G_{n,i} \\ & + (n-i+1)\rho G_{n,i-1} - (n-i)\rho G_{n,i}\end{aligned}$$

Dynamics at the nodes level

$$\frac{dS_m}{dt} = \mu(g_m - S_m) - m\textcolor{brown}{r}S_m$$

Mean-field quantities

$$\rho(t) = r(t) \frac{\sum_m m(m-1)S_m}{\sum_m mS_m} ; \quad r(t) = \frac{\sum_{n,i} (n-i)\Theta_{n,i,\beta} G_{n,i}}{\sum_{n,i} (n-i)G_{n,i}}$$

Global prevalence

$$I(t) = \sum_m [g_m - S_m(t)]$$

The model and its mathematical description

$r(t)$: mean infection rate from a random group to which a random node belongs

$\rho(t)$: mean infection rate from all *other* groups to which a random node in a random group belongs

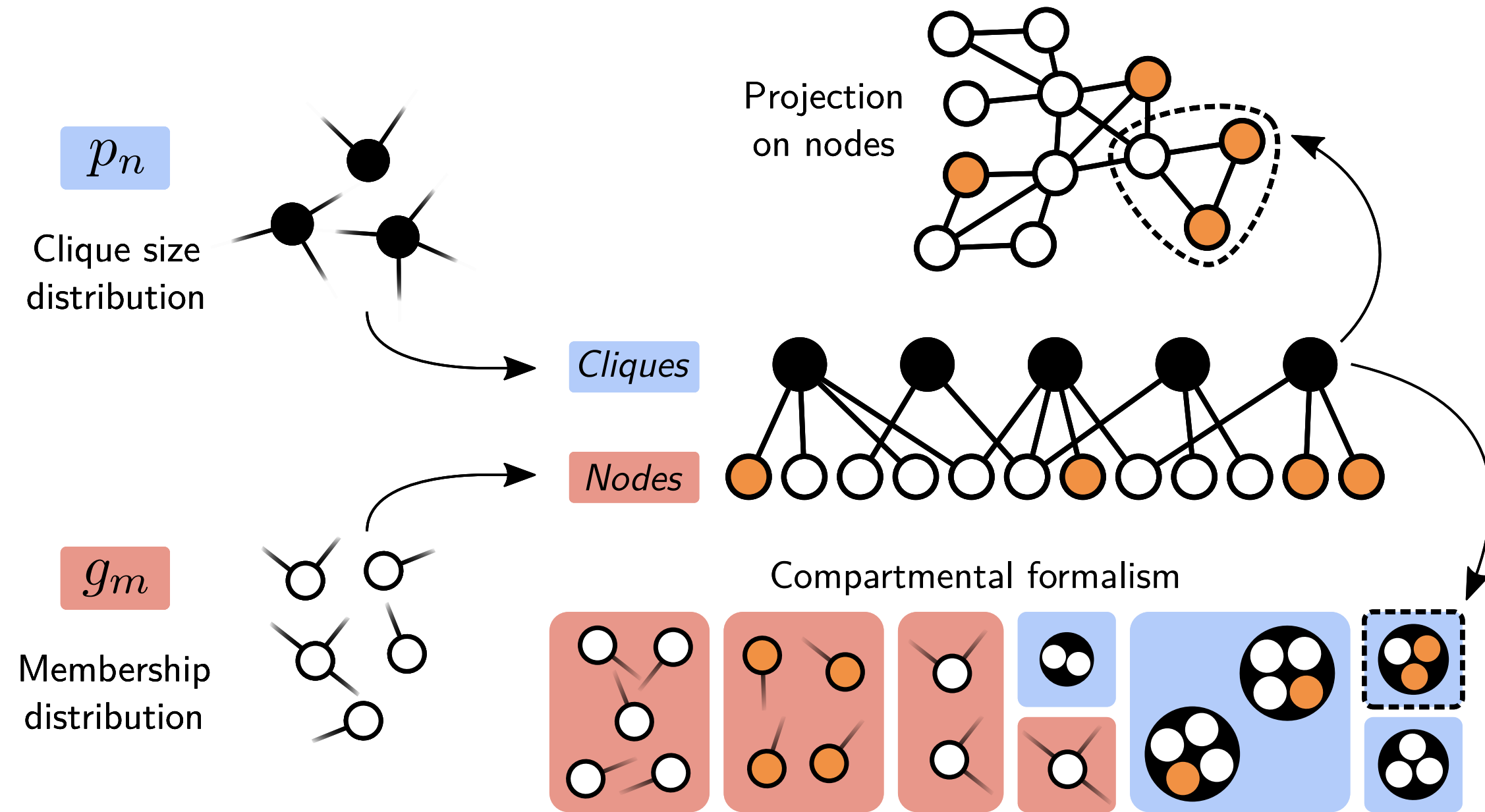
S_m : fraction of nodes with membership m and that are susceptible

$G_{n,i}$: fraction of group of size n and that contain i infectious nodes

g_m : fraction of nodes with membership m

p_n : fraction of groups of size n

The model and its mathematical description



g_m : fraction of nodes with membership m

p_n : fraction of groups of size n

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Dynamics at the nodes level

$$\frac{dS_m}{dt} = \mu(g_m - S_m) - m r S_m$$

$r(t)$: mean infection rate from a random group to which a random node belongs

$\rho(t)$: mean infection rate from all *other* groups to which a random node in a random group belongs

Mean-field quantities

$$\rho(t) = r(t) \frac{\sum_m m(m-1)S_m}{\sum_m m S_m} ; \quad r(t) = \frac{\sum_{n,i} (n-i)\Theta_{n,i,\beta}G_{n,i}}{\sum_{n,i} (n-i)G_{n,i}}$$

Global prevalence

$$I(t) = \sum_m [g_m - S_m(t)]$$

The model and its mathematical description

Stationary state

$$S_m^* = \frac{g_m}{1 + mr}$$
$$\mu(i + 1)G_{n,i,\beta}^* = [\mu i + (n - i)(\Theta_{n,i,\beta} + \rho)] G_{n,i,\beta}^* - (n - i + 1)(\Theta_{n,i-1,\beta} + \rho) G_{n,i-1,\beta}^*$$

Epidemic threshold

$$\left. \frac{dF}{d\rho} \right|_{\rho \rightarrow 0} > 1$$

where

$$F(\rho) \equiv r(\rho) \frac{\sum_m m(m - 1) S_m(\rho)}{\sum_m m S_m(\rho)}$$