

Disease Transmission in Layered Networks

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Part I: Final size relations for deterministic models

The Kermak-McKendrick Epidemic Model (1927)

$$v(t) = S(t) \int_0^{\infty} \beta(\tau) v(t - \tau) d\tau$$

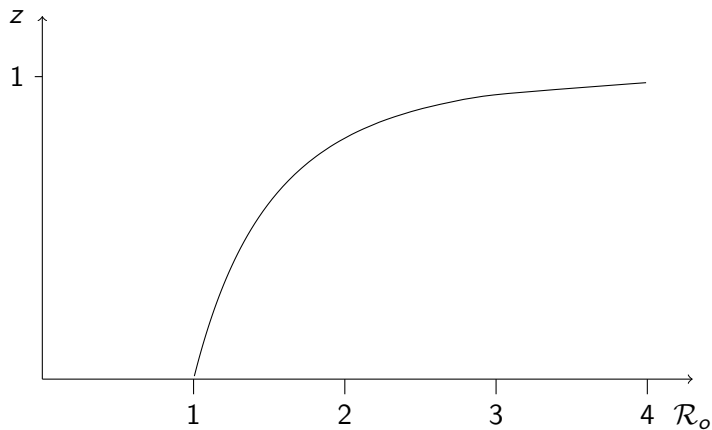
$$S'(t) = -v(t)$$

The final size, $z = 1 - \frac{S(\infty)}{N_0}$, is a solution of

$$\mathcal{R}_o z + \ln(1 - z) = 0$$

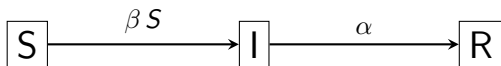
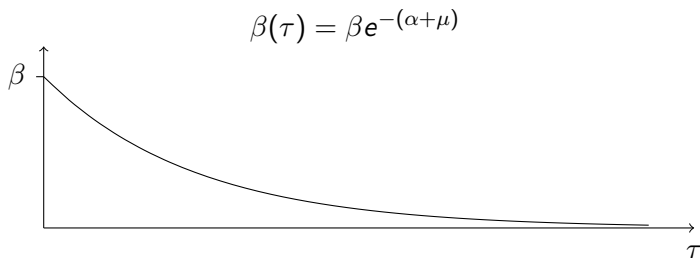
$$\text{where } \mathcal{R}_o = N_0 \int_0^{\infty} \beta(\tau) d\tau$$

The final size relation



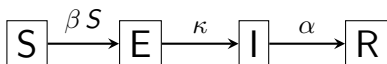
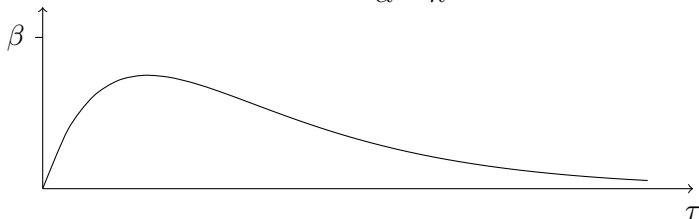
$$\mathcal{R}_o z + \ln(1 - z) = 0$$

Example 1: The SIR model

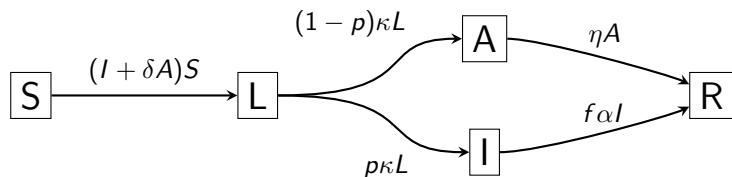


Example 2: The SEIR model

$$\beta(\tau) = \beta \frac{\kappa(e^{-\kappa\tau} - e^{-\alpha\tau})}{\alpha - \kappa}$$



Example 2: The SLIAR model



$$S' = -S(I + \delta A)$$

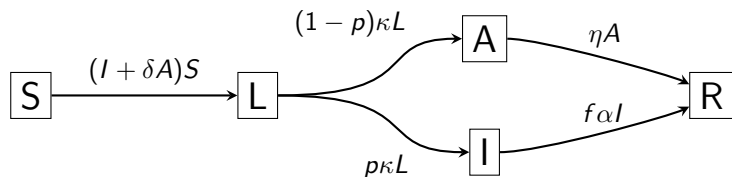
$$L' = S(I + \delta A) - \kappa L - \varphi_L L + \theta_L L_T$$

$$I' = p\kappa L - \alpha I - \varphi_I I + \theta_I I_T$$

$$A' = (1 - p)\kappa L - \eta A - \varphi_A A + \theta_A A_T$$

$$R' = f\alpha I + f_T \alpha_T I_T + \eta A + \eta_T A_T$$

Example 2: The SLIAR model



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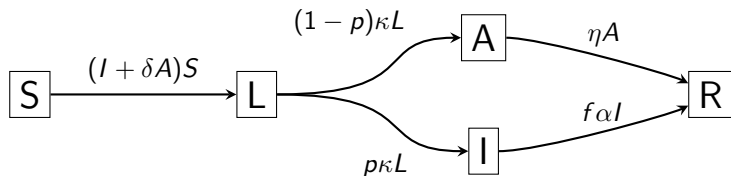
$$A' = (1 - p)\kappa L - \eta A - \varphi_A A + \theta_A A_T$$

$$R' = f\alpha I + f_T \alpha_T I_T + \eta A + \eta_T A_T$$

$$S' = -S\beta^T x$$

$$x' = S\beta^T x Q - Vx$$

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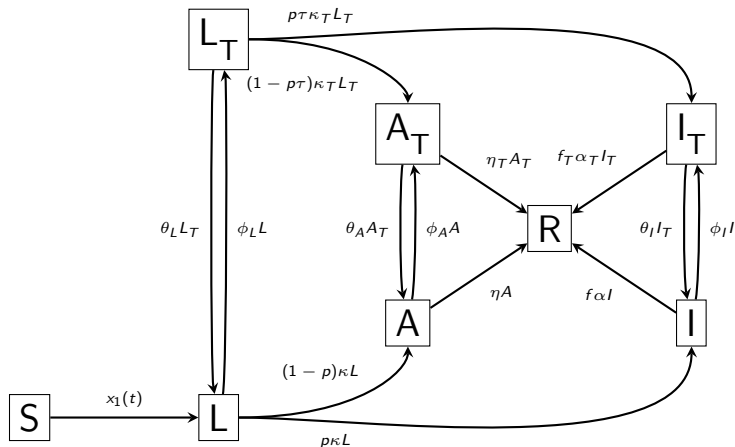
$$R' = f\alpha I + f_T \alpha_T I_T + \eta A + \eta_T A_T$$

$$S' = -S\beta^T x$$

$$x' = S\beta^T x Q - Vx$$

$$\beta(\tau) = \beta^T e^{-V\tau} Q$$

Example 3: The SLIAR model with treatment



$$\beta(\tau) = (1 - \phi(\tau))\beta_U(\tau) + \phi(\tau)\beta_T(\tau)$$

The multi-type Epidemic Model

$$v_i(t) = S_i(t) \sum_{j=1}^m f_{ij}(N(t)) \int_0^\infty \beta_j(\tau) v_j(t - \tau) d\tau$$

$$S_i'(t) = -v_i(t)$$

The final size, $z_i = 1 - \frac{S_i(\infty)}{N_{i0}}$, is a solution of

$$\boxed{K_{ij} z_j + \ln(1 - z_i) = 0} \quad \text{where } K_{ij} = N_{0i} f_{ij}(N^*) \int_0^\infty \beta_j(\tau) d\tau.$$

This equation has a unique solution in $(0, 1)^m$ if and only if $\mathcal{R}_o = \rho(K) > 1$.

Example 4: Age structured epidemic

Population: $N_i = (60^2 - (40 - i)^2)/266667$, $0 < i < 100$,

$$\text{Contact rates: } a_i = \begin{cases} 1.46 & 0 \leq i \leq 12, \\ .66 & 12 < i \leq 18, \\ .16 & 18 < i \leq 100, \end{cases}$$

$$\text{Infectivity: } \int_0^{\infty} \beta_j(\tau) d\tau = A_j = 3, \quad 0 < j < 100.$$

The final size is

$$z_i = \begin{cases} 941.50 & 0 \leq i \leq 12, \\ 722.86 & 12 < i \leq 18, \\ 267.35 & 18 < i \leq 100, \end{cases}$$

in infections per 1000 for the three age groups.

Example 4: Age structured epidemic

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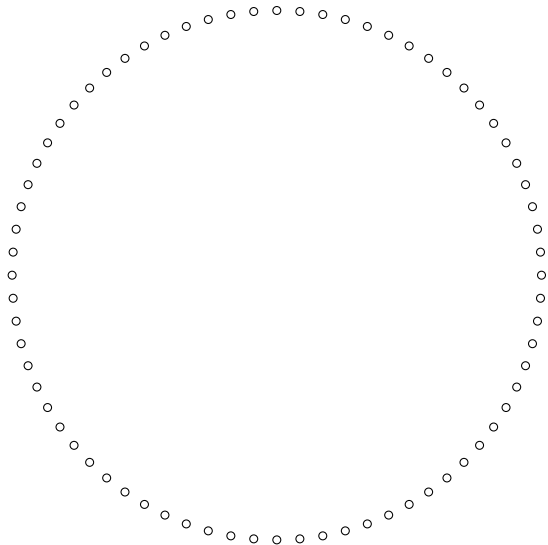
$$\text{Susceptibility: } b_i = \begin{cases} 1 & i < 77, \\ 5 & i > 77 \end{cases}$$

$$\text{The final size is } z_i = \begin{cases} 951.35 & 0 \leq i \leq 12, \\ 745.03 & 12 < i \leq 18, \\ 282.01 & 18 < i \leq 77, \\ 863.00 & 77 < i \leq 100 \end{cases}$$

in infections per 1000 for the three age groups.

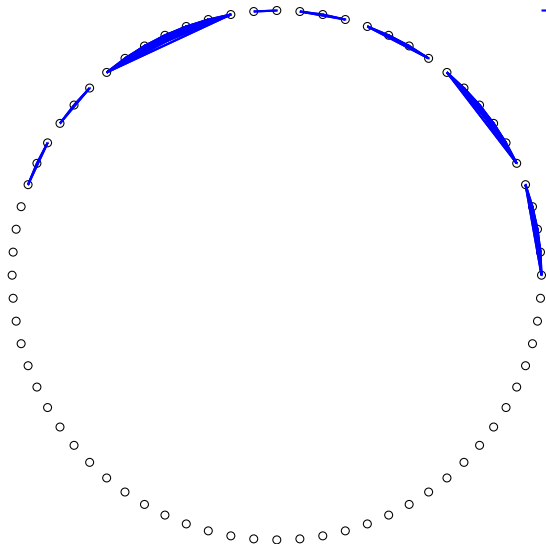
Part II: Layered Networks

A layered network

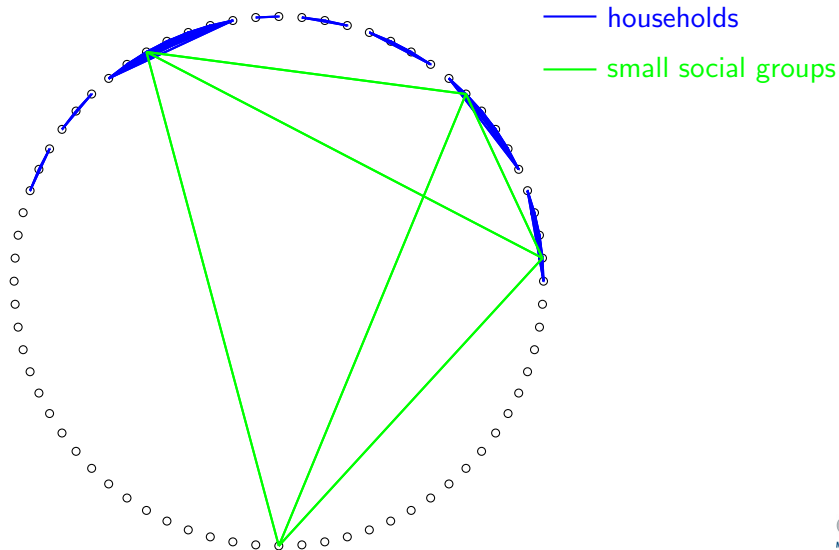


A layered network

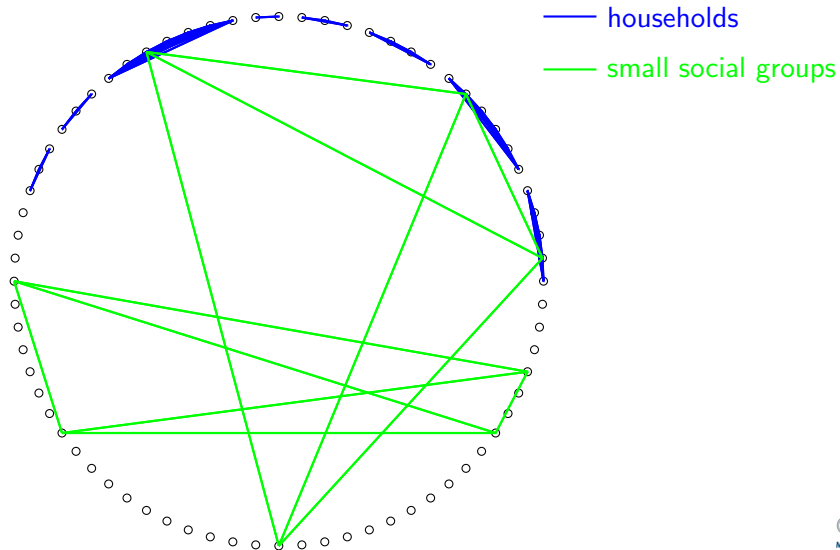
— households



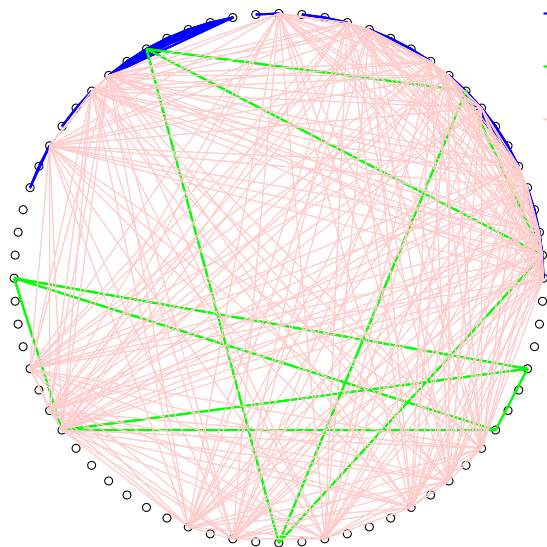
A layered network



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A layered network

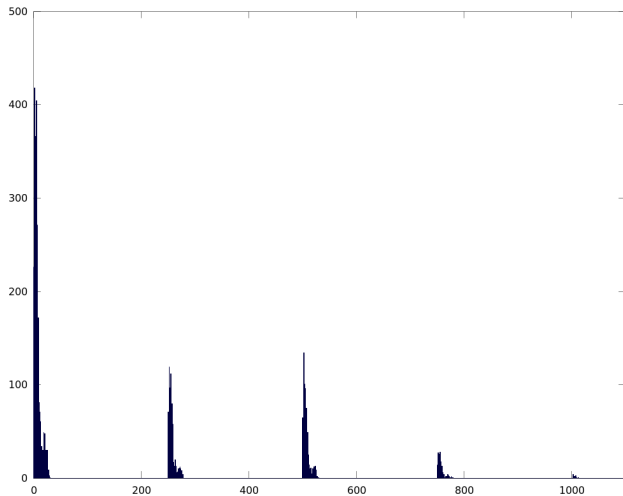


— households

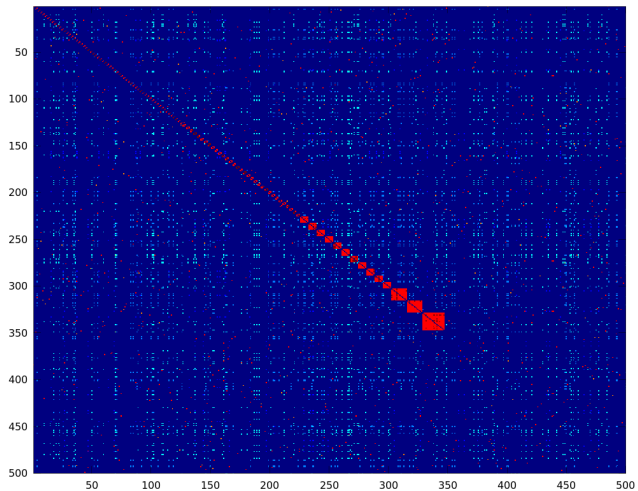
— small social groups

— large asocial network

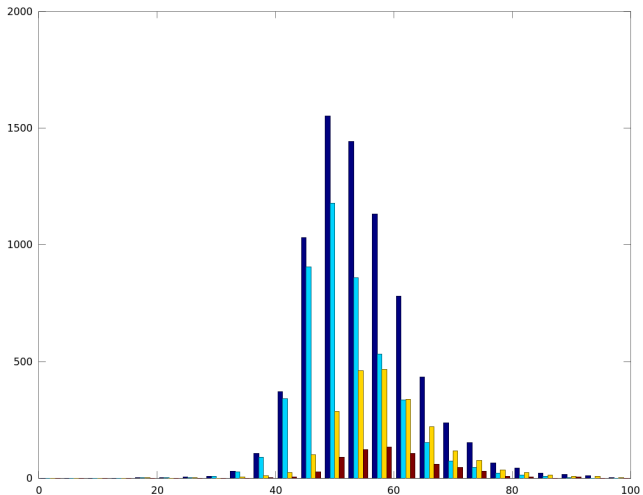
The Degree Distribution



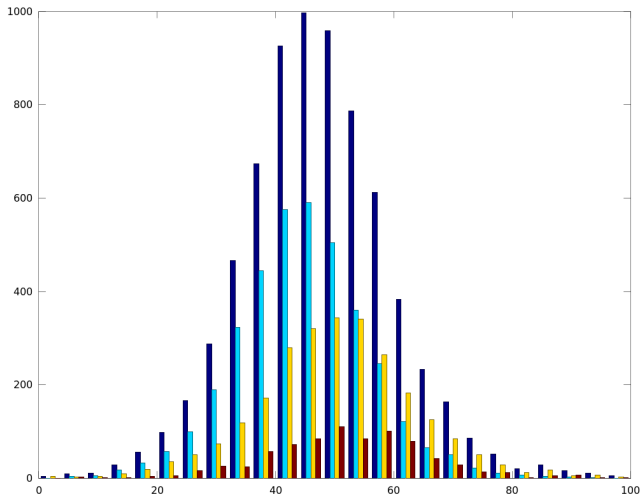
The Contact Matrix



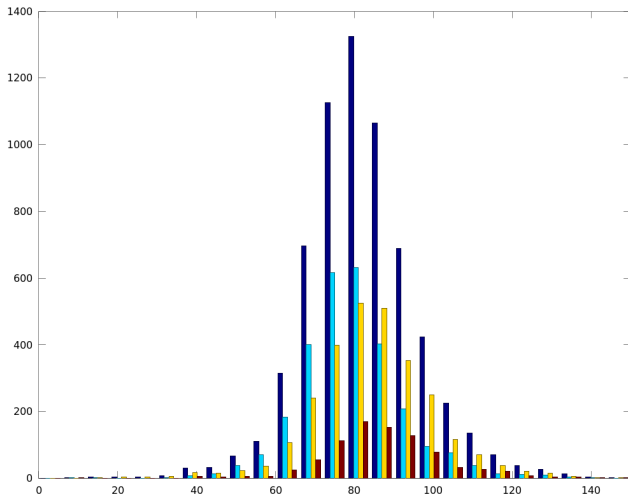
The Incidence Curves I



The Incidence Curves II



The Incidence Curves III



The Incidence Curves III

