

```

function [Sh, Ih, Rh] = simulate_absir(M, Iv0, T, infection_rate, recovery_rate)
% Simulate agent-based transmission model. Uses a graph to represent social
% connectivity between agents.
%
% Note: You may find it helpful to summarize the state history matrices via
% summation. For instance sum(Ih, 1) will return the total number of
% infected persons at each timestep in the simulation.
%
% Inputs
% M (square matrix): Adjacency matrix
% Iv0 (column vector): Initial infection state
% T (integer): Number of timesteps to simulate
% infection_rate (float): Infection rate (probabilistic)
% recovery_rate (float): Recovery rate (probabilistic)
%
% Returns
% Sh (matrix): Susceptible state history
% Ih (matrix): Infected state history
% Rh (matrix): Recovered state history

% Setup
dim = length(Iv0); % Dimensions of initial state
Ih = zeros(dim, T); % Infection history
Ih(:, 1) = Iv0; % Record the initial state to infection history
Rh = zeros(dim, T); % Recovery history

% Construct an "action" helper function
function [I, R] = action(I, R)
    % Compute infection probabilities based on the social graph
    v_eff = infection_rate * M * I;
    v_pr = v_eff ./ (1 + v_eff);

    % Draw random values
    v_infect = rand(dim, 1) <= v_pr;
    v_recover = rand(dim, 1) <= recovery_rate;

    % Infect non-recovered individuals
    I = I | v_infect & (~R);
    % Recover infected individuals
    R = R | (I & v_recover);
    I = I & (~R);
end

% Run simulation
for i = 2:T
    [Ih(:, i), Rh(:, i)] = action(Ih(:, i-1), Rh(:, i-1));
end

% Compute susceptible history
Sh = ones(dim, T) - Ih - Rh;
end

```