

19CSE337 Social Networking and Security

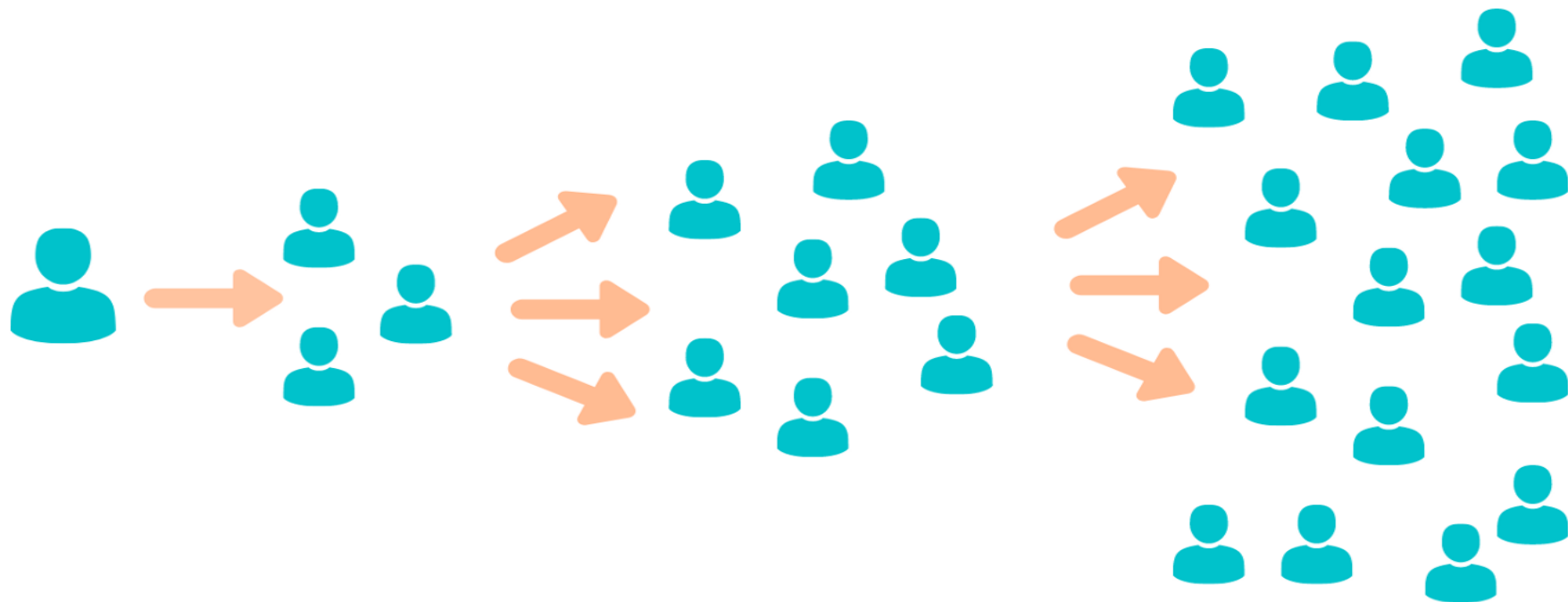
Lecture 16

A vertical sidebar on the left side of the slide, featuring a dark blue background with a grid of various white and light blue icons. These icons include a television, a camera, a lightbulb, a hand, a speech bubble, a padlock, a shopping cart, a smartphone, a person, a Twitter bird, and a lowercase 't' (Tumblr).

Topics to Discuss

- Contagion

Contagion





Contagion

- Contagion means spreading of diseases.
- In social network perspective, it is information spreading from person to person.
- Also termed as information diffusion or simply diffusion.
- Can be used to model infection spread, economic flow etc.



Contagion

- Social networks are prominent tool for information diffusion in society.
- Modelling and predicting information diffusion is very important.
- It is very useful in predicting, marketing, advertising, political campaigns etc.
- Tie strength is very important in information diffusion.



Social Contagion

- Phenomena or various processes that depend on individual propensity to adopt and diffuse knowledge, ideas and information.
- The spread of behaviours, attitudes and effect through crowds and other types of social aggregates from one member to other.



Simple Contagion

- Single exposure is sufficient for transmission.
- Eg; disease, information etc
- Various epidemic models are there to model simple contagion.
- Eg; SIR, SIS etc



Complex Contagion

- Single exposure is not sufficient for a transmission.
- Multiple exposures are needed.
- The behaviour is risky or technology requires coordination.
- Complex contagion can be modelled using threshold model, generalized epidemic model, diffusion percolation etc.
- Most of the real world networks are described by complex contagion models (spread of technology, trends in technology etc).



Simple Contagion Model

SIR Model

- S-Susceptible (groups under observation).
- I-Infected (already done).
- R-Recovered (recovered from the effect).
- S-I-R model is capable of predicting the number of people in S,I,R stages respectively.
- It is a compartmental model introduced in 1927.



SIR Model

- It is a differential equation model.
- There are dependent variables and independent variables to model disease situation.
- Dependent variables: people in each group, total population.
- Independent variable: Time t , measured in days/hrs.



SIR Model

- Susceptible individuals may become infected.
- This measures rate of transfer from susceptible to infected.
- The rate of transfer depends on number of individuals in each compartments and frequent contacts.
- The rate of contact which converts susceptible to infected is β .
- This β value can be reduced through quarantine, lockdowns etc.



SIR Model

- Rate of recovery: infected to recovered.
- Also used as rate of mortality.
- R_0 : basic reproduction number. The mean number of infections caused by a single infected individual over the course of their illness.
- R_0 is the ratio between β and γ .
- R_0 also known as contacts per infection.
- A decrease in β reduces R_0 .
- If $R_0 > 1$, spread will increase in the upcoming days else reduced.

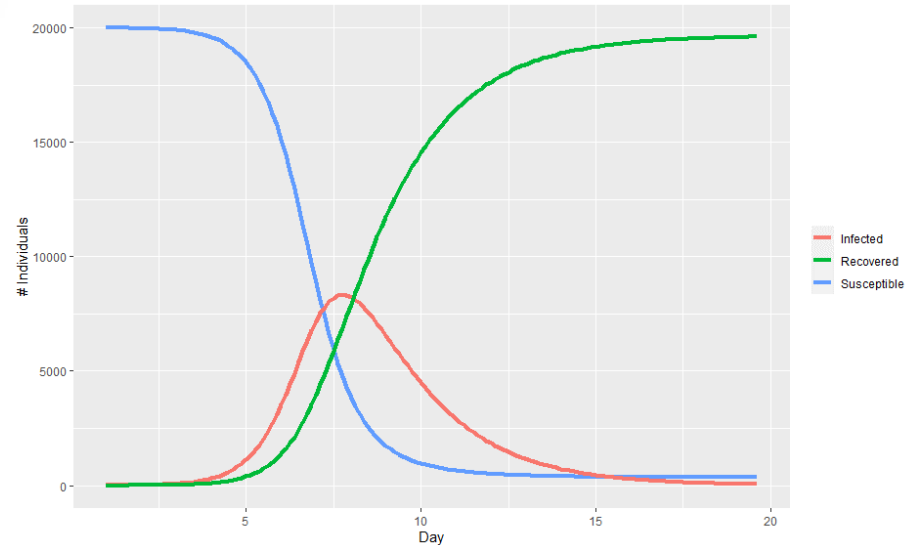


SIR Model

- Expressing S , I , R , in terms of t as $S(t), I(t), R(t)$.
- The total population $N = S(t) + I(t) + R(t)$.
- To form a differential equation, express S , I and R as rate of change of S, I, R with respect to time t as: $dS/dt, dI/dt, dR/dt$.

SIR Model

- The equations are:
 - $dS/dt = -\beta SI$ (-ve value as susceptible change to infected).
 - $dI/dt = \beta SI - \gamma I$
 - $dR/dt = \gamma I$



SIR Model

- From above equations:

- $dS = -\beta SI * dt$

- $dI = (\beta SI - \gamma I) dt$

- $dR = \gamma I * dt$

- Change in S, $dS = S_{i+1} - S_i$

- $S_{i+1} - S_i = -\beta S_i I_i * dt$

- $S_{i+1} = S_i - (\beta S_i I_i) dt$

SIR Model

- Similarly,

$$I_{i+1} = I_i + (\beta S I_i - \gamma I_i) dt$$

$$R_{i+1} = R_i + \gamma I_i dt$$



Limitations

- Not including any other parameters other than S,I,R (other models include carriers, vaccinated etc).
- Assume homogenous mixing of population (SxI). In real world, human networks are highly local.



Problem

- Consider the statistics of an epidemic in country X. $S=1000$, $I=10$, $R=0$. The time t is measured in days. Each infected person would make infecting contacts every two days. The average recovery period is 3 days. Find S , I , and R on the second day of epidemic.



Thanks.....