

# Lab 2 Summary

**Consequences of Heterogeneity** 

Roger Ying MMED 2018

### Assumptions

- 1. Everyone is the same except for their contact rates, which are gamma distributed
- 2. Disease transmission occurs one at a time (similar to event-driven model from yesterday)
- 3. Disease recovery rate is exponentially distributed



### Step 1 – Low variance (homogeneous)



contact rate (1/day)

cumulative # infected (final size)

Beta.mean	2
Beta.var	0.001
Runs	5
Pop.size	100



### Step 2 – Increase number of runs



Beta.mean	2
Beta.var	0.001
Runs	50
Pop.size	100



### Step 3 – Change variance





outbreak size distribution



2 Beta.mean 0.01 Beta.var 30 Runs Pop.size 100

cumulative # infected (final size)

#### outbreak size distribution



Beta.mean	2
Beta.var	5
Runs	30
Pop.size	100

cumulative # infected (final size)





time series







### Step 4 – Larger population size





outbreak size distribution



Beta.mean	2
Beta.var	0.01
Runs	30
Pop.size	500

cumulative # infected (final size)

#### outbreak size distribution



Beta.mean	2
Beta.var	5
Runs	30
Pop.size	500

cumulative # infected (final size)





30

20

10

0

0

Frequency





contact rate (1/day)

6

une senes

8 10

# Step 5 – Change mean beta $\rightarrow$ 0.7





outbreak size distribution



cumulative # infected (final size)



distribution of average R

8

6

15

10

5

0

0

Frequency

time series



outbreak size distribution



Beta.mean	0.7
Beta.var	10
Runs	30
Pop.size	100

cumulative # infected (final size)



## Summary

• Heterogeneous populations have:





500





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