

Back of the Envelope
Modelling of Infectious
Disease Transmission
Dynamics for
Veterinary Students

Back of the Envelope Modelling of Infectious Disease Transmission Dynamics for Veterinary Students

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For my family.

R

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SIR

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use

1.1 Introduction

1.2 Draw a flow chart



Fig. 1-1 *The beginnings of a basic flow chart consisting of three host states: susceptible (S), infected (I) and recovered (R).*

S

S

is

R

S I I
S I

λ

I R

I R

δ

-
-

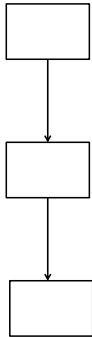


Fig. 1-2 A basic flow chart consisting of three host states: susceptible (*S*), infected (*I*) and recovered (*R*). Hosts in state *S* can move into state *I* as the result of **transmission** (the rate of transmission is represented by λ). Recovery is represented by a **transition** from box *I* to box *R* (the arrowed labeled δ).

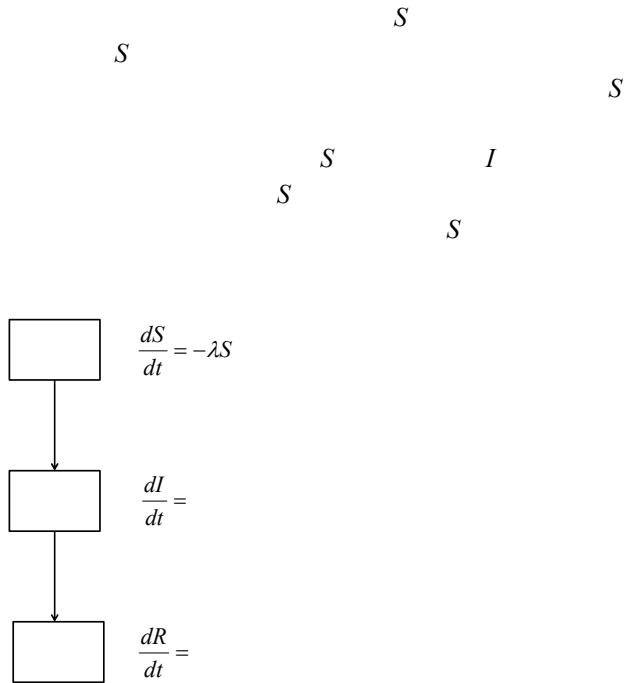
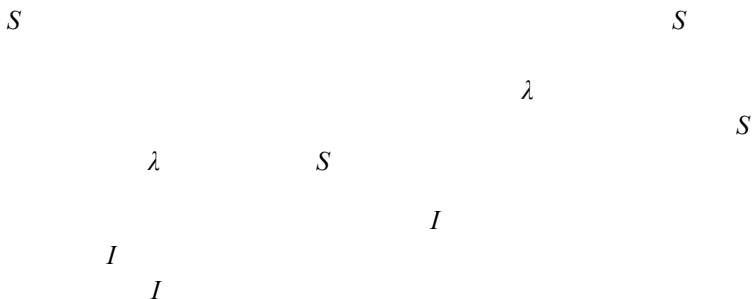


Fig. 1-4 The right-hand side of the equation for S is constructed by first noting that there is a single arrow leaving box S . This means that there is only one term to the right of the equals sign and it must be negative. We supply the term by multiplying the arrow label (λ) by S , which is the label of the box from which the arrow originates.



λ S λS
 $\delta,$

$I.$

$$\frac{dI}{dt} = +\lambda S - \delta I$$

$$\frac{dS}{dt} = -\lambda S$$

$$\frac{dI}{dt} = \lambda S - \delta I$$

$$\frac{dR}{dt} =$$

S S $I.$ λS
 S λS I
 S I λS λI
 λS $S.$ λI I
 R R R R I δI
 δ

$$\frac{dS}{dt} = -\lambda S$$

$$\frac{dI}{dt} = \lambda S - \delta I$$

$$\frac{dR}{dt} = \delta I$$

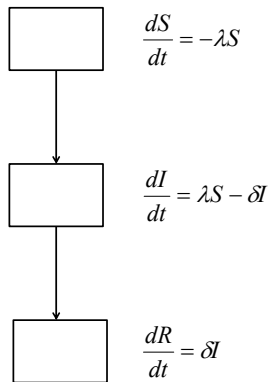


Fig. 1-5 A complete SIR model of avian influenza in a broiler house. S = number of susceptible birds in a single broiler house, I = number of infected birds, R = number of recovered birds, λ = the force of infection, δ = the recovery rate.

1.4 Using the recipe for other systems

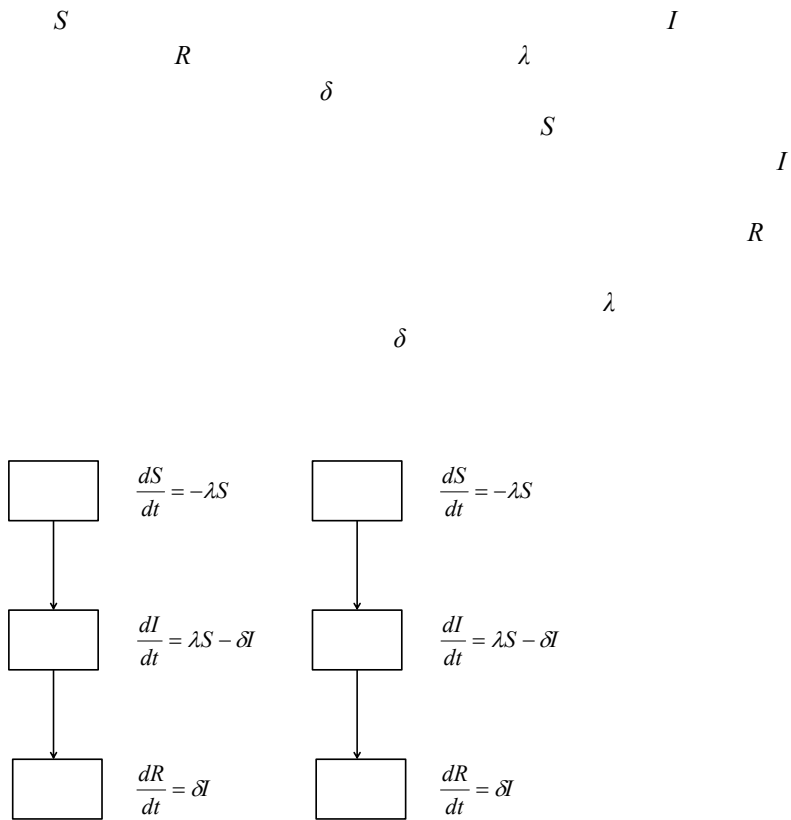


Fig. 1.6 *SIR* model of how avian influenza infection spreads between broilers in a single broiler house; *SIR* model of how avian influenza spreads between broiler houses.

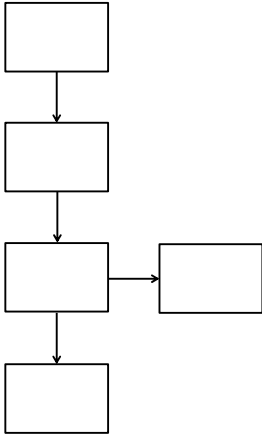
1.5 Adding new boxes

E

D

I

E *I* *E*
 I



$$\frac{dS}{dt} = -\lambda S$$

$$\frac{dE}{dt} = \lambda S - \sigma E$$

$$\frac{dI}{dt} = \sigma E - \alpha I - \delta I$$

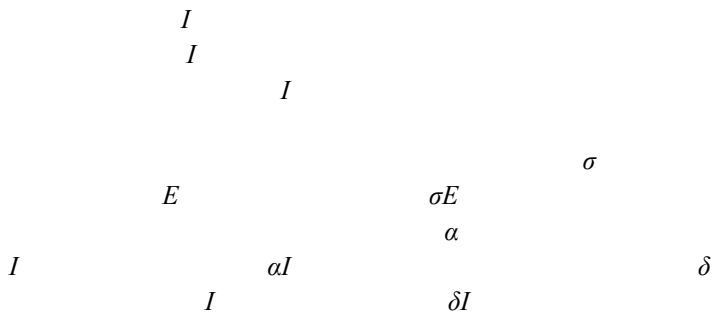
$$\frac{dD}{dt} = \alpha I$$

$$\frac{dR}{dt} = \delta I$$

Fig. 1-7 An SEIR model of avian influenza in a broiler house. *S* = number of susceptible birds in a single broiler house, *E* = number of infected-but-not-yet-infectious birds, *I* = number of infectious birds, *R* = number of recovered birds, λ = the force of infection, σ = rate at which infected birds become infectious, δ = the recovery rate.

E

E



$$\frac{dI}{dt} = \sigma E - \alpha I - \delta I$$

1.6 Suggested further reading

et al.

1.7 Summary

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-
-
-

1.8 References

- Dynamics and Control*) *Infectious Diseases of Humans.*
- Infectious Diseases: Epidemiology and Ecology.* : *Parasitic and*
- Diseases in Humans and Animals.* *Modelling Infectious*
- to Infectious Disease Modelling.* *An introduction*
- et al*

2.1 Introduction

transition

