

Modelling Hepatitis C virus infection and treatment impact through serological surveillance data

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1 A Global Health Challenge

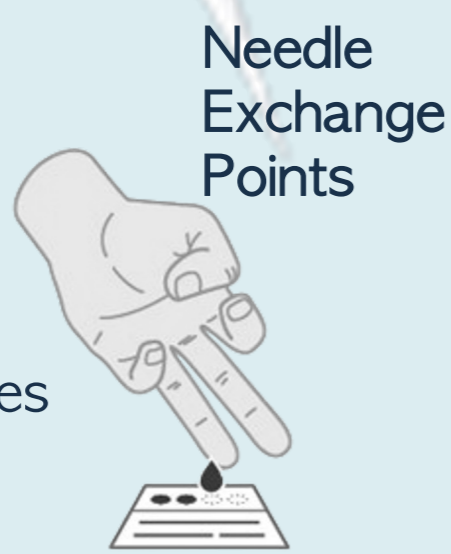
- Hepatitis C is a blood-borne virus that can cause **life-threatening liver damage** over time.
- In the UK, the Hepatitis C virus (HCV) is primarily transmitted through the sharing of drug-injection equipment, making **people who inject drugs (PWID)** a key at-risk group.
- From 2015 to 2022, the number of people living with chronic HCV infection in England has fallen dramatically in the general adult population by 51.6% and is now estimated at **62,600** (1).

2 New Treatment, New Hope

- The advent of **effective new treatments** for HCV in 2015 has transformed the landscape for HCV control.
- This has enabled the formulation of an **elimination strategy** led by the World Health Organisation (WHO) based on a **reduction in HCV incidence**.

3 Evidencing Elimination

- The **unlinked anonymous (UAM)** survey annually monitors blood-borne viruses among PWID in England, Wales and Northern Ireland.
- PWID are recruited at **needle exchange points** to participate in the UAM survey – this involves taking a **dried blood-spot (DBS)** sample and completing a **biobehavioural questionnaire**.



- DBS samples are tested for **antibody (Ab)** and **ribonucleic acid (RNA)** status – this can be leveraged to **differentiate between current and past HCV infections**.
- This information facilitates the development of a **multi-state model** characterising HCV transmission dynamics.

UK Health Security Agency Labs



Monitoring incidence directly is challenging – cross-sectional biobehavioural serosurveys, which capture information on infection status and duration of exposure, provide a **means to estimate incidence indirectly** and can be conducted serially to track changes over time.

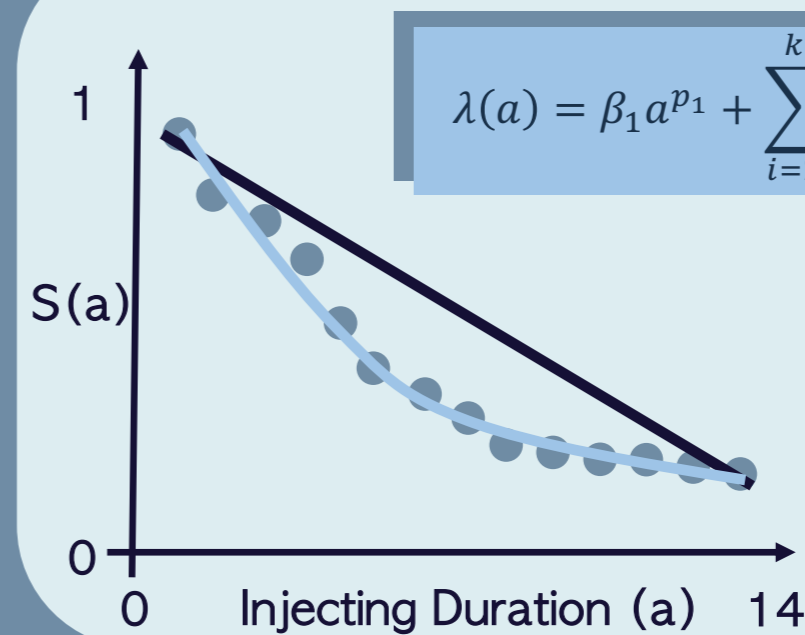
Incidence is the rate at which new cases of a disease occur in a specified population during a defined time period – it measures the risk of contracting the disease.

Primary incidence rates among recent initiates rose between 2011 and 2022, from 0.21 (95% CrI: 0.20–0.22) to 0.32 (95% CrI: 0.30–0.33), coinciding with increased needle-sharing behaviour – conversely, **rates among long-term injectors declined**.

Multi-state models serve as robust tools to **monitor progress toward WHO elimination goals** and **inform public health strategies**.

- Transition parameters are modelled over **time at risk (a)** and **calendar time (t)** – information obtained from the biobehavioural questionnaire.

Aim 1: Flexibly model transition parameters



- Fractional polynomials** (2) are chosen from a suite of flexible model parametrisations (3) to model the transition parameters.



Spontaneous clearance rates are higher than widely reported – consistent with studies including RNA-negative individuals.

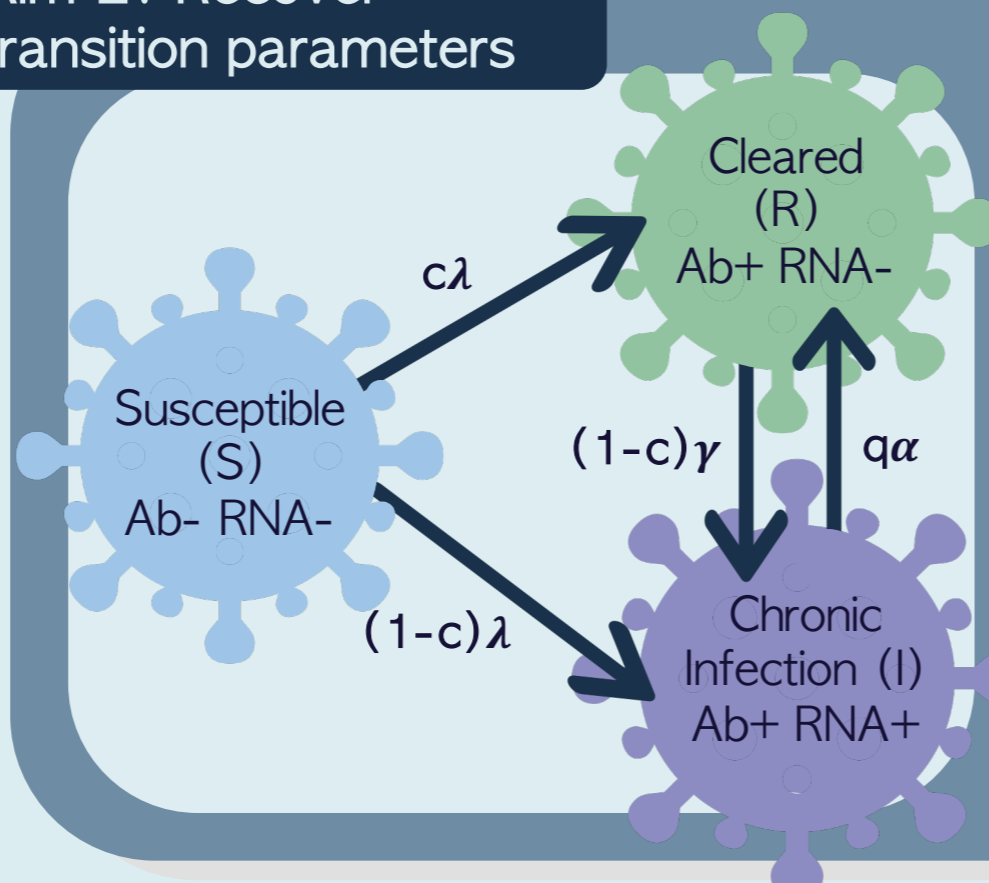
Key Findings

HCV Working Groups



Aim 2: Recover transition parameters

- The multi-state model is implemented within a **Bayesian framework** – integrating external information to inform our prior beliefs about the unknown parameters.



The multi-state model is governed by a system of ordinary differential equations.

$$\begin{aligned} \frac{dS}{dt} &= -\lambda S \\ \frac{dI}{dt} &= (1-c)(\lambda S + \gamma R) - q\alpha I \\ \frac{dR}{dt} &= c\lambda S + q\alpha I - (1-c)\gamma R \end{aligned}$$

λ : primary incidence rate
 γ : reinfection rate
 c : spontaneous clearance rate
 α : treatment uptake rate
 q : sustained virologic response rate

Biostatistician (me)

