

# Health Analytics Workshop

## **MODELLING NEGLECTED TROPICAL DISEASES IN INDIA**

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# DAY 2



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## Day 2

9:30 am	Plenary talk by Dr Tanu Jain, NCVBDC
10:15 am	Plenary talk by Dr Vinod P. Choudhary, NCVBDC
11:00 am	Tea break
11:15 am	Recap of Day 1 and Introduction to Day 2
11:45 am	Break-out session
1:00 pm	Lunch
2:00 pm	Break-out session (continued)
3:15 pm	Tea break
3:30 pm	Interactive review of challenges
5.00 pm	Close

11:15 – 11:45 am  
**RECAP DAY 1**





# Recap of Day 1

1. Introduction to the epidemiology and control of VL
2. Draft initial research question in pairs
3. Development of compartmental models + template VL model
4. Choose and refine research question + design conceptual model in groups
5. Data needs + strategies to address lack of data

Questions about:

- Any of the workshop sessions on VL so far
- The template VL model
- Parameter values in the hand-out





## Goal of Day 2

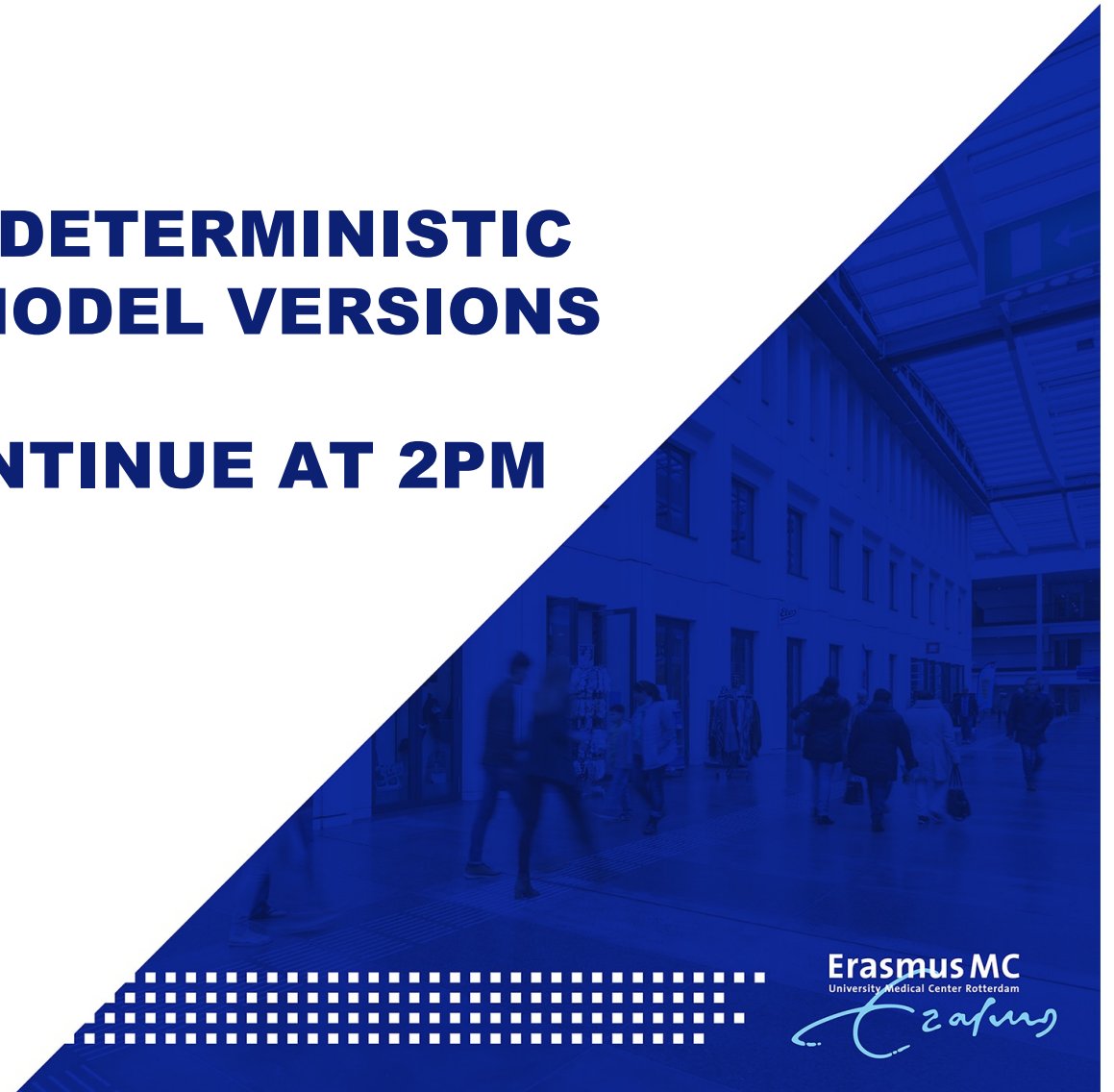
- Implement two versions of your group's conceptual model:
  - Deterministic
  - Stochastic (if time allows, prioritise the deterministic version)
- Get some hands-on experience working with the *pomp* package in R
  - Based on a template VL model that you can adjust yourself
- Draft a visualisation of the model-predicted answer to the research question
  - Discuss results and encountered challenges



11:45 am – 3:15 pm

# **BREAK-OUT: BUILD DETERMINISTIC AND STOCHASTIC MODEL VERSIONS**

**LUNCH AT 1PM, CONTINUE AT 2PM**



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# Link between deterministic and stochastic model versions

Deterministic model:

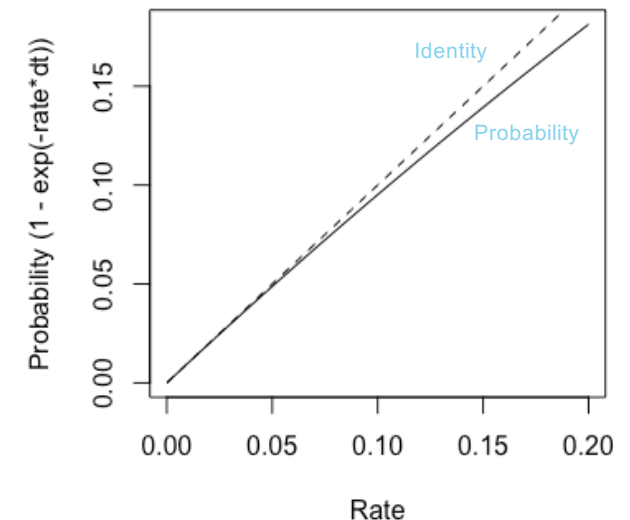
- Fractions of population transition between compartments

Stochastic model:

- Discrete number of individuals transition between compartments
  - Multinomial draws for number of transitions per  $\Delta t$
- Define transition probability  $P_{X \rightarrow Y}$  from compartment X to Y, given all exit rates  $\rho_{X \rightarrow i}$  from X:

$$P_{X \rightarrow Y} = \frac{\rho_{X \rightarrow Y}}{\sum_i \rho_{X \rightarrow i}} \cdot (1 - \exp[-\Delta t \cdot \sum_i \rho_{X \rightarrow i}])$$

- For VL model here, we take fixed time step size  $\Delta t$  (Euler steps) of 1 day
  - Chosen such that  $\Delta t \times$  the highest rate in the human part of the model is  $\ll 0.1$ , and  $\ll 1$  in the sandfly model (fly dynamics are almost instantaneous compared to humans in this VL model)







# Notes on sandfly sub-model

The stochastic model is actually a hybrid model:

- Sandfly model is implemented deterministically
- Little point in simulating individual sandflies
  - Dynamics are almost instantaneous, compared to humans
  - We have no idea how many sandflies there are in absolute terms and how many of them are relevant for transmission





# Pomp

Package for Partially Observed Markov Processes (<https://kingaa.github.io/pomp/>)

- Convenient to develop model code that will be translated to C and compiled
- A lot of additional functionality that we will not use

## Ingredients

- C-snippets of model code (strings that will be translated to C code and compiled)
  - Function to set initial values
  - Function to calculate derivatives
  - “Accumulator” variables (stochastic model only) that keep track of how many individuals transitioned (useful for calculating incidences)
- Vectors of state names (i.e., compartments) and parameter names
- Named vector of fixed parameter values (i.e., these have the same value throughout a simulation)
  - Including initial values (must have same names as states, but with suffix “\_0”)
- Optional: table of time-varying parameter values

Stochastic model only:

- Vector of accumulator names





**Go through example script**





# Break-out instructions

Code repository: <https://github.com/Ananthu89/NDMC-VL-Workshop-IITB-2023>

- Run the example script
  - Have a look at the C-snippets and see if you understand what's going on
- Develop code in R for a deterministic (and stochastic?) version of your group's conceptual model
  - Make a copy of what you want to work on and adapt the copy, leaving the original intact
- Perform sense checks and debug any discrepancies between the two model versions
  - Refer to readme.md at the above web address
- Produce a plot that illustrates the answer to your group's research question
- Prepare 2 slides to guide discussion about your model during the next session (send to trainers)
  - Results / plot + interpretation
  - Encountered challenges





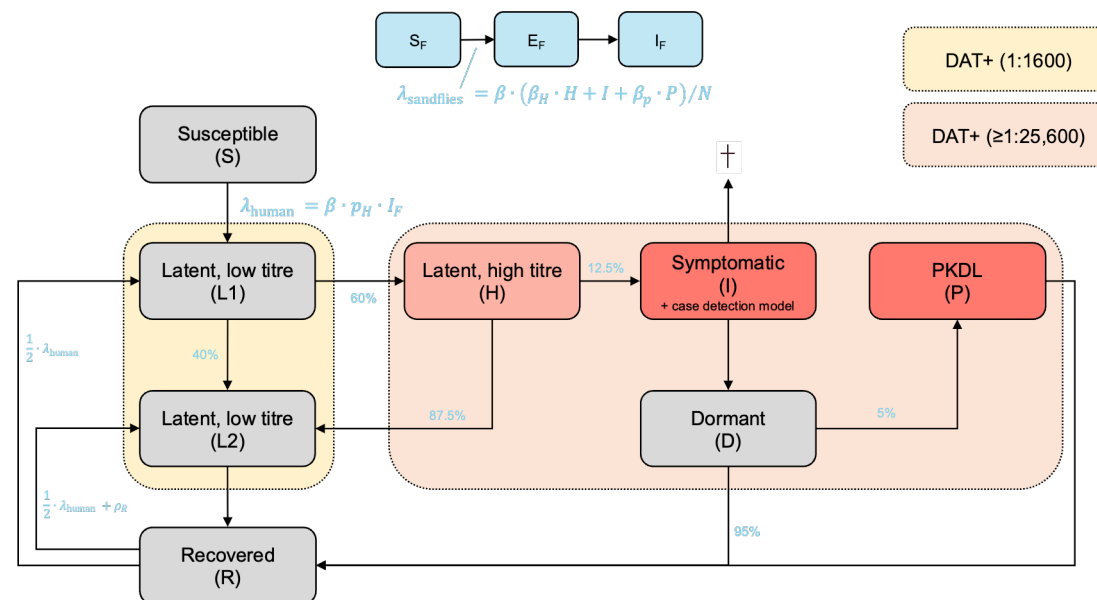
# Sense checks to perform

- Does the population size remain stable?
- Do the deterministic and stochastic model agree?
- ...



# Debugging strategy

- Temporarily simplify the two model versions and compare them again
  - Set a strategically chosen rate to zero, e.g., rate from H to I (but make sure that H is at least somewhat infective towards sandflies then)

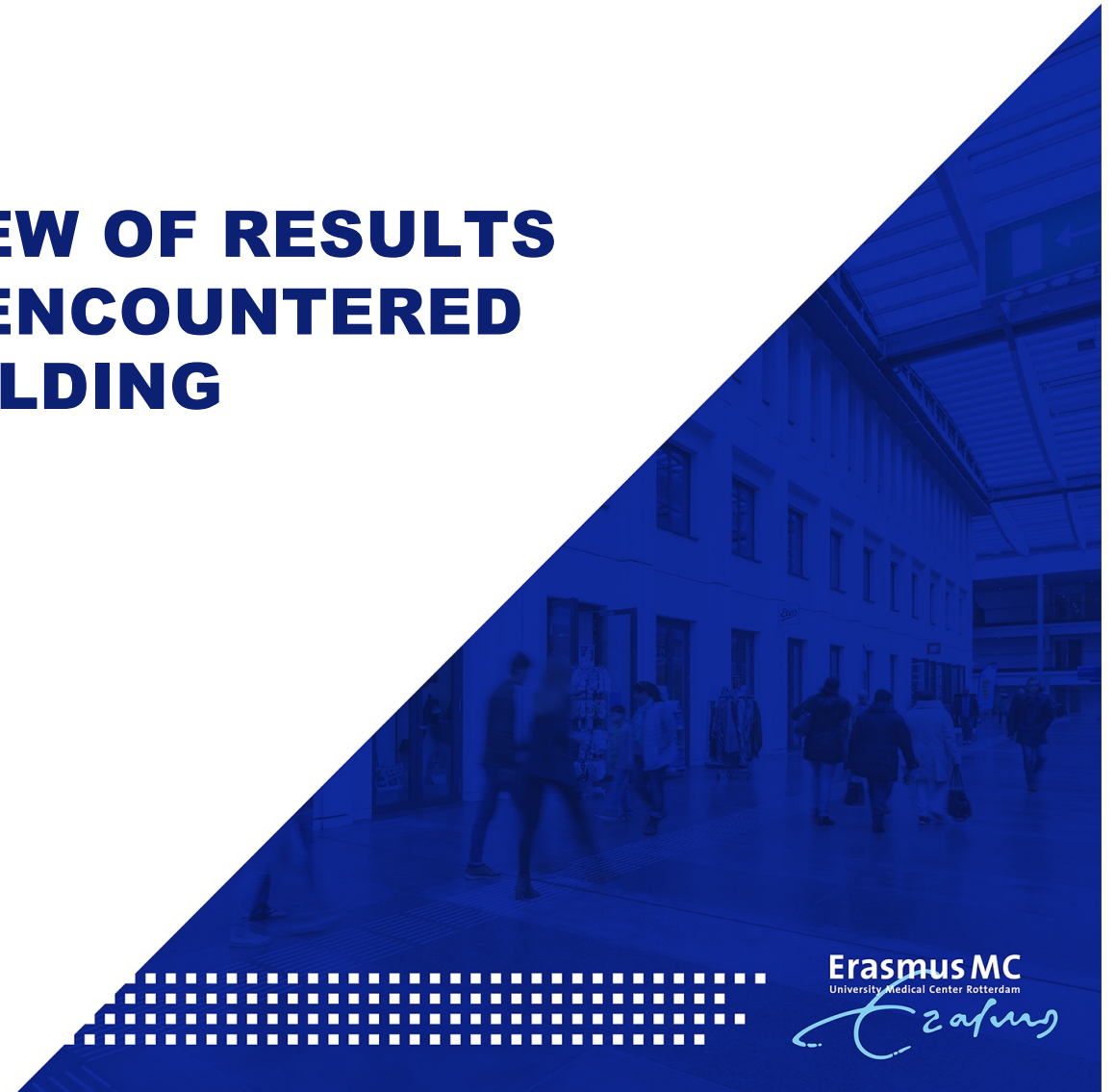


3:15 pm – 3:30 pm  
**TEA BREAK**



3:30 – 5:00 pm

# **INTERACTIVE REVIEW OF RESULTS AND CHALLENGES ENCOUNTERED DURING MODEL BUILDING**



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