



SACEMA

Real-world example: Initial spread of COVID-19 in South Africa

Juliet Pulliam, PhD

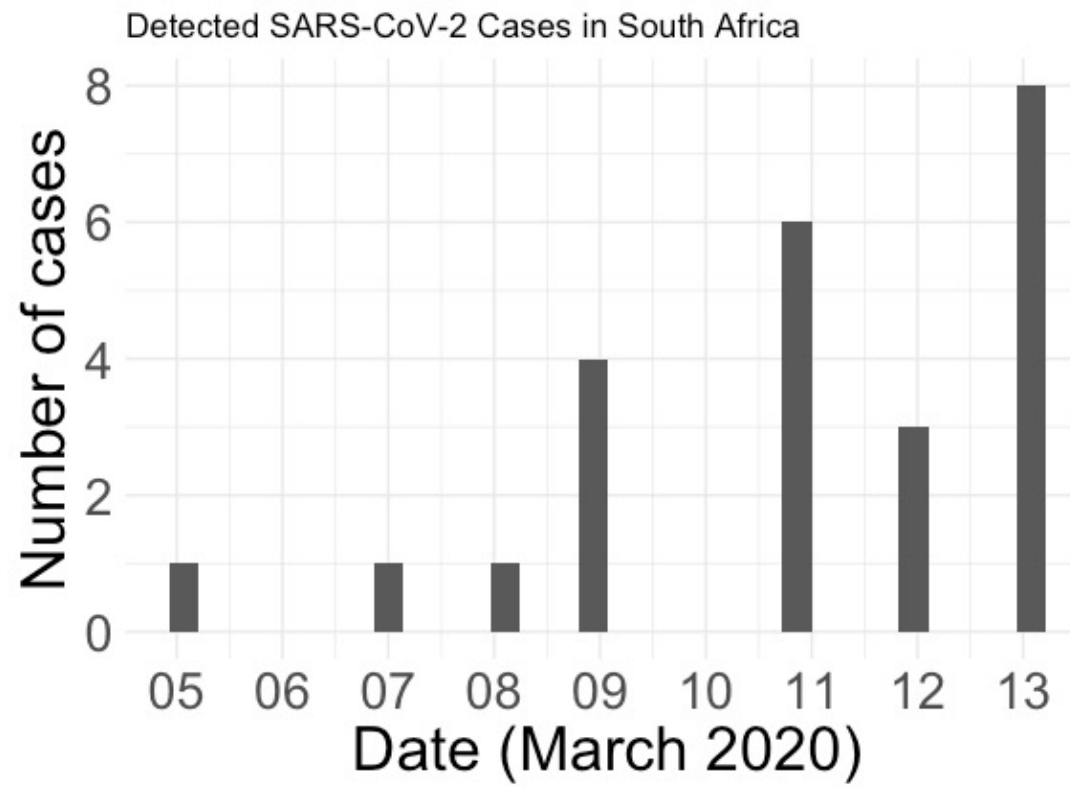
SMART Meeting

2022-02-15



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As of 13 Mar 2020:

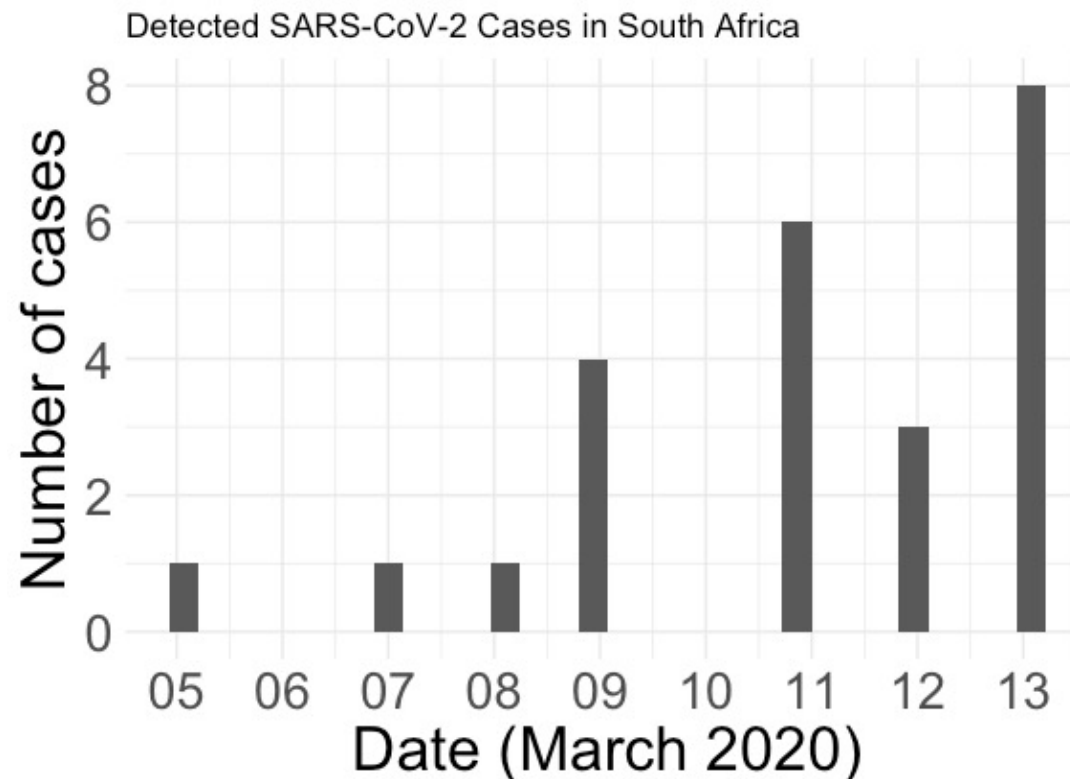




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Question (13 Mar):

“When do we expect to see the first 1,000 and 10,000 cases?”





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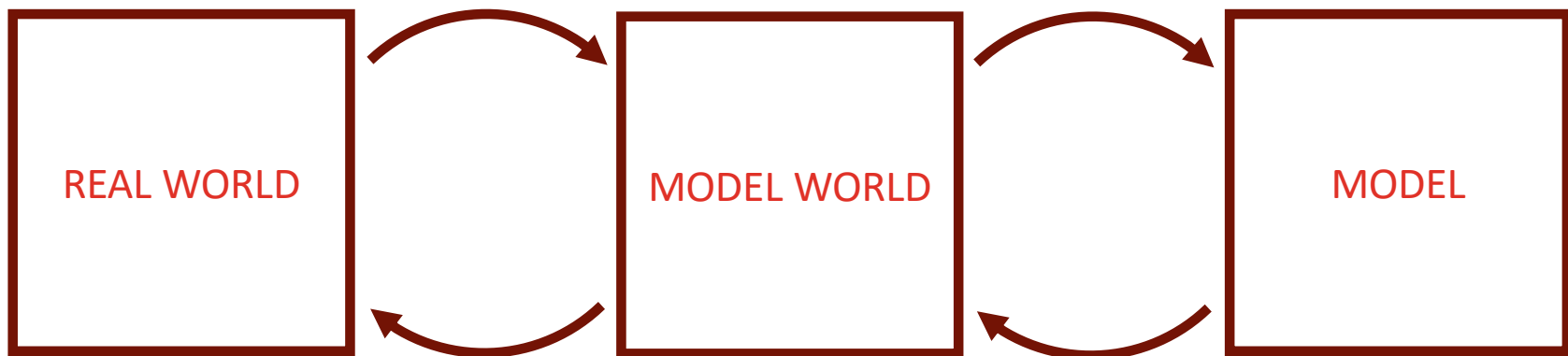


Figure courtesy of the International Clinics on Infectious Disease Dynamics and Data (ICI3D) Program
<https://doi.org/10.6084/m9.figshare.5044606.v3>



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Question (13 Mar):

“When do we expect to see the first 1,000 and 10,000 cases?”

DEVELOPING A MODEL WORLD

Data-limited situations require simple models

What was known?

Initial estimates available of the basic reproduction number and time between successive cases (serial interval)

No interventions had been put in place at the time

The virus was introduced in SA in early March (24 cases detected to date)



SMART, 2022-02-15

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DEVELOPING A MODEL WORLD

Data-limited situations require simple models

Approach

Branching process model

Seed introductions on the same timeline as known infections

Assume constant detection probability, with about half of infections being detected

Assume the epidemic trajectory continued without substantial intervention



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STRUCTURE OF THE REPORT

Projection of early spread of COVID-19 in South Africa

Prepared by Carl A.B. Pearson (SACEMA, LSHTM), Sheetal Silal (UCT), and Juliet Pulliam (SACEMA)

Last updated: 2020-03-14



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STRUCTURE OF THE REPORT

Summary paragraph with results

Figure

Assumptions and parameters

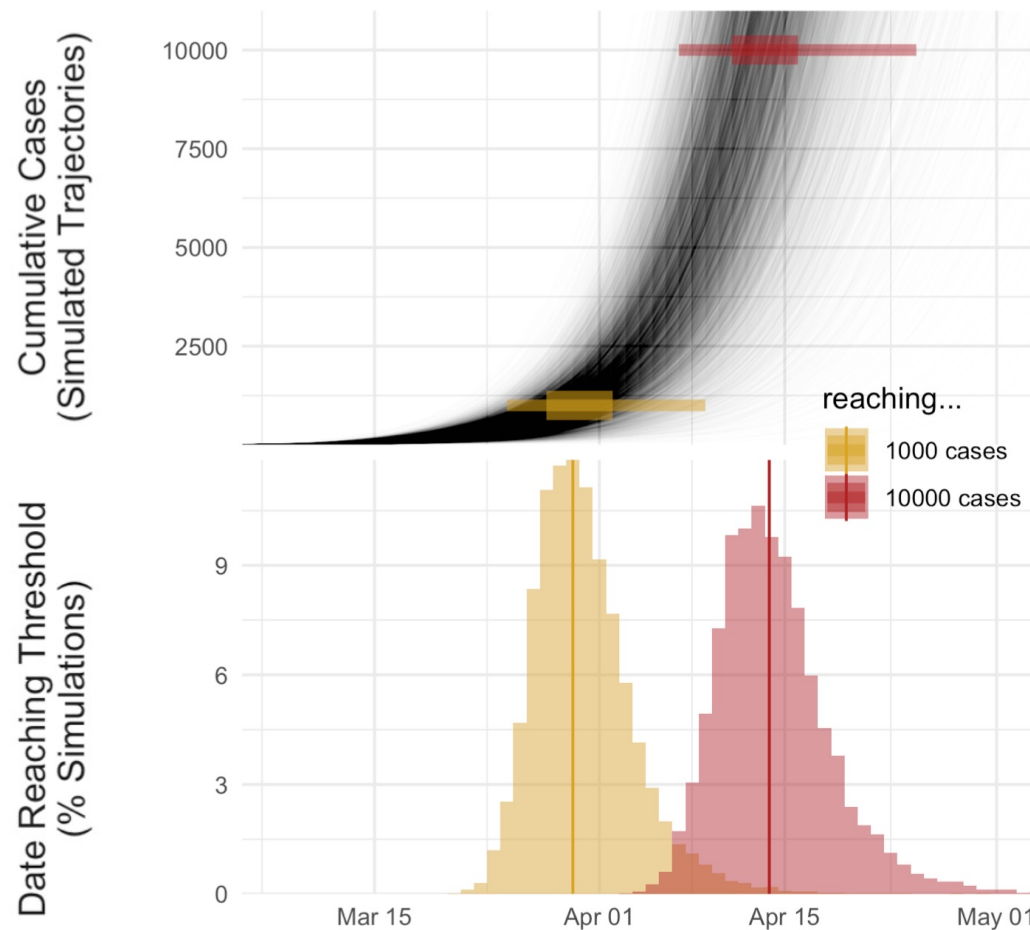
Contact

References



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IN AN ABSENCE OF INTERVENTIONS THAT CHANGE THE EPIDEMIC TRAJECTORY:

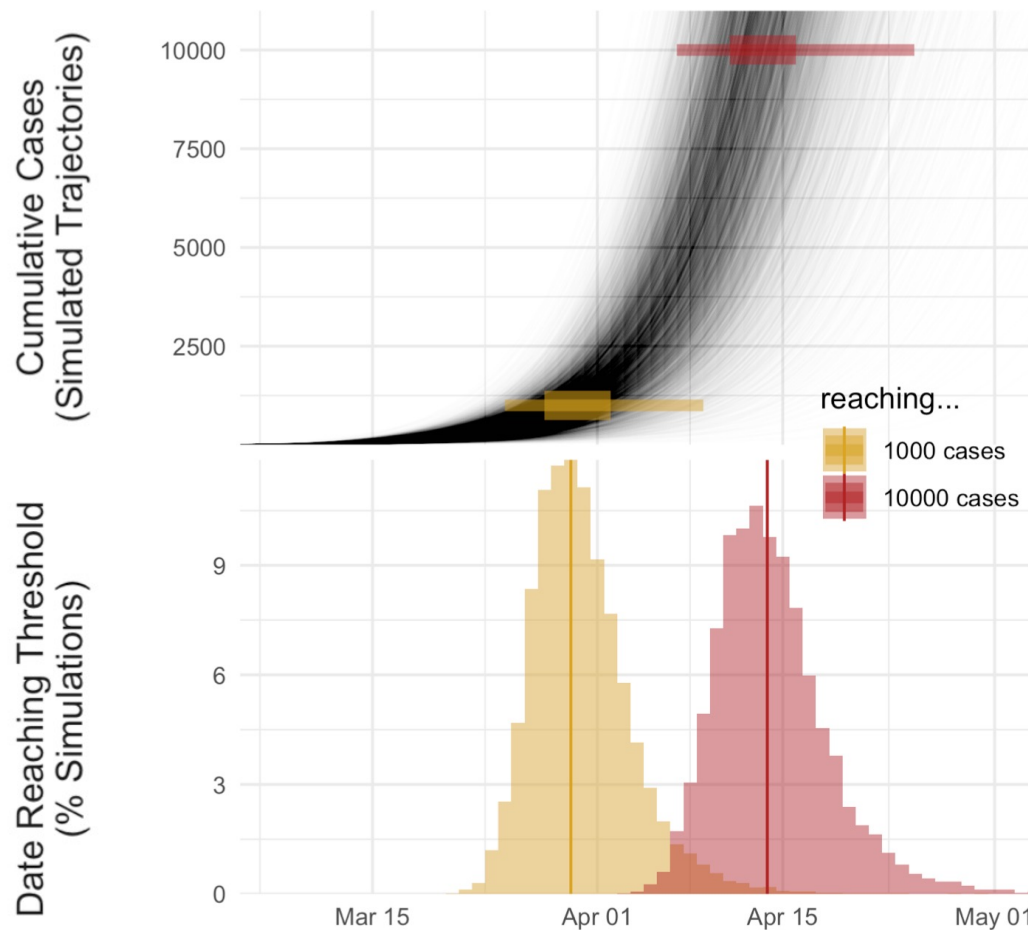


cumulative case count is...



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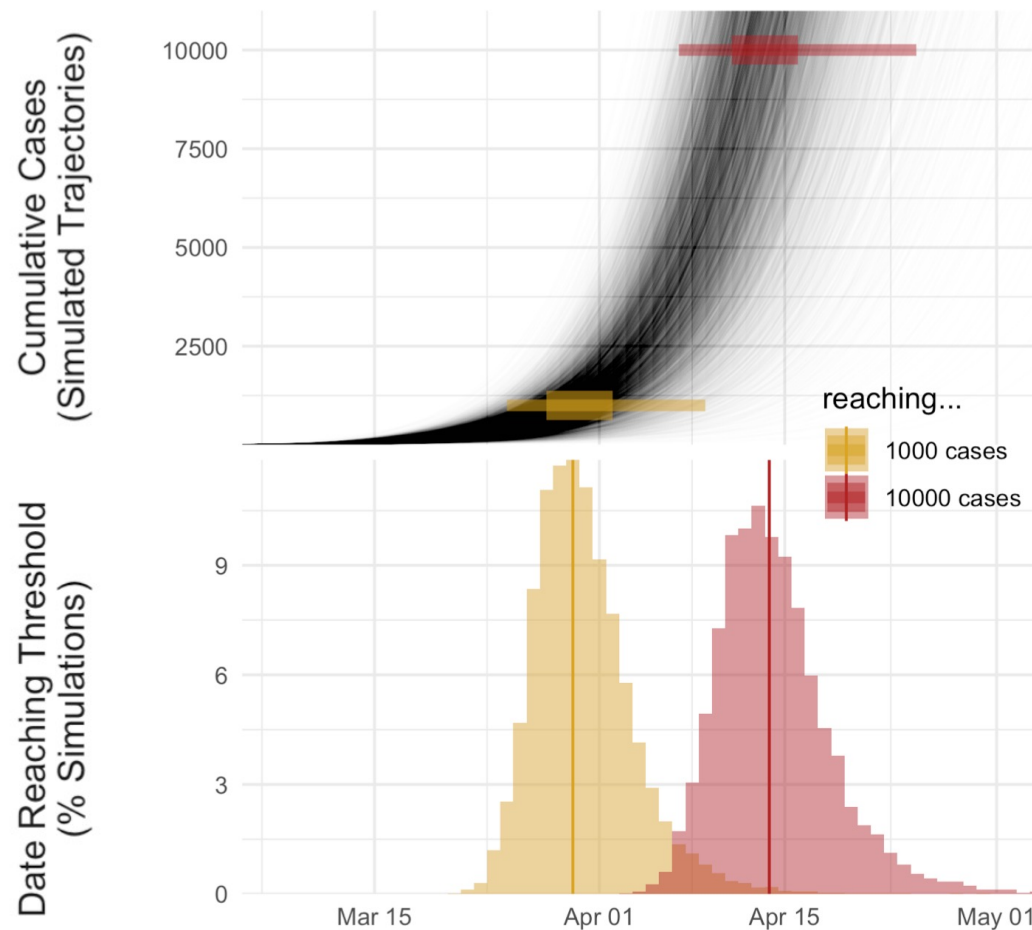
cumulative case count is...

likely to reach 1,000 cases
between 28 March and 2 April
(95% range: 25 March - 9 April)



SMART, 2022-02-15

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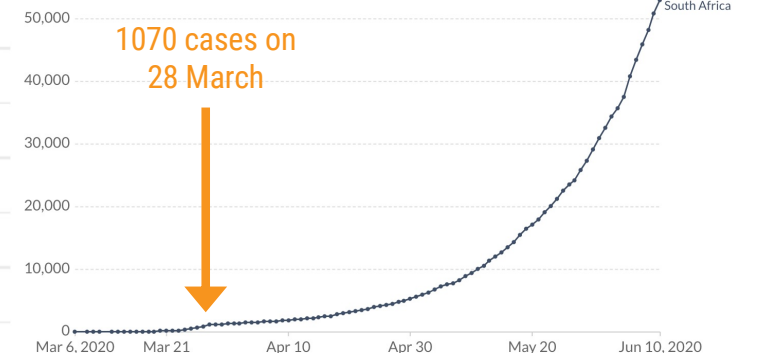
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between 28 March and 2 April
(95% range: 25 March - 9 April)

Total confirmed COVID-19 cases

The number of confirmed cases is lower than the number of total cases. The main reason for this is limited testing.

[LINEAR](#) [LOG](#)



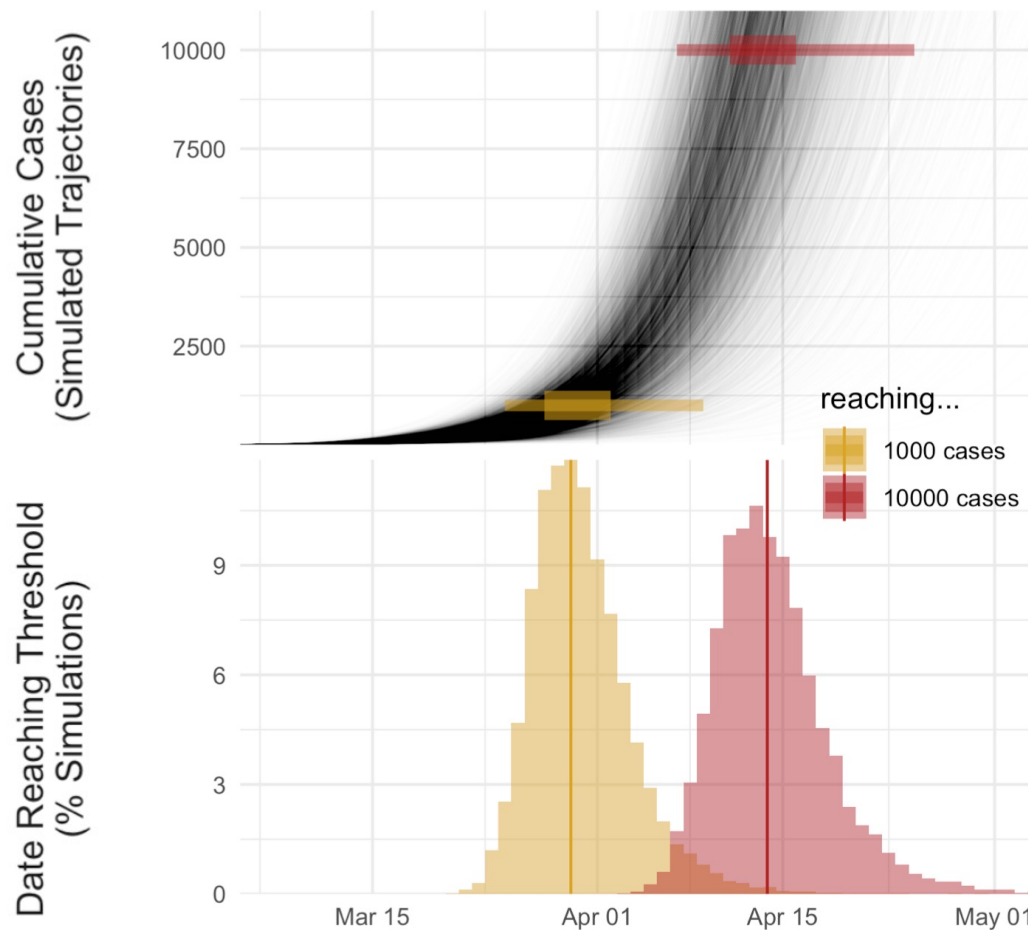
Source: European CDC - Situation Update Worldwide - Last updated 10th June, 12:00 (London time)

CC BY



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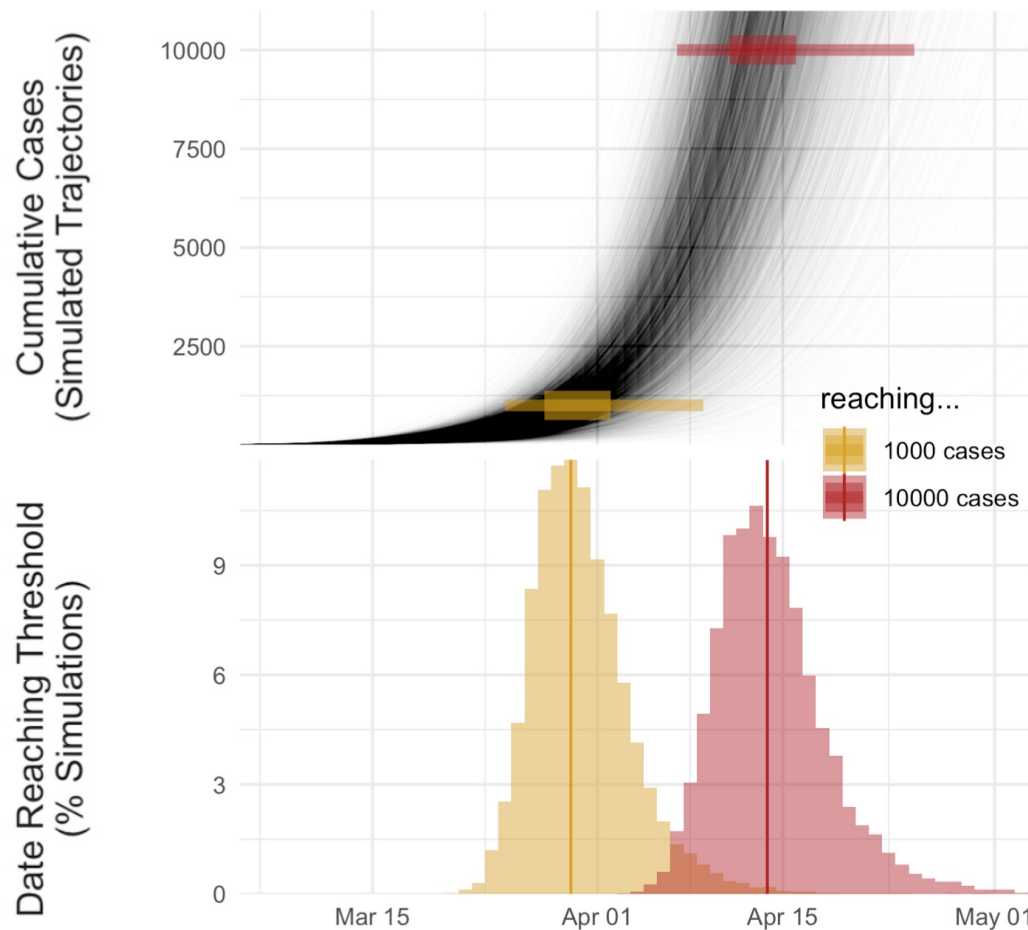
likely to reach 1,000 cases
between 28 March and 2 April
(95% range: 25 March - 9 April)

likely to reach 10,000 cases
between 11 April and 16 April
(95% range: 7 April - 25 April)



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IN AN ABSENCE OF INTERVENTIONS THAT CHANGE THE EPIDEMIC TRAJECTORY:



cumulative case count is...

likely to reach 1,000 cases
between 28 March and 2 April
(95% range: 25 March - 9 April)

likely to reach 10,000 cases
between 11 April and 16 April
(95% range: 7 April - 25 April)

10,015 on 11 May

(lockdown from 27 March)



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PACKAGE USED

<https://github.com/sbfknk/bpmodels>

CODE

<https://github.com/SACEMA/COVID10k>

SPECIFIC COMMITS FOR REPORTS

Projection of early spread of COVID-19 in South Africa:
328b02b6b841ab742e4a594cd170d2397bc3c24e

Early spread and hospital load for COVID-19 in South Africa:
d92a62d8502f8a7c6eac68f687845d0e5cf16aab



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STRENGTHS

Provides probabilistic view of the likely epidemic trajectory **in the short term**

Clear assumptions and quick implementation

LIMITATIONS

Assumes a constant epidemic trajectory

Does not account for potential interventions or behaviour change

Assumes a constant detection probability



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LESSONS

Start with a clear question

Build on what others have done (bpmodels package by Seb Funk, parameter estimates from Abbot, Bi, and Nishiura papers)

Decision-makers need clear, concise, and quick results