

Testing for COVID-19

Date Thursday 4 June 12.30pm

Presenters
Mr Robert Skeen
Prof Dominic Dwyer
Ms Ruth Luppino



This activity has been developed in partnership with Aboriginal Health & Medical Research Council of NSW and NSW Health



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Acknowledgement of Country

We recognise the traditional custodians of the land and sea on which we live and work.

We pay our respects to Elders past and present.



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Learning Outcomes

By the end of this webinar, participants will:

- Understand the importance of testing for COVID-19
- Understand the role of different types of testing in the diagnosis and management of cases with COVID-19
- Understand how to access testing and support for test interpretation



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Presenters



Robert Skeen
CEO, AH&MRC



Professor Dominic Dwyer
Director of Public Health Pathology, New
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Ruth Luppino
Practice Manager at Coonamble Aboriginal
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Robert Skeen

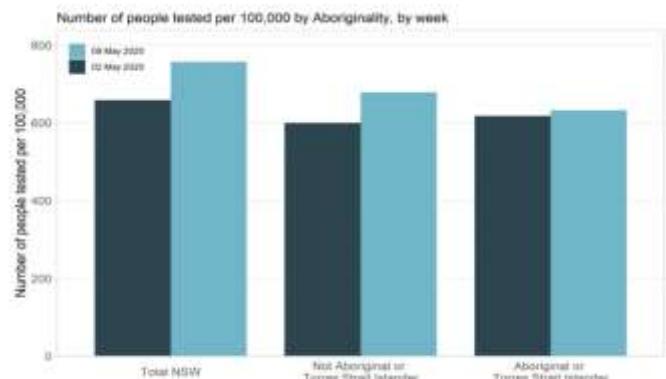
WELCOME & UPDATE



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Current Situation in NSW

- 30 cases = 1% of all cases in NSW
- No deaths
- Comparable rates of testing



*Total rates include people with unknown Aboriginality status.



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Testing criteria is now symptom based

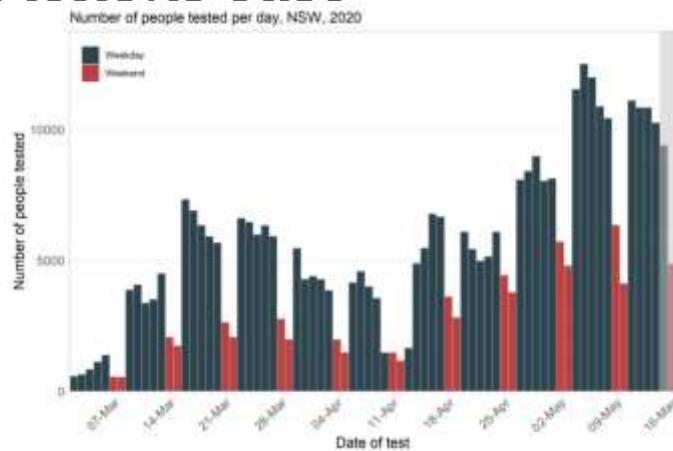
Test everyone with:

- Respiratory symptoms OR
- Unexplained fever



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High testing rates in NSW are needed as restrictions ease

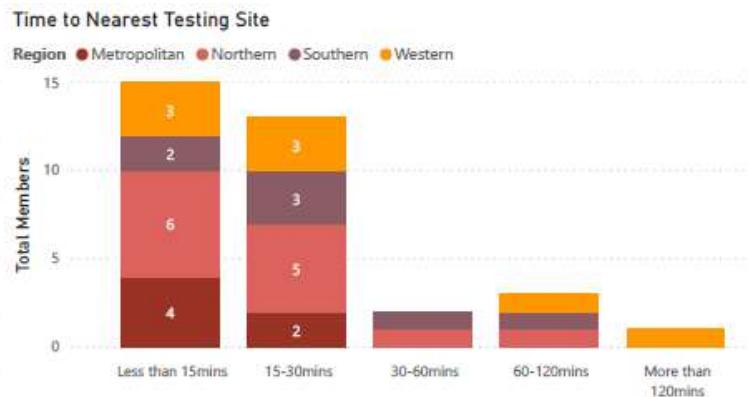


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Supporting Community Testing

Barriers include:

- Access to transport
- Limited hours at GP Respiratory clinics
- Long distances for regional members



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Dominic Dwyer

TESTING FOR COVID-19



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A new coronavirus (SARS-CoV-2) causing a new illness (COVID-19)

C ⓘ Not secure | virological.org/t/novel-2019-coronavirus-genome/319



Novel 2019 coronavirus genome

Novel 2019 coronavirus



edward_holmes

6 Jan 10

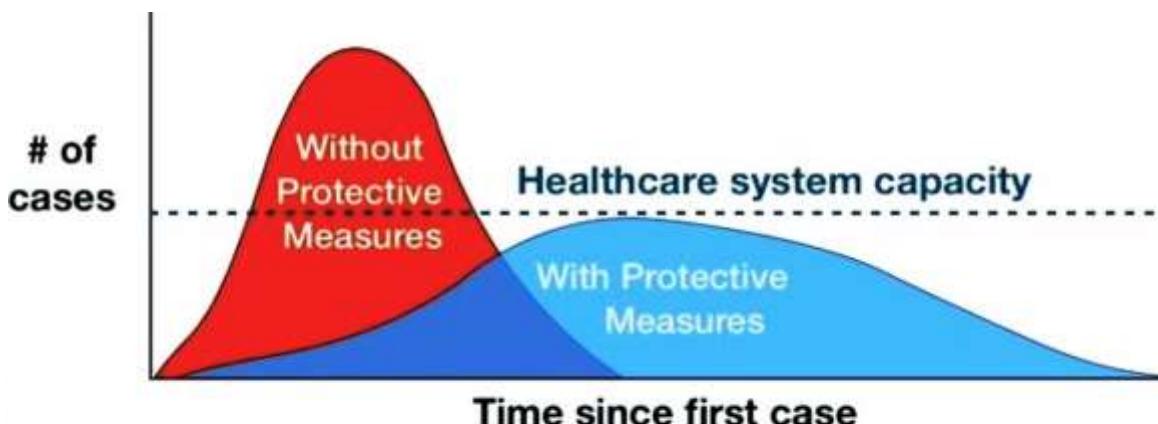
10th January 2020

This posting is communicated by Edward C. Holmes, University of Sydney on behalf of the consortium led by Professor Yong-Zhen Zhang, Fudan University, Shanghai

profession.australia.

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Interventions to slow disease spread



Adapted from CDC / The Economist

Laboratory tests for COVID-19 disease

- Nucleic acid testing
- Serology
- Virus isolation (or virus culture)
- Whole genome sequencing (WGS)
- Tests for disease management



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Developing SARS-CoV-2 nucleic acid detection tests

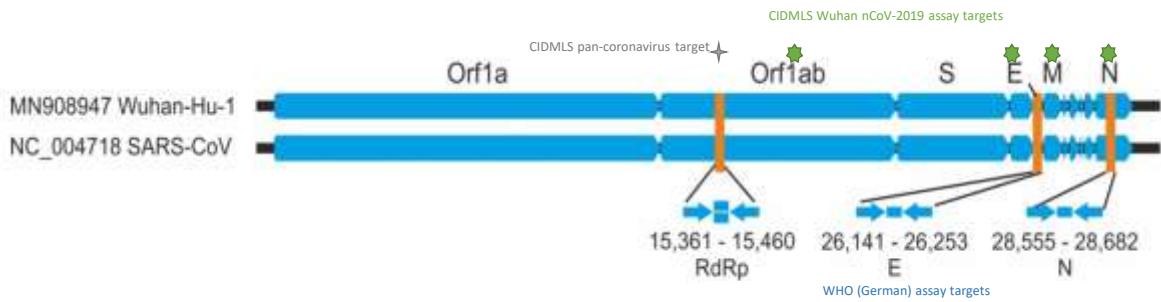


Figure 1 relative positions of amplicon targets on SARS-CoV ad Wuhan-CoV genome. N: nucleocapsid; ORF: open reading frame; RdRp: RNA-dependent RNA polymerase. Numbers below amplicon are genome positions according to SARS-CoV, NC_004718.

Nucleic acid testing for COVID-19 disease

- 'NAT' or 'PCR' or 'molecular tests'
- Multiple high throughput platforms
- Turnaround times vary and depend on the definition
 - Clinical TATs ~12-40 hours
 - In-laboratory TATs ~6 hours
- Rapid individual PCR tests eg Genexpert
 - <1 hour in-lab



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Nucleic acid testing for COVID-19 disease

- Issues around sensitivity and specificity
- Variation between platforms
- Sample collection
 - Methods of swabbing the upper respiratory tract
 - Swab types
 - Transport media
- Reagent shortages
- Clinical indications for testing



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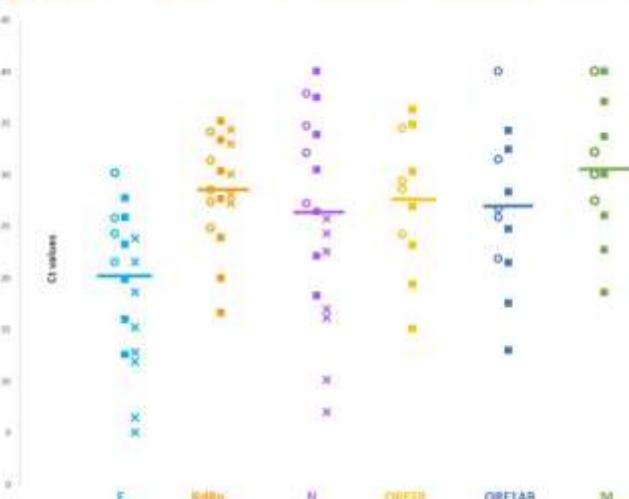


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40,000 tests in NSW Health Pathology later..

Figure 1 Cycle threshold values of clinical samples (○), cell culture supernatant (■) and synthetic positive controls (X) for nucleic acid test assays targeting different regions of the SARS-CoV-2 genome.



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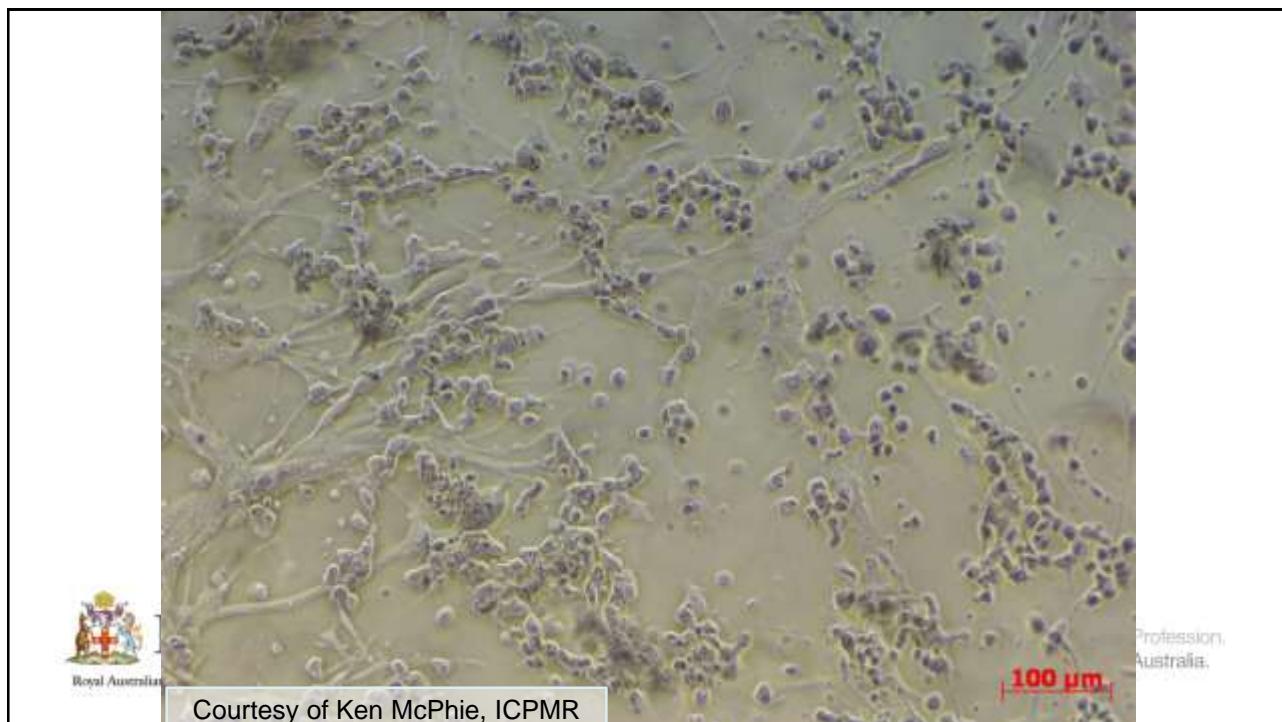
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Virus isolation or culture

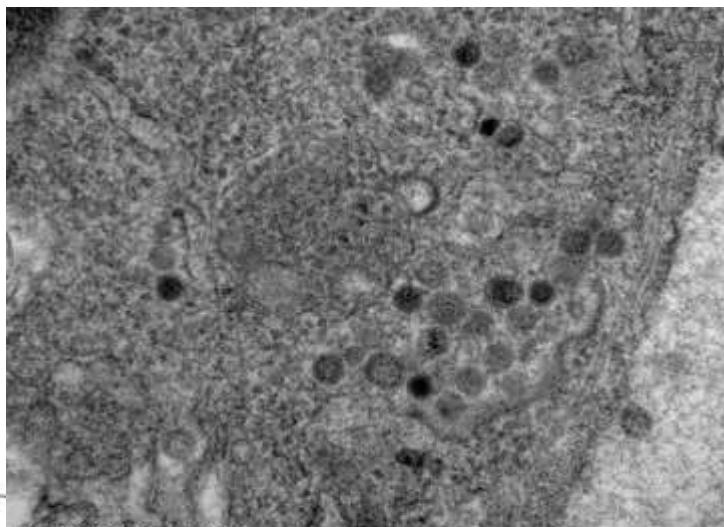
- Risk group 3 pathogen
 - Cultures performed in PC3/4 laboratory
- Use cell lines - Vero-E6 cells
- Observe for cytopathic effect (CPE) and confirm by PCR
- Studies in persistently NAT positive patients, including in healthcare workers



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Electron microscopy of SARS-CoV-2



ER-20-131 CMDS_K_022 [B] INFECTED!!!
[B] INFECTED
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TEM Mode: Imaging
Microscopist: LN

200 nm
HV=60kV
Direct Mag: 70000 x
ICPMR WESTMEAD

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PCR positivity vs culture positivity



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Indications for serology (SARS-CoV-2 antibodies)

- Making a **retrospective diagnosis** in individuals who have recovered from infection prior to testing
- Identifying cases where **false negative nucleic acid testing** has occurred, either because of sampling issues or mutations at primer/probe binding sites
- Confirmation of **unexpected positive nucleic acid tests**: especially important in settings of low incidence
- Identifying **asymptomatic infection**, especially in close contacts of cases or healthcare workers
- Determining the extent of infection in a population through **serosurveys**



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SARS-CoV-2 serology at NSWHP–ICPMR Westmead

- First results 20th February using immunofluorescence (IFA)
- SARS-CoV-2-specific IgG, IgM and IgA
- IgG is a marker of past infection; IgM and IgA are markers of more recent infection
- Virus neutralisation is available as another specific test for SARS-CoV-2-specific antibody



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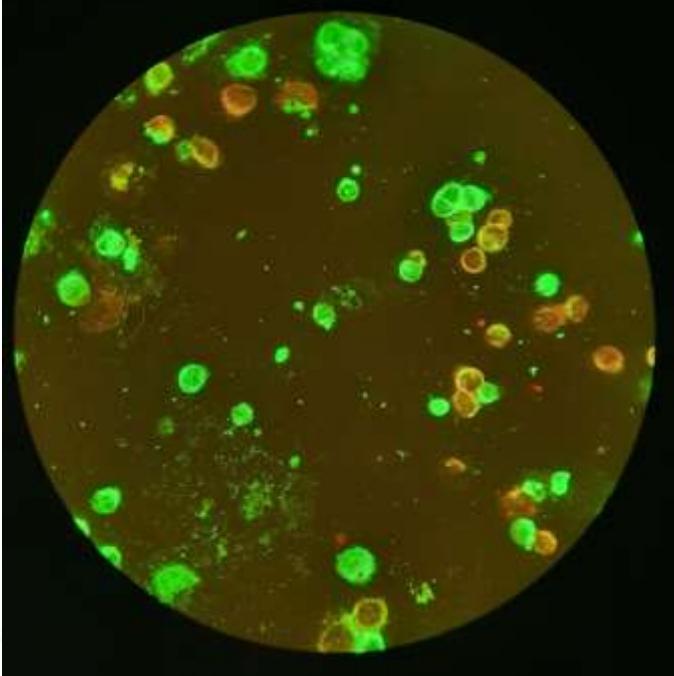


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SARS-CoV-2 infected cells
showing cytoplasmic
immunofluorescent
staining

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Courtesy of Linda Hueston, ICPMR

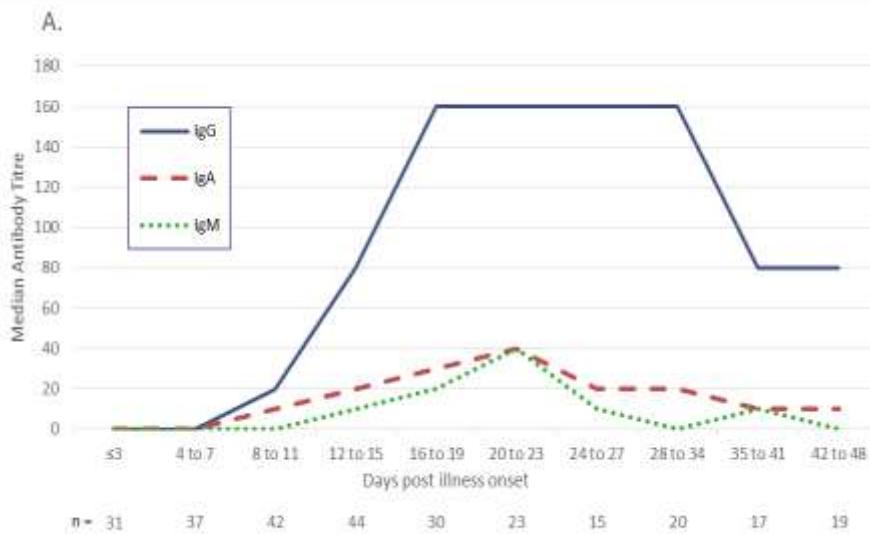
SARS-CoV-2 serology at NSWHP-ICPMR, Westmead

- >6500 samples tested
 - Routine diagnosis
 - Outbreak investigations (schools, ACFs, Ruby Princess crew)
 - Population serosurveys
- Evaluations of commercial platforms
 - Point-of-care antibody tests
 - Euroimmun ELISA, Abbott Architect CMA....
- In-house EIA in development



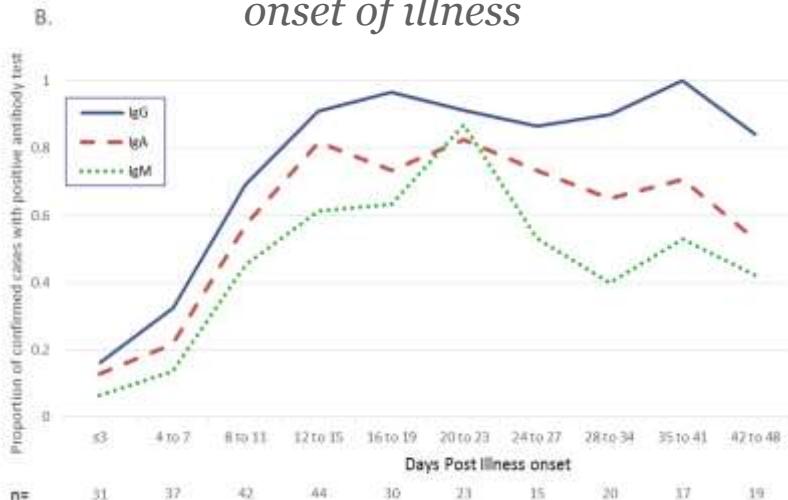
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Kinetics of the SARS-CoV-2 antibody response



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Detection of SARS-CoV-2 antibodies at different times after onset of illness



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RAPID COMMUNICATION

Estimating the asymptomatic proportion of coronavirus disease 2019 (COVID-19) cases on board the Diamond Princess cruise ship, Yokohama, Japan, 2020

Kenji Mizumoto^{1,2,3}, Katsushi Kagaya^{2,4}, Alexander Zarebski⁵, Gerardo Chowell³

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4. Seto Marine Biological Laboratory, Field Science, Education and Research Center, Kyoto University, Shirahama-cho, Nishimuro-gun, Wakayama, Japan

5. Department of Zoology, University of Oxford, Oxford, United Kingdom

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Citation style for this article:
Mizumoto Kenji., Kagaya Katsushi
cases on board the Diamond Prin
E5, 2020, 25, 10, 2000180

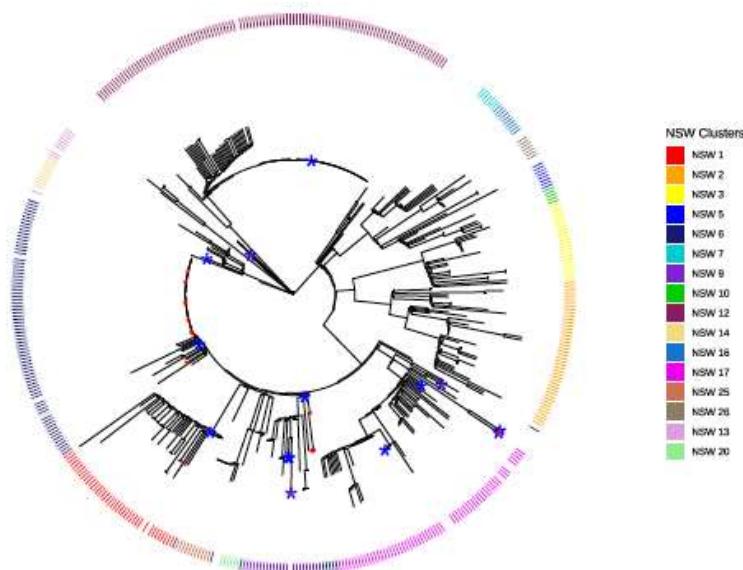
The estimated asymptomatic
proportion was
17.9%

95% credible interval: 15.5–20.2%

ise 2019 (COVID-19)
B07/1560-7917

/ published on 12 Mar 2020

Whole Genome Sequencing – infection clusters in NSW



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Sintchenko V et al. WGS SitRep 29/05/20



Coronavirus Disease 2019 (COVID-19) CDNA National Guidelines for Public Health Units

<https://www1.health.gov.au/internet/main/publishing.nsf/Content/cdna-song-novel-coronavirus.htm>

Confirmed case

A person who:

- I. tests positive to a validated specific SARS-CoV-2 nucleic acid test;
- OR
- II. has the virus isolated in cell culture, with PCR confirmation using a validated method;
- OR
- III. undergoes a seroconversion to or has a significant rise in SARS-CoV-2 neutralising or IgG antibody level (e.g. four-fold or greater rise in titre).¹

Probable case

A person who:

- I. has not been tested, with fever ($\geq 38^\circ\text{C}$)² or history of fever (e.g. night sweats, chills) **OR** acute respiratory infection (e.g. cough, shortness of breath, sore throat) **AND** is a household contact (refer to [Contact definition](#) below) of a confirmed or probable³ case of COVID-19;
- OR
- II. has detection of SARS-CoV-2 neutralising or IgG antibody⁴ **AND** has had a compatible clinical illness **AND** is a close contact (refer to [Contact definition](#) below) of a confirmed or probable³ case of COVID-19.



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COONAMBLE'S EXPERIENCE



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Useful resources

- NSW Health Website – Testing Advice for General Practitioners
<https://www.health.nsw.gov.au/Infectious/covid-19/Pages/case-definition.aspx>
- NSW Health Website – Advice for ACCSs
<https://www.health.nsw.gov.au/Infectious/covid-19/Pages/aboriginal-services.aspx>
- AH&MRC Website
<https://www.ahmrc.org.au/coronavirus/>



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Any questions....

Thank you



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