

Emerging Evidence Search Request

Please note: Our “Emerging Evidence” feature is based on the existing literature at the time the question is asked to provide the most informed answers possible to questions about COVID-19, and the science continues to evolve. As with any research, studies need to be replicated and there needs to be a strong body of evidence before any absolute conclusions can be made.

Questions about UHN policy or practice are best directed to the appropriate department (e.g. IPAC, Occupational Health).

Question:

How long is a person contagious after they contract Covid? What is the best evidence for determining people are no longer infectious?

Additional Results from an Updated Search: November 3, 2020

<1>

Unique Identifier

33104776

Title

Repeat COVID-19 Molecular Testing: Correlation of SARS-CoV-2 Culture with Molecular Assays and Cycle Thresholds.

Source

Clinical Infectious Diseases. 2020 Oct 27.

VI 1

Status

Publisher

Authors

Gniazdowski V; Morris CP; Wohl S; Mehoke T; Ramakrishnan S; Thielen P; Powell H; Smith B; Armstrong DT; Herrera M; Reifsnyder C; Sevdali M; Carroll KC; Pekosz A; Mostafa HH.

Authors Full Name

Gniazdowski, Victoria; Morris, C Paul; Wohl, Shirlee; Mehoke, Thomas; Ramakrishnan, Srividya; Thielen, Peter; Powell, Harrison; Smith, Brendan; Armstrong, Derek T; Herrera, Monica; Reifsnyder, Carolyn; Sevdali, Maria; Carroll, Karen C; Pekosz, Andrew; Mostafa, Heba H.

Institution

Gniazdowski, Victoria. Johns Hopkins University School of Medicine, Department of Pathology, Division of Medical Microbiology, Baltimore, Maryland, USA.

Morris, C Paul. Johns Hopkins University School of Medicine, Department of Pathology, Division of Medical Microbiology, Baltimore, Maryland, USA.

Morris, C Paul. National Institutes of Allergy and Infectious Disease, Bethesda, Maryland, USA.

Wohl, Shirlee. Johns Hopkins Bloomberg School of Public Health, Baltimore, Maryland, USA.

Mehoke, Thomas. Johns Hopkins University Applied Physics Laboratory, Laurel, Maryland, USA.

Ramakrishnan, Srividya. Johns Hopkins University, Department of Computer Science, Baltimore, Maryland, USA.

Thielen, Peter. Johns Hopkins University Applied Physics Laboratory, Laurel, Maryland, USA.

Powell, Harrison. Bio-Rad Laboratories, Pleasanton, California, USA.

Smith, Brendan. Bio-Rad Laboratories, Pleasanton, California, USA.

Armstrong, Derek T. Johns Hopkins University School of Medicine, Department of Pathology, Division of Medical Microbiology, Baltimore, Maryland, USA.

Herrera, Monica. Bio-Rad Laboratories, Pleasanton, California, USA.

Reifsnnyder, Carolyn. Bio-Rad Laboratories, Pleasanton, California, USA.

Sevdali, Maria. Bio-Rad Laboratories, Pleasanton, California, USA.

Carroll, Karen C. Johns Hopkins University School of Medicine, Department of Pathology, Division of Medical Microbiology, Baltimore, Maryland, USA.

Pekosz, Andrew. W. Harry Feinstone Department of Molecular Microbiology and Immunology, The Johns Hopkins Bloomberg School of Public Health, Baltimore, Maryland, USA.

Mostafa, Heba H. Johns Hopkins University School of Medicine, Department of Pathology, Division of Medical Microbiology, Baltimore, Maryland, USA.

Abstract

BACKGROUND: Repeat COVID-19 molecular testing can lead to positive test results after negative tests and to multiple positive test results over time. The association between positive tests and infectious virus is important to quantify.

METHODS: A two months cohort of retrospective data and consecutively collected specimens from COVID-19 patients or patients under investigation were used to understand the correlation between prolonged viral RNA positive test results, cycle threshold (Ct) values and growth of SARS-CoV-2 in cell culture. Whole genome sequencing was used to confirm virus genotype in patients with prolonged viral RNA detection. Droplet digital PCR (ddPCR) was used to assess the rate of false negative COVID-19 diagnostic tests.

RESULTS: In two months, 29,686 specimens were tested and 2,194 patients received repeated testing. Virus recovery in cell culture was noted in specimens with SARS-CoV-2 target genes' Ct value average of 18.8 +/- 3.4. Prolonged viral RNA shedding was associated with positive virus growth in culture in specimens collected up to 20 days after the first positive result but mostly in individuals symptomatic at time of sample collection. Whole genome sequencing provided evidence the same virus was carried over time. Positive tests following negative tests had Ct values higher than 29.5 and were not associated with virus culture. ddPCR was positive in 5.6% of negative specimens collected from COVID-19 confirmed or clinically suspected patients.

CONCLUSIONS: Low Ct values in SARS-CoV-2 diagnostic tests were associated with virus growth in cell culture. Symptomatic patients with prolonged viral RNA shedding can also be infectious. Copyright © The Author(s) 2020. Published by Oxford University Press for the Infectious Diseases Society of America.

Publication Type

Journal Article

Year of Publication

2020

Link to the Ovid Full Text or citation:

[Click here for full text options](#)

Link to UHN Full Text:

[Serial Solutions](#)

<2>

Unique Identifier

Library staff make every effort to provide accurate and complete database search results. However, we assume no liability for information retrieved, its interpretation, applications or omissions.

33077007

Title

Understanding Viral Shedding of SARS-CoV-2: Review of Current Literature.

Source

Infection Control & Hospital Epidemiology. 1-35, 2020 Oct 20.

VI 1

Status

Publisher

Authors

Fontana L; Villamagna AH; Sikka MK; McGregor JC.

Author NameID

McGregor, Jessina C; ORCID: <https://orcid.org/0000-0002-6093-5444>

Authors Full Name

Fontana, Lauren; Villamagna, Angela Holly; Sikka, Monica K; McGregor, Jessina C.

Institution

Fontana, Lauren. Division of Infectious Diseases, Department of Medicine, School of Medicine, Oregon Health & Science University; Portland, OR.

Villamagna, Angela Holly. Division of Infectious Diseases, Department of Medicine, School of Medicine, Oregon Health & Science University; Portland, OR.

Sikka, Monica K. Division of Infectious Diseases, Department of Medicine, School of Medicine, Oregon Health & Science University; Portland, OR.

McGregor, Jessina C. Department of Pharmacy Practice, College of Pharmacy, Oregon State University; Portland, OR.

Abstract

OBJECTIVE: Transmission of SARS-CoV-2 has significant implications for hospital infection prevention and control, discharge management, and public health. We reviewed available literature to reach an evidenced-based consensus on the expected duration of viral shedding.

DESIGN: We queried four scholarly repositories/search engines for studies reporting SARS-CoV-2 viral shedding dynamics by PCR and/or culture available through September 8, 2020. We calculated the pooled median duration of viral RNA shedding from respiratory and fecal sources.

RESULTS: Seventy-seven studies on SARS-CoV-2 were included. All studies reported PCR-based testing and 12 also included viral culture data. The overall pooled median duration of RNA shedding from respiratory sources was 18.4 days (95% CI: 15.5 days - 21.3 days; I²=98.87%, p<0.01) among 28 studies. When stratified by disease severity, the pooled median duration of viral RNA shedding from respiratory sources was 19.8 days (95% CI: 16.2 days - 23.5 days; I²=96.42%, p<0.01) among severely ill patients and 17.2 days (95% CI: 14.0 days - 20.5 days; I²=95.64%, p<0.01) in mild/moderate illness. Viral RNA was detected up to 92 days after symptom onset. Viable virus was isolated by culture from -6 days to 20 days relative to symptom onset.

CONCLUSIONS: SARS-COV-2 RNA shedding can be prolonged, yet high heterogeneity exists. Detection of viral RNA may not correlate with infectivity since available viral culture data suggests shorter durations of shedding of viable virus. Additional data is needed to determine the duration of shedding of viable virus and the implications for risk of transmission.

Publication Type

Journal Article.

Year of Publication

2020

Link to the Ovid Full Text or citation:

[Click here for full text options](#)

Link to UHN Full Text:

[Serial Solutions](#)

Library staff make every effort to provide accurate and complete database search results. However, we assume no liability for information retrieved, its interpretation, applications or omissions.

<3>

Unique Identifier

33049331

Title

The duration of infectiousness of individuals infected with SARS-CoV-2. [Review]

Source

Journal of Infection. 2020 Oct 10.

VI 1

Status

Publisher

Authors

Walsh KA; Spillane S; Comber L; Cardwell K; Harrington P; Connell J; Teljeur C; Broderick N; de Gascun CF; Smith SM; Ryan M; O'Neill M.

Authors Full Name

Walsh, Kieran A; Spillane, Susan; Comber, Laura; Cardwell, Karen; Harrington, Patricia; Connell, Jeff; Teljeur, Conor; Broderick, Natasha; de Gascun, Cillian F; Smith, Susan M; Ryan, Mairin; O'Neill, Michelle.

Institution

Walsh, Kieran A. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland. Electronic address: kiwalsh@hiqa.ie.

Spillane, Susan. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland.

Comber, Laura. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland.

Cardwell, Karen. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland; Health Research Board Centre for Primary Care Research, Department of General Practice, Royal College of Surgeons in Ireland, 123 St. Stephens Green, Dublin 2, Ireland.

Harrington, Patricia. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland.

Connell, Jeff. National Virus Reference Laboratory, University College Dublin, Belfield, Dublin 4, Ireland.

Teljeur, Conor. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland.

Broderick, Natasha. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland.

de Gascun, Cillian F. National Virus Reference Laboratory, University College Dublin, Belfield, Dublin 4, Ireland.

Smith, Susan M. Health Research Board Centre for Primary Care Research, Department of General Practice, Royal College of Surgeons in Ireland, 123 St. Stephens Green, Dublin 2, Ireland.

Ryan, Mairin. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland; Department of Pharmacology & Therapeutics, Trinity College Dublin, Trinity Health Sciences, James Street, Dublin 8, Ireland.

O'Neill, Michelle. Health Information and Quality Authority, Smithfield, Dublin 7, Ireland.

Abstract

OBJECTIVES: To summarise the evidence on the duration of infectiousness of individuals in whom SARS-CoV-2 ribonucleic acid is detected.

METHODS: A rapid review was undertaken in PubMed, Europe PubMed Central and EMBASE from 1 January 2020 to 26 August 2020.

RESULTS: We identified 15 relevant studies, including 13 virus culture and 2 contact tracing studies. For 5 virus culture studies, the last day on which SARS-CoV-2 was isolated occurred within 10 days of symptom onset. For another 5 studies, SARS-CoV-2 was isolated beyond day 10 for approximately 3% of included patients. The remaining 3 virus culture studies included patients with severe or critical disease; SARS-CoV-2 was isolated up to day 32 in one study. Two studies identified immunocompromised patients from whom SARS-CoV-2 was isolated for up to 20 days. Both contact tracing studies, when close contacts were first exposed greater than 5 days after symptom onset in the index case, found no evidence of laboratory-confirmed onward transmission of SARS-CoV-2.

CONCLUSION: COVID-19 patients with mild-to-moderate illness are highly unlikely to be infectious beyond 10 days of symptoms. However, evidence from a limited number of studies indicates that patients with severe-to-critical illness or who are immunocompromised, may shed infectious virus for longer.

Copyright © 2020. Published by Elsevier Ltd.

Publication Type

Journal Article. Review.

Year of Publication

2020

Link to the Ovid Full Text or citation:

[Click here for full text options](#)

Link to UHN Full Text:

[Serial Solutions](#)

Databases Searched:

Ovid MEDLINE(R) ALL <1946 to November 02, 2020>

Search history sorted by search number ascending

# Searches	Results	Type
1 Virus Shedding/	3465	Advanced
2 ((viral or virus) adj2 shed*).mp.	6639	Advanced
3 1 or 2	6639	Advanced
4 limit 3 to covid-19	411	Advanced
5 limit 4 to dt="20201006-20201103"	47	Advanced
6 limit 5 to english language	47	Advanced
7 remove duplicates from 6	47	Advanced
8 from 7 keep 9, 17, 19	3	Advanced

Search Completed By:

Ani Orchanian-Cheff