

COVID-19 Pathogenesis

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Jonathan Corum & Carl Zimmer *The New York Times*, April 3rd 2020



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Speaker Disclosure

Speakers are required to disclose any commercial relationships before today's presentation.



Contents

- Basics of COVID-19 disease
- Appearance and spread
- Current status of the pandemic
 - Recent developments in testing, vaccines, and treatment
- Emergence from an animal reservoir
 - Likelihood of other coronavirus emergence



Kevin Olival, EcoHealth Alliance





emergency symptoms seek medical attention

inability to wake or stay awake difficulty breathing bluish lips or face chest pain or pressure loss of speech or movement new confusion

sources: US Centers for Disease Control & Prevention, World Health Organisation, UK NHS

Basics of COVID-19 disease



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COVID-19 and SARS-CoV-2

- <u>Coronavirus Disease 2019 (COVID-19) is the disease</u>
- SARS-Coronavirus type 2 (SARS-CoV-2) is the virus that causes COVID-19
- SARS-CoV-2 is closely related to SARS-CoV, the virus that caused <u>Severe Acute Respiratory Syndrome</u> (SARS) in 2002
- SARS-CoV and SARS-CoV-2 are also related to MERS-CoV, the virus that causes Middle Eastern Respiratory Syndrome (MERS) in 2012/2013
- All are a type of virus called Coronaviruses



Coronaviruses

- Spike proteins form a crown shape (or corona) on virus surface
- Prior to SARS outbreak in 2002, coronaviruses were thought to be harmless in humans
- One of the causes of the common cold (15%)
- 3 human outbreaks
 - SARS (2002) 8,098 cases, 774 deaths (10% CFR)
 - MERS (2013) 2,519 cases, 866 deaths (34% CFR)
 - COVID-19 (2020) 16.5 million cases, 650,000 deaths* (4% CFR, likely an overestimate as we do not have reliable case counts)

*as of 07/28/2020, data from Johns Hopkins University COVID-19 dashboard



Science vol 339 pp1269-1273 (2013)



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Especially Those with Existing Conditions

% of deceased with serious ailments



study of 44,672 confirmed cases in Mainland China sources: China Center for Disease Control & Prevention, Statista



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How Contagious & Deadly is It?

We don't fully know yet but it's in this range

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Infection with a virus

- In the first step of infection, any virus has to recognize and interact with a specific receptor protein on the target cell's surface
- Each type of virus recognizes a different receptor
- Different cells have different proteins on their cell surface
- So, each virus can infect a specific set of cells, and not others
- SARS-CoV and SARS-CoV-2 both use the same receptor: <u>ACE2</u>

Infectious mechanism used by SARS-CoV-2

- ACE2 is found of pneumocytes in the lung; also enterocytes in the gut
- The spike protein of SARS-CoV and SARS-CoV-2 is cleaved by the human enzyme TMPRSS2
 - the drug <u>camostat mesylate</u> inhibits this
- This drug is already licensed in Europe & Japan (as a treatment for pancreatitis), and is now in a clinical trial for COVID-19

Cell vol 181 271-280.e8 (2020)

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COVID-19 subversion of the immune response

- Viral infection normally induces type I interferon response in immune cells
- COVID-19 proteins block interferon α/β expression and signaling
- Viral clearance is delayed; robust B cell, T cell and cytokine response develops
- "Cytokine storm" results in lung tissue damage

Cell vol 181 pp1046-1045e9 (2020)

Infection and disease in COVID-19

- Respiratory infection, primarily through droplets and contact with contaminated surfaces
- The extent of aerosol infection is not clear
- Many infections are mild or asymptomatic, so it is hard to determine the true case count
- Severe acute respiratory distress syndrome (ARDS) and extensive lung damage

Dr. Keith Mortman, George Washington University

COVID-19 distinguishing features

- Spread by droplet infection (like SARS, MERS, and also influenza), BUT
- COVID-19 patients produce infectious virus well before they feel ill (if they feel ill at all)
 - Spreads in the general population
 - Many asymptomatic (but infectious) infections
- SARS and MERS patients only shed virus after the onset of symptoms
 - Secondary infections were most common in family members and healthcare workers

Nature (2020) https://doi.org/10.1038/s41586-020-2488-1

Superspreaders in COVID-19

- Individual person who infects a large number of others
- Daegu, South Korea, Feb 2020; 70 cases linked to "patient 31" at the Shinceonji Church of Jesus
- New Rochelle NY, USA, March 2020; one person spread the virus to at least 20 individuals, creating a cluster of infections that ultimately exceeded 100
- Rheda-Wiedenbrück, Germany, March-April 2020; one asymptomatic worker at a meat packing plant infected 60% of other workers within a 25-foot radius

Prevention is better than cure

Lots of science-y folks are posting this graph. But if there is one thing I have learned from being on the internet, it is this:

Data/graphs: Not compelling to many.

Kitties: Compelling to many.

So I present: #Catteningthecurve.

#scicomm #epitwitter

Replying to @amdar1ing

Awesome! Here's a live re-enactment, courtesy of my two.

Oreo and Spooky showing how to #FlattenTheCurve #EpiCats

7:57 AM · Mar 12, 2020 · Tweetbot for iOS	
11:28 AM · Mar 12, 2020 · Twitter Web App	
II View Tweet activity	
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Masks work

Masks reduce airborne transmission

Infectious aerosol particles can be released during breathing and speaking by asymptomatic infected individuals. No masking maximizes exposure, whereas universal masking results in the least exposure.

Science vol 368 pp1422-1424 (2020)

- SARS-CoV-2 replicates three times faster than other coronaviruses
- Can spread rapidly to the pharynx and be exhaled before the innate immune response becomes activated and the infected person shows symptoms

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Impact of COVID-19

Infection & Fatality Rates Vary by Country

	confirmed cases			deaths	% cases who have died	
USA	4,233,923	+1.3%		146,935		3.5%
Brazil	2,419,09	9 1 1.0%	+4.7%	87,00	4	3.6%
India	1,435,616	+3.6%	+4.1%	32,771		2.3%
Russia	811,073		-1.2%	13,249	- •	1.6%
South Africa	445,433	+2.6%	-1.2%	6,769	- •	1.5%
Mexico	390,516	+1.4%		43,680	• · · · ·	11.2
Peru	375,961		-2.5%	17,843		4.7%
Chile	345,790			9,112		2.6%
UK	301,020		-1.4%	45,837	•	15.2
Iran	291,172			15,700	•	5.4%
Pakistan	273,113		-6.7%	5,822		2.1%
Spain	272,421		+10.4%	28,432		10.4
Saudi Arabia	266,941		-2.2%	2,733	•	1.0%
Italy	246,118		+3.0%	35,107	•	14.3
Colombia	240,795			8,269		3.4%
Turkey	226,100			5,613		2.5%
Bangladesh	223,453	+1.0%	-1.3%	2,928		1.3%
France	217,801		+6.0%	30,195	•	13.9
Germany	206,667		+4.9%	9,124		4.4%
Argentina	162,526	+2.6%	+4.4%	2,939		1.8%
					AVERAGE 5.3%	

CASE FATALITY RATE	
CFR is unreliable during an outb	

	cases per million people
Chile	19,152
USA	13,049
Peru	11,688
Brazil	11,559
Saudi Arabia	8,104
Sweden	7,971
South Africa	7,854
Israel	7,445
Belarus	7,090
Bolivia	6,282
UAE	6,267
Spain	5,877
om. Republic	5,843
Belgium	5,777
Russia	5,633
Kyrgyzstan	5,428
Colombia	4,908
Portugal	4,856
Ecuador	4,854
Kazakhstan	4,566

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updated 27 July 2020

deaths per million people •=9	
Belgium	
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Spain •••••••613	
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sources: Johns Hopkins University

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The New York Times, July 27th 2020

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COVID-19 in Allegheny County

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Recent updates in cases, treatment, and vaccines

Figure from The New York Times, July 20th 2020

COVID-19 detections are rising. It's not due to increased testing.

How Coronavirus Cases Compare With Expectations

The charts show how the number of reported cases compares with the expected count based on expanded testing. The gap for states with more cases than expected is highlighted. Each state is on its own scale.

+8,743

The New York Times, July 24th 2020

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Seroprevalence of antibodies to SARS-CoV-2 in the USA

- Convenience sample of sera collected for routine clinical testing, March – May 2020
- 16,025 samples tested
- Most had no evidence of SARS-CoV-2 antibodies
 - Proportion of reactive sera ranged from 1-7%
- But the proportion of reactive sera was much greater than the number of reported cases in each area
 - Between 6 and 24 times greater

JAMA Internal Medicine (2020) doi:10.1001/jamainternmed.2020.4130

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Coronavirus Vaccine Tracker

By Jonathan Corum, Denise Grady, Sui-Lee Wee and Carl Zimmer Updated July 27, 2020

The New York Times, July 27th 2020

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PHASE III

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Immunity to coronaviruses after vaccination or infection

- There is no COVID-19 vaccine yet, although many are now beginning to undergo clinical trials, and COVID-19 survivors have only been infected a few months ago at most
- In the case of SARS, many infected-recovered people showed waning immunity a few years afterwards
- Responses to experimental coronavirus vaccines also decline after one year
- Will we have lifelong immunity to COVID-19?

Months after disease onset

Journal of Immunology vol **186** pp7264-7268 (2011)

Current state of COVID-19 therapies

- The antiviral drug <u>Remdesivir</u> shortens recovery time by 4 days on average but has no impact on mortality, compared to placebo [NEJM 2020; DOI:10.1056/NEJMoa2007764]
- The only drug shown to date to affect mortality is the steroid dexamethasone
 - Effect is greatest for patients needing oxygen, or mechanical ventilation
 - Little to no effect in less severe cases
- Other drug discovery pipelines
 - Repurposing of existing drugs [Nature 2020; doi:10.1038/s41586-020-2577-1]
 - Protein-protein interaction screening [Nature 2020; doi:10.1038/s41586-020-2286-9]

The RECOVERY study: impact of dexamethasone on mortality

"Scientifically robust and ethically sound clinical research remains the quickest and most efficient pathway to effective treatment and prevention strategies for patients with Covid-19"

-accompanying editorial by Lane & Fauci

New England Journal of Medicine advance publication DOI: 10.1056/NEJMoa2021436

Are there any human genes that affect COVID-19 severity?

New England Journal of Medicine, advance publication DOI: 10.1056/NEJMoa2020283

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Where did SARS-CoV-2 come from, and are there others?

Nature vol 583 pp366-368 (2020)

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Multiple early introductions of COVID-19 into the USA from China

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Washington state and California cases

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How do coronaviruses infect humans?

Trends in Microbiology vol 25 pp35-48 (2017)

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Zoonotic transfer of coronavirus in SARS

Horseshoe bat

Coronavirus 88-92% similarity to human

Palm civet

Wet market

Coronavirus 99.8% similarity to human

After the initial zoonotic transfer to humans, almost all infections were due to human-to-human spread

Zoonotic transfer of coronavirus in MERS

Lancet Infectious Diseases vol **15** pp495-497 (2015)

Direct or indirect transmission to human beings

Lancet vol 386 pp995-1007 (2015)

Zoonotic transfer of coronavirus in COVID-19?

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The Huanan Seafood and Wildlife Market in Wuhan, China

Cell vol 181, pp223-227 (2020)

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Coronavirus adaptation to humans

- The interaction between the spike protein, especially the receptor-binding domain (RBD) with ACE2, is critical to infection
- Early in the SARS outbreak, the SARS-CoV RBD underwent two small genetic changes that increased its binding affinity for human ACE2 by 1,000 fold
- This allowed for extensive human-to-human transmission after the initial species jump
- Similar small changes seen in MERS

Nature Microbiology vol 5 pp562-569 (2020)

How are SARS-CoV and SARS-CoV-2 different?

- SARS case fatality rate was 10%; COVID-19 is much lower (1-2% current estimate)
- COVID-19 patients have virus in the upper respiratory tract; SARS patients did not
- ACE2 levels are lower on cells of the upper respiratory tract
- SARS-CoV-2 spike protein has an extra region that SARS-CoV does not. This polybasic furin-type cleavage site increases fusion activity and may allow the virus to enter cells with lower levels of ACE2

Wuhan	23481	CTAATGTTTTTCAAACACGTGCAGGCTGTTTAATAGGGGCTGAACATGTCAACAACTCAT	23540
RaTG13	23463	CTAATGTTTTCAAACACGTGCAGGTTGTTTAATAGGGGCTGAACATGTCAATAACTCGT	23522
Wuhan	23541	ATGAGTGTGACATACCCATTGGTGCAGGTATATGCGCTAGTTATCAGACTCAGACTAATT	23600
RaTG13	23523	ATGAGTGTGACATACCTATTGGTGCAGGAATATGCGCCAGTTATCAGACTCAAACTAATT	23582
Wuhan	23601	CTCCTCGGCGGGCACGTAGTGTAGCTAGTCAATCCATCATTGCCTACACTATGTCACTTG	23660
RaTG13	23583	CACGTAGTGTGGCCAGTCAATCTATTATTGCCTACACTATGTCACTTG	23630
Wuhan	23661	GTGCAGAAAATTCAGTTGCTTACTCTAATAACTCTATTGCCATACCCACAAATTTTACTA	23720
RaTG13	23631	GTGCAGAAAATTCAGTTGCTTATTCTAATAACTCTATTGCCATACCTACAAATTTTACTA	23690
Wuhan	23721	TTAGTGTTACCACAGAAATTCTACCAGTGTCTATGACCAAGACATCAGTAGATTGTACAA	23780
RaTG13	23691	TTAGTGTGACCACTGAAATTCTACCTGTGTCTATGACAAAGACATCGGTAGACTGTACAA	23750

...but it's not an engineered virus (unless they did it really badly)

http://virological.org/t/tackling-rumors-of-a-suspicious-origin-of-ncov2019

BatRaTG13 Wuhan

SARS-CoV-2 is mutating like other coronaviruses

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Is COVID-19 evolving to become more deadly?

By Jonathan Corum | Source: Lizhou Zhang et al., Scripps Research

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The New York Times, June 12th 2020

Nextstrain

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Why do bats carry so many viruses that are fatal to humans?

- SARS, MERS, COVID-19, Nipah, Hendra, Ebola, Marburg, etc
- Bats roost together in large numbers, often with members of other species
- Bats harbor these viruses but do not get sick and die – how?
 - Bats have evolved a complex series of immune responses that enable them to tolerate the presence of many viruses
- Bats are flying mammals and have a high body temperature
 - viruses that can survive this can survive the high fever in humans that is intended to kill pathogens

Cell vol 173 pp1098-1110 (2018)

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Will COVID-19 be the last new coronavirus infection?

- There are many Coronaviruses circulating in many bat species
- 6.5% of bats surveyed harbor one or more coronaviruses (Tang et al (2006), J. Virol vol 80 pp7481-7490)
- EcoHealth Alliance data: ≈400 novel coronaviruses in bats in China; multiple "spillover" events where these have infected humans, but not spread

Viruses vol **11**, 210 (2019) Each red block represents one CoV in one bat species

The global virome project

- Around 263 viruses from 25 viral families are currently known to infect humans
- Estimated that ~1.67 million yet-to-be-discovered viral species from key zoonotic viral families exist in mammal and bird hosts
- Reasonable expectation is that between 631,000 and 827,000 of these unknown viruses have zoonotic potential
- GVP launched in 2018, with goal of identifying this viral threat

Science vol 359 pp872-874 (2018)

GVP targeting strategy

The project will capitalize on economies of scale in viral testing, systematically sampling mammals and birds to identify currently unknown, potentially zoonotic viruses that they carry.

Of these 111 viral families, the GVP will target **25** containing viruses known to infect (or to have substantial risk of infecting) people.

In these 25 families, an estimated **1.67 million** unknown viruses exist in mammals and birds—hosts that represent 99% of the risk for viral emergence.

Of these 1.67 million viruses, an estimated **631,00 to 827,000** likely have the capacity to infect people.

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Summary

- COVID-19 is a severe disease caused by infection with SARS-CoV-2
- Coronaviruses are a widespread family of viruses that typically do not cause severe disease
- SARS-CoV-2 infects cells after its spike protein interacts with ACE2
 - mutations in the spike protein facilitate infection in humans
- The long-term immune response to COVID-19 is unknown
- There are MANY more coronaviruses out there

MidAtlantic AIDS Education and Training Center - Contact Information

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