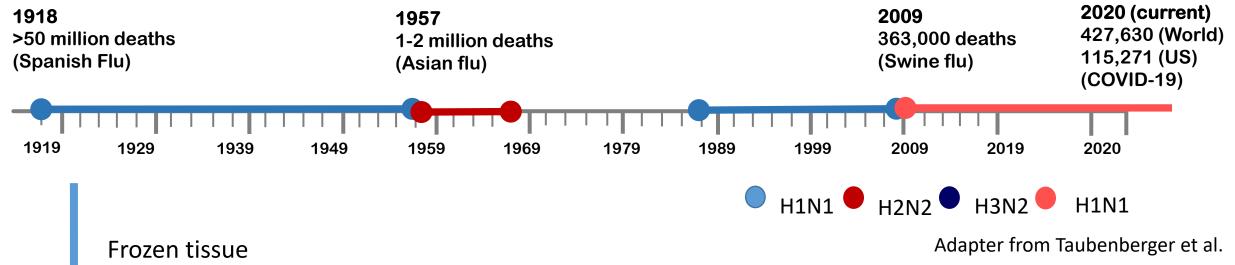
Development, Validation and Interpretation of Testing for SARS-CoV-2

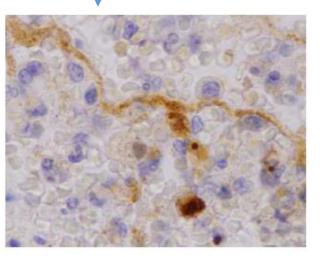
> David R Hillyard MD Professor, Pathology University of Utah School of Medicine Director Molecular Infectious Disease Testing ARUP Laboratories

Outline

- Pandemics: lessons from the past
- The virus and its challenges
- Viral infection, symptoms, shedding and transmission
- Tests and their comparative performance
- Viral evolution and the role of sequencing



Multiple viral and bacterial sequences Reconstruction of 1918 virus Test in many animal models



Highest incidence in crowded populations Virtually all deaths were due to secondary bacterial pneumonia High mortality in young adults due to higher case incidence

> COVID-19 is a different virus Mitigation is similar process

Taubenberger et al., Sci. Transl. Med. 11, (2019)

WHO Global Flu Strategy 2019-2030 (Prevent. Control. Prepare.)

Announced March 2019



<u>Pillars of Flu Preparedness</u> (national-community-health care system-physician)

Viral Surveillance and Risk Assessment Early disease recognition, Diagnostics and Drugs Vaccines

Infrastructure Preparedness "non-pharmaceutical"

Non-pharmaceutical Recommendations

During a Pandemic: Limit the Spread of Germs and Prevent Infection

- Avoid close contact with people who are sick.
- When you are sick, keep your distance from others to protect them from getting sick too.
- Cover your mouth and nose with a tissue when coughing or sneezing. It may prevent those around you from getting sick.
- Washing your hands often will help protect you from germs.
- Avoid touching your eyes, nose or mouth.
- **Practice other good health habits**. Get plenty of sleep, be physically active, manage your stress, drink plenty of fluids, and eat nutritious food.

• Before a Pandemic:

- Store a two week supply of water and food.
- Periodically check your regular prescription drugs to ensure a continuous supply in your home.
- Have any nonprescription drugs and other health supplies on hand, including pain relievers, stomach remedies, cough and cold medicines, fluids with electrolytes, and vitamins.
- Get copies and maintain electronic versions of health records from doctors, hospitals, pharmacies and other sources and store them, for personal reference. Get help accessing electronic help records.
- Talk with family members and loved ones about how they would be cared for if they got sick, or what will be needed to care for them in your home.





Guidelines to Prevent Pandemic Influenza through Non-pharmaceutical Interventions (NPIs) and Community Engagement MMWR / April 21, 2017 / Vol. 66 / No. 1

Social Distancing

CDC recommendations

Social distancing measures: Even though the evidence base for the effectiveness of some of these measures is limited, CDC might recommend the simultaneous use of multiple social distancing measures to help reduce the spread of influenza in community settings (e.g., schools, workplaces, and mass gatherings) during severe, very severe, or extreme influenza pandemics while minimizing the secondary consequences of the measures. Social distancing measures include the following:

- Increasing the distance to at least 3 feet (98) between persons when
 possible might reduce person-to person transmission. This applies
 to apparently healthy persons without symptoms. In the event of a
 very severe or extreme pandemic, this recommended minimal
 distance between people might be increased.
- Persons in community settings who show symptoms consistent with influenza and who might be infected with (probable) pandemic influenza should be separated from well persons as soon as practical, be sent home, and practice voluntary home isolation.

Environmental cleaning

CDC recommendations

Environmental surface cleaning measures: CDC recommends environmental surface cleaning measures in all settings, including homes, schools, and workplaces, to remove influenza viruses from frequently touched surfaces and objects. Use of these measures might help prevent transmission of various infectious agents, including seasonal and pandemic influenza (https://www.cdc.gov/nonpharmaceutical-interventions/environmental/ index.html; https://www.cdc.gov/oralhealth/infectioncontrol/questions/ cleaning-disinfecting-environmental-surfaces.html).

Additional guidance is available from CDC for health care facilities (https:// www.cdc.gov/hicpac/pdf/guidelines/eic_in_HCF_03.pdf), schools (https:// www.cdc.gov/flu/school/cleaning.htm), and airline, travel, and transportation industries (https://www.cdc.gov/flu/pandemic-resources/archived/ transportation-planning.html).

Use of Face Masks

CDC recommendations

Use of face masks by ill persons: CDC might recommend the use of face masks by ill persons as a source control measure during severe, very severe, or extreme influenza pandemics when crowded community settings cannot be avoided (e.g., when adults and children with influenza symptoms seek medical attention) or when ill persons are in close contact with others (e.g., when symptomatic persons share common spaces with other household members or symptomatic postpartum women care for and nurse their infants). Some evidence indicates that face mask use by ill persons might protect others from infection.

Use of face masks by well persons: CDC does not routinely recommend the use of face masks by well persons in the home or other community settings as a means of avoiding infection during influenza pandemics except under special, high-risk circumstances (https://www.cdc.gov/flu/ professionals/infectioncontrol/maskguidance.htm). For example, during a severe pandemic, pregnant women and other persons at high risk for influenza complications might use face masks if unable to avoid crowded settings, especially if no pandemic vaccine is available. In addition, persons caring for ill family members at home (e.g., a parent of a child exhibiting influenza symptoms) might use face masks to avoid infection when in close contact with a patient, just as health care personnel wear masks in health care settings.

School Closure

CDC recommendations

School closures and dismissals: CDC might recommend the use of preemptive, coordinated school closures and dismissals during severe, very severe, or extreme influenza pandemics. This recommendation is in accord with the conclusions of the U.S. Community Preventive Services Task Force (https://www.thecommunityguide.org/findings/emergencypreparedness-and-response-school-dismissals-reduce-transmissionpandemic-influenza), which makes the following recommendations:

- The task force recommends preemptive, coordinated school dismissals during a severe influenza pandemic.
- The task force found insufficient evidence to recommend for or against preemptive, coordinated school dismissals during a mild or moderate influenza pandemic. In these instances, jurisdictions should make decisions that balance local benefits and potential harms.



Absent from WHO and CDC Pandemic Plans:

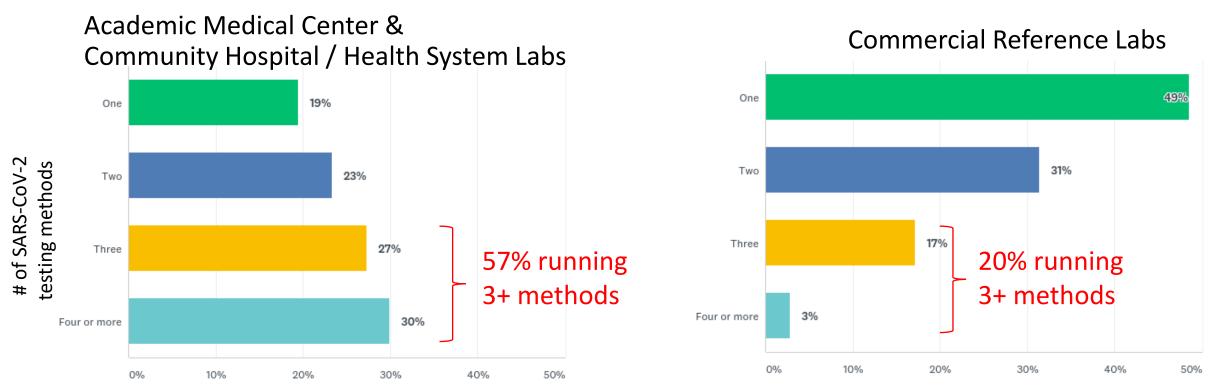
How to get ready for high capacity, rapid and sensitive testing

resulting in ...

Use: <u>whatever you have</u> Add: <u>whatever you can get</u> Experience:

extreme shortfalls in supply chain tests and collection reagents

Due to supply shortages and uncertainties, laboratories are deploying multiple testing methodologies



Why did you choose this SARS-CoV-2 testing method?

"Whatever reagents were able to receive" "Independent supply chain" "Limited kit availability" "Supply chain issues are a major hurdle currently, which is preventing us from moving forward with this as a primary instrument." "We use the [company name's] extraction reagents and they are hard to get and the shortage affects our 24 other LDTs." "We are concerned with this test and have it as a back-up for increased capacity if it needs to be deployed. The supply chain for this test has been very un-reliable." "Next door (Virology Lab) is offering COVID19 testing on three platforms to minimize the risk of inventory shortage."

©2020 AMP COVID-19 Molecular Testing Survey www.amp.org/COVID19 8

All US-based labs: top 10 primary testing methods

SARS-CoV-2 Molecular Testing Methods	Primary*	Secondary	Tertiary
SARS-COV-2 Molecular resting Methods	(n=112)	(n=88)	(n=59)
Laboratory developed testing procedure (LDP / LDT) with EUA submission	21%	8%	<mark>5%</mark>
Roche Molecular Systems cobas SARS-CoV-2	17%	5%	0%
Abbott Molecular RealTime SARS-CoV-2 assay	16%	6%	7%
Cepheid Xpert Xpress SARS-CoV-2 test	8%	19%	25%
Hologic Panther Fusion SARS-CoV-2	<mark>6%</mark>	1%	0%
Quidel Corporation Lyra SARS-CoV-2 Assay	5%	2%	2%
Thermo Fisher Scientific TaqPath COVID-19 Combo Kit	<mark>5%</mark>	<mark>3</mark> %	7%
CDC 2019-nCoV Real-Time RT-PCR Diagnostic Panel	4%	9%	7%
DiaSorin Molecular Simplexa COVID-19 Direct assay	<mark>4</mark> %	13%	9%
Abbott Diagnostics ID NOW COVID-19	<mark>3</mark> %	2%	9%

* Data sorted by the primary testing method from largest to smallest



Performance of COVID tests done in a condensed timeframe

Gaps in analysis of tests due to urgency for testing Rapidly assembled quantified validation and control materials not carefully standardized to each other.

Universal control materials late to arrive and sometimes not well standardized

Requirements for EUA validation of LOD

30 positive and 30 negative clinical specimens

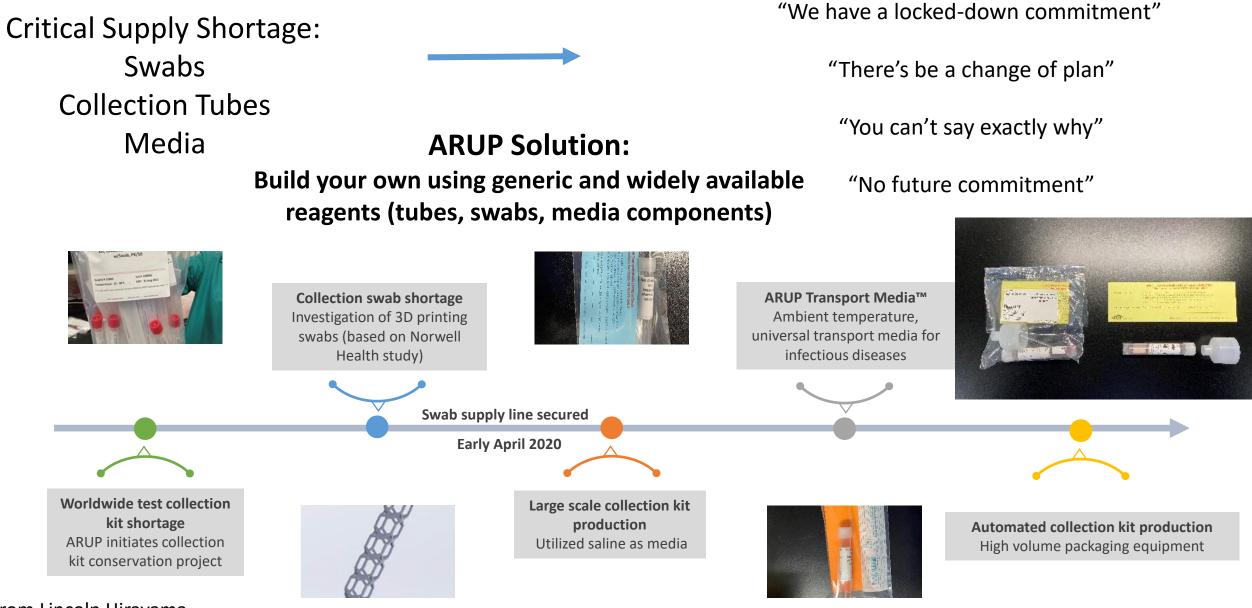
Dilution series of 3 replicates per concentration with inactivated virus on actual patient specimen matrix Confirmation at the final concentration of 20 replications with 19/20 required to claim that concentration as LOD

> **EUA Bridging studies** Altered application of COVID test, now discontinued

Highly accurate comparisons of COVID-19 test sensitivities not available for all tests (<u>https://www.fda.gov/media/135659/download</u>)

Company	Test	LOD	EUA date
Becton, Dickinson & Company	BioGX SARS-CoV-2 Reagents for BD MAX System	40 copies/mL	4/2/20
Abbott	Abbott RealTime SARS-CoV-2 assay	100 copies/mL	3/18/20
Abbott	ID NOW COVID-19	125 copies/mL	3/27/20
Quest Diagnostics	Quest SARS-CoV-2 rRT-PCR	136 copies/mL	3/17/20
Cepheid	Xpert Xpress SARS-CoV-2 test (lab test)	250 copies/mL	3/20/20
Cepheid	Xpert Xpress SARS-CoV-2 test (point of care test)	250 copies/mL	3/20/20
bioMerieux	BioFire COVID-19 Test	330 copies/mL	3/23/20
Qiagen	QIAstat-Dx Respiratory SARS-CoV-2 Panel	500 copies/mL	3/30/20
DiaSorin	Simplexa COVID-19 Direct assay	500 copies/mL	3/19/20
Quidel	Lyra SARS-CoV-2 Assay	800 copies/mL*	3/17/20
lpsum	COV-19 IDx Assay	850 copies/mL*	4/2/20
CDC	CDC 2019-nCoV Real-Time RT-PCR Diagnostic Panel (CDC)	3,160 copies/mL; 1,000 copies/mL*	2/4/20
Co-Diagnostics	Logix Smart Coronavirus Disease 2019 (COVID-19) Kit	4,290 copies/mL	4/3/20
Luminex	NxTAG CoV Extended Panel Assay	5,000 copies/mL	3/27/20
LabCorp	COVID-19 RT-PCR Test	6,250 copies/mL*	3/16/20
Luminex	ARIES SARS-CoV-2 Assay	75,000 copies/mL	4/3/20
GenMark	ePlex SARS-CoV-2 Test	100,000 copies/mL	3/19/20
PerkinElmer	PerkinElmer New Coronavirus Nucleic Acid Detection Kit	3 copies/ reaction	3/24/20
Gnomegen	Gnomegen COVID-19 RT-Digital PCR Detection Kit	8 copies/ reaction	4/6/20
Thermo Fisher	TaqPath COVID-19 Combo Kit	10 copies/ reaction	3/13/20
New York State Department of Public Health	New York SARS-CoV2 Real-time Revers Transcriptase (RT)-PCR Diagnostic Panel	25 copies/ reaction	2/29/20
Mesa Biotech	Accula SARS-CoV-2 Test	200 copies/ reaction	3/23/20
Roche	Cobas SARS-CoV2	0.009 TCID50/ mL	3/12/20
Hologic	Panther Fusion SARS-CoV-2	0.01 TCID 50/mL	3/16/20

Sample Collection: An <u>unexpected?</u> supply chain pinch point



From Lincoln Hirayama

COVID-19 Saliva Collection

- High 'invalid' rate at first 15%!
- Viscosity/higher order protein complexes possibly interfering
- All saliva samples are freezethawed → invalid rate now <1%
- Correct amount very important

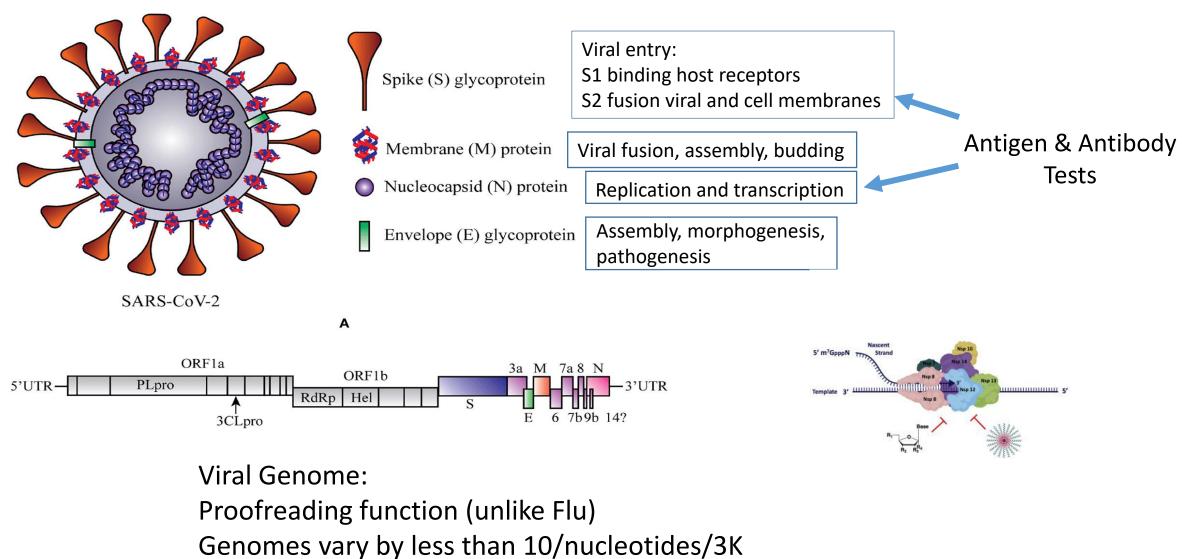




Unacceptable Collection: An under filled saliva collection can generate poor results such as "<u>False</u> Negatives"



Unacceptable Collection: Even a small amount of overfilled saliva can create too much viscosity in the sample and generate <u>"In-valid"</u> results



Multiple highly conserved targets for NAT tests ORFab, RdRp, S, E and N genes

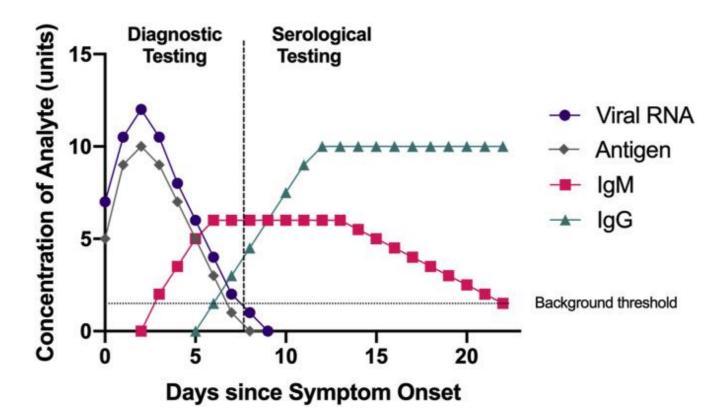
Islam et al. Natureresearch Aug 2020 | https://doi.org/10.1038/s41598-020-70812-6

COVID-19 A Respiratory and Vascular Infection

- Respiratory failure Acute respiratory distress syndrome (ARDS)
- Secondary infections -respiratory infections and bacteremia
- Inflammatory complications -exuberant inflammatory response, with persistent fevers, elevated inflammatory markers, elevated proinflammatory cytokines
- Cardiac and cardiovascular complications Other complications arrhythmias, acute cardiac injury, and shock
- Thromboembolic complications Pulmonary embolism and acute stroke
- Neurologic complications Encephalopathy (common), stroke, movement disorders, motor and sensory deficits, ataxia, and seizures

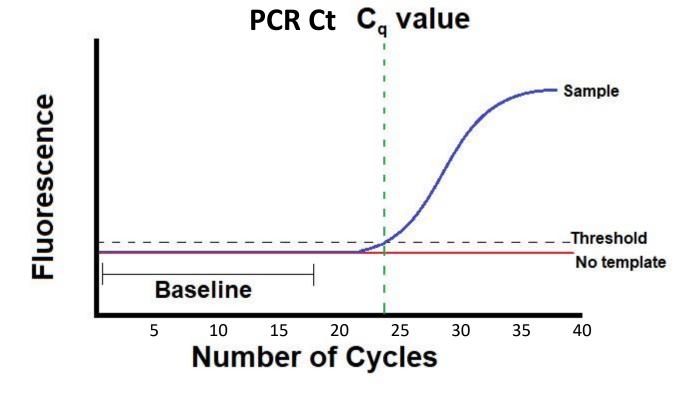
CDC extended symptoms: https://www.cdc.gov/mmwr/volumes/69/wr/mm6930e1.htm

COVID-19 biomarkers for testing



Diagnostic Tests

- Antigen
 - BinaxNow
 - DiaSorin
- NAT high throughput & sensitive
 - Roche PCR (closed box)
 - Hologic fusion PCR (closed box)
 - Hologic Panther TMA RLU signal (closed box)
 - ThermoFisher PCR (open platform)
- NAT POC low throughput-rapid
 - Abbott
 - Cephiad
 - Biofire
- Differentiating clinical vs analytical sensitivity is critical!



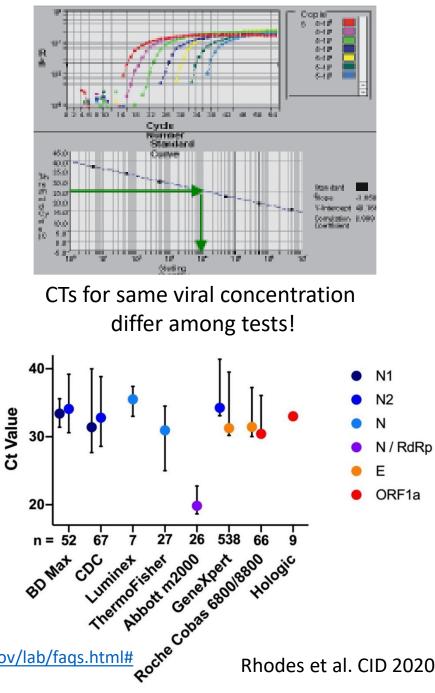
Crossing thresholds (Cts) provide a good approximation of viral concentration in a liquid sample

Cts do not accurately measure viral burden in host

Cts vary among different instruments!

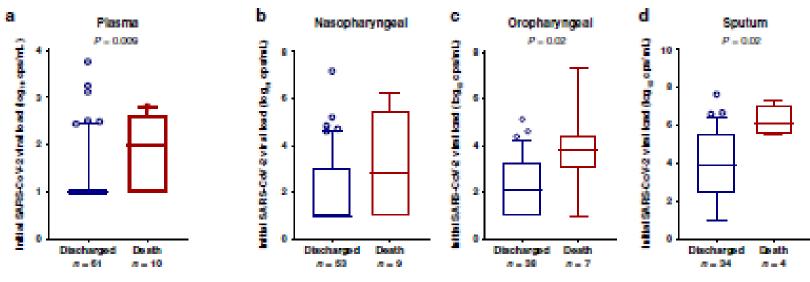
8 fold variability in CTs in 26 lab survey https://www.publichealthontario.ca/-/media

https://www.cdc.gov/coronavirus/2019-ncov/lab/faqs.html# Interpreting-Results-of-Diagnostic-Tests



Utility PCR Crossing Threshold Analysis/reporting

- Predict rising tide of infections at a given phase of pandemic
- Understand dynamics of viral shedding
- Release patients with still detectable virus from quarantine
- Indicator disease severity and likelihood of death
- Stratification of patient risk
- Crossing thresholds vary with sample type.

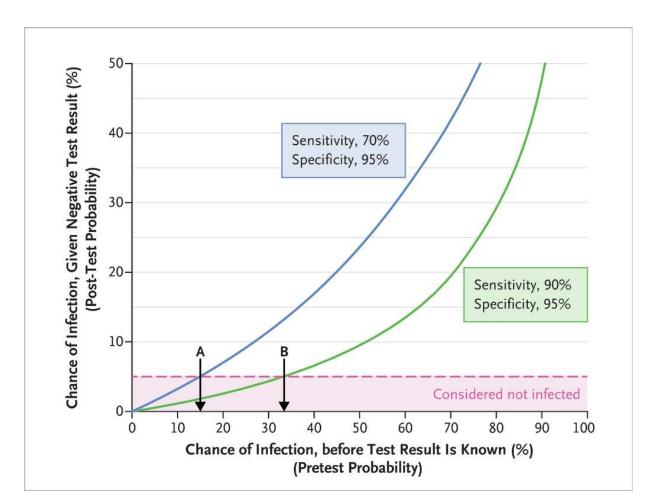


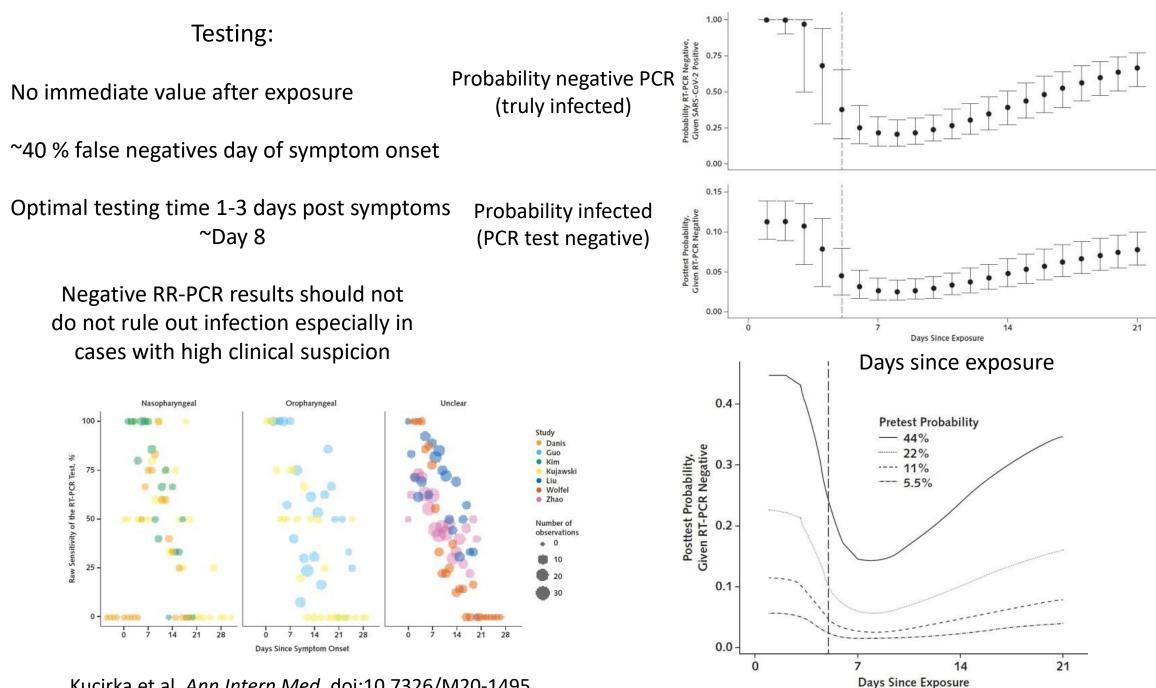
NATURE COMMUNICATIONS | (2020) 11:5493 |

Bryan et al. Open Forum Infect Dis. 2020 Nov 3;7(12)

Analytic vs Clinical Sensitivity and Specificity

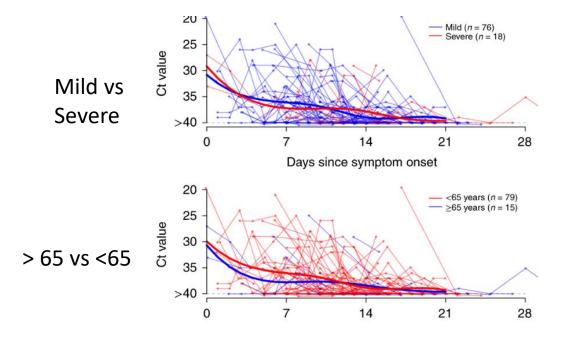
- Traditional test performance validation:
 - Clinical or contrived specimens
 - Comparison to reference test
- Sensitivity:
 - % positivity patients with disease
 - No absolute reference for disease status
 - Reference material lacking
 - <u>COVID EUA</u> allows establishing agreement with results positive material from symptomatic patients or contrived material
 - Swabs and saliva miss infected material
- Specificity:
 - High and reliable for NAT tests



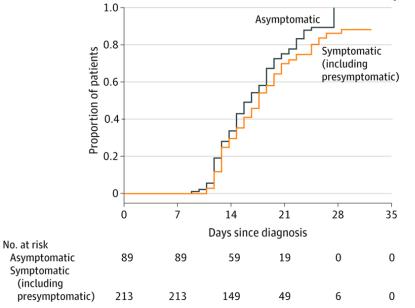


Kucirka et al. Ann Intern Med. doi:10.7326/M20-1495

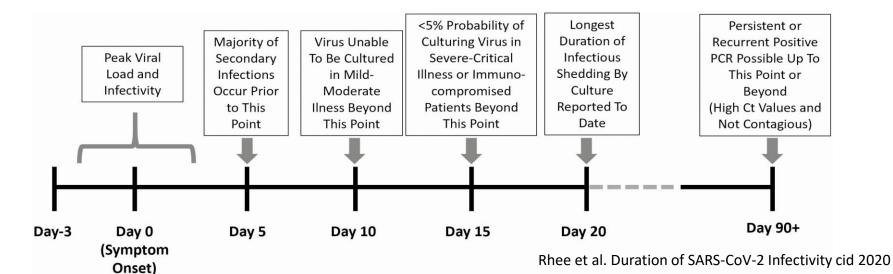
COVID-19 Viral Dynamics and Shedding



He et. Nature Medicine | VOL 26 | 672 MAY 2020 | 672–675 |



Lee et al. JAMA Internal Medicine November 2020 Volume 180, Number 11



Symptomatic vs Asymptomatic

Approaches to test comparisons and power to predict % positives and % missed

- Traditional FDA requirements
 - Rigorous LOD
 - Large population studies
 - Examples: HIV, HCV HPV, Flu
- Comparative testing of samples of different viral concentration identified by PCR crossing thresholds (CTs).
 - Across the spectrum viral concentration
 - Within the spectrum of low viral concentration
- Prediction based on established test LOD and relative CTs, +/- confirmatory testing

Comparative Assay Testing of a Sampled Population

Comparison of Two High-Throughput RTPCR Systems for the Detection of COVID-19

Kraner et al. JCM May 2020

Characteristics of discordant specimens

RT-PCR Result					
Target 1Target 2Ct ValueCt Value					
35.26	37.69				
N/A	N/A				
N/A	N/A				
N/A	N/A				
N/A	N/A				
N/A	38.26 °				
N/A	N/A				
35.37	N/A				
N/A	37.64 c				
35.84	38.05				
N/A	N/A				
N/A	N/A				
36.13	N/A				

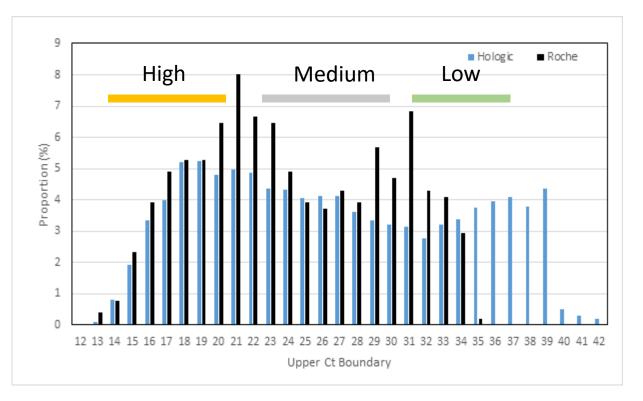
cobas 6800 SARS-CoV02

Panther Fusion SARS-CoV-2 RT-PCR Result

CT Value	
N/A	
36.5	
38.1	
38.1	
36.9	
N/A	
38	
N/A	
N/A	
N/A	
N/A	
38	
38.1	

N/A

Distribution CTs Hologic and Roche (~ 5% Positive Symptomatic Utah Population)

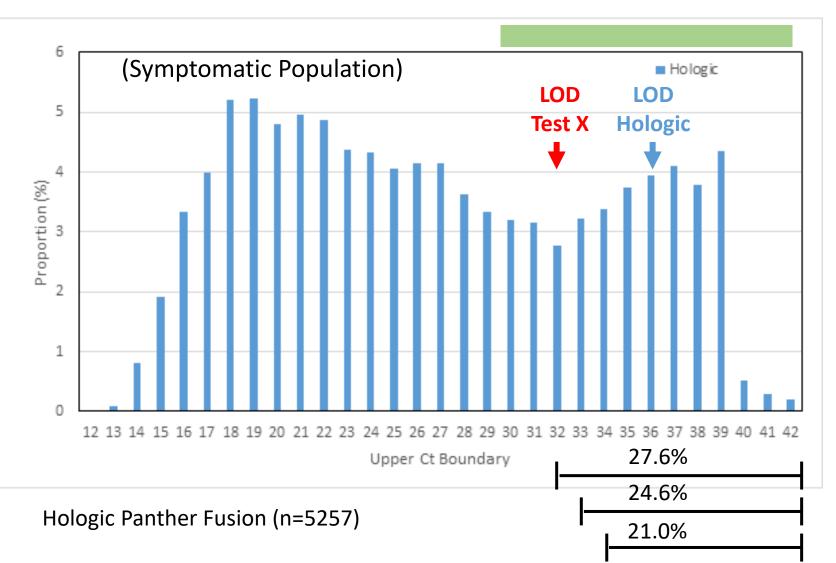


Hologic Panther Fusion (n=5257)

Roche cobas SARS-CoV-2 Assay (n=511)

Berry et al. JCM.00743-20

Comparative testing within a spectrum of low viral concentrations

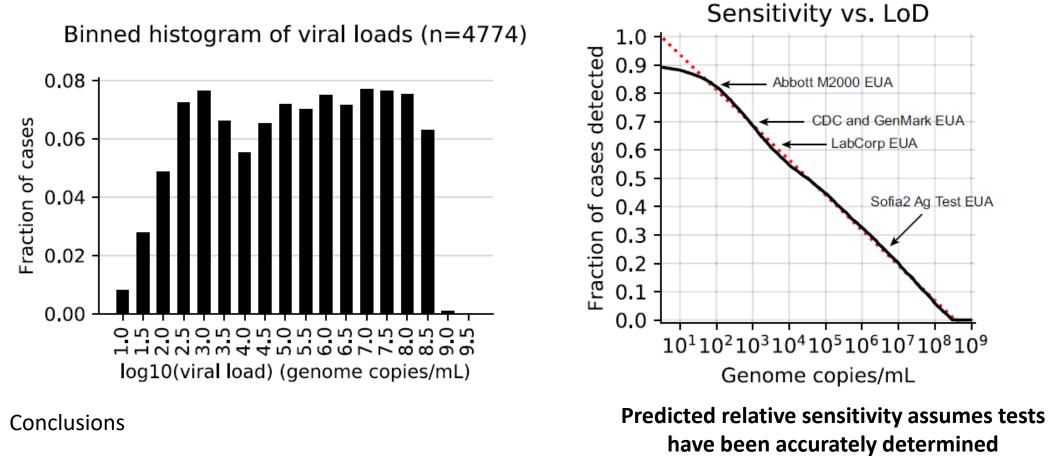


Selectively Test Samples In "Low Range"

Advantage: test many low positives more accurate predictions

Calculate percentage of samples "likely" missed based on comparative CTs and knowledge of LODs

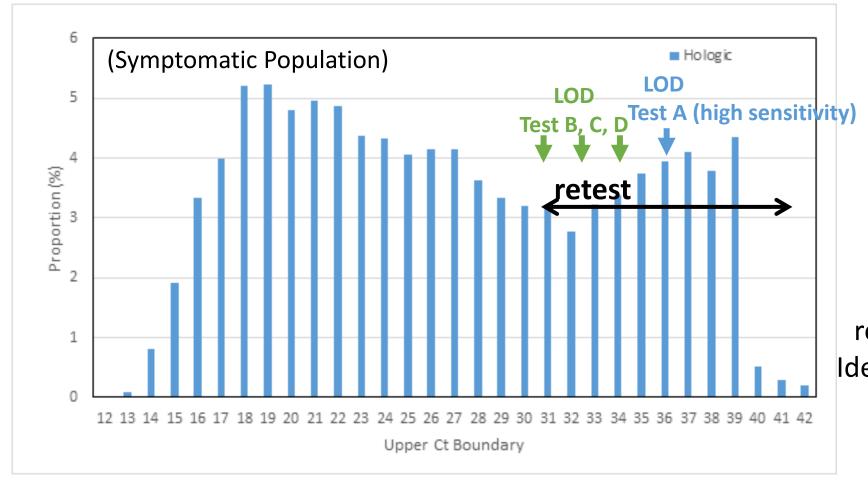
Calculate expected CT based on Anchor CT and LOD of Less Sensitive Test Use of CTs to determine numbers of low positives in a population and predict negatives among different assays



Arnaout et al. bioRxiv preprint doi: https://doi.org/10.1101/2020.06.02.131144

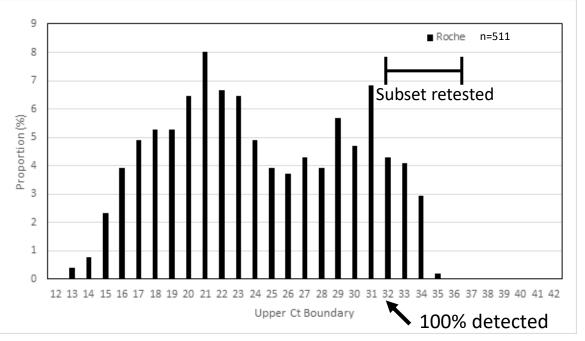
Limits of Detection (LOD)

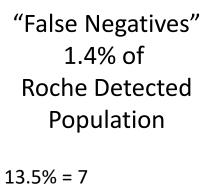
Prediction based on established test LOD and relative CTs with Confirmation



y) Calculate percentage of samples "likely" missed based on comparative CTs of several tests and knowledge of LODs with confirmation by retesting low positive samples Identified by high sensitivity test

Hologic Panther Fusion (n=5257)





samples missed

CDC assay results for 52 low positive Roche samples

	_			
Roche Ct	Positive	Negative	Inconclusive	Row Total
31-31.9	100% (18)	-	-	100% (18)
32-32.9	85.0% (17)	10.0% (2)	5.0% (1)	100% (20)
33-33.9	61.5% (8)	30.8% (4)	7.7% (1)	100% (13)
≥34	-	100% (1)	-	100% (1)
Grand Total	82.7% (43)	13.5% (7)	3.8% (2)	100% (52)

CDC assay results for 140 low positive Hologic samples

Row Total

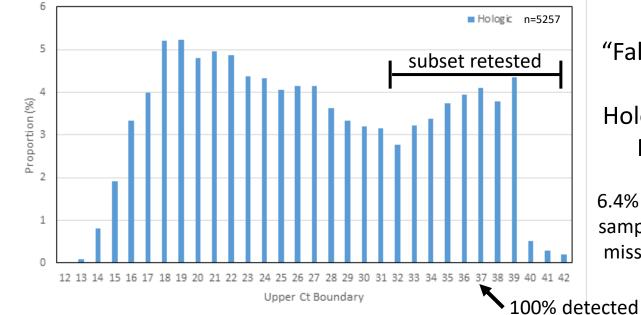
100% (17) 100% (29)

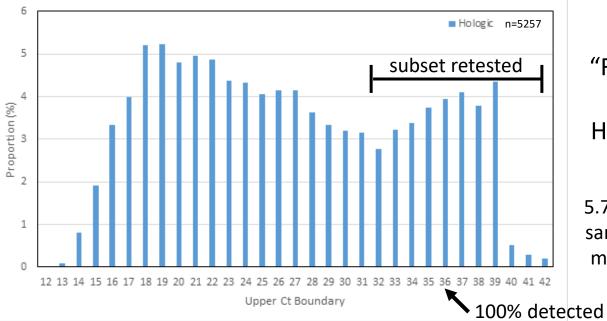
100% (21)

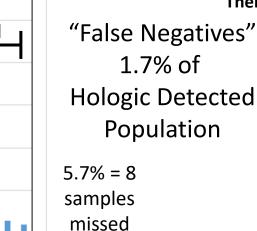
100% (13) 100% (15) 100% (19)

100% (20) 100% (6) 100% (140)

						_		-		-	-	-
			E F	lo logic	n=5257				C	DC Assay Res	ult	
								Hologic Ct	Positive	Negative	Inconclusive	
subset retested		"False Negatives"	32-32.9	94.1% (16)	-	5.9% (1)*						
			_				1.7% of	33-33.9	100% (29)	-	-	
								34-34.9	100% (21)	-	-	
ι.	_	a I					Hologic Detected	35-35.9	100% (13)	-	-	
Ħ	н.	HH	н				Population	36-36.9	100% (15)	-	-	
							Population	37-37.9	94.7% (18)	5.3% (1)	-	
H				_		_		38-38.9	65.0% (13)	30.0% (6)	5.0% (1)	
							6.4% = 9	>39	33.3% (2)	33.3% (2)	33.3% (2)	
H		\square			-	_	samples	Grand Total	90.7% (127)	6.4% (9)	2.1% (3)	
					LL.		missed	*Not retested – Insu	ıfficient RNA remainiı	ıg		
28 29	30 31 32 3	3 34 35	36 3	7 38 3	9 40 41 4	12						



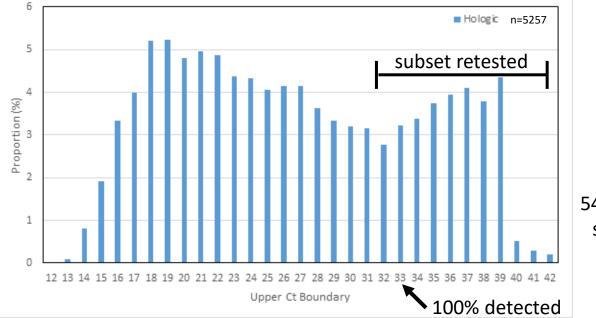




ThermoFisher assay results for 140 low positive Hologic samples

	Thermo			
Hologic Ct	Positive	Negative	Inconclusive	Row Tota
32-32.9	100% (17)	-	-	100% (17)
33-33.9	100% (29)	-	-	100% (29)
34-34.9	100% (21)	-	-	100% (21
35-35.9	100% (13)	-	-	100% (13
36-36.9	100% (15)	-	-	100% (15
37-37.9	94.7% (18)	5.3% (1)	-	100% (19
38-38.9	75.0% (15)	15.0% (3)	10.0% (2)	100% (20)
>39	66.7% (4)	33.3% (2)	-	100% (6)
Grand Total	94.3% (132)	4.3% (6)	1.4% (2)	100% (140

Quidel assay results for 140 low positive Hologic samples



"False Negatives" 14.6% of
Hologic Detected Population
F4 20/ - 76

54.3% = 76
samples
missed

	Qui	ult		
Hologic Ct	Positive	Negative	Invalid	Row Total
32-32.9	100% (17)	-	-	100% (17)
33-33.9	89.7% (26)	6.9% (2)	3.4% (1)	100% (29)
34-34.9	38.1% (8)	57.1% (12)	4.8% (1)	100% (21)
35-35.9	23.1% (3)	61.5% (8)	15.4% (2)	100% (13)
36-36.9	13.3% (2)	73.3% (11)	13.3% (2)	100% (15)
37-37.9	-	100% (19)	-	100% (19)
38-38.9	-	95.0% (19)	5.0% (1)	100% (20)
>39	-	83.3% (5)	16.7% (1)	100% (6)
Grand Total	40.0% (56)	54.3% (76)	5.7% (8)	100% (140)

Assessing Sensitivity COVID-19 Antigen Test (BinaxNOW)

BinaxNOW False N

8+ ASx

BinaxNOW True Pos

Adults

Days Post Symptom Onset

Sensitivity 84.6% 🔺

Specificity 100%

Symptomatic Population

1380 Adults 928 Children

Cycle Threshold

8+ ASx

30-

20-

10-

0

Asymptomatic Population 2,645 **College Students** p < 0.0001 **40** ThermoFisher TaqPath 30. Ct Value 20 10 Positive Negative **BinaxNOW COVID-19 Ag Card** All positive RT-PCR Discordant negative BinaxNOW Ct 20 **Co-collected Nasal Swabs** 46 (1.7 %) NAT positive 24 (0.9%) Antigen positive 53.3 % sensitive, 100% specific

All group sensitivity & Crossing Threshold 99.3% Ct <25 95.8% Ct < 30 81.2% Ct <35.

Children

Days Post Symptom Onset

Sensitivity 95%

Specificity 100%

40-

30-

20-

10-

2 3

Cycle Threshold

Pollock et al. https://doi.org/10.1101/2021.01.09.21249499;

Perchetti et al. J Clin Microbiol doi:10.1128/JCM.02880-20

Okoye/Barker/Pearson et al. Accepted JCM Jan 2021

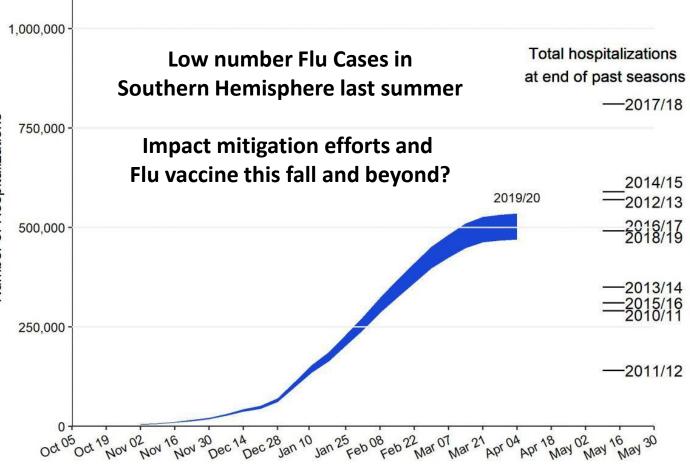
Ct Value



Co-Circulation COVID and Other Respiratory Viruses (Flu A, Flu B, RSV AB 2020-21? 39,000,000 - 56,000,000 flu **illnesses** 1,000,000 Number of Hospitalizations 750,000 18,000,000 - 26,000,000 flu medical visits 500,000 410,000 - 740,000 flu hospitalizations 250.000

24,000 - 62,000

flu deaths



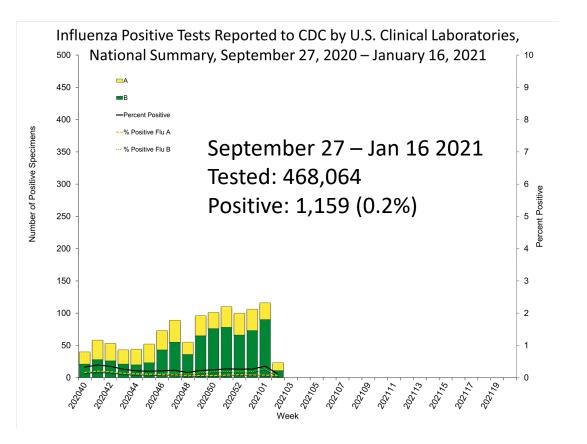
https://www.cdc.gov/flu/about/burden/preliminary-in-season-estimates.htm

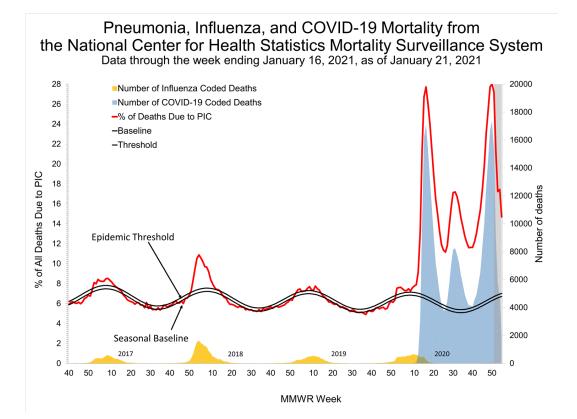
COVID/Flu Twindemic?

Issues

- Shared symptoms
- Co-infections more lethal?
- Competition testing resource
- Availability of high throughput Co-tests

- Commitment to a specific % Co-test reagents
- Adapt to COVID testing process
- Deal with unused reagents in case of low Flu season







Centers for Disease Control and Prevention "Because some of the symptoms of flu and COVID-19 are similar, it may be hard to tell the difference between them based on symptoms alone, and testing may be needed to help confirm a diagnosis."

Co-Testing

Attractive to symptomatic patient

Detect and differentiate COVID, Flu, RSV For diagnosis, treatment, tracking (provides a "final" diagnosis)

Potential to improve operational efficiency and cost for patients, labs, and public health

Symptom	Influenza	COVID
Fever/Chills	\checkmark	\checkmark
Cough	\checkmark	\checkmark
Sore throat	\checkmark	\checkmark
Nasal congestion	\checkmark	\checkmark
Body aches	\checkmark	\checkmark
Headache	\checkmark	\checkmark
Fatigue	\checkmark	\checkmark
Vomiting/diarrhea	\checkmark	\checkmark
Loss of taste or smell		\checkmark
Shortness of breath		\checkmark

RSV: Disease Burden and Impacted **Populations**

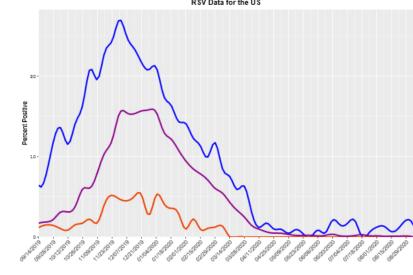
Each year in U.S. :

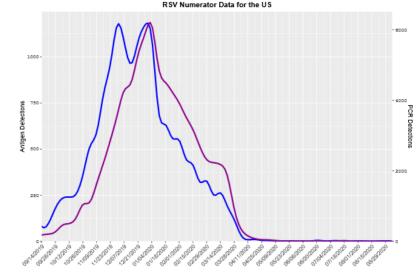
•2.1 million outpatient visits among children younger than 5 years old¹

•57,527 hospitalizations among children younger than **5 years old**¹ •177,000 hospitalizations among adults older than 65 years² "Revalue of the among addition of the second with a disease burden similar to that of non-pandemic influenza A in a population in which the prevalence of vaccination for influenza is high" Falsey et al.

Testing for RSV is appropriate for adults over 65 as well as children as well as transplant patients of any age

RSV in a COVID/Flu not indicated in uncompromised population 5-65





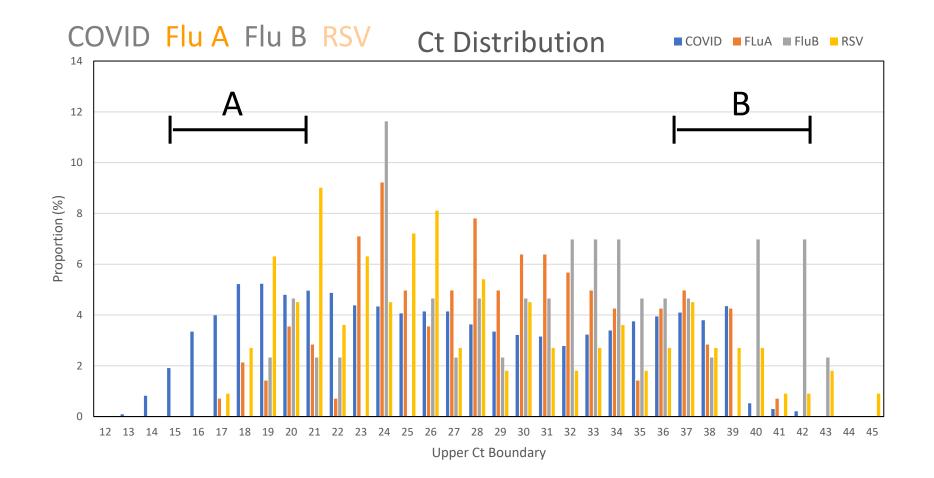
Hall CB et al. New Engl J Med. 2009;360(6):588-98 The burden of respiratory syncytial virus infection in young children Falsey AR et al. New Engl J Med. 2005;352(17):1749-59 Respiratory syncytial virus infection in elderly and high-risk adults

RSV Co-infections in Hospitalized Young Children

From: American Academy of Pediatrics https://www.aappublications.org/news/2020/06/10/coronavirusbronchiolitis061020

Researchers studied 1,880 children hospitalized with bronchiolitis from two multicenter cohorts — a group under 2 years from 2007-'10 and a group under 1 year from 2011-'14. Children were tested for 18 viruses, including four endemic coronaviruses (CoV), which are not the newest CoV identified (SARS-CoV-2).

Roughly 12% of the children had a coronavirus. **Of those, 85% also had another infection, most commonly respiratory syncytial virus (RSV),** according to "Severe Coronavirus Bronchiolitis in the Pre-COVID-19 Era," (Mansbach JM, et al. *Pediatrics*. June 10, 2020 <u>https://doi.org/10.1542/peds.2020-1267</u>). What is the impact of Population A viruses on Population B detectability for Co-Positive Samples?



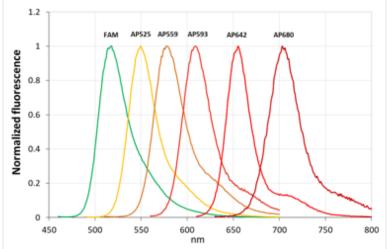
Data: Panther fusion

Co-testing Analytic Challenges (single well, high throughput assays)

- Optimize tests for high sensitivity and specificity for multiple viral targets
- Single well multiplex Co-detection
 - 4 targets: COVID, IC, Flu A, Flu B
 - 5 targets: COVID, IC, Flu A, Flu B, RSV (AB)
 - 6 targets: COVID, IC, Flu A, Flu B, RSV (AB), host target

• ISSUE: Fluorescence bleed through <u>Impacts Test Specificity</u>

- Current Limitation:
 - Max out at 5-6 targets
 - Multi-well or extract once amplify many
- Current Fluor Limitations:
 - Max out at 5-6 targets
 - Alternative: Multi-well or extract once amplify many
- Co-positive samples: Potential for suppression of signal of minor viral population by major population?
 - Impacts Test Spensitivity



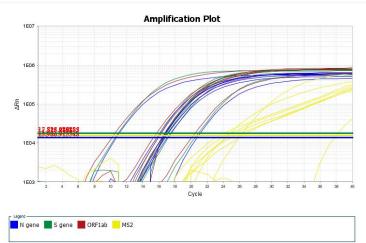
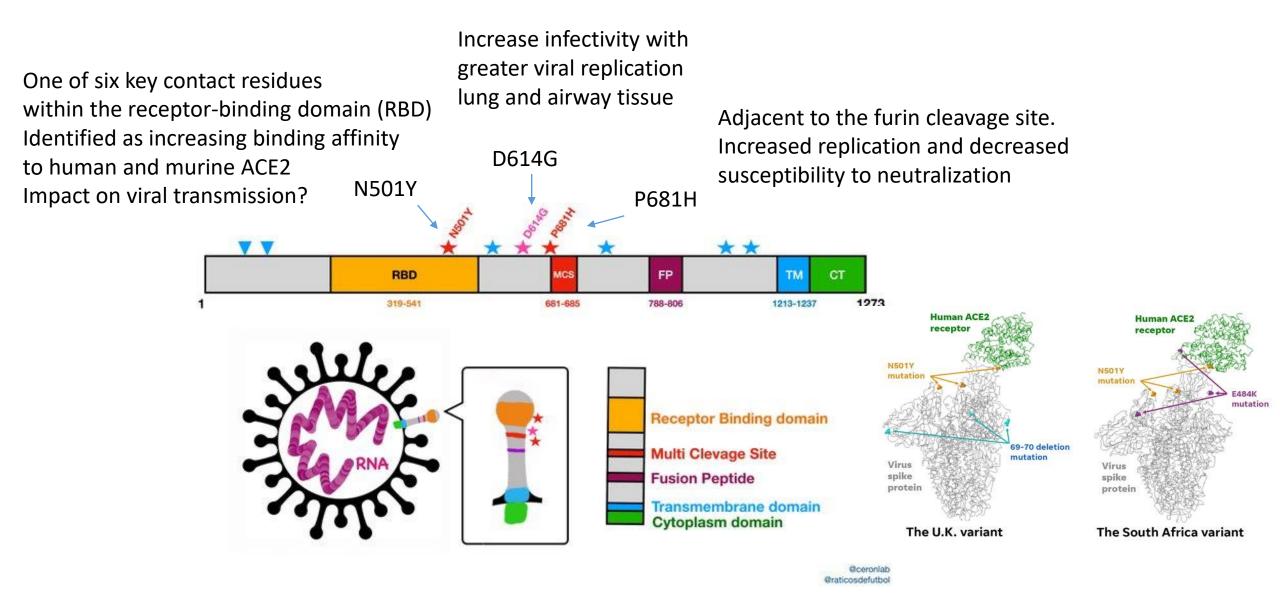


Image courtesy Walt Mahoney Elitec Group

ARUP study:Roche, Hologic, and Chromacode assays are resilient to bleed through and suppression!

COVID-19 Mutations and Their Consequences

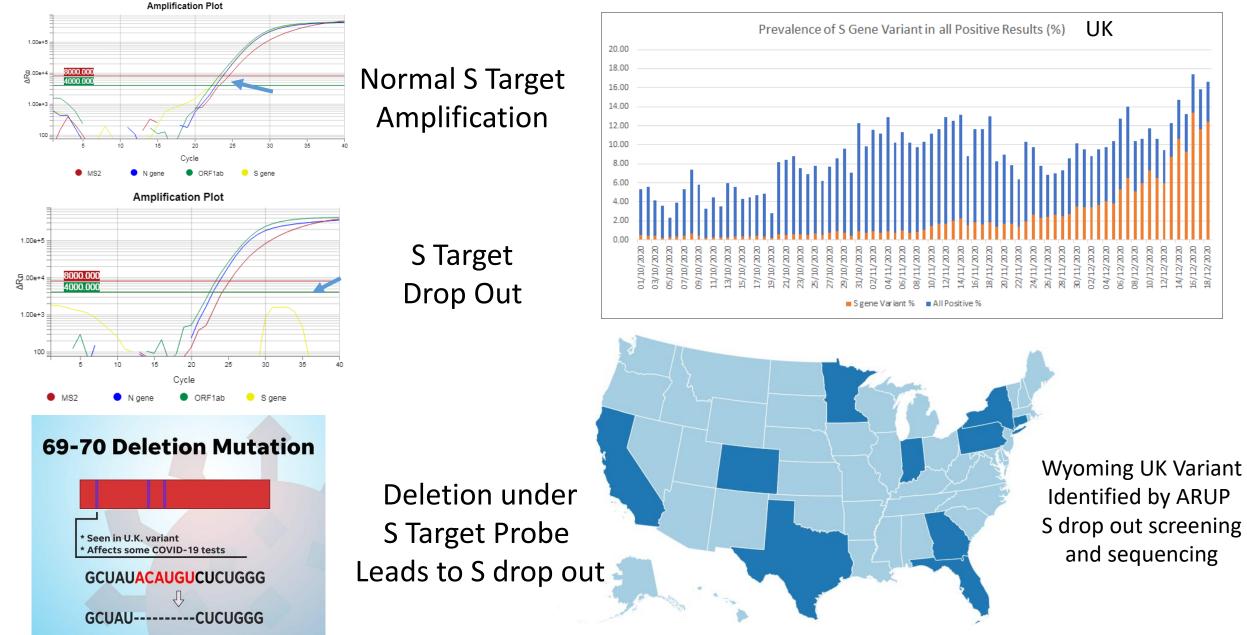


Nature | Vol 585 | 10 September 2020 | Corrected 16 September

COVID-19 Mutations and Timeline

- D614G variant
 - Emerged in late Jan. or early Feb. 2020
 - Replaced the initial SARS-CoV-2 strain identified in China
 - Increased infectivity and transmission
- "Denmark Mink" variant
 - Emerged mink farm sector June 2020
 - Variant "cluster 5" Nov 5 in 12 human cases with new mutations.
 - Worry of reduced viral neutralization and vaccine effectiveness not confirmed
- VOC 202012/01, lineage B.1.1.7, "UK" variant
 - Reported Kent England Sept 2020
 - Spread rapidly to be dominant English strain
 - Clear capacity to spread more quickly
 - Multiple mutations including D614G, P681H and deletion 69-70.
- 501Y.V2 lineage B.351 variant "South Africa"
 - Reported Dec 2020 with combination of mutations
 - Worry more rapid spreading, vaccine resistant

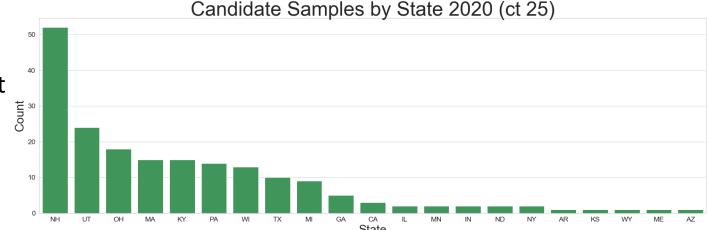
Identification and Tracking "UK" Variant: Screen for S Target Drop Outs



https://www.cdc.gov/coronavirus/2019-ncov/transmission/variant-cases.html

Issues "UK" Variant screening by S Target Drop Out Identification

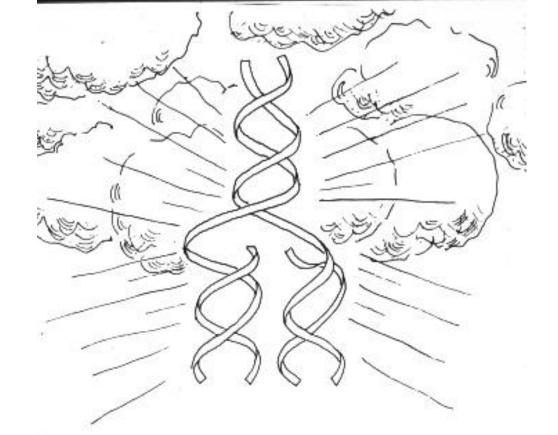
- Most tests don't signal S target drop out.
 - Will current tests be modified to avoid drop out?
- S target drop out requires confirmatory identification by sequencing
 - Low prevalence population presents high burden for sequencing
 - U.S. COVID sequencing capacity is lacking (compare to UK)
- Solutions
 - Expand sequencing capacities (federal, academic, commercial)
 - Engage labs with high "sequence to diagnosis" capacities (Ginkgo model)
 - Validate Multiplex mutation detection assavs



>5,000 Positive Samples screened for S drop out ~ 150 drop outs identified and sequenced Jan 8-15 4 UK variants identified

Unmet Needs in Testing

- Super NAT Tests
- Home sample collection conveniently and safely linked to testing
- Affordable and scalable sequencing
- Better informatics for communicating with test populations



ARUP, University of Utah, and Utah State Teams

