



Pandemic lessons learned from the mitochondrial disease community

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2022 MitoAction

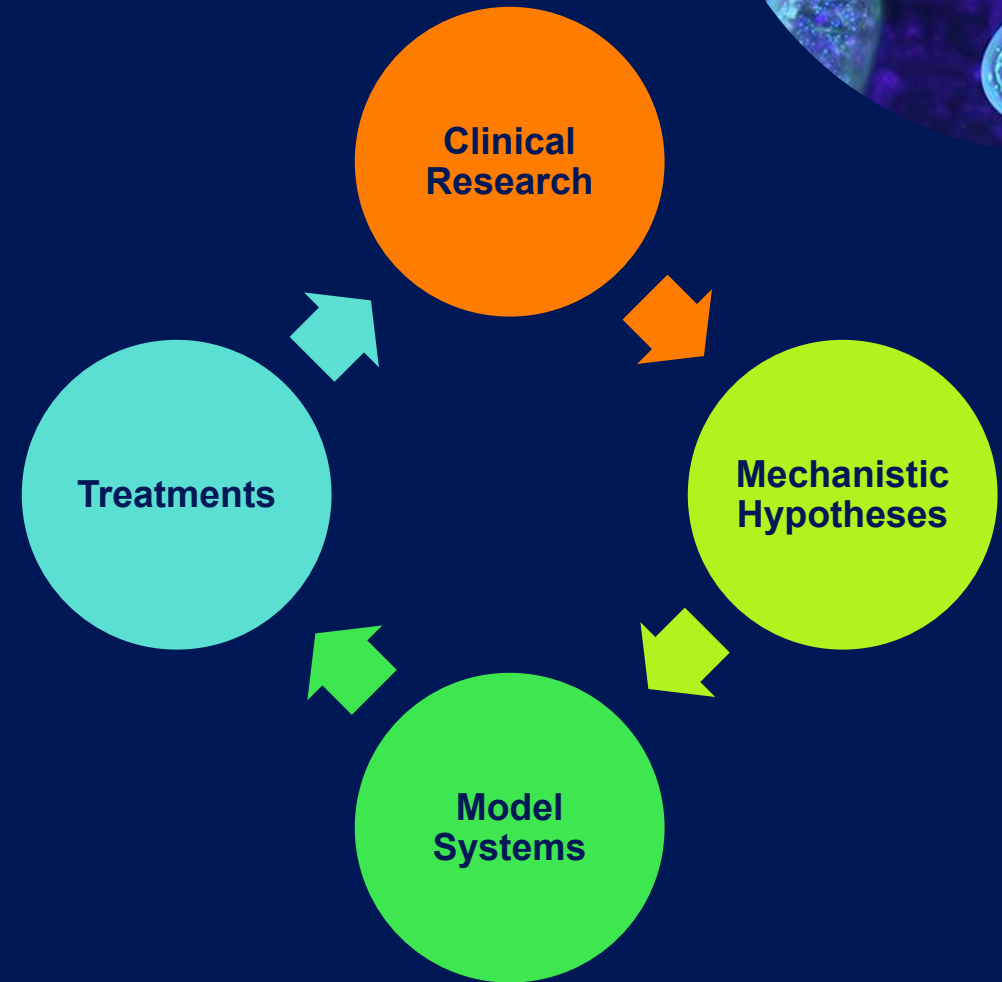


National Human Genome
Research Institute



MINI Section Mission

“The Metabolism, Infection and Immunity (MINI) Section aims to define the risk factors and mechanisms involved in infection-related clinical decline in children with mitochondrial disease.”



Why do we care about infection and mitochondrial disease?

Infection and mitochondrial disease (MD)

- Sepsis and pneumonia are two most common causes of death in children with MD (Eom et al., 2017).
- Sepsis is one of the top five admitting diagnoses for pediatric patients with MD (McCormack et al., 2017) .
- Up to 80% of children with MD experience recurrent infections, mostly respiratory (Tarasenko et al., 2017) .
- Intercurrent infection is a leading cause of episodic neurodegeneration in mitochondrial disease (Edmonds et al., 2002).
- COVID-19 pandemic represents a threat to patients with mitochondrial disease.

A photograph of a neonatal intensive care unit (NICU). In the foreground, a newborn baby is lying in a hospital bed, covered with a patterned blanket. The baby's head is visible, and they are connected to various medical tubes and wires. To the right of the bed, there is a large medical machine, likely a ventilator or a pump, with a digital display showing red numbers. Above the machine, a large, orange, star-shaped balloon with a smiling face is attached. In the background, there are several other medical monitors and equipment, including a Philips monitor displaying vital signs. The room has a window with white blinds. A large, semi-transparent blue circle is overlaid on the left side of the image, containing the title and a list of bullet points.

Metabolic decompensation

- *In extremis* (life-threatening)
 - Bioenergetic failure
 - Lactic acidosis
 - Disease progression
 - Organ failure (e.g. liver failure)
 - Encephalopathy
 - Metabolic “stroke”
 - Sequelae
- Extensive ICU care
- Viral infections
- Treatment is limited

**Therefore, one of the goals of the
MINI Section is to keep patients with
mitochondrial disease healthy**

Dec. 31, 2019



China alerts World Health Organization (WHO) to several cases of pneumonia with no known cause in Wuhan. The disease goes on to be named COVID-19.

Jan. 7



WHO officials announce they have identified a new virus named SARS-CoV-2 that causes COVID-19. It belongs to the coronavirus family, which includes viruses that cause SARS, MERS and the common cold.

Jan. 11



China announces the first death linked to COVID-19.

Jan. 13



WHO reports the first case outside of China in Thailand.

Feb. 26



National Institutes of Health (NIH) begin the first clinical trial in the U.S. for a potential COVID-19 treatment, remdesivir, an antiviral drug originally developed to treat Ebola.

Feb. 29



The FDA took steps to expand novel coronavirus testing to hospital clinical microbiology laboratories.

Mar. 11



WHO declares COVID-19 a pandemic, with more than 100,000 cases and 4,000 deaths in 114 countries.

Apr. 2

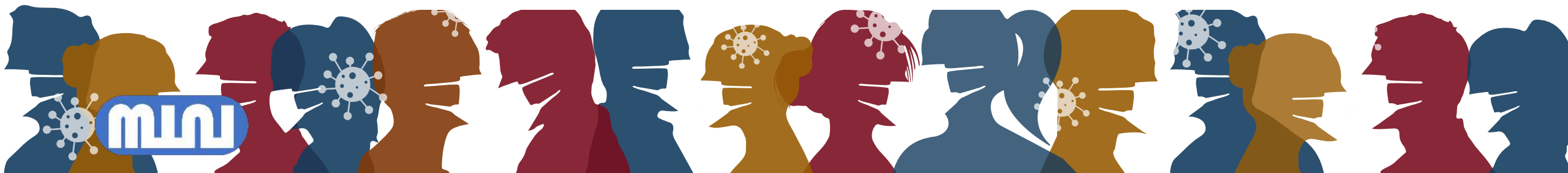


Confirmed cases of COVID-19 top 1 million worldwide.

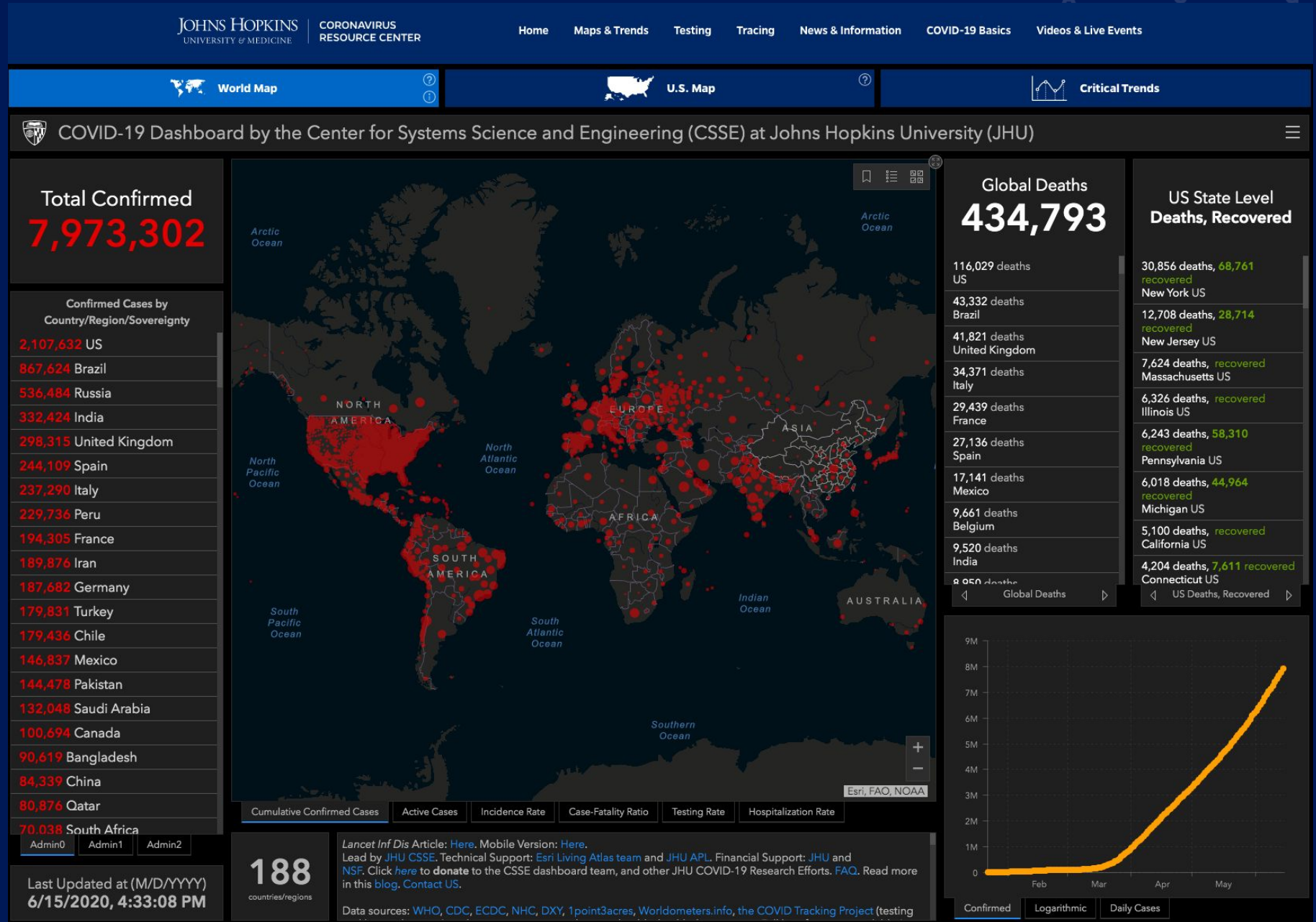
Apr. 10



Global deaths due to COVID-19 top 100,000.



June 15, 2020



Addressing COVID19 and beyond

Feb-Mar 2020
UMDF
COVID19
Position Statement

May 2020
Phase 2: Acute Infection in
Mitochondrial Disease: An
Observational Prospective
Natural History Study of
Metabolism, Infection and
Immunity During the
COVID19 Pandemic
(Actively recruiting)

Late Summer/Fall 2020
Phase 3: SARS-CoV-2
Exposure in Children
with Mitochondrial
Disease

2021-
Phase 5: Viral History
and Mitochondrial
Disease



NIH MINI Study:

- Study of infection and immunity in mitochondrial disease
- Instituted remote enrollment/samples for participants
- Invited to NIH Clinical Center at a later date

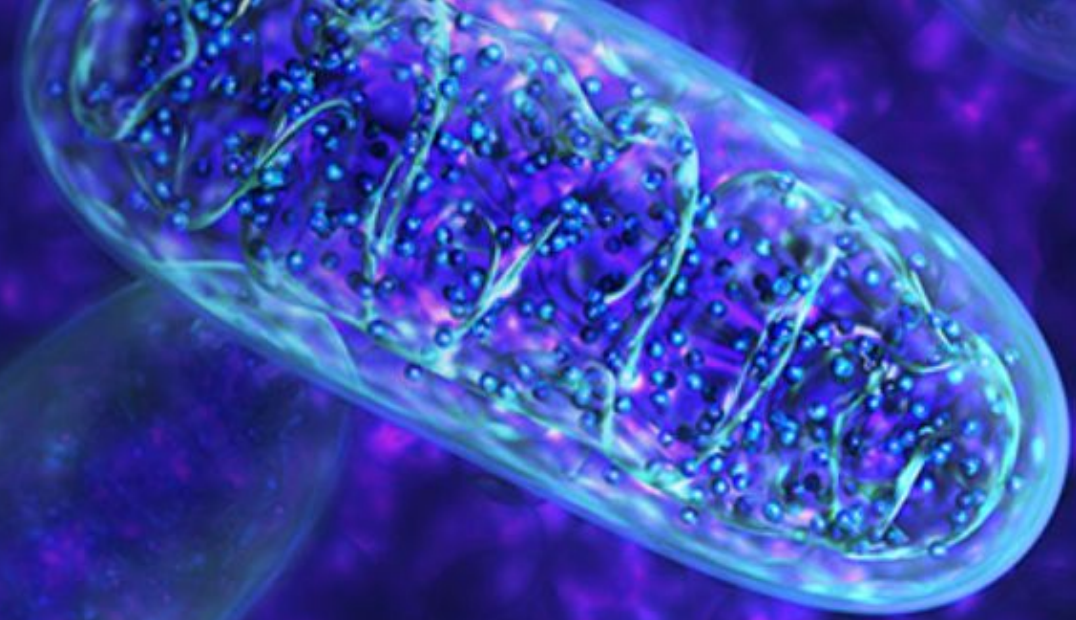
Apr 2020
Phase 1: Understanding
the Experience of the
Mitochondrial Disease
Community During the
COVID19 Pandemic
(>450 participants)

Late Apr 2020
Ask the Mito Doc:
COVID19

June 2020
Phase 1: Understanding
the Experience of the
Mitochondrial Disease
Community During the
COVID19 Pandemic
(Open now!)

2021-2022
Phase 4: SARS-CoV-2
Vaccination and
Mitochondrial Disease

Metabolism, Infection and Immunity (MINI) Study



Understanding why people with mitochondrial disease decline during infection is the first critical step to improving their well-being.



Dr. Peter McGuire



Dr. Eliza Gordon-Lipkin



Ms. Shannon Kruk

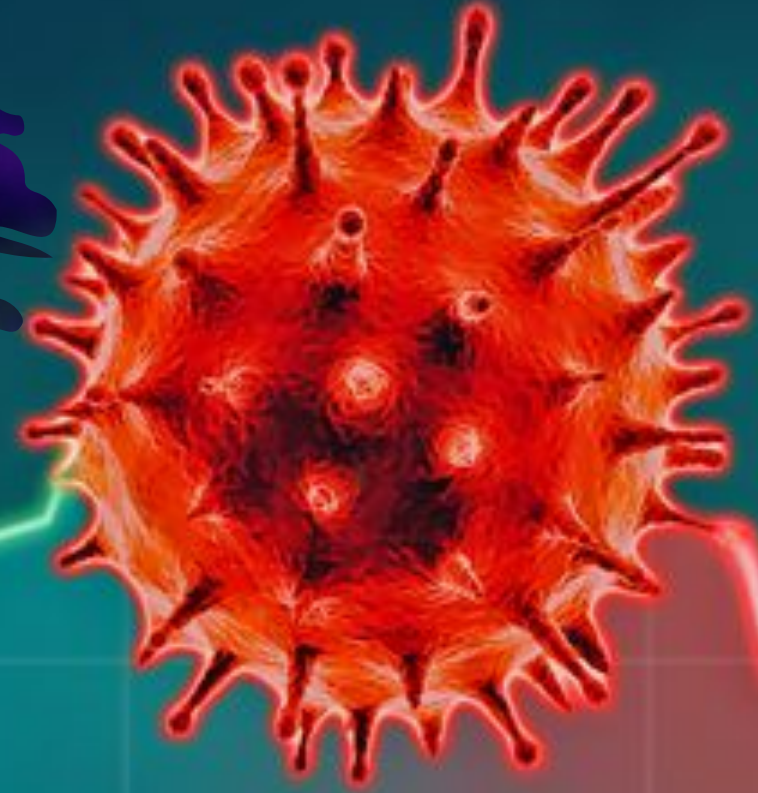
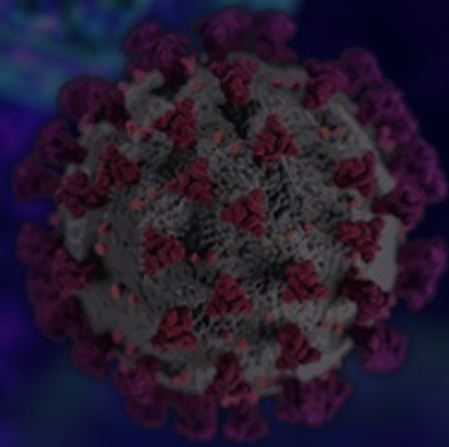


NIH MINI Study: Metabolism, Infection and Immunity (NCT01780168)



- Natural history study of infection and immunity in children with MD
 - ✓ Infection history
 - ✓ Immune function
 - ✓ Disability

**Understanding the
Experience of the
Mitochondrial
Disease
Community During
the COVID-19
Pandemic**



Online surveys



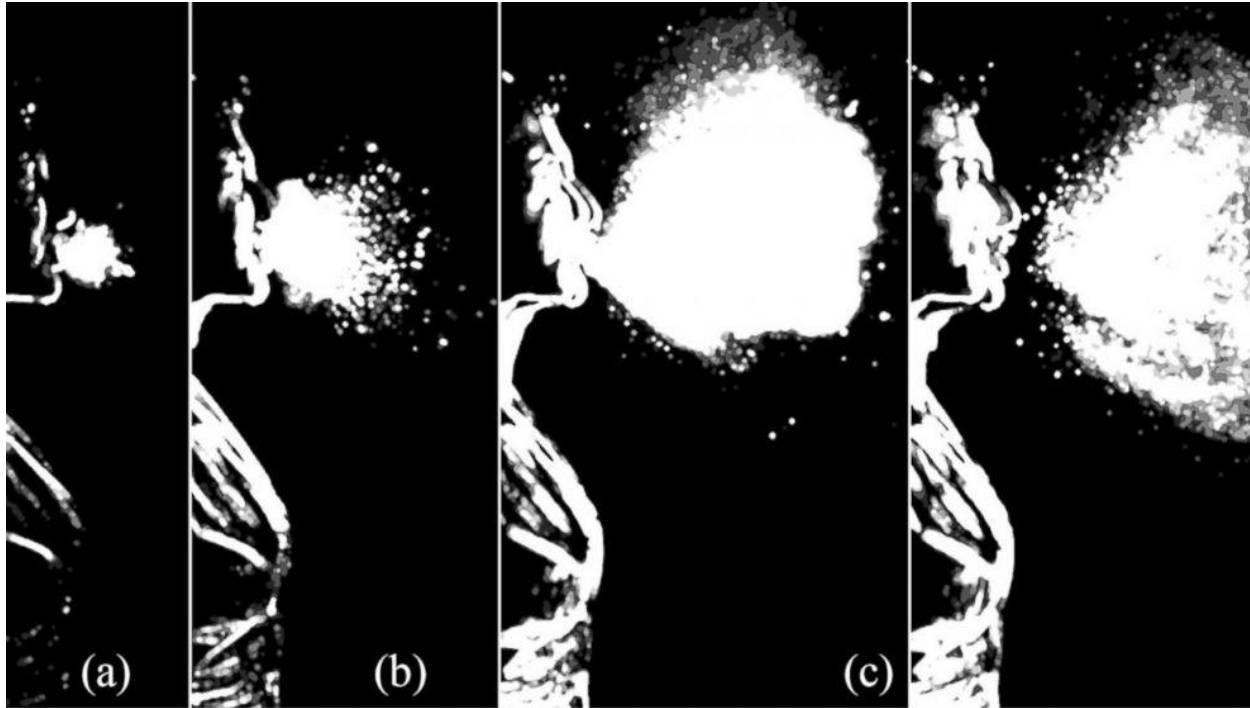
- April 2020, June 2020
- 688 responses; 82% completion rate.
- 30% pediatric MD patients.
- Diagnoses: Mitochondrial Disease Not Otherwise Specified (33%), Mitochondrial Myopathy (28%) and Leigh Syndrome (11%).
- 62% known pathogenic variant
- 5 positive COVID-19 cases
- 68 requested a SARS-CoV-2 test, 14 (21%) were *unable* to receive testing.



Symptoms That Overlap with COVID19

Symptoms that overlap with COVID19 occurred frequently:

Fever	23%
New Cough	15%
New Shortness of Breath	14%
Pneumonia	3%



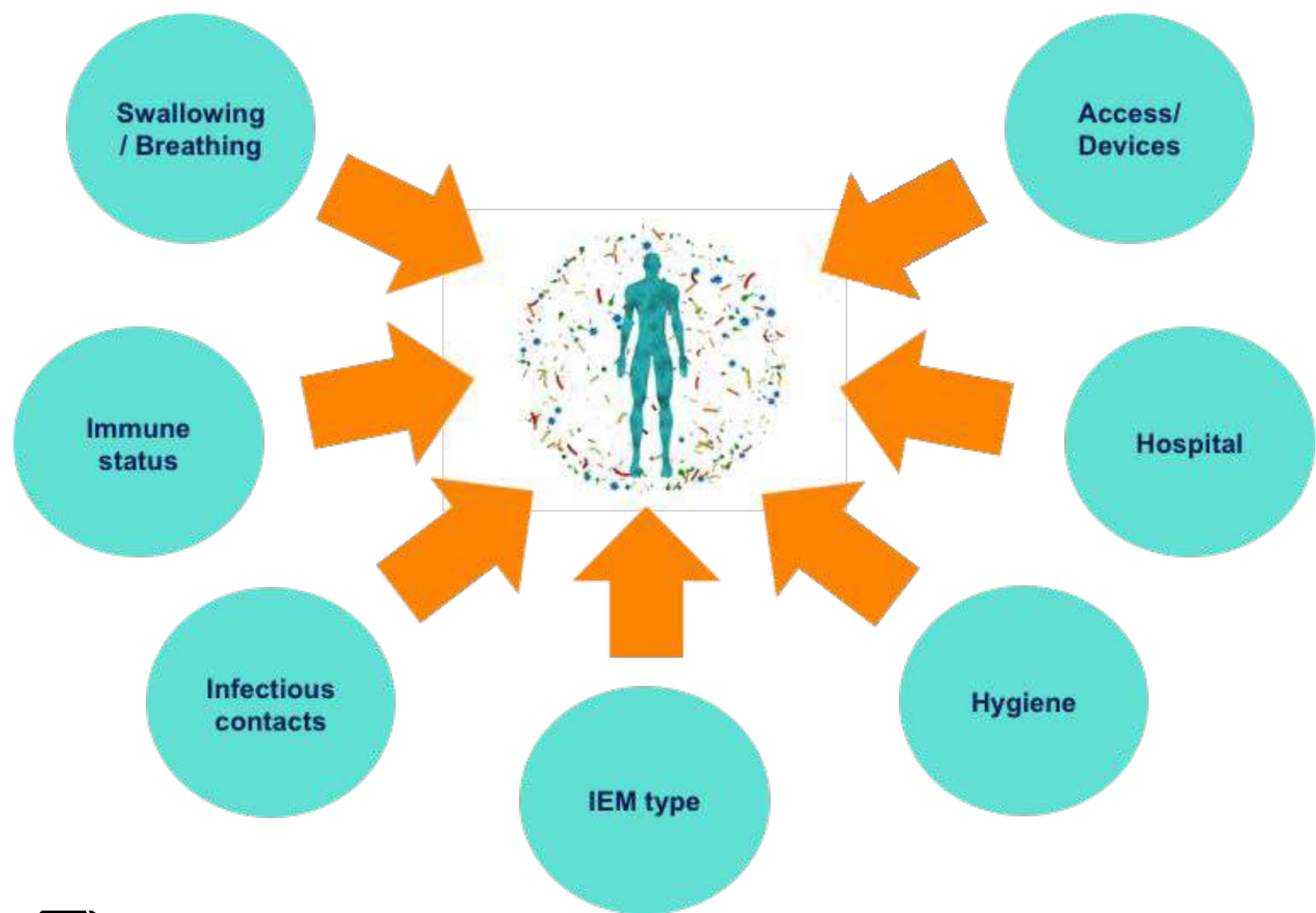
Risk Factors for Exposure to SARS-COV2

Exposure in Healthcare Settings

- **55% doctor's visit**
- 14% ER visit
- 12% hospitalized




Public Contact During COVID19

- **38% have a household member who is an essential worker**
- 12% of adult patients are essential workers



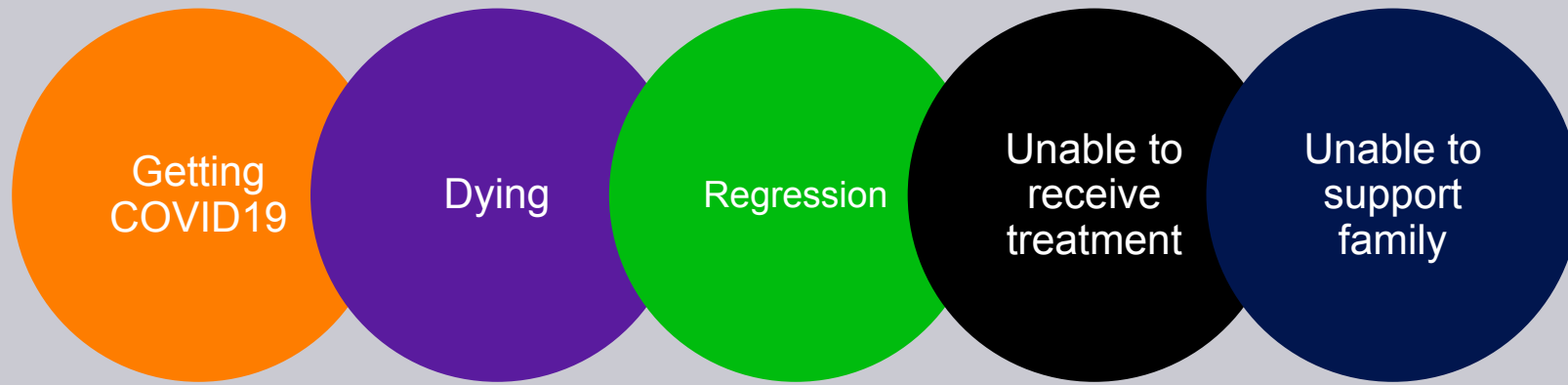
Risk Factors for Severe COVID19

73% reported at least one condition recognized by CDC as risk factors for severe COVID19, including:

- 35% Respiratory muscle weakness 
- 31% Immuno-deficiency 
- 25% Asthma 

“What is your greatest concern about COVID-19?”

“[We are terrified of] our child with Leigh Syndrome contracting COVID19, or requiring hospitalization for some other reason, experiencing a metabolic crisis, losing developmental skills and [having our lives] changed forever.”



SLOW THE SPREAD OF COVID-19

cdc.gov/coronavirus



Wear a cloth face covering
in public spaces



Stay at least 6 feet
from other people

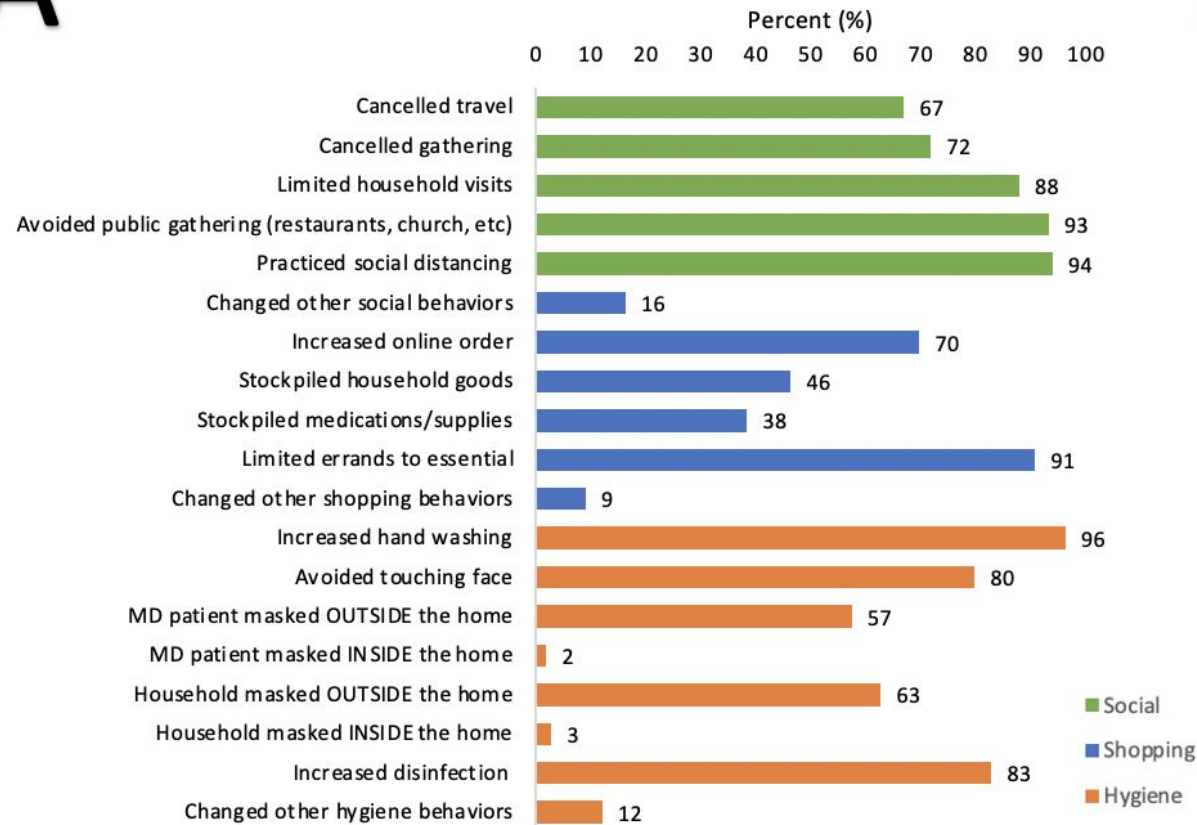


Frequently wash
your hands

Risk mitigation behaviors and mitochondrial disease

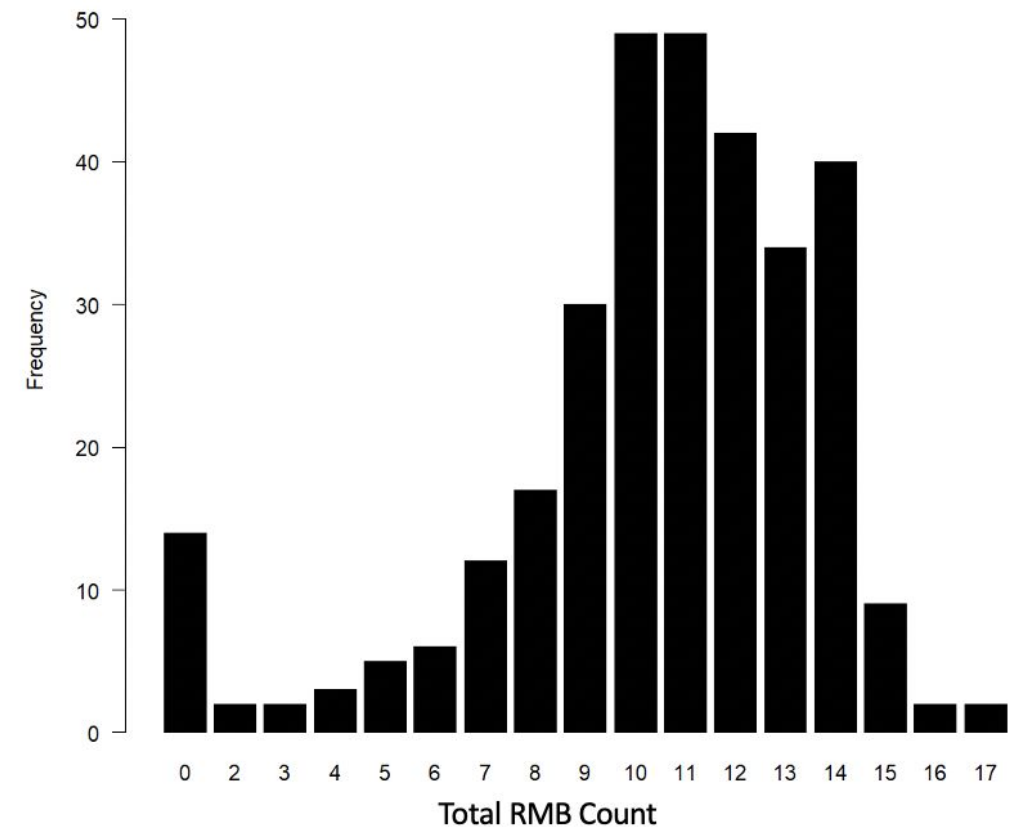
A

Proportion of Respondents Who Implemented Each RMB



B

Distribution of Total Number of RMBs Implemented By Each Respondent



Household Viral Exposure in Children with Mitochondrial Disease During COVID-19 Study

A study to learn more about the role of viral infection and biomarkers of immunity in mitochondrial disease using new technology with Neoteryx™ fingerstick at-home sampling.





SARS-CoV-2 antibody status in households with children with mitochondrial disease

Household members as risk factors for SARS-CoV-2 transmission to children with MtD

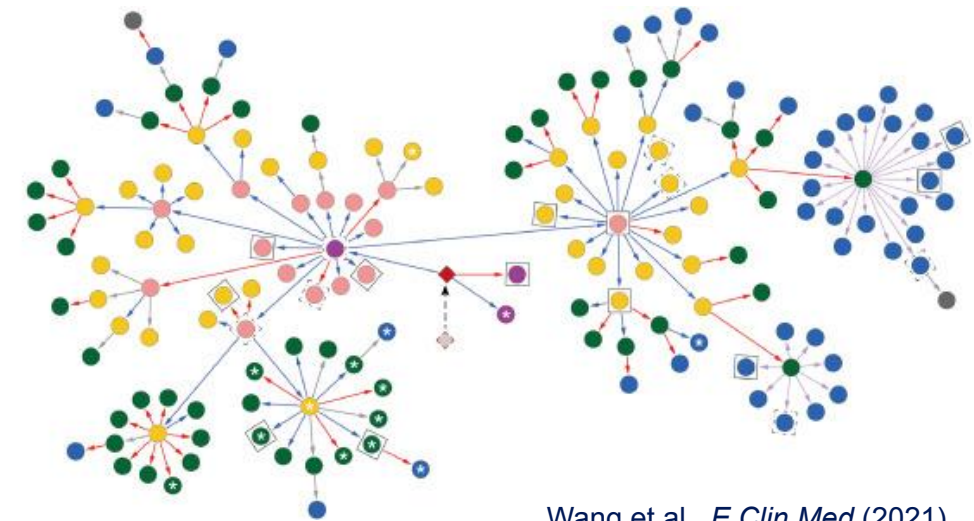
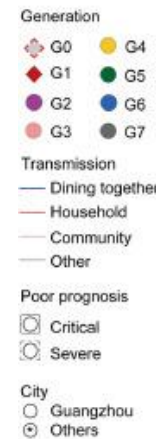


Original Investigation | Global Health

Household Transmission of SARS-CoV-2 A Systematic Review and Meta-analysis

Zachary J. Madewell, PhD; Yang Yang, PhD; Ira M. Longini Jr, PhD; M. Elizabeth Halloran, MD, DSc; Natalie E. Dean, PhD

- Meta-analysis: 54 studies, 77,258 participants
- Household secondary attack rate 16.6%



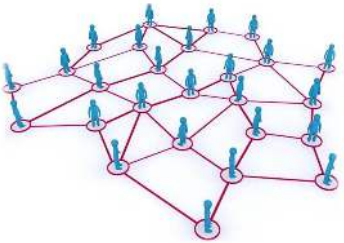
Wang et al., *E Clin Med* (2021)



Chris Marcum, PhD



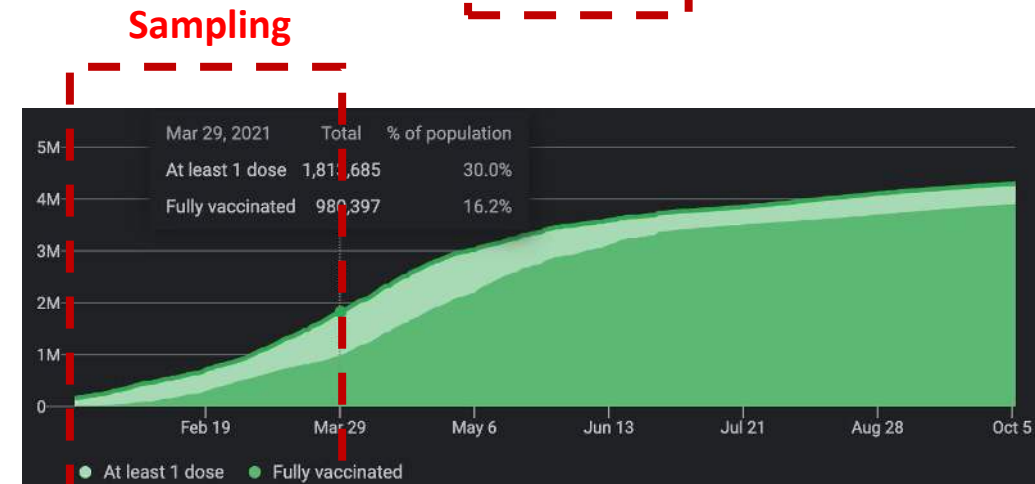
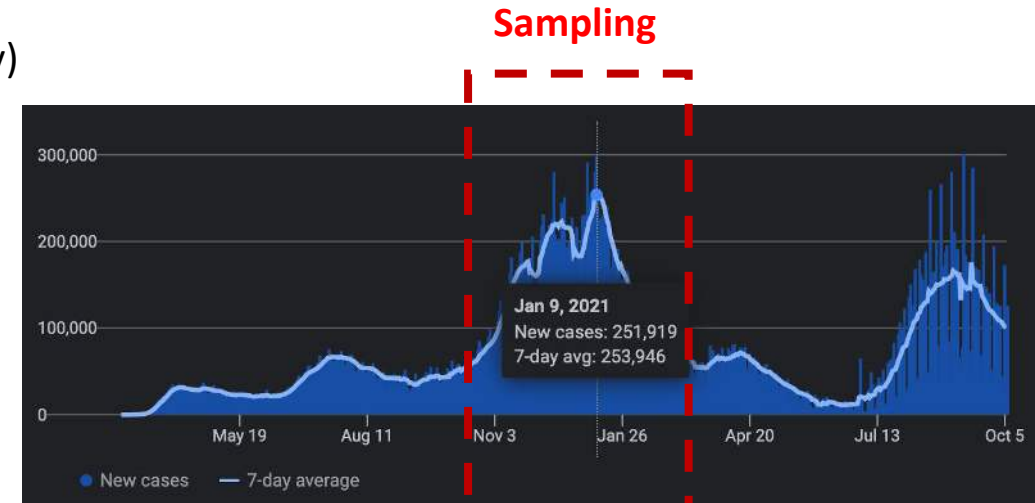
Neoteryx Mitra
collection system



Risk determination using
network scale up estimators

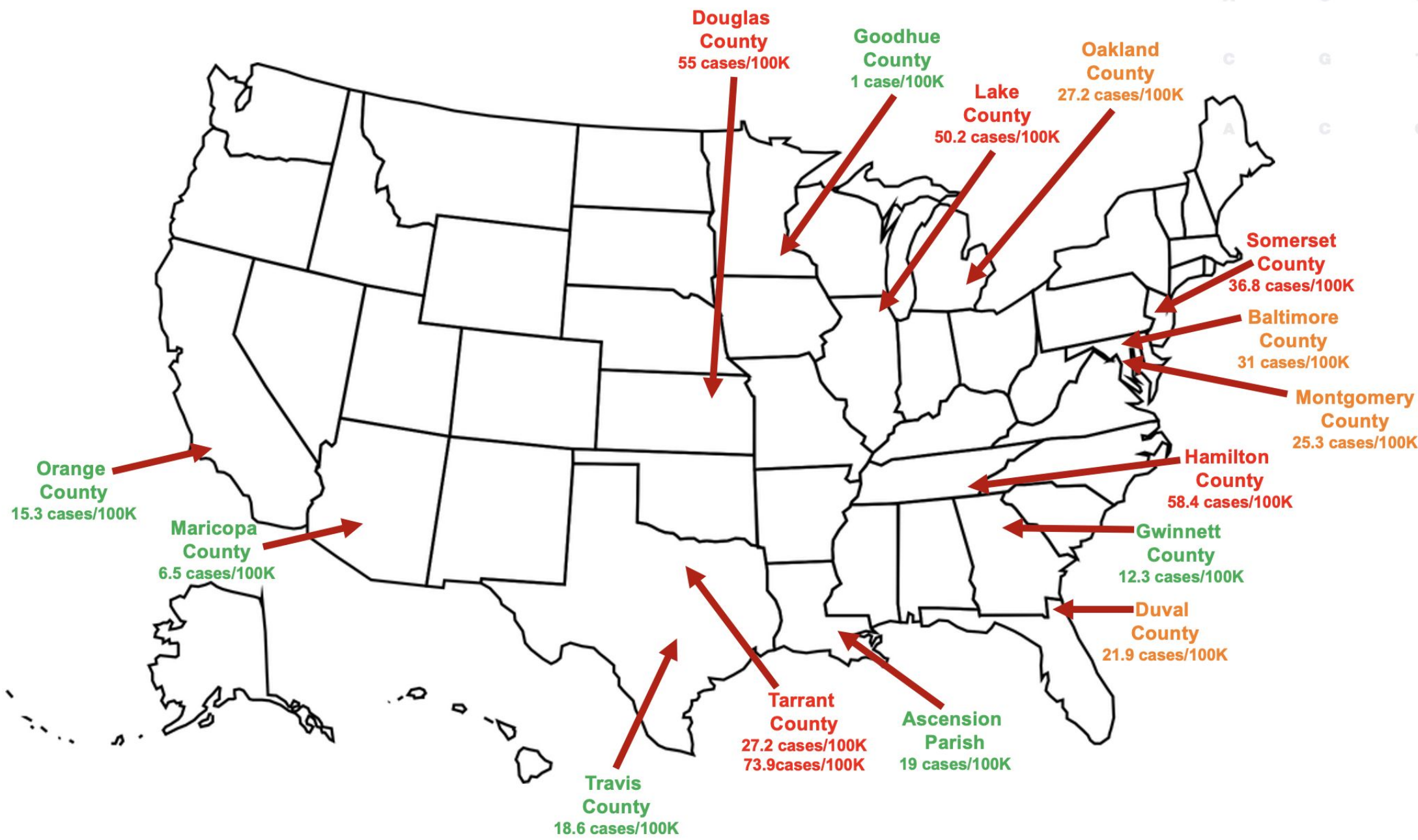
Ongoing SARS-CoV-2 family studies: interim analysis

- Families with children with MtD
 - N=20 families
 - N=83 samples (~4 members/family)
 - Confirmed dx of MtD
- Remote sampling methods
 - October 2020 – March 2021
- SARS-CoV2 antibodies
 - Infection rate
 - Vaccination rate
- Other data collected
 - Network scale up estimators
 - Symptomatology
 - Local case count
- Post-vaccination surveillance



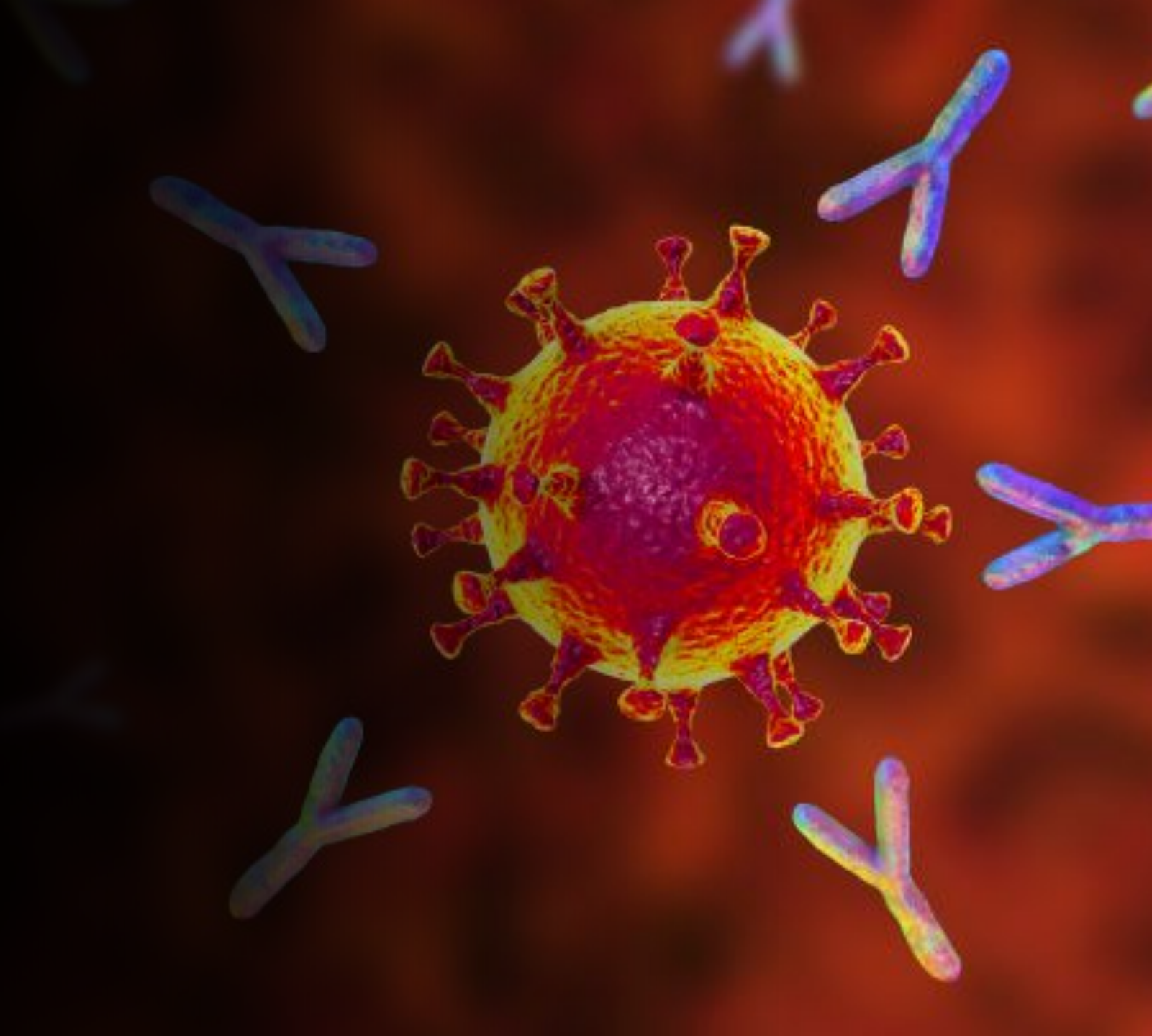
Participants		MtD (N=22)	Household (> 16 y/o, N=49)
		Number/total (%/Std dev)	
Mean age		8.8 (4.4)	38.8 (11.2)
Sex			
	<i>Males</i>	11/22 (50%)	24/49 (49%)
	<i>Females</i>	11/22 (50%)	25/49 (51%)
Job/school setting			
	<i>On site</i>	2/22 (9.1%)	8/49 (16.3%)
	<i>Hybrid</i>	2/22 (9.1%)	9/49 (18.4%)
	<i>Remote</i>	18/22 (81.8%)	23/49 (46.9%)
	<i>Not applicable</i>	0/22 (0.0%)	9/49 (18.4%)
COVID19 exposure/testing/diagnosis			
	<i>Exposed</i>	7/22 (31.8%)	15/49 (30.6%)
	<i>Tested</i>	14/22 (63.6%)	23/49 (46.9%)
	<i>Diagnosed</i>	1/22 (4.2%)	1/49 (2.0%)
COVID19 symptoms			
	<i>Fever or chills</i>	6/22 (27.2%)	6/49 (12.2%)
	<i>New or worsening cough</i>	2/22 (8.3%)	10/49 (20.4%)
	<i>New or worsening shortness of breath</i>	1/22 (4.2%)	5/49 (10.2%)
	<i>Pneumonia</i>	2/22 (8.3%)	1/49 (2.0%)
	<i>Muscle or body aches</i>	0/22 (0.0%)	10/49 (20.4%)
	<i>Vomiting or diarrhea</i>	0/22 (0.0%)	5/49 (10.2%)
	<i>Loss of taste or smell</i>	0/22 (0.0%)	1/49 (2.0%)

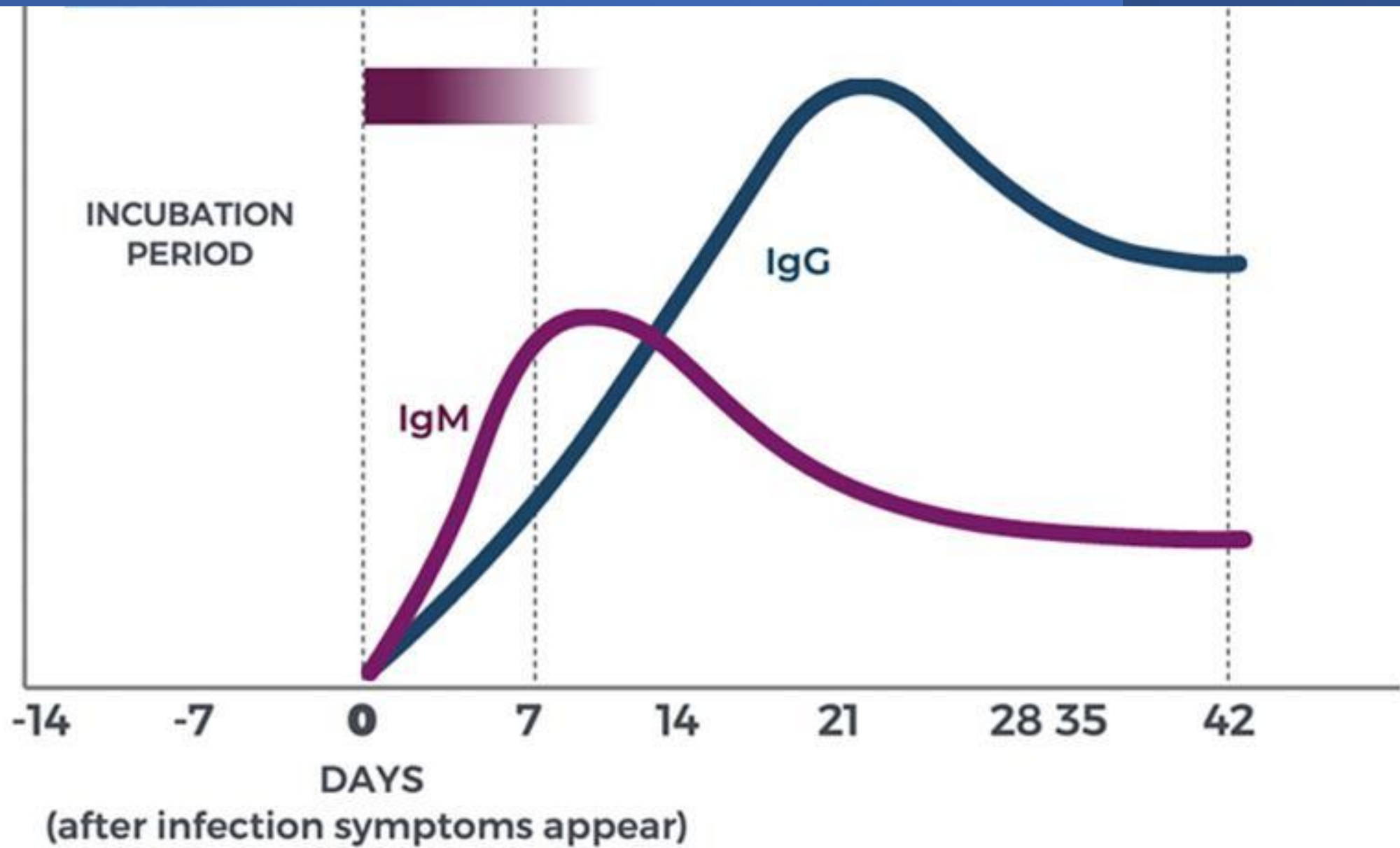
A C G
C G T
C G

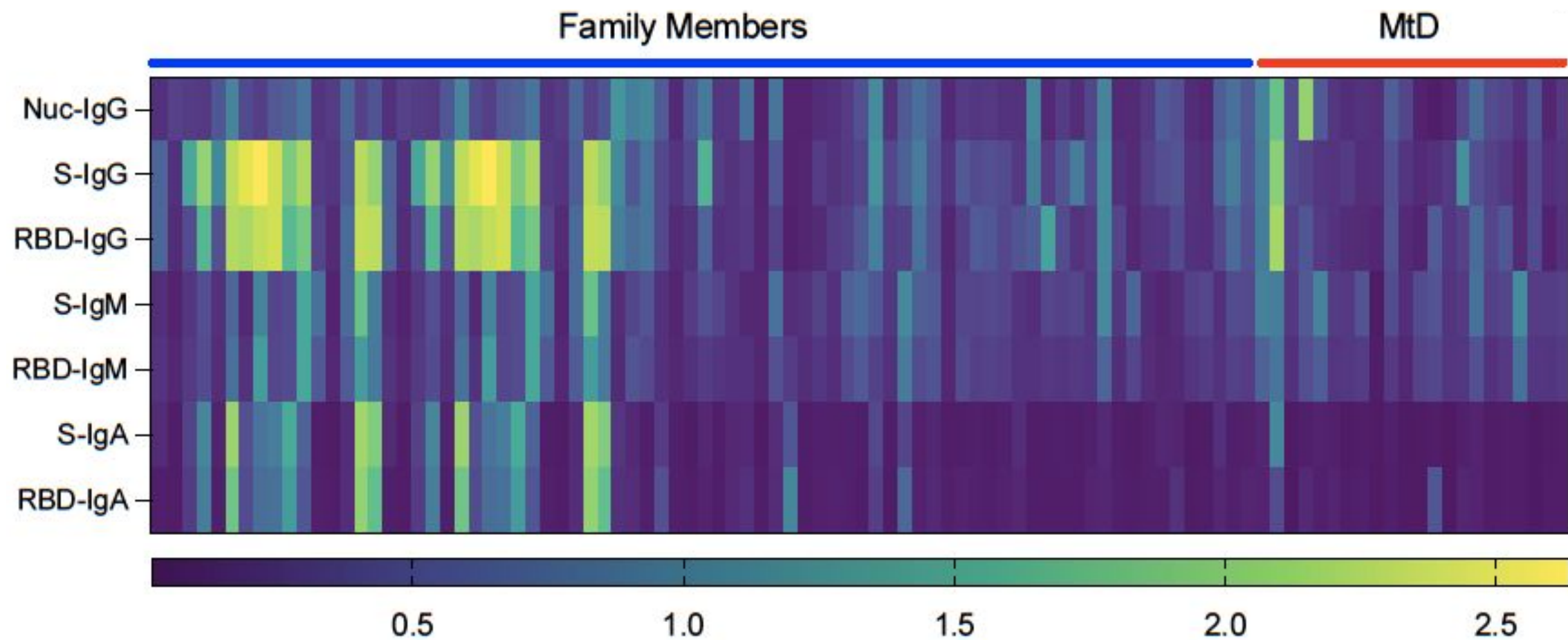


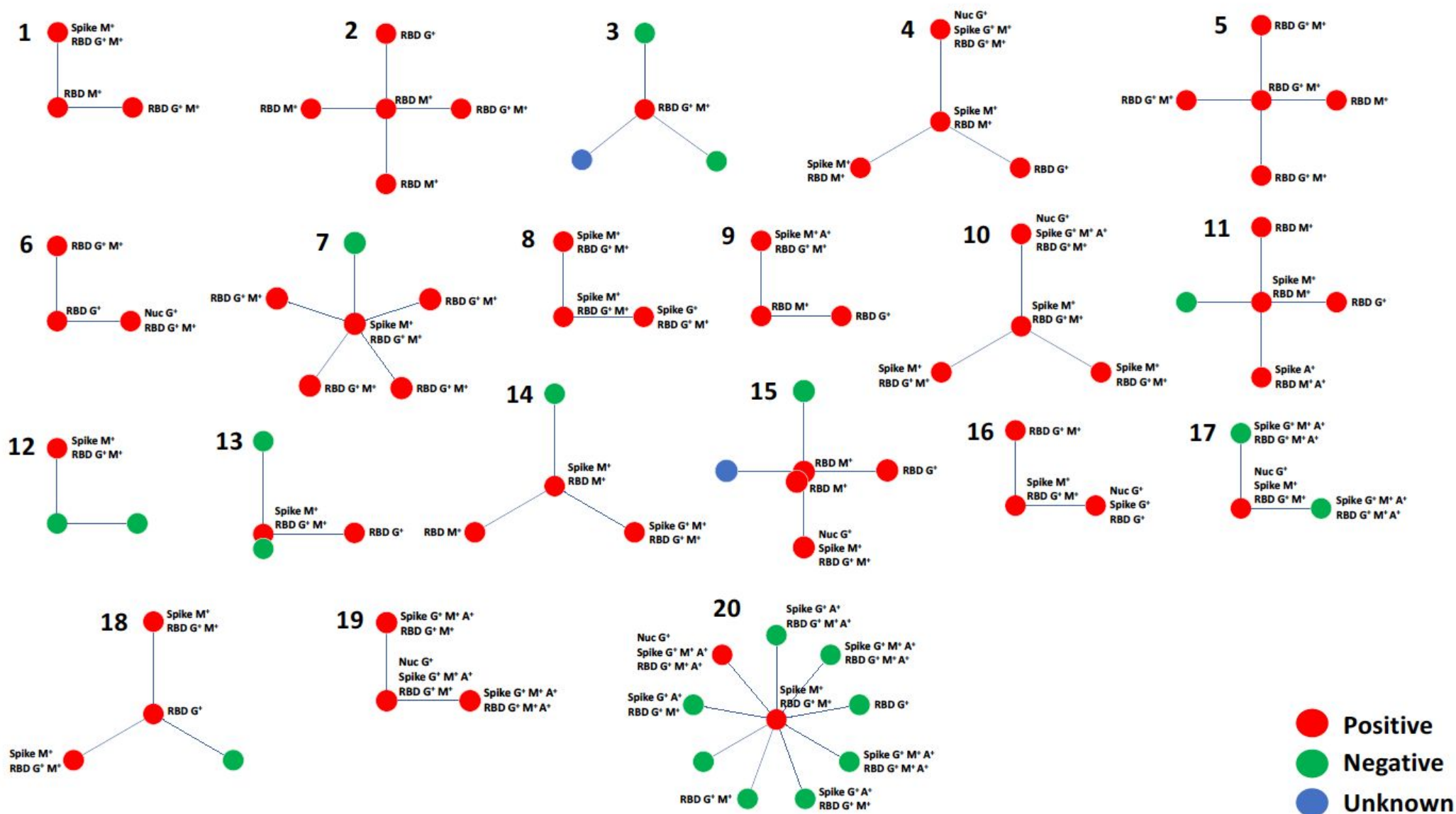
Antibodies against SARS-CoV-2

- Nucleocapsid
- Spike protein
- Receptor binding domain

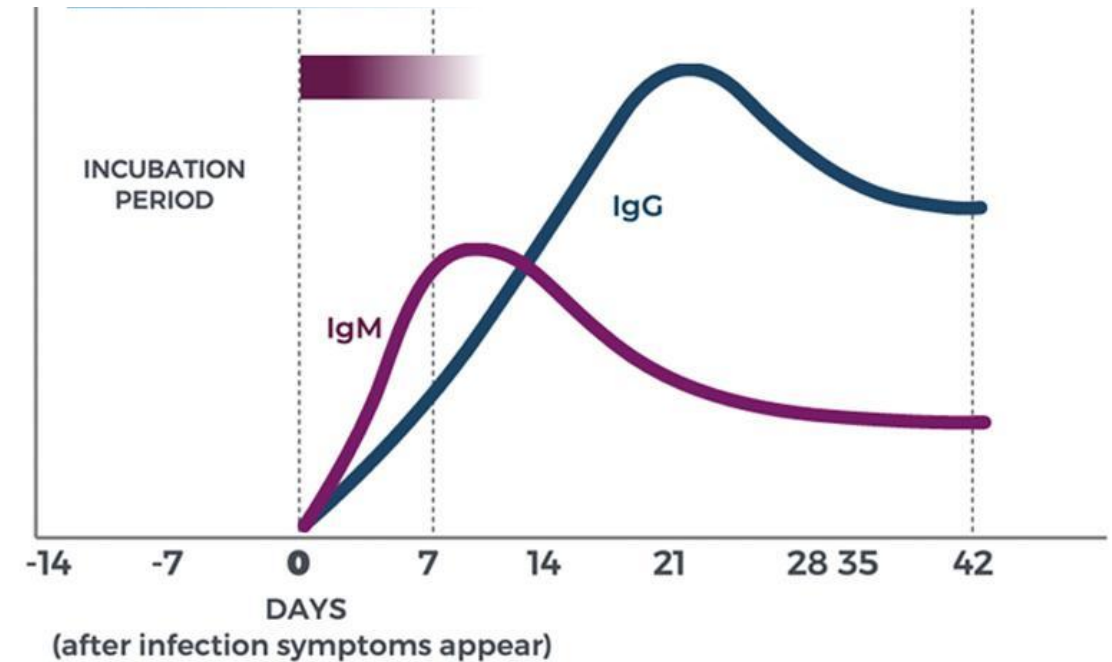
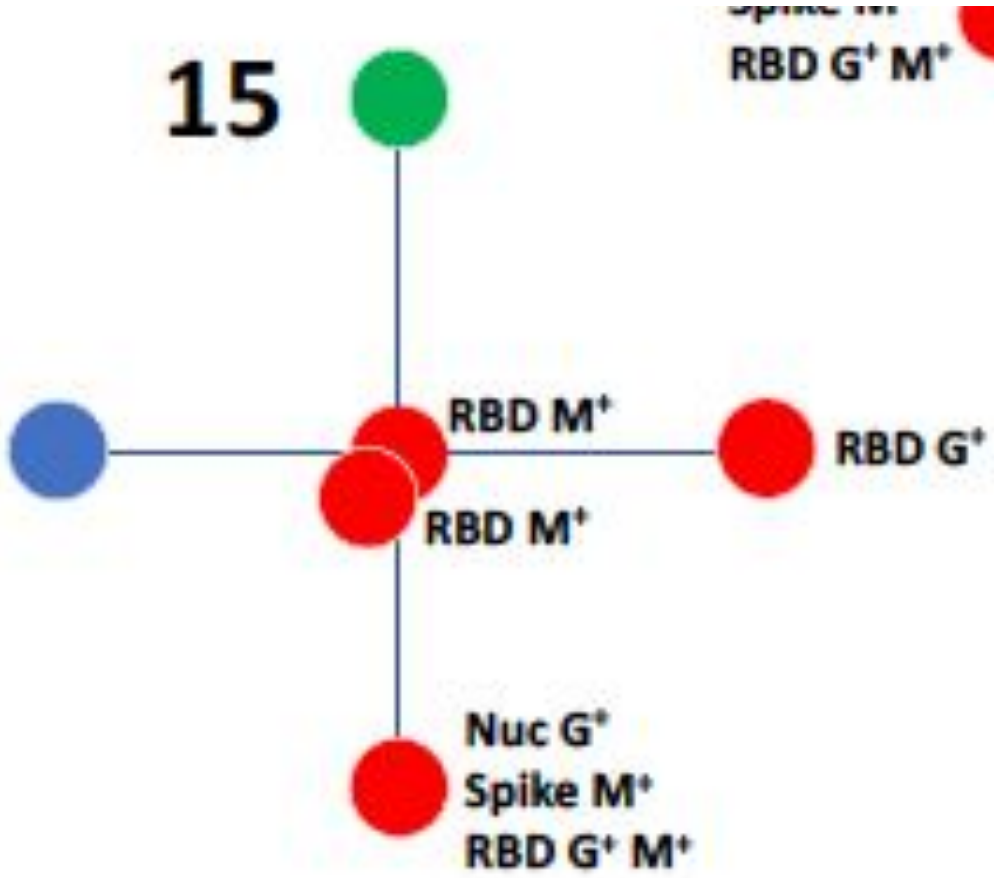








Transmission from other household members?

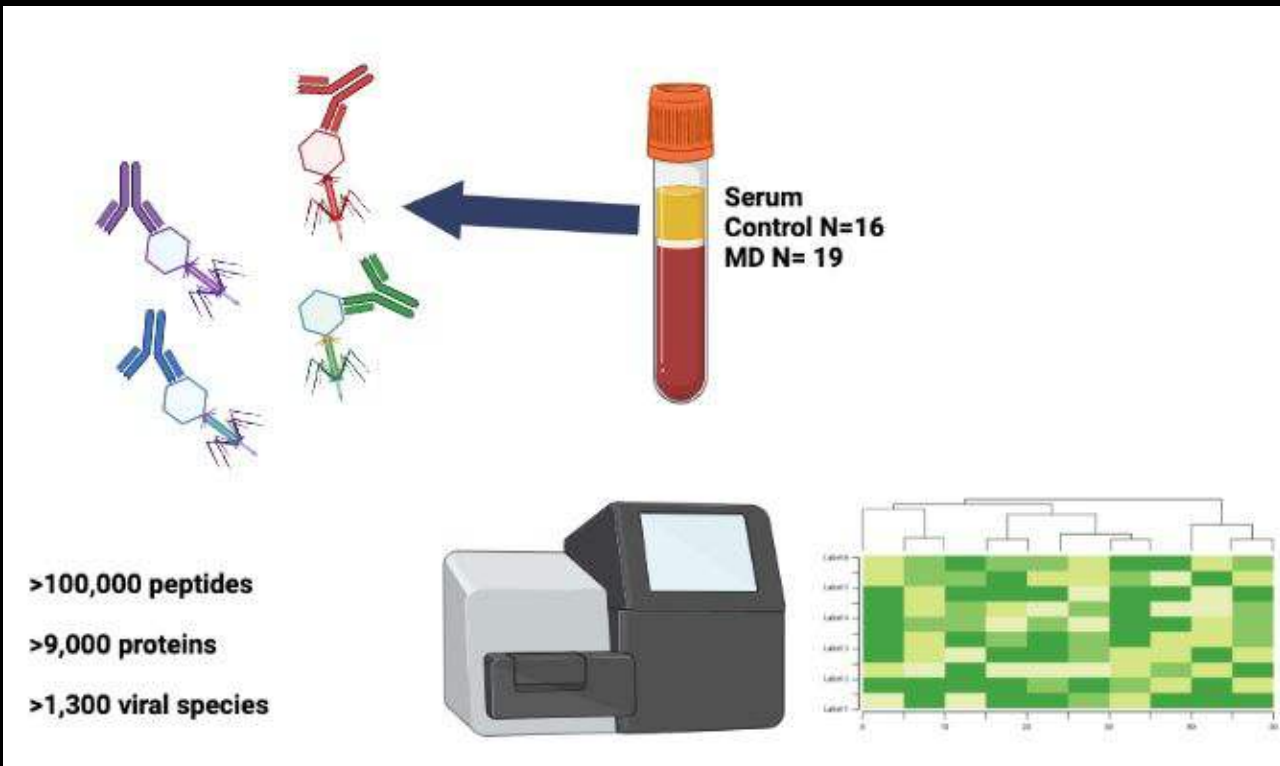


Household Viral Exposure in Children with Mitochondrial Disease During COVID-19 Study

A study to learn more about the role of viral infection and biomarkers of immunity in mitochondrial disease using new technology with Neoteryx™ fingerstick at-home sampling.



Can we assess the entire antiviral antibody repertoire?



VirScan articles

RESEARCH

RESEARCH ARTICLE

VIRAL IMMUNOLOGY

Comprehensive serological profiling of human populations using a synthetic human virome

George J. Xu,^{1,2,3,4,*} Tomasz Kula,^{3,4,5,*} Qikai Xu,^{3,4} Mamie Z. Li,^{3,4} Suzanne D. Vernon,⁶ Thumbi Ndung'u,^{7,8,9,10} Kiat Ruxrungtham,¹¹ Jorge Sanchez,¹² Christian Brander,¹³ Raymond T. Chung,¹⁴ Kevin C. O'Connor,¹⁵ Bruce Walker,^{8,9} H. Benjamin Larman,¹⁶ Stephen J. Elledge^{3,4,6,†}

VIRAL IMMUNOLOGY

Measles virus infection diminishes preexisting antibodies that offer protection from other pathogens

Michael J. Mina^{1,2,3,*†}, Tomasz Kula^{1,2}, Yumei Leng¹, Mamie Li², Rory D. de Vries⁴, Mikael Knip^{5,6}, Heli Siljander^{5,6}, Marian Rewers⁷, David F. Choy⁸, Mark S. Wilson⁸, H. Benjamin Larman⁹, Ashley N. Nelson^{10,†}, Diane E. Griffin¹⁰, Rik L. de Swart⁴, Stephen J. Elledge^{1,2,11,†}



Temporal virus serological profiling of kidney graft recipients using VirScan

Pierre Isnard^{a,b,1}, Tomasz Kula^{c,d,1}, Véronique Avettand Fenoel^{e,f}, Dany Anglicheau^{a,b,f}, Fabiola Terzi^a, Christophe Legendre^{a,b,f}, Stephen J. Elledge^{c,d}, and Guillaume Canaud^{a,b,f,2}

^aINSERM U1151, Institut Necker Enfants Malades, Hôpital Necker-Enfants Malades, 75015 Paris, France; ^bService de Néphrologie Transplantation Adultes, Hôpital Necker-Enfants Malades, 75015 Paris, France; ^cDivision of Genetics, Department of Medicine, Howard Hughes Medical Institute, Brigham and Women's Hospital, Boston, MA 02115; ^dDepartment of Genetics, Harvard University Medical School, Boston, MA 02115; ^eLaboratoire de Virologie, Hôpital Necker-Enfants Malades, 75015 Paris, France; and ^fUniversité Paris Descartes, Sorbonne Paris Cité, Hôpital Necker-Enfants Malades, 75006 Paris, France

nature
medicine

LETTERS

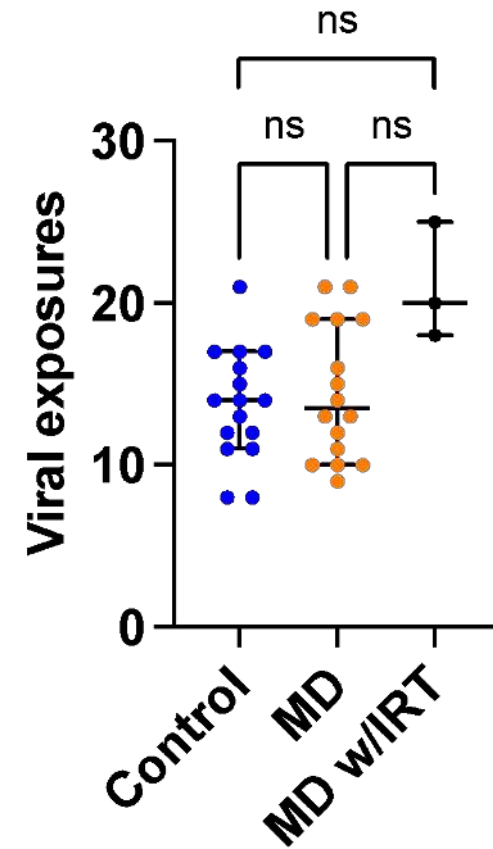
<https://doi.org/10.1038/s41591-019-0392-8>

The repertoire of maternal anti-viral antibodies in human newborns

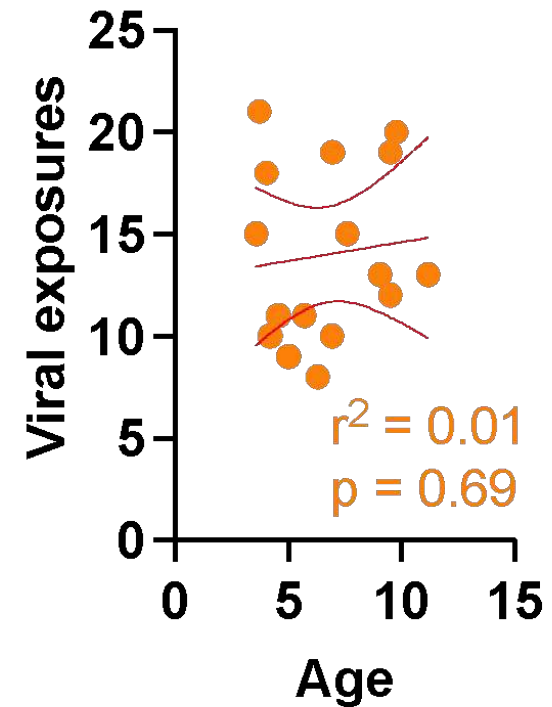
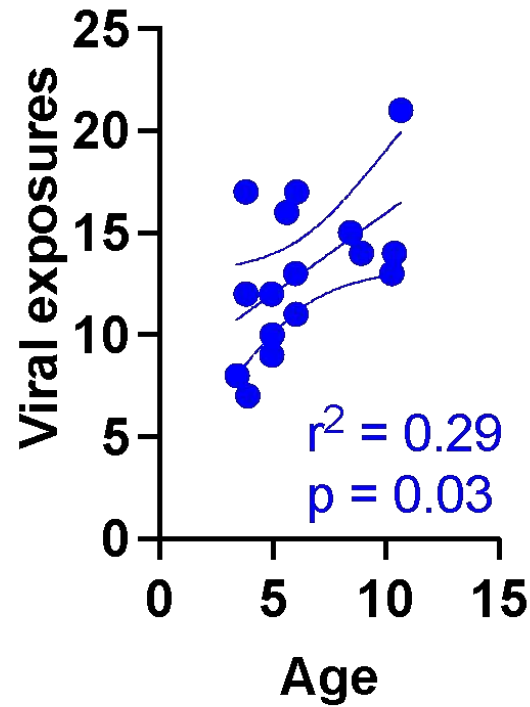
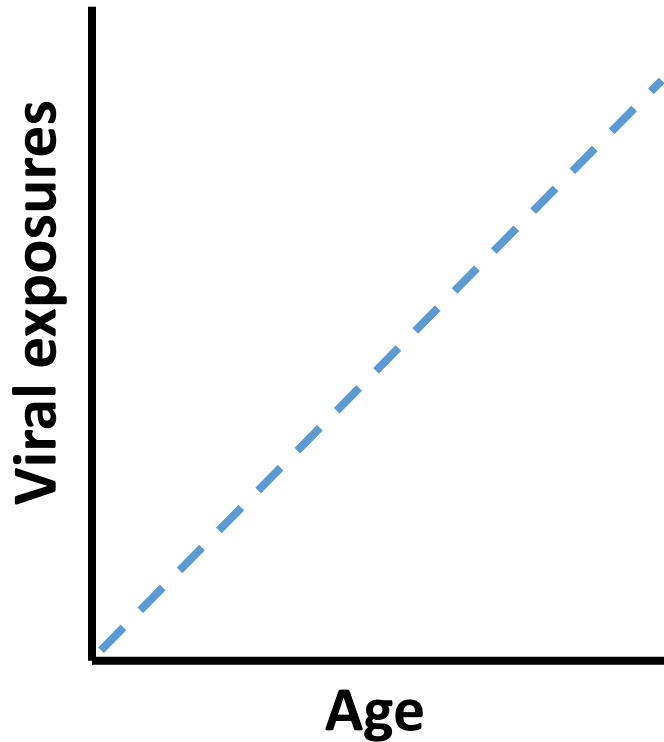
Christian Pou^{1,4}, Dieudonné Nkulikiyimfura^{1,4}, Ewa Henckel^{2,3}, Axel Olin¹, Tadeally Lakshmikanth¹, Jaromir Mikes¹, Jun Wang¹, Yang Chen¹, Anna Karin Bernhardsson^{1,3}, Anna Gustafsson^{2,3}, Kajsa Bohlin^{2,3} and Petter Brodin^{1,3*}

Viral exposures via the AntiViral Antibody Response Deconvolution Algorithm (Monaco et al., bioRxiv, 2018)

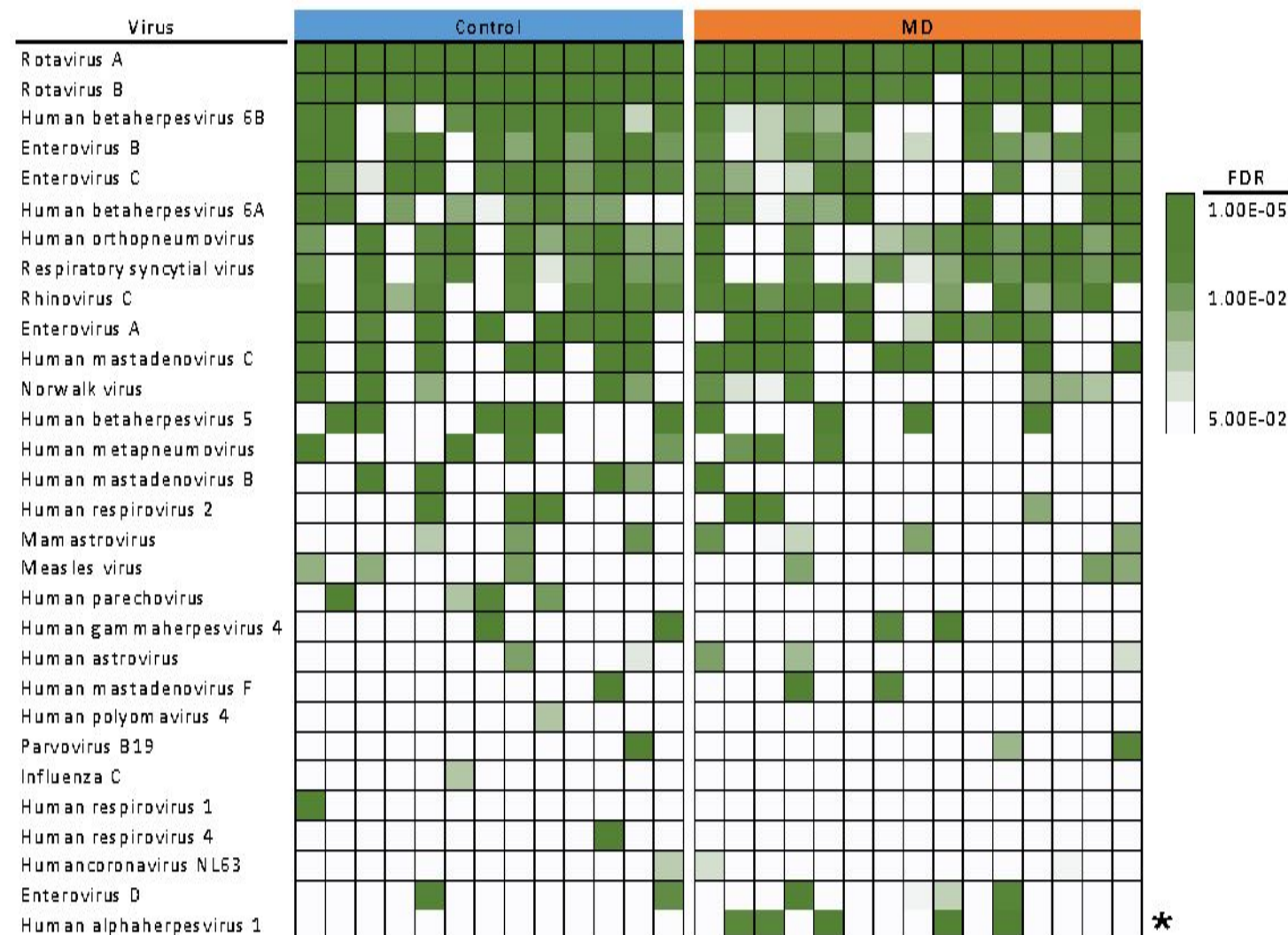
- Disproportionate representation of viruses
- Antibody cross-reactivity for related viruses



Viral exposures via the AntiViral Antibody Response Deconvolution Algorithm (Monaco et al., bioRxiv, 2018)



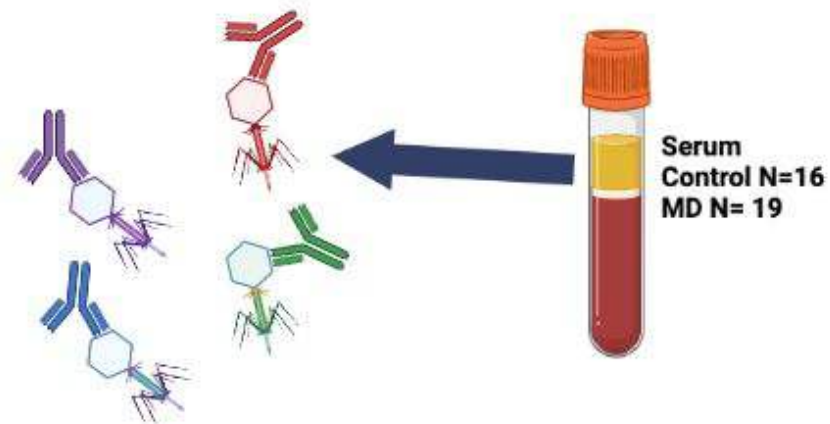
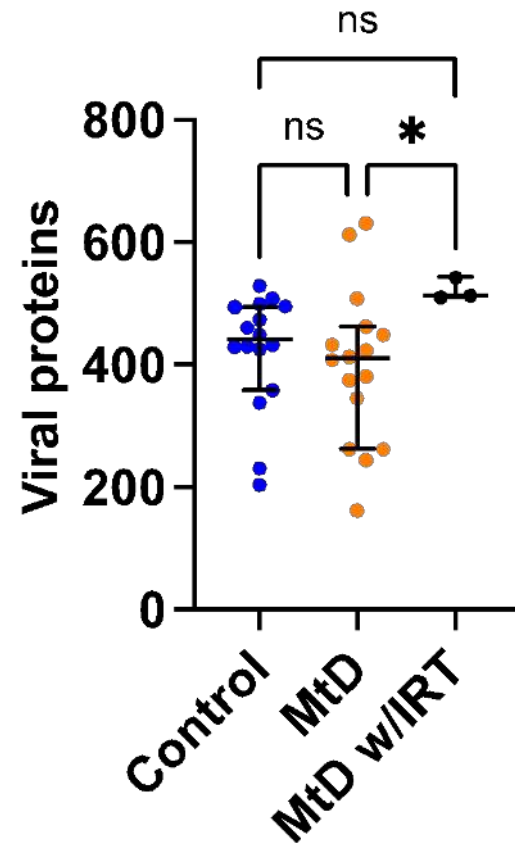
Viral exposures via the AntiViral Antibody Response Deconvolution Algorithm (Monaco et al., bioRxiv, 2018)



Viral exposures via the AntiViral Antibody Response Deconvolution Algorithm (Monaco et al., bioRxiv, 2018)

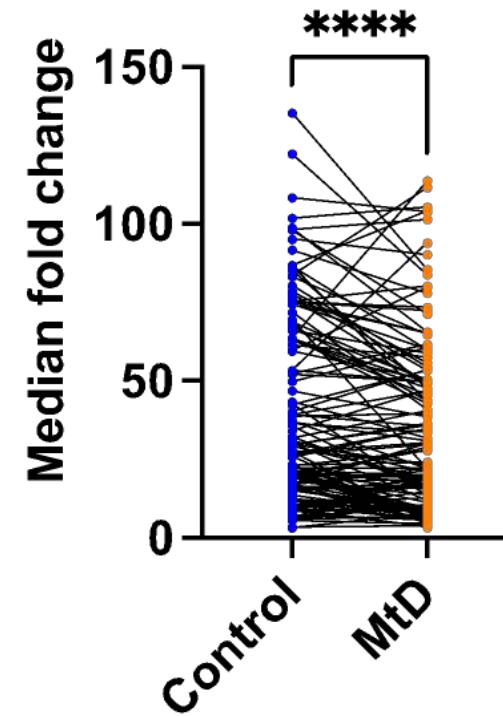
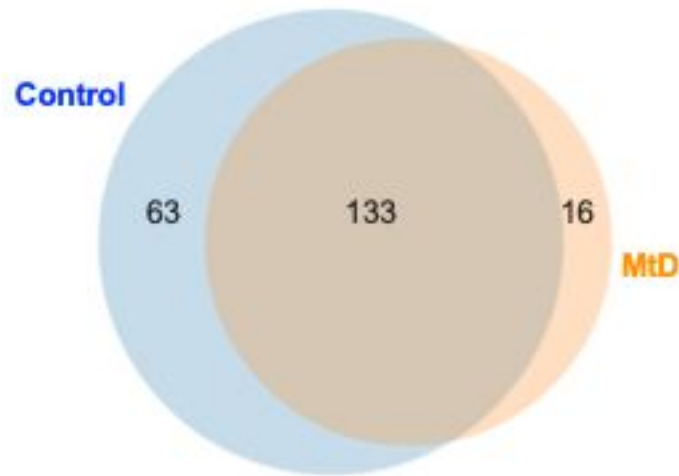
Control		MD	
Virus	% positive	Virus	% positive
Rotavirus A	100	Rotavirus A	100
Rotavirus B	100	Rotavirus B	93
Enterovirus C	92	Human beta herpesvirus 6B	80
Human beta herpesvirus 6B	85	Enterovirus B	80
Enterovirus B	85	Respiratory syncytial virus	80
Respiratory syncytial virus	77	Human orthopneumovirus	73
Human orthopneumovirus	77	Rhinovirus C	73
Rhinovirus C	69	Enterovirus C	67
Human beta herpesvirus 6A	69	Human beta herpesvirus 6A	60
Enterovirus A	62	Enterovirus A	60

Number of viral proteins per individual

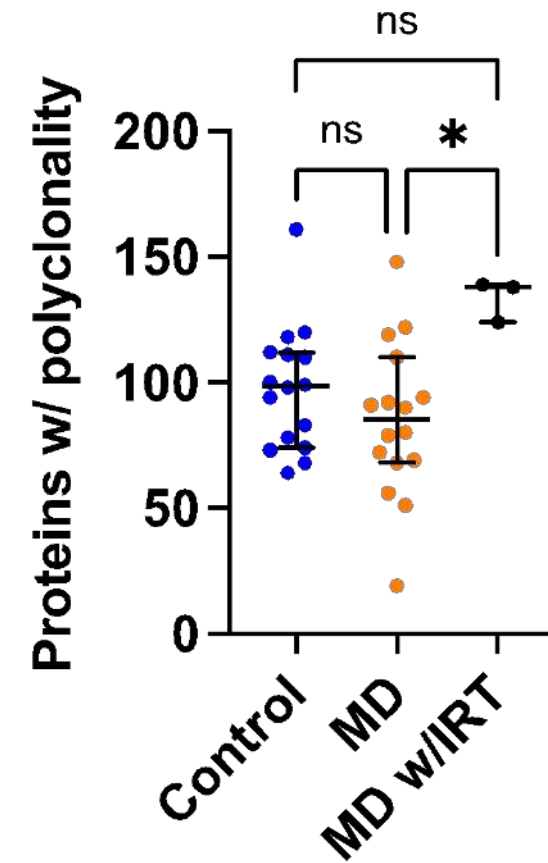
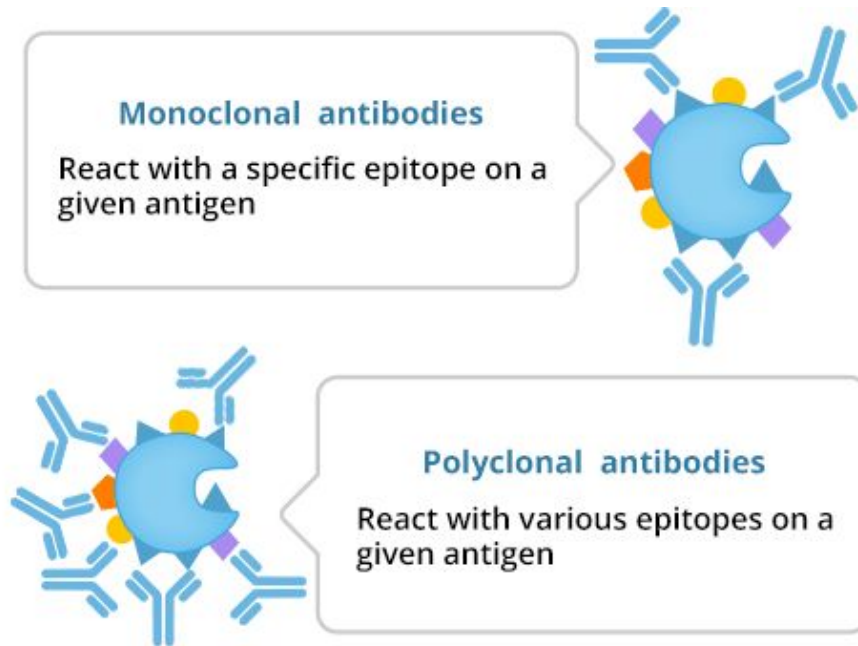


Controls mean age = 6.4 years old
MtD mean age = 6.6 years old

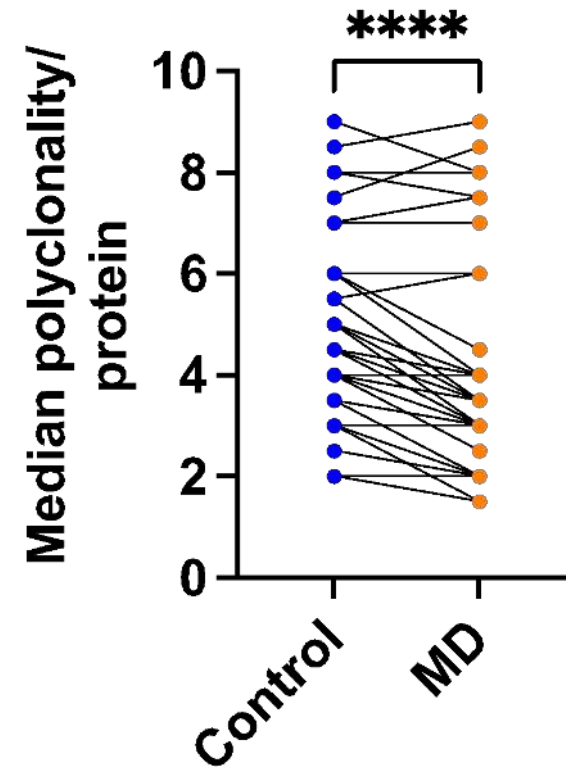
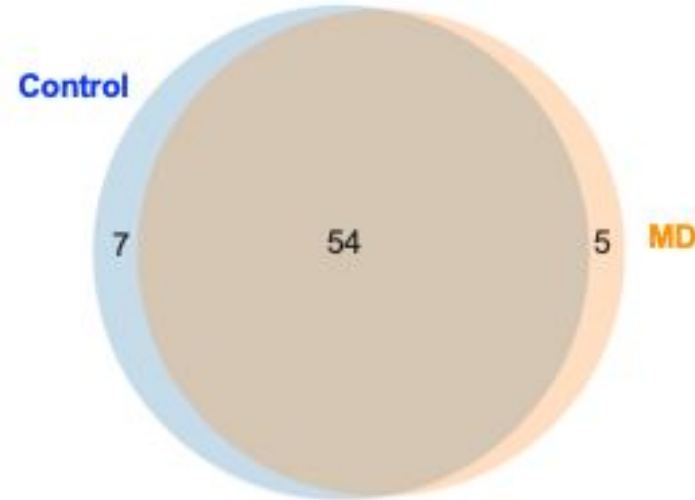
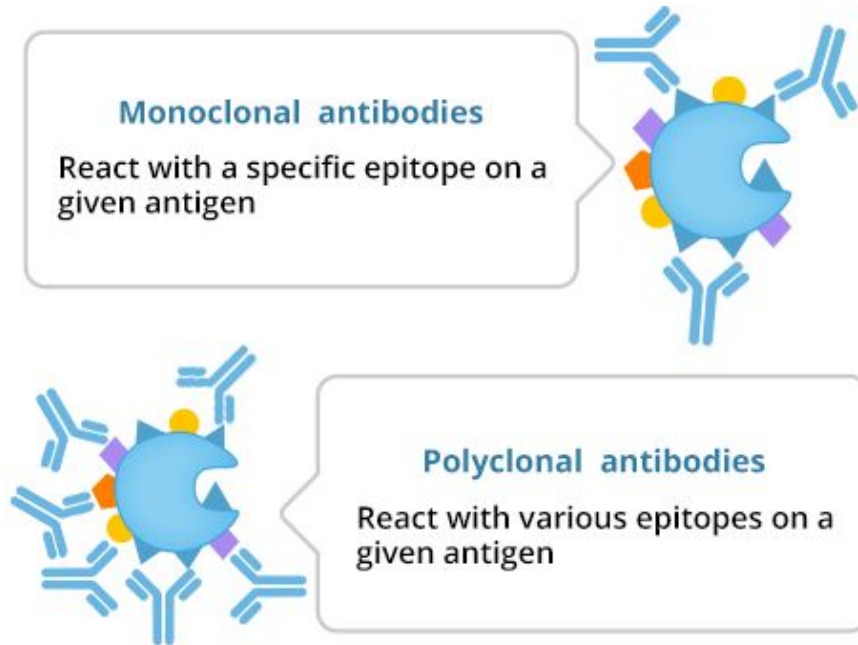
Shared/unique proteins



Polyclonality for individuals



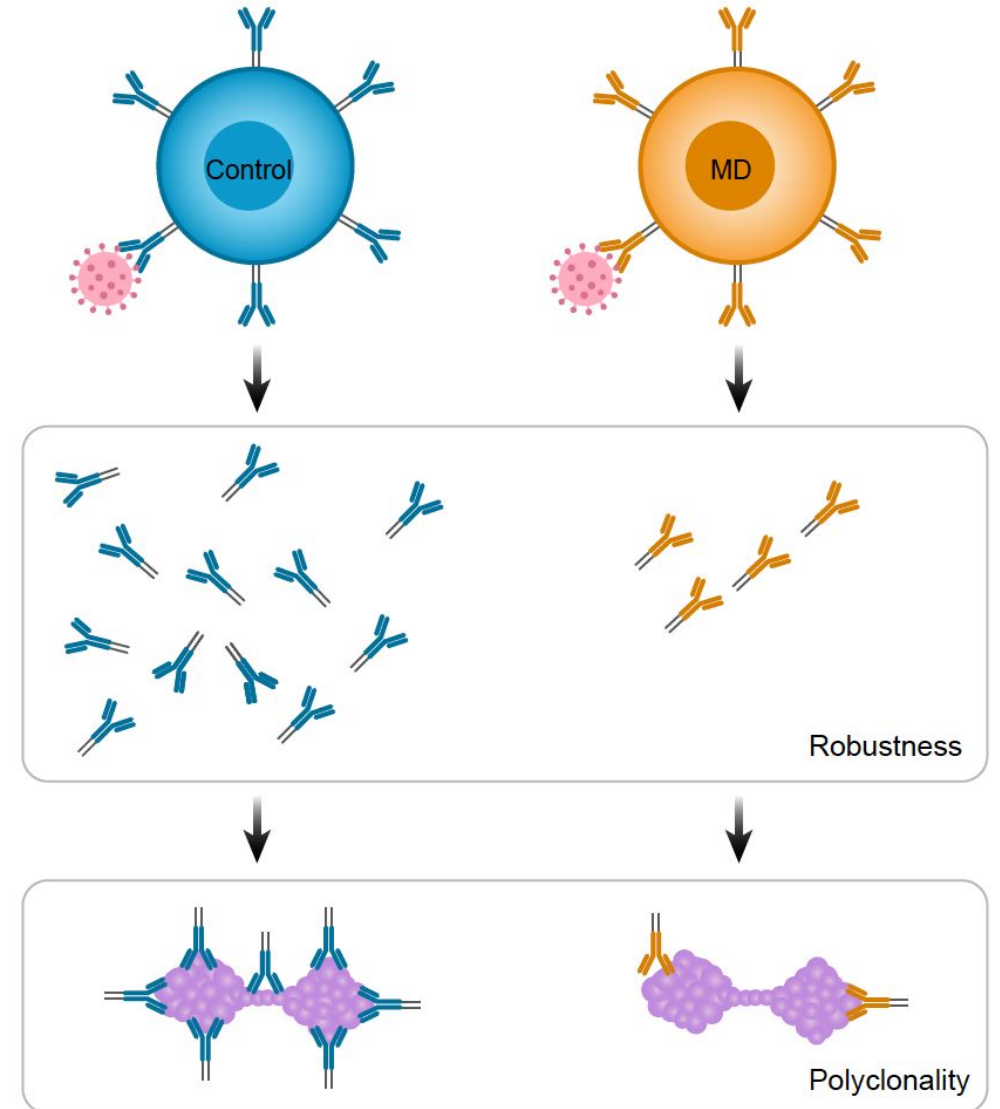
Polyclonality for shared/unique proteins



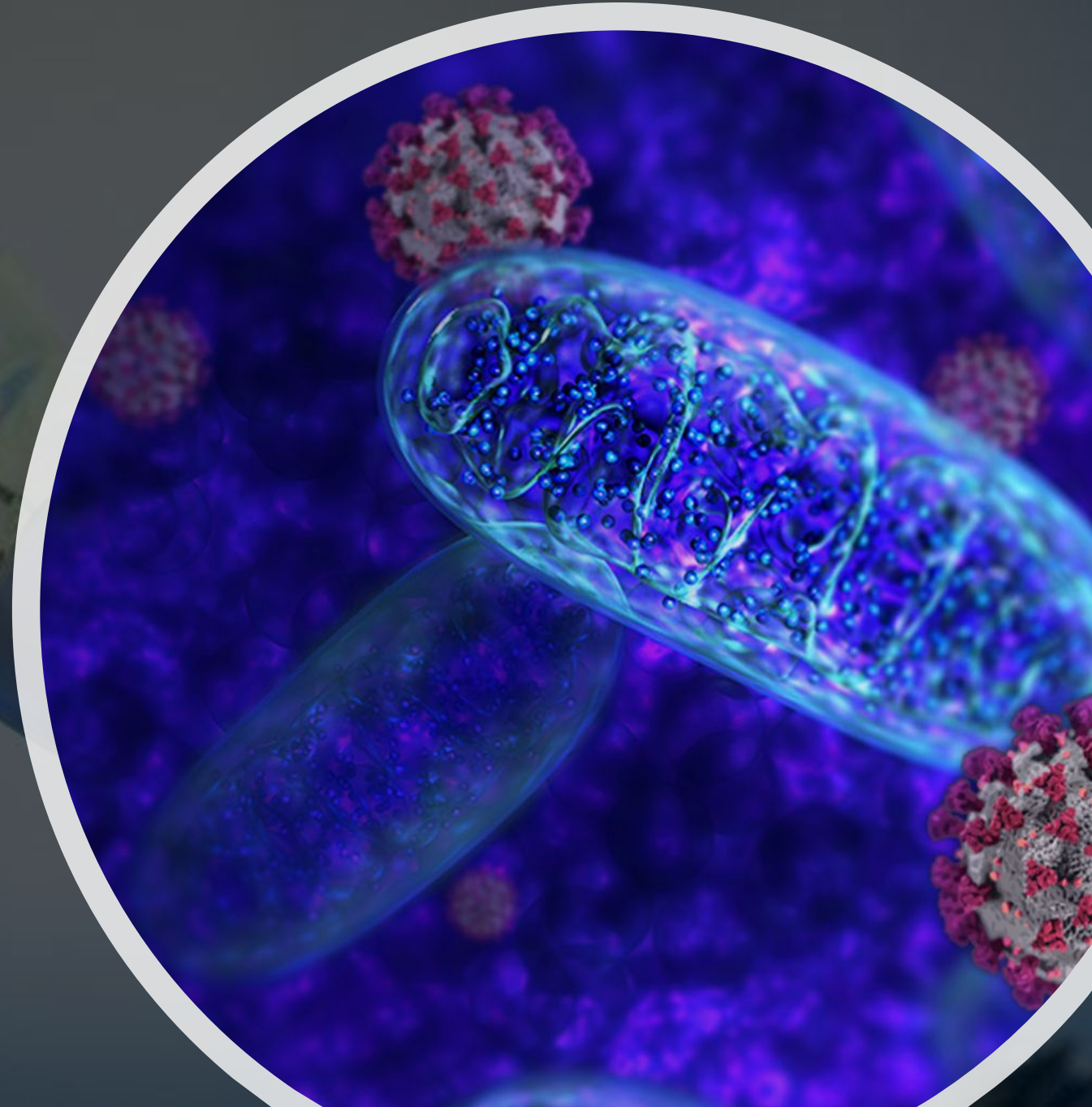
Summary

Children with MtD:

- Infection is inevitable
 - Viral exposure signatures are similar to healthy children
- Limitations in the antibody repertoire
- Asynchronous relationship with viral exposure
 - ↑↑ exposure at earlier age
- HSV-1 exposures earlier
 - ?natural host
 - Chronic neurodegeneration
- Necessary to start building a compendium of viral exposures in MtD

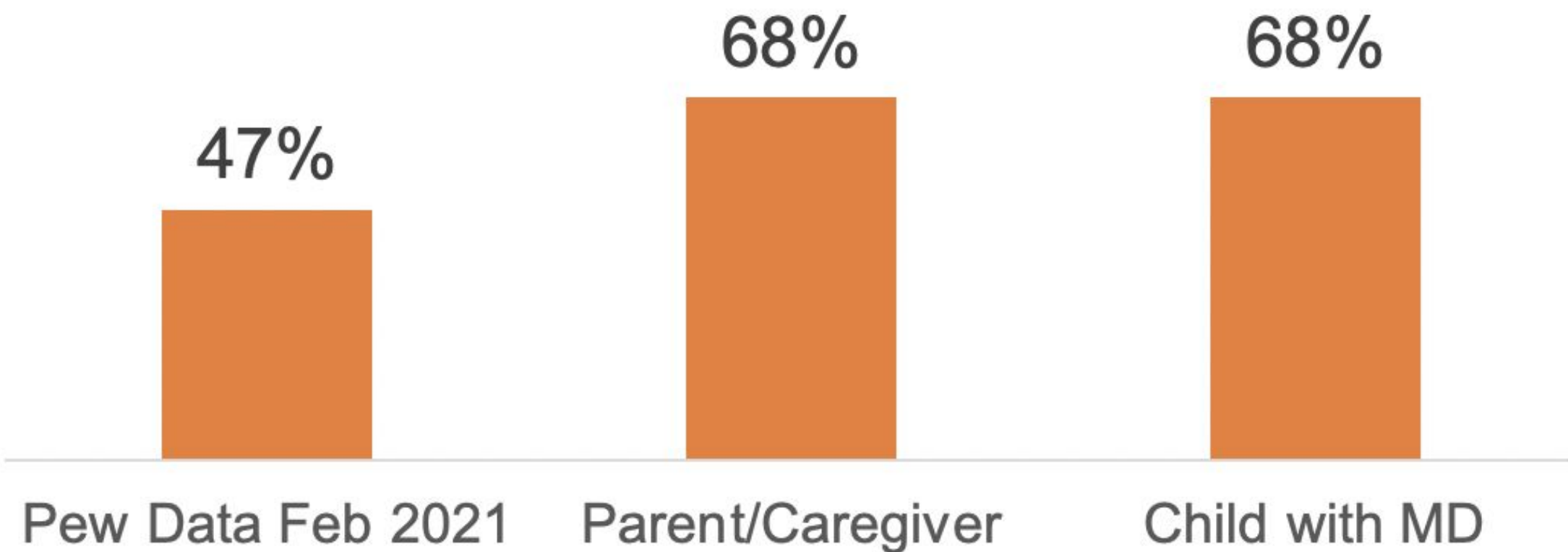


Vaccine hesitancy and the COVID-19 vaccine



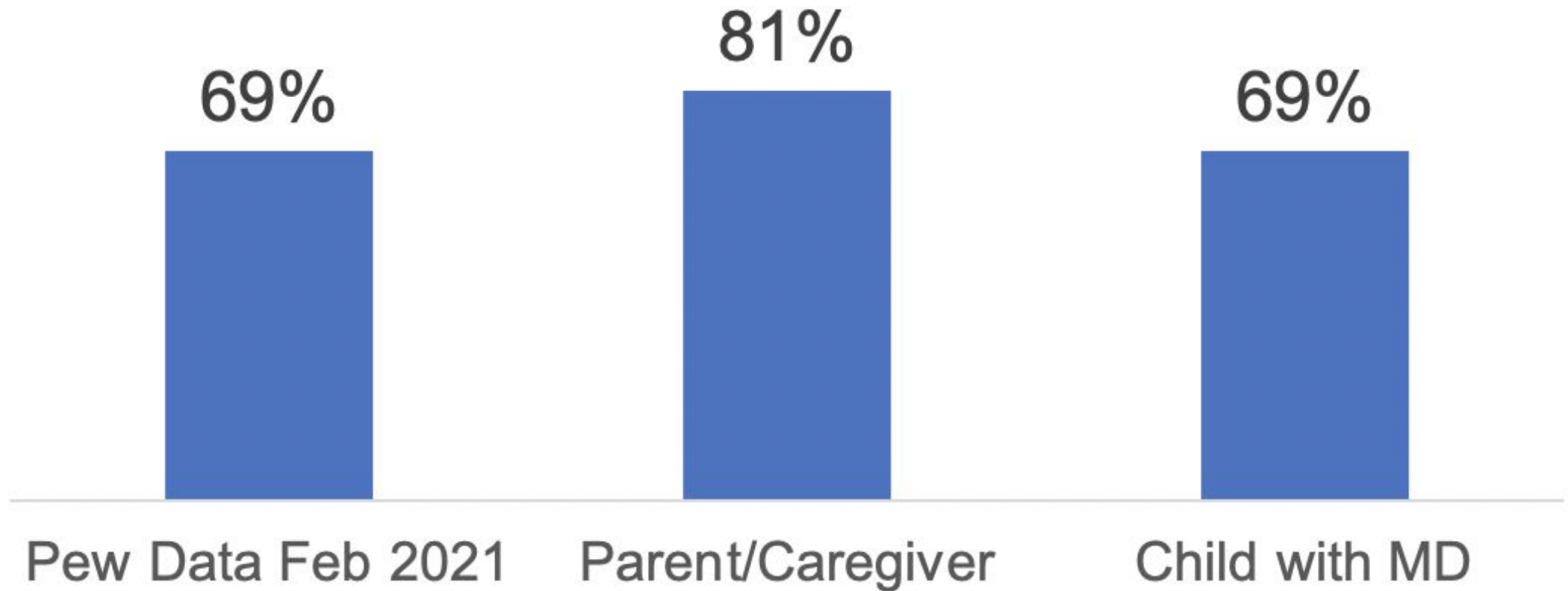
Receives the annual flu shot

■ Yes



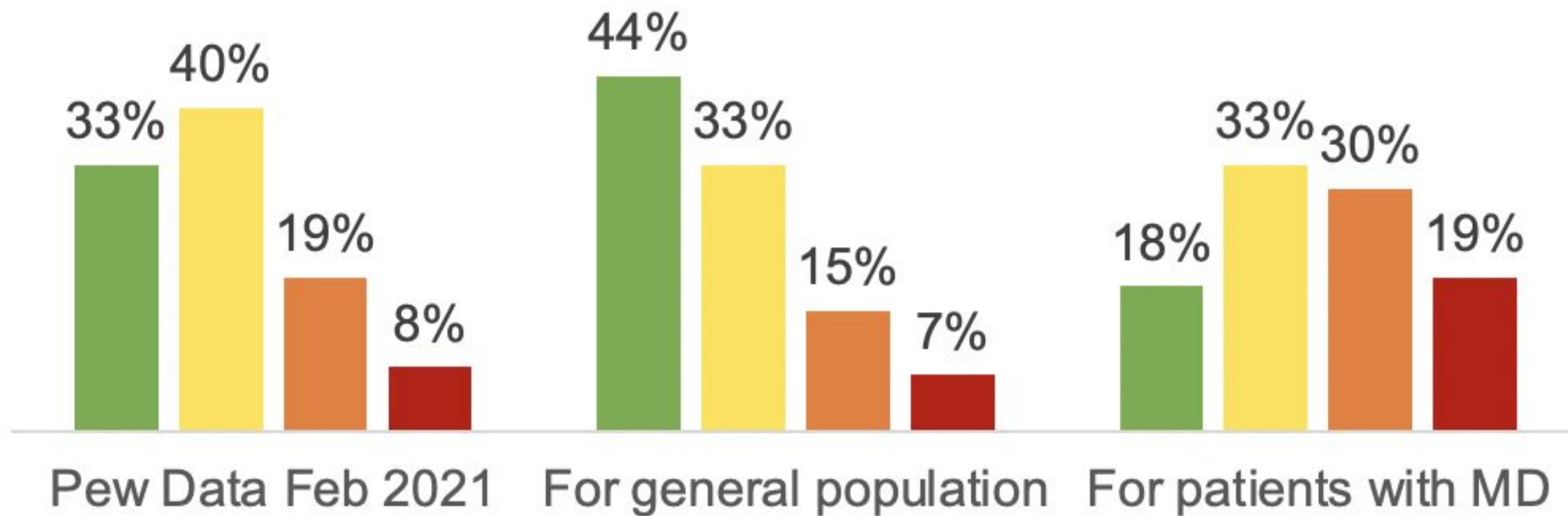
Intends to get COVID19 Vaccine

■ Probably/Definitely Yes

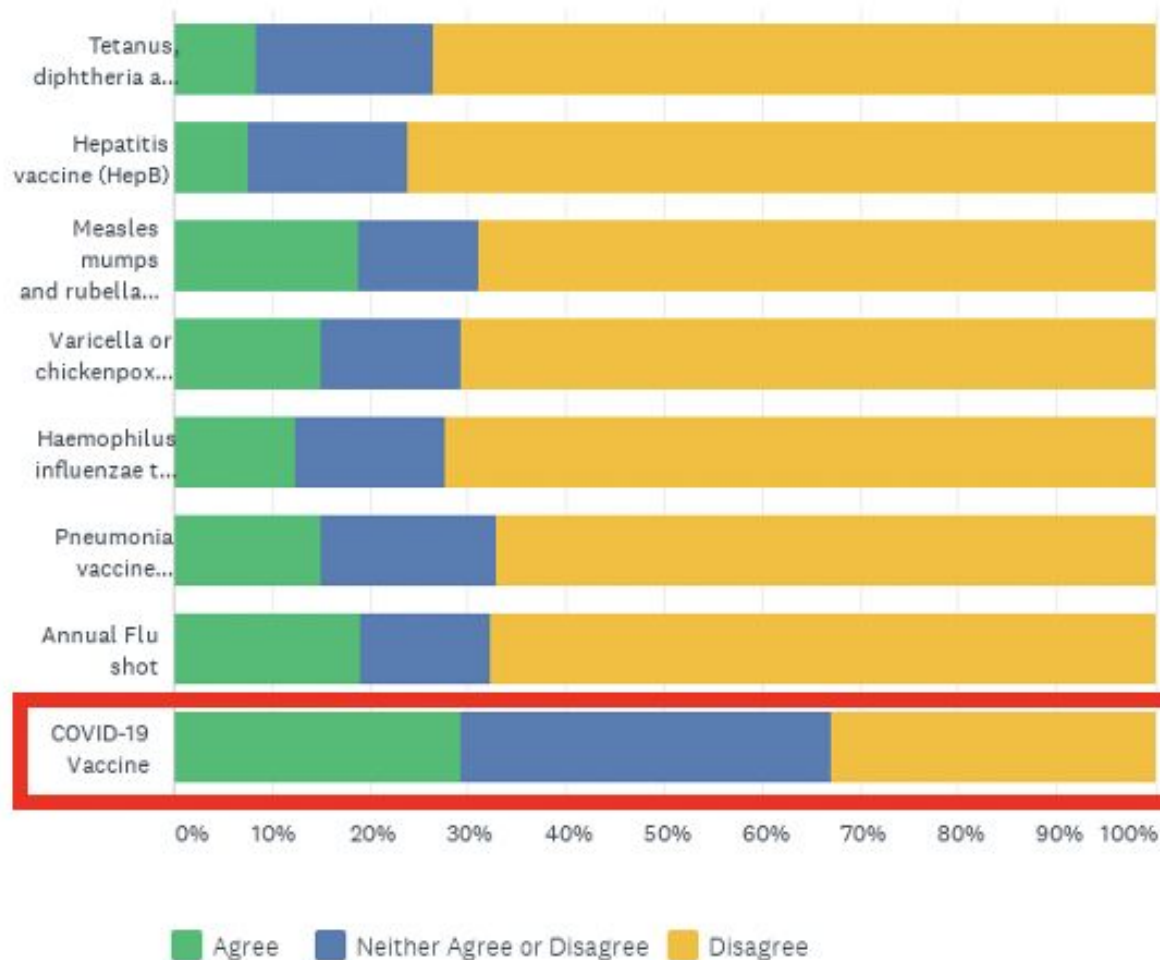


Confidence in Safety/Efficacy of COVID19 Vaccine

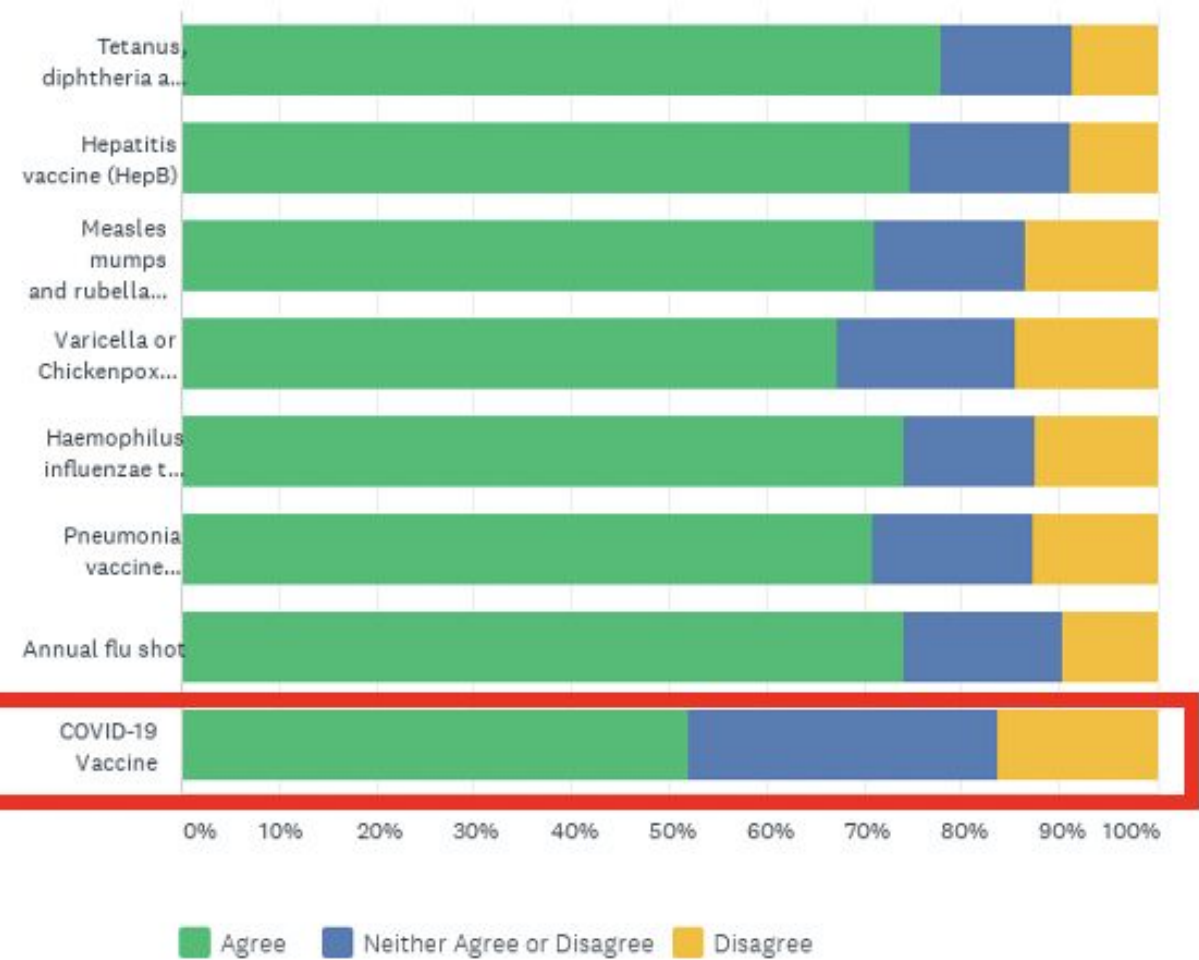
■ Great Deal ■ Fair Amount ■ Not too much ■ None

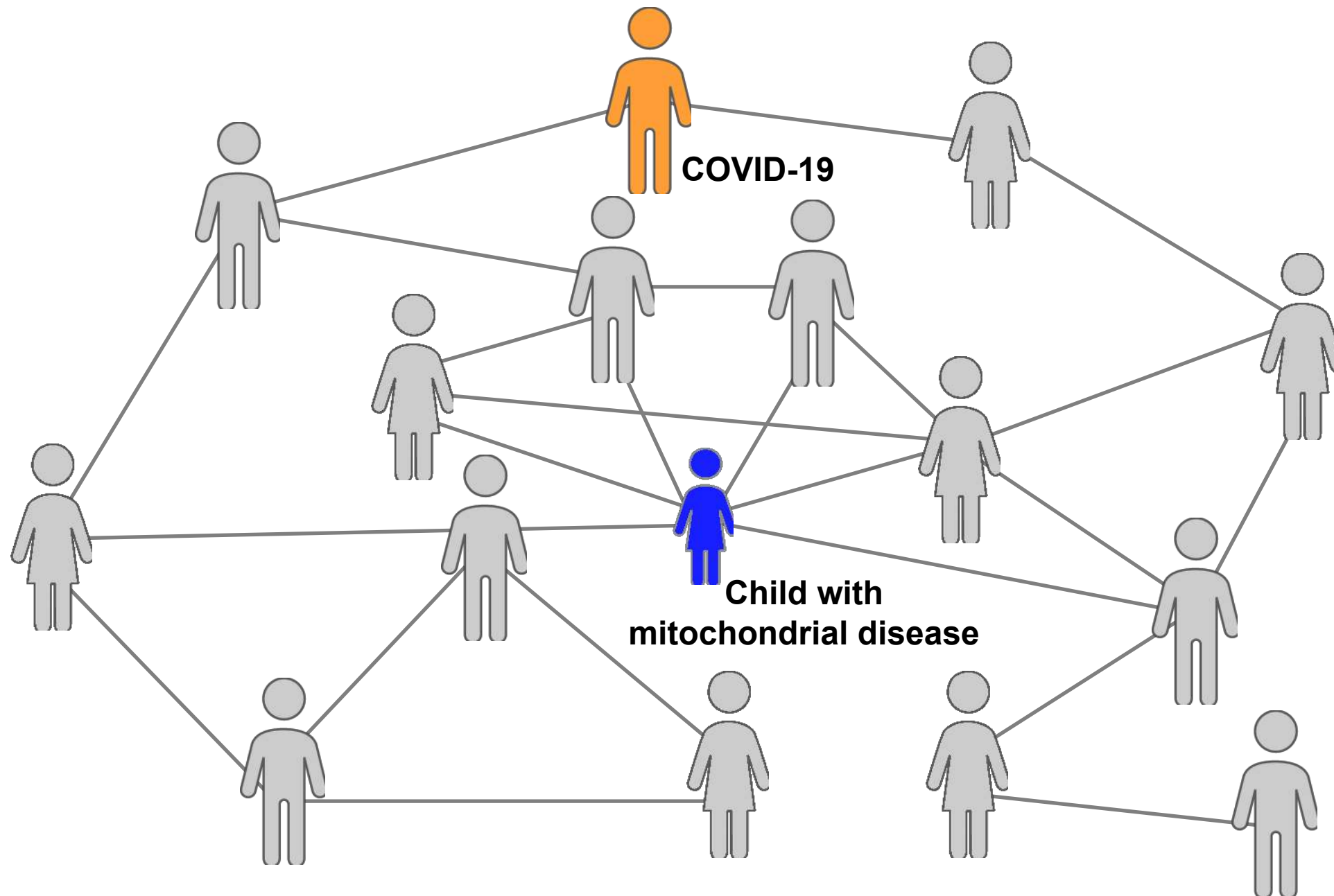


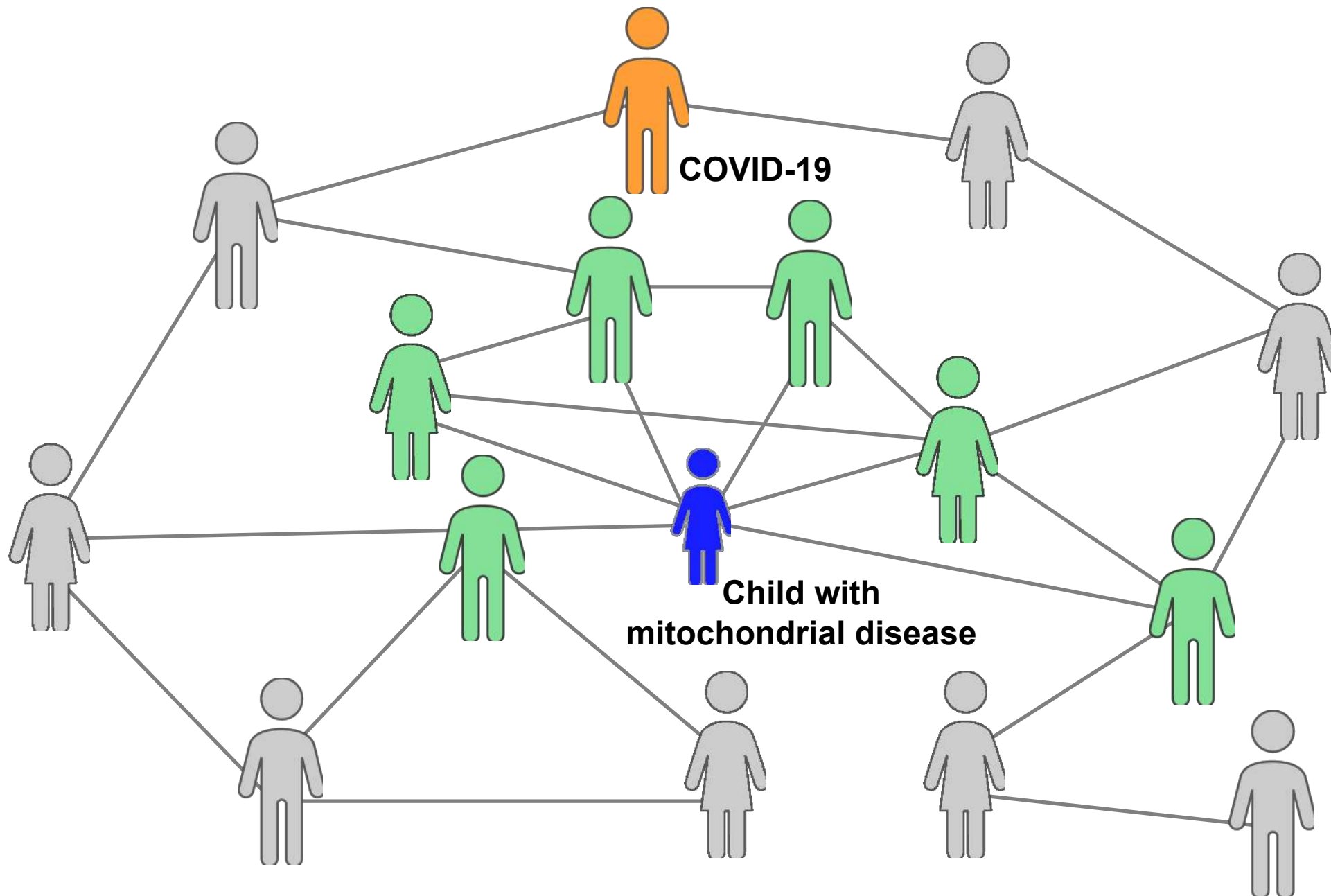
“I am concerned that my child with MtD will become sick or deteriorate after the following vaccinations:”



“The benefits of the following vaccines outweigh the risks in children with MtD.”

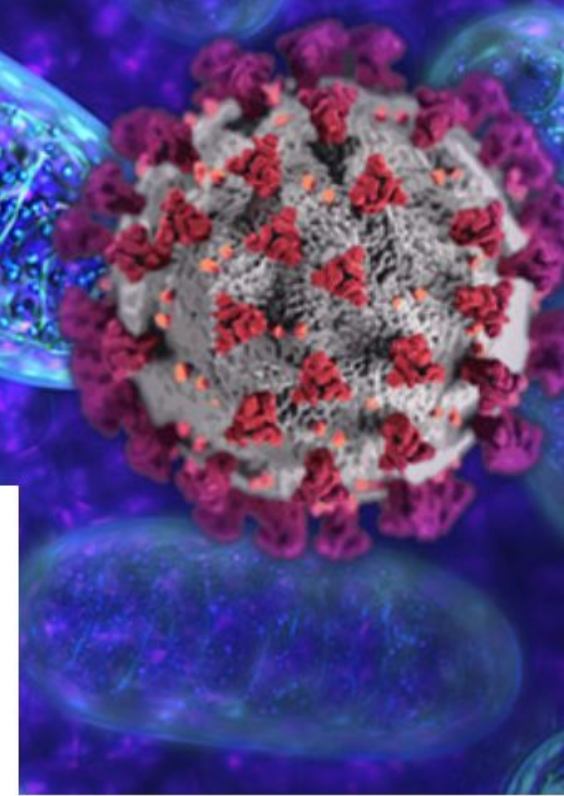






Post COVID-19 Vaccine Registry for Patients with Mitochondrial Disease

The NIH Metabolism, Infection and Immunity (MINI) Section invites people with mitochondrial disease who have already received the COVID-19 vaccine to participate in our new initiative: **the Post COVID-19 Vaccine Registry**.



Interim data from registry

- 66 patients/caregivers
 - 14 children
- Diagnoses: MtD not otherwise specified (23%), mitochondrial myopathy (23%), and MELAS (13%).
- 20% immunodeficiency/immunocompromised
- 12% previous adverse reaction
- Side effects: injection site reaction (62%), fatigue (47%), aches (26%), headache (24%), weakness (18%), fever (10%), allergic reaction (1.5%), no symptoms (15%)

Summary

- Individuals with MtD have risk factors for adverse outcomes with COVID-19
- Families with MtD are highly adherent to RMBs
- COVID-19 was widespread and underdiagnosed in MtD households
- RMBs may have worked
- Vaccine hesitancy occurs in the MtD community
- Ring vaccination is important
- Studies are ongoing: viral exposures, acute infection, vaccine registry

Thank you



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