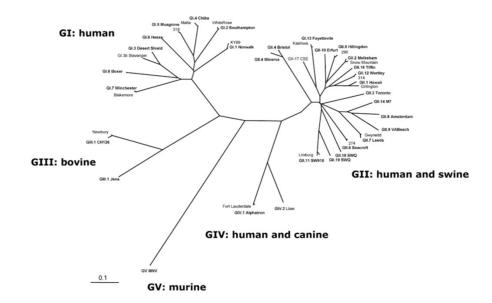
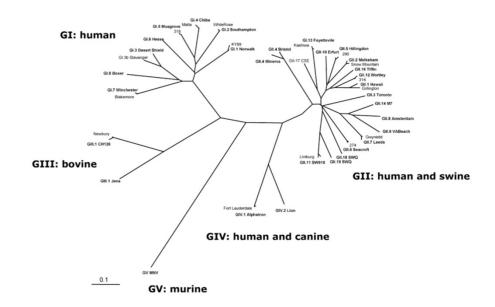


- Not traditionally (prior to 2014) cultivable in vitro
- Low infectious dose
- Survives on surfaces
- Disinfection difficult
- Asymptomatic shedding and carriers
- Diversity and rapid evolution
- No long-lasting immunity



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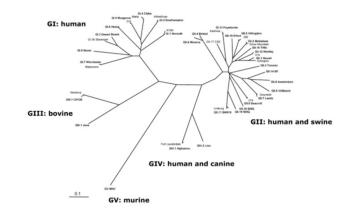


Not traditionally (prior to 2014)
 cultivable in vitro
 Note: Genogroup (II) → Genotype (4) → Strain

Low infectious dose

Survives on surfaces

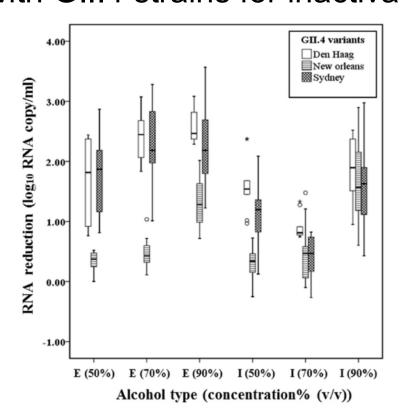
- Disinfection difficult
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(Sydney): GII.4 Sydney

Diversity of Susceptibility

Differences with GII.4 strains for inactivation



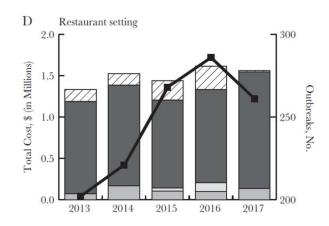
(Park et al. 2016)

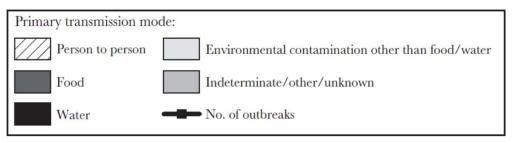
Challenges

- Resistance to inactivation
 - At currently allowed levels, many agents not effective enough
 - Even heat: 72°C did not remove signal until >10 min

Melting Temperatures Using Aptamer M6-2		Melting Temperatures Using HBGA	
Strain	Melting Temperature (°C)	Strain	Melting Temperature (°C)
SYV	73.10 ± 0.43	SYV	71.71 ± 0.49
SMV	75.03 ± 0.80	SMV	N/A
HOV	68.88 ± 1.10	HOV	67.69 ± 0.30

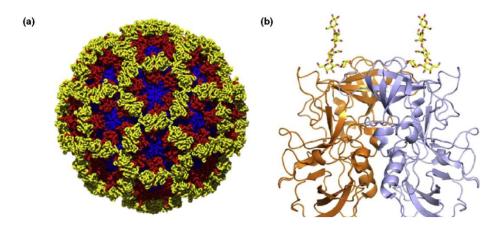
- Alleviate a major burden
 - 900 lives
 - 109,000 hospitalizations
 - 465,000 emergency department visits
 - 2.3 million urgent care visits
 - \$430-740 million in healthcare costs alone
 - \$10.6 billion overall annually





(Burke et al. 2020; Bartsch et al. 2020)

- Numerous challenges:
 - Antigenic and biological diversity
 - General lack of immune cross-protection with other genotypes
 - Can be re-infected with same genotype after a few years
 - Immune correlate of protection still a little unclear



(Esposito et al. 2020; Prasad et al. 2016)

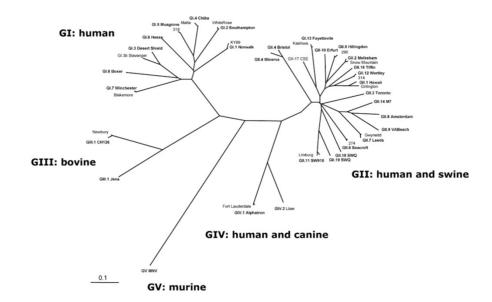
- Virus-like particle (VLP) based vaccines
 - Viral capsid without nucleic acid inside
 - GI.1, GII.3, GII.4 genotypes
 - Some combined with other virus targets
 - Preclinical through Phase II
 - A number of Phase II trials with positive preliminary results
- P particle vaccines
 - The "Protruding" domain of the major capsid protein
 - Outermost portion
 - Involved in receptor binding and major antigenic target
 - Mostly preclinical

(Esposito et al. 2020)

- Recombinant adenovirus vaccine
 - Uses adenovirus to deliver norovirus major capsid protein
 - Orally administered
 - GI.1 and GII.4
 - Currently in Phase II
 - Positive results for safety as well as robust immune response in animal model

(Esposito et al. 2020)

- Not traditionally (prior to 2014) cultivable in vitro
- Low infectious dose
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- In vitro cultivation
 - Over 40 years
 - Mid-life crisis: 3D cell culture model

- In vitro cultivation
 - Over 40 years
 - Mid-life crisis: 3D cell culture model

In Vitro Cell Culture Infectivity Assay for Human Noroviruses

Timothy M. Straub,* Kerstin Höner zu Bentrup,† Patricia Orosz-Coghlan,‡ Alice Dohnalkova,* Brooke K. Mayer,* Rachel A. Bartholomew,* Catherine O.Valdez,* Cynthia J. Bruckner-Lea,* Charles P. Gerba,‡ Morteza Abbaszadegan,§ and Cheryl A. Nickerson†

(Straub et al. 2007)

- In vitro cultivation
 - Over 40 years
 - Mid-life crisis: 3D cell culture model

OPEN ACCESS Freely available online



Challenges of Culturing Human Norovirus in Three-Dimensional Organoid Intestinal Cell Culture Models

Efstathia Papafragkou^{1,2}, Joanne Hewitt³, Geun Woo Park¹, Gail Greening³, Jan Vinjé¹*

1 Division of Viral Diseases, Center for Disease Control and Prevention, Atlanta, Georgia United States of America, 2 Center for Food Safety and Applied Nutrition, Division of Molecular Biology, Food and Drug Administration, Laurel, Maryland, United States of America, 3 Institute of Environmental Science and Research Ltd, Kenepuru Science Centre, Porirua, New Zealand

(Papafragkou et al. 2013)

- In vitro cultivation
 - Over 40 years
 - Mid-life crisis: 3D cell culture model
 - Complicates study
 - Infectivity dilemma

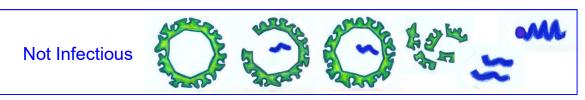
Infectivity Dilemma

- Lack of widely available/ideal in vitro cultivation
 - Cannot follow/examine behavior of infectious particles
 - Implications for inactivation methods, detection, and fundamental study of solution of viral particles
- Genome amplification techniques overestimate infectious particles
 - Free RNA
 - RNA from damaged/fatally mutated capsids
 - RNA with fatal mutations/damage



Theoretically Infectious





(Images courtesy Rebecca Goulter)

Cultivable Surrogates

- Utilize related cultivable surrogate viruses that are genetically and/or structurally related/similar
- Feline calicivirus, Tulane virus, murine norovirus (MNV), bacteriophage MS2
- However, differences in susceptibility
 - Possibility they are weaker than human norovirus
- Murine norovirus widely used for pathogenesis
 - In vivo and in vitro
- Differences in biological presentation

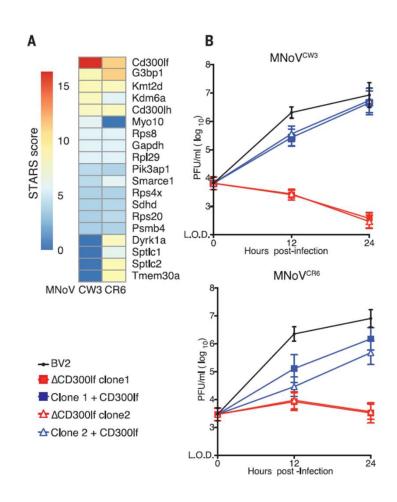
(Richards 2012; Wobus 2018; Karst & Tibbetts 2016)

VIROLOGY

Discovery of a proteinaceous cellular receptor for a norovirus

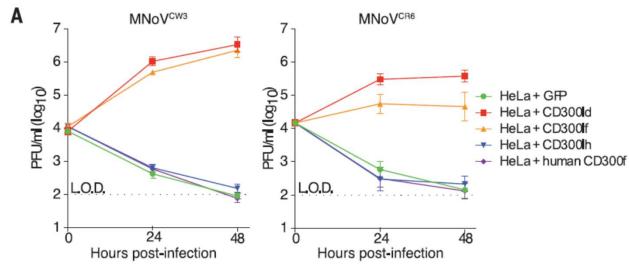
Robert C. Orchard,^{1*} Craig B. Wilen,^{1*} John G. Doench,² Megan T. Baldridge,¹ Broc T. McCune,¹ Ying-Chiang J. Lee,¹ Sanghyun Lee,¹ Shondra M. Pruett-Miller,³ Christopher A. Nelson,¹ Daved H. Fremont,¹ Herbert W. Virgin¹†

 Identify proteinaceous receptors, CD300lf and CD300ld, necessary for cell permissiveness to MNV infection



(Orchard et al. 2016)

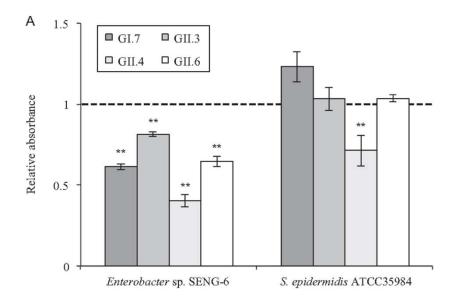
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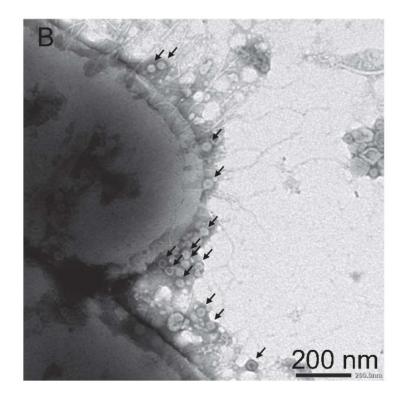
(Orchard et al. 2016)

- Identify functional proteinaceous receptors, CD300lf and CD300ld, necessary for cell permissiveness to MNV infection
- Makes human cells permissive, not vice versa
- Evidence for a small (<5kDa) nonproteinaceous cofactor to facilitate binding
- Not necessarily comparable to human norovirus

- In vitro cultivation
- Enteric bacteria –Potentially a co-factor?
 - HuNoV found to bind Enterobacter cloacae



(Miura et al. 2013)



(Miura et al. 2013)

 Enteric bacteria as potential co-factor for infection in human B cells

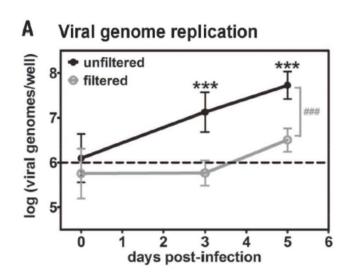


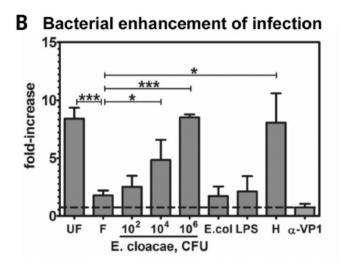
Enteric bacteria promote human and mouse norovirus infection of B cells

Melissa K. Jones, ^{1*} Makiko Watanabe, ^{1*} Shu Zhu, ¹ Christina L. Graves, ^{2,3} Lisa R. Keyes, ¹ Katrina R. Grau, ¹ Mariam B. Gonzalez-Hernandez, ⁴ Nicole M. Iovine, ⁵ Christiane E. Wobus, ⁴ Jan Vinjé, ⁶ Scott A. Tibbetts, ¹ Shannon M. Wallet, ^{2,3} Stephanie M. Karst ¹

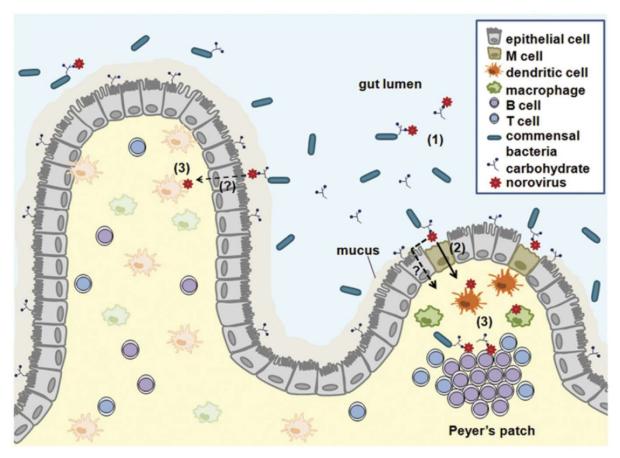
(Jones et al. 2014)

 Enteric bacteria as potential co-factor for infection in human B cells





(Jones et al. 2014)



(Karst & Wobus 2015)

- Enteric bacteria as potential co-factor for infection in human B cells
 - However, can have issues in replicating
 - < 3 log₁₀ production → sensitivity a challenge
 - Requirement of additional co-factors could confound inactivation study
- Antibiotic treatment reduces viral titer in mouse model

(Jones et al. 2015; Almand et al. 2017)

Enteric Bacteria and Norovirus

- Some evidence binding increases stability of virus
 - Though some conflicting reports
- Some bacteria down-regulate by enhancing virusspecific antibody titer
- Others promote viral adherence and binding
- Bacterially modified bile acids (secondary bile acids) may enhance viral infection
 - Can enhance binding to receptor

(Moore & Jaykus 2018; Almand et al. 2017; Neu & Mainou 2020; Jones et a. 2016; Sullender & Baldridge 2018)

Enteric Bacteria and Norovirus

- Commensal bacteria can enhance production of secretory immunoglobulins
 - Actually promotes MNV infection
- Potential bacterial effect on tropism of virus
- Can alter gut microbiota, promote higher Firmicutes to Bacteroidetes ratio (dysbiosis)
 - How long that lasts unclear
- Native microbiota on produce may aid attachment

(Moore & Jaykus 2018; Almand et al. 2017; Neu & Mainou 2020; Jones et a. 2016; Sullender & Baldridge 2018)

Enter the Enteroids

- Another cell culture model released from group at Baylor
- Involves creation of human intestinal enteroids (HIEs)

Replication of human noroviruses in stem cell-derived human enteroids

Khalil Ettayebi,¹* Sue E. Crawford,¹* Kosuke Murakami,¹* James R. Broughman,¹ Umesh Karandikar,¹ Victoria R. Tenge,¹ Frederick H. Neill,¹ Sarah E. Blutt,¹ Xi-Lei Zeng,¹ Lin Qu,¹ Baijun Kou,¹ Antone R. Opekun,²,³,⁴ Douglas Burrin,³,⁴ David Y. Graham,¹,²,⁵ Sasirekha Ramani,¹ Robert L. Atmar,¹,² Mary K. Estes¹,²†

(Ettayebi et al. 2016)

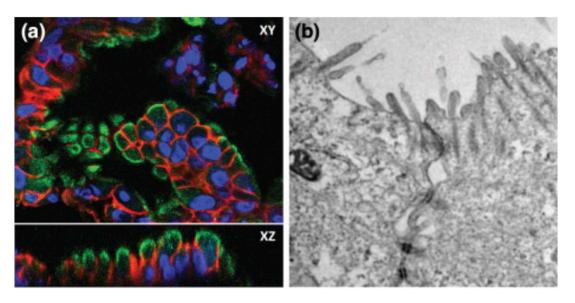
The Enteroid Strikes Back

HIE Recipe:

Isolate stem cells from human intestinal crypts

Provide growth factors and nutrients to develop into differentiated "mini

guts"



(Kovbasnjuk et al. 2013)

The Enteroid Strikes Back

- HIE Recipe:
 - Isolate stem cells from human intestinal crypts
 - Provide growth factors and nutrients to develop into differentiated "miniguts"
 - Behave like intestine, multiple intestinal epithelial cell types:
 enterocytes, goblet, enteroendocrine, Paneth cells
 - Can be 3D or grown as monolayer

(Ettayebi et al. 2016; Kovbasnjuk et al. 2013)

Enteroids for Inactivation

- Aged green tea; heat; chlorine; ethanol
- Real value in comparison to surrogates

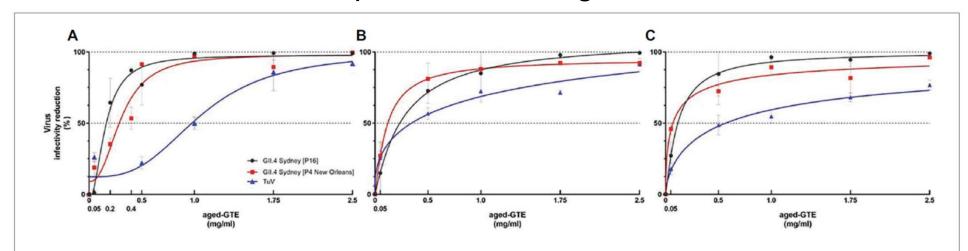
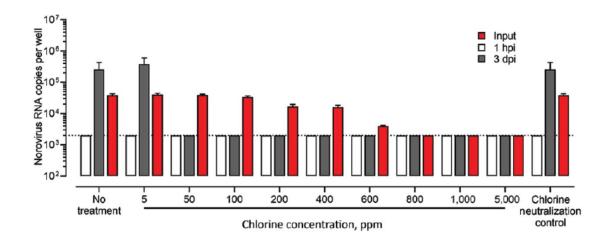


FIGURE 4 | Effective concentration (EC₅₀) of aged-GTE that inhibits human norovirus and Tulane virus replication. Viruses were exposed to aged-GTE for 1 h at 37°C **(A)**, 21°C **(B)**, and 7°C **(C)**. A nonlinear regression (curve fit) function on data sets of three experiments with three technical replicates for each treatment, time point, and virus were used to obtain EC₅₀ values. The data is based on genomic copies for human norovirus and on TCID₅₀ for Tulane virus.

(Ettayebi et al. 2016; Constantini et al. 2018; Randazzo et al. 2020)

Enteroids for Inactivation

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- 60°C for 15 min; 50 ppm chlorine for 1 min



(Ettayebi et al. 2016; Constantini et al. 2018; Randazzo et al. 2020)

Enteroids for Inactivation

- Aged green tea; heat; chlorine; ethanol
- Real value in comparison to surrogates
- 60°C for 15 min; 50 ppm chlorine for 1 min
- Still utilize RT-qPCR
- Not much sensitivity (~2 log reduction)
- Great model for antivirals/therapeutics

(Ettayebi et al. 2016; Constantini et al. 2018; Randazzo et al. 2020)

HIE Model: Ready for Our Purposes?

- Major breakthrough, both models game-changers
- Replicated in multiple labs
- Great for studying HuNoV biology/infection mechanisms
- Not the highest production
 - Not high enough to provide ideal stock sensitivity for testing inactivation agents
- Bottom line for applied purposes: Stay tuned

Noroviruses Can Potentially have Good Effects?

Some evidence in germ-free and antibiotic-treated mice with MNV



doi:10.1038/nature13960

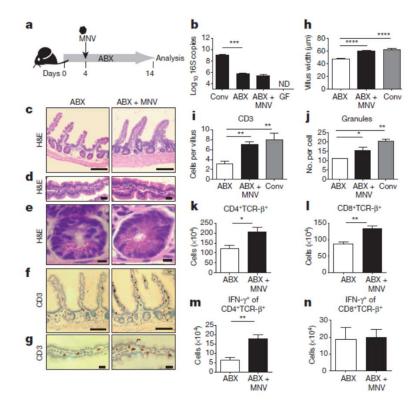
An enteric virus can replace the beneficial function of commensal bacteria

Elisabeth Kernbauer^{1,2}, Yi Ding^{3,4} & Ken Cadwell^{1,2}

Noroviruses Can Potentially have Good Effects?

Improved intestinal morphology and mucosal immune

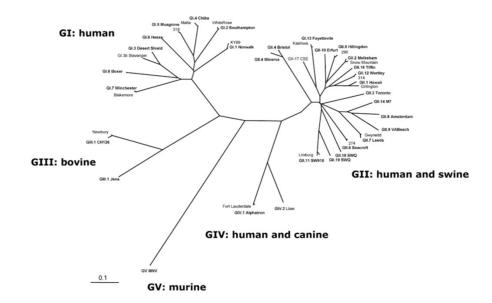
function



(Kernbauer et al. 2014)

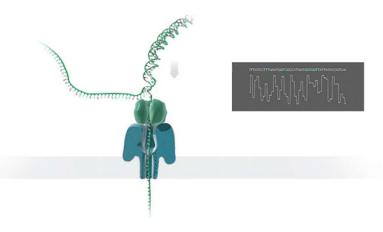
Human Norovirus

- Not traditionally (prior to 2014) cultivable in vitro
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- Disinfection difficult
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- Diversity and rapid evolution
- No long-lasting immunity



(Vinje 2014; Bull et al. 2010; Patel 2009)

- Whole genome sequencing
- Nanopore sequencing: MINion
- How it works
- Size of graphing calculator, 90g
- Powered by USB to computer
- Real-time reads to laptop
- Reads up to 60 kb (NoV= 7.5 kb)
- Can read straight RNA
- \$1,000 to get started



(Hoenen et al. 2016; Wang et al. 2015; Oxford Nanopore Technologies)

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(Hoenen et al. 2016; Wang et al. 2015; Ibtimes.com)

- Has been used for other viruses
 - Influenza
 - Ebola
 - Poxviruses
 - Lambda bacteriophage
- Usually requires amplification
- Not yet used for foodborne viruses
- Challenges: base calling accuracy, data analysis



(Hoenen et al. 2016; Wang et al. 2015; The Atlantic)

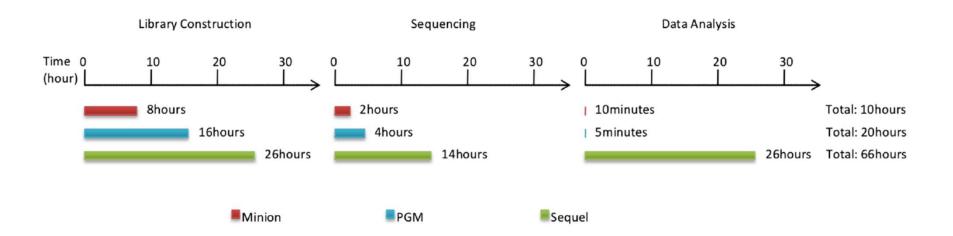
 Able to detect GII norovirus from fecal samples using 3rd generation sequencing technology (MinION and PacBio Sequel)

Comparison of third-generation sequencing approaches to identify viral pathogens under public health emergency conditions

Yang Li¹ · Xiao-zhou He¹ · Ming-hui Li² · Bo Li³ · Meng-jie Yang¹ · Yao Xie² · Yi Zhang¹ · Xue-jun Ma^{1,4}

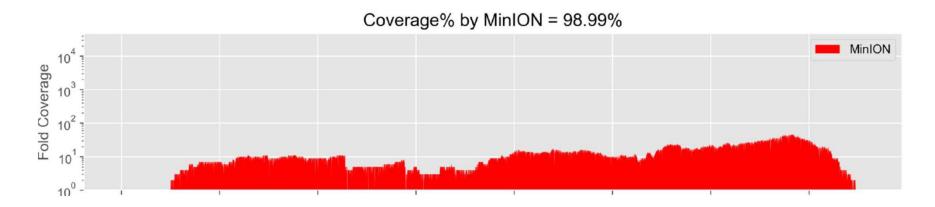
(Li et al. 2020)

- Able to detect GII norovirus from fecal samples
- Comparatively quicker sample-to-result (~10 hours)



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(Li et al. 2020)

- Able to detect GII norovirus from fecal samples
- Comparatively quicker sample-to-result (~10 hours)
- Not amazing depth, but enough coverage to be valuable
- Only applicable for clinical samples at this time
- Implications for identification of outbreaks

Conclusions

- Norovirus imposes huge burden globally and in US
- Seemingly small and simple structurally
- Many challenges
 - Infectious dose, diversity, multiple transmission routes, stability, requirement for concentration, asymptomatic shedding
 - Big recent developments: cell culture models; vaccine development, potential beneficial effects, 3rd-generation sequencing
- Major take-aways from this presentation:
 - Noroviruses present MANY challenges, but a lot of discoveries are likely to occur in the next 5 years
 - Even with breakthroughs in fundamental understanding, challenges will still be difficult in our field

Acknowledgements

- Everyone attending!
- NEHA
- Dr. Clyde Manuel
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- Terryn Laird
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- Hao-Yuan Hsu
- Minji Kim
- Pragathi Kamarasu
- Louisa Bachman
- Melina Demokritou
- Nicholas Holt
- Sloane Stoufer

UMassAmherst



Questions?

