Performing the GI Virus Panel by Real-Time PCR Procedure

Purpose: This procedure provides instructions for how to perform the GI Virus Panel by Real-Time Polymerase Chain Reaction (q-PCR) to detect Sapovirus(Sapo), Rotavirus(Rota), Adenovirus(Adv) and Astrovirus(Ast) from patient's fecal and vomitus samples using a standardized methodology.

I. Samples: Nucleic acid from patient fecal and vomitus samples extracted using the bioMérieux easyMAG. See PHGI_101_1600 for acceptable sample types and collection containers.

II. Materials:

A. Media: N/A

B. Primers and Probes

Target	Primer	Primer Sequence (5'- 3')		Manufacturer
	Sav1F	TTG GCC CTC GCC ACC TAC	Junction of	
Sapovirus	Sav5F	TTT GAA CAA GCT GTG GCA TGC TAC		Life
Sapovirus	Sav124F	GAY CAS GCT CTC GCY ACC TAC	polymerase and capsid	Technologies
	Sav124R	CCC TCC ATY TCA AAC ACT A	Capsiu	reciliologies
Potoviruo Tyro	NSP3-F	ACC ATC TWC ACR TRA CCC TCT ATG AG	Non structure	
Rotavirus Type A	NSP3-R	GGT CAC ATA ACG CCC CTA TAG C	Protein 3	Life Technologies
Adenovirus	AdV-F	GCC TGG GGA ACA AGT TCA G		
40/41	AdV-R	ACG GCC AGC GTA AAG CG	Hexon	Life Technologies
	Ast-F	AAG CAG CTT CGT GAR TCT GG	Junction of	
Astrovirus	Ast-R	GCC ATC RCA CTT CTT TGG TCC	polymerase and capsid	Life Technologies

Target	Probe	Sequence(5'-3')	Reporter	Quencher	Manufacturer
	Sav124TP	CCR CCT ATR AAC CA			
Sapovirus	Sav5TP	TGC CAC CAA TGT ACC A	FAM	MGB-NFQ	Life Technologies
Rotavirus Type A	NSP3-P	AGT TAA AAG CTA ACA CTG TCA AA	VIC	MGB-NFQ	Life Technologies
Adenovirus 40/41	AdV-P	ACC CAC GAT GTA ACC AC	NED	MGB-NFQ	Life Technologies
Astrovirus	Ast-P	CAC AGA AGA GCA ACT CCA TCG CAT TTG	Су5	Tao-IBDRQ	Integrated DNA Technologies

C. Other Reagents, Supplies and Equipment:

Reagent	Supplies	Equipment
Applied Biosystems TaqMan® Fast Virus 1-Step Master Mix(cat# 4444434)	Applied Biosystems Fast Optical 96-Well Plates or MicroAmp Optical 8-well strips	ABI 7500 Real-Time PCR System with Fast Block

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IDTE 1x TE Buffer pH8.0	MicroAmp Optical Adhesive Film (P/N4311971) and Applicator or MicroAmp Optical 8-well strip caps	Biological safety cabinet- 6ft, Nuaire, Class II Type A/B3
UltraPURE DNAse/RNAse Free water	Sterile Barrier Tips for pipets	Pipettes (2.5 μL, 10 μL, 20 μL, 100 μL and 1000 μL)
DNA AWAY, Molecular BioProducts	Disposable Powder-free Gloves and Gowns	Mini-Micro Vortex Mixer
gBlocks® control oligos (see table C)	Under pad, disposable	96-well plate centrifuge
Custom primers and probes (see table B)		

D. gBlocks Control Oligos (Integrated DNA Technologies)

D11-	0		
gBlock	Sequence		
Human adenovirus	5'-		
type 40/41_gBlock	GCCTGGGGAACAAGTTCAGAAACCCCACCGTGGCTCCGACCCACG		
	ATGTAACCACAGACAGGTCACAGCGACTGACGCTGCGATTCGTGCC		
	CGTCGACCGCGAGGAAACCGCCTACTCTTACAAAGTGCGCTTTACG		
	CTGGCCGT - 3'		
Human astrovirus	5'-		
gBlock	TGGAGCACTGCCTCTCACGGACTGCAAAGCAGCTTCGTGACTCTGG		
gblock	CCTTCCA		
	GCCAGACTCACAGAAGAGCAACTCCATCGCATTTGGAGGGGAGGA		
	CCAAAGAAGTGTGATGGCTAGCAAGTCCAATA - 3'		
Human rotavirus	5'-		
type A gBlock	ACTATGAATATGCATATGAGTAGTCACATAATAAAGAATGTTCACCAT		
	CTACACA		
	TGACCCTCTATGAGCACAATAGTTAAAAGCTAACACTGTCAAAAACC		
	TAAATGGCTATAGGGGCGTTATGTGACC - 3'		
Human sapovirus	SaV124		
Polymerase/Capsid	5'-		
Junction gBlock®	TAACACCAACTATGACCAGGCTCTCGCCACCTACAATGCTTGGTTCA		
Juliction applock	TAGGTGGTACAGTACCTGA		
	CCCAGTGGGTTACACTGAAGGAACCCACAAAATAGTGTTTGAGATG		
	GAGGGCAATGGCTCCAAC - 3'		
	SaV1		
	5'-		
	TGAATACAAATTTTGATTTGGCCCTCGCCACCTACAATGCCTGGTTC		
	ATAGGTGGTACAGCTCCA		
	GATCCAGAGCGCCCCACTGAAGGTGCACCCAAATTAGTGTTTGAGA		
	TGGAGGGCAATGGCTCCA - 3'		
	SaV5		
	5'-		
	GTTAATACCAACTTTGAACAAGCTGTGGCATGCTACAACAGCTGGTA		
	CATTGGTGGCACCACACC		
	AGAAATGCCCACTACCAATGAAGGCTGTGGGCTATTAGTGTTTGAG		
	ATGGAGGCAATGGCTCCC - 3'		

III. Quality Control:

A. Positive control

IDT gBlocks® or suitable extracts of known positive targets diluted such that the Ct value is between 29 and 31 cycles when the threshold is set at a Δ Rn value of 0.05. Currently the controls are set to be:

Astrovirus gBlock	12500 copies per reaction	
Adenovirus gBlock	1000 copies per reaction	
Rotavirus gBlock	1000 copies per reaction	
Sapovirus RNA Extract	Aim for CT value between 27-31	

B. Negative (no template control -NTC)

The negative PCR control is DEPC water substituted for sample.

C. Extraction control (Mean + 2SD)

DEPC water follows the entire extraction protocol and should be used as an extraction control.

D. Inhibition control

All fecal extracted and first tested for Norovirus which includes an exogenous internal control West Nile Virus. All samples should pass the detection of -WNV before carrying on to the GI panel test.

WNV should be detected in all samples . Any sample that is negative for WNV is either inhibited or insufficient and must be repeated.

IV. Procedure:

A. Running the q-PCR on the TaqMan 7500

	. Kuming	Related Documents Title Number					
	Note: This assay was specifically designed to detect Human sapovirus, Human rotavirus type A, Enteric adenovirus types 40/41 and Human astrovirus Types 1-8 all associated with gastroenteritis. The assay has not been designed or validated to detect other types that cause infection in other body sites.				BCCDC Laboratory Safety Manual		
1.		mple extraction file into the Norofast PCR System.	st run template on	the ABI 7500 Fast			
2.	extraction	e BSCs in the PCR reagent clean clean room, assure stable function with DNA Away prior to starting wor	ality and wipe dov		BCCDC Safety Manual, Section F-BSCs		
4.	Note: The 20x primer/probe mix should be prepared ahead of time and QC'd before use in a sample run. See Appendix A for recipe.						
		Reagent	Vol./reaction (µl)				
		PCR grade Water	9				
		TaqMan® Fast Virus 1-Step Mix (4X)	5				
		20x GI panel multiplex primer/probe mix	1				
	Note: Reco	ord the Lot Numbers for all PCR Maste eet.	er Mix reagents in th	ne spaces provided on			
5.	Remove an Applied Biosystems Fast Optical 96-Well Plate and or MicroAmp Optical 8-well strip from the box, inside the BSC, and place it in a MicroAmp 96-Well Plate Holder to ensure that the bottom of the plate has no contact with anything that could fluoresce.						
6.	Note: <u>Do not</u> use permanent marker on optical plates, the ink will fluoresce. Aliquot 15µL of the GI panel cocktail to the wells being used for the assay in the plate/strips.						
	Note: Only use ABI Fast Optical 96-Well Reaction Plates or MicroAmp Optical 8-well strips in the fast block enabled TaqMan 7500.						
7.	Transport the qPCR plate/strips in a ziploc bag to the genomic preparation room. Do not remove the plate from the bag until you are ready to add the sample extracts.						
8.	Place the	qPCR plate/strips in the plate holde	er in BSC.				
9.	Add 5µL o	f control or sample to the appropria by the extraction/plate layout sheet	ate well of the 96-v	well plate/strip as			

		Related Documents Title Number				
	Note: This assay was a type A, Enteric adenov gastroenteritis. The assause infection in other	BCCDC				
10.	Seal the plate with a the MicroAmp optica			optical 8-well strips witl pplicator.	h	
	Note: Be careful not cont as ink, dust or fingerprints		tical surface with anyth	ing that could fluoresce suc	h	
11.	•	•		des and bottom of each wn in a plate centrifuge		
	Note: Place the plate/strip contaminated with dust. spin down the plate if this	Do not run a plate with re		down so it is not underside of the optical film,		
12.	Place the sealed plat	te/strips back in the z	rip-loc bag to trans	port it to the ABI 7500.		
	Note: transporting the prior to loading on the		bag prevents contai	mination with dust etc.		
13.	Load the plate/strips Note: Use the appro			block.	PHGI_283_2500 Using the 7500 Fast Real-Time PCR System	
14.		on the ABI Fast 750	0 using the following	ng program stored as G	SI .	
	Fast Run Protocol:					
	Step	Temperature (°C)	Time (min:sec)	cycles		
	UNG activation* (or RT)	50	5:00	1		
	Taq activation	95	0:20	1		
	Amplification	95	0:03	40		
	7	60	0:30			
	*Note: Published UNG activation for TaqMan Fast Advance 2x Master Mix is 2 minutes. Holding the reactions at 50°C for 5 minutes does not adversely affect master mix performance.					
	Confirm that the cycling program is correct before starting the instrument and the all detectors are selected for each sample					
15.	Initiate the run by clic					
16.	Confirm that the instr	rument is running bef	fore you leave.			

B. Results Analysis

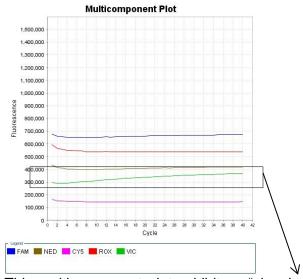
	Action				
	Note: Use Universal Precautions and appropriate PP according to the Waste Management Policy. Process handling occurs in the PCR Genomic Extraction Clear PCR Reagent Clean Room	BCCDC Laboratory Safety Manual			
1.	Open the ANALYSIS setting window and entargets is set to 0.05.	isure that the C_TT	hreshold for all		
2.	Analyze the q-PCR results of the controls for If	r each of the prime	er sets;		
			no the positive	_	
	the CT value for the negative control is Undetermined	Proceed to analy controls	se the positive		
	the CT value for the negative control is NOT Undetermined	Check the multice	omponent plot		
		If	Then		
		there is actually a rise in fluorescence relative to the baseline ROX	 The run has failed due to contamination. Consult your supervisor and repeat the run with reagents that have passed QC. 		
		there is No rise in fluorescence relative to the baseline ROX	 There may be background fluorescence in the wells or the plate. Inform your supervisor 		
3.	Check the CT value for all positive controls				
	If]			
	The Positive Control has a C_T value within its quality control range for all targets	The run is valid a can be analyzed.			
	NOTE: See QC Stat Tools for details	C Stat Tools for details			
	The Positive Control does not have a C_T value within its quality control range.	The positive cont Repeat the run us new control.			

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Analyze the q-PCR results of the patient samples				
If Then	Then			
Ct values are not detected for the target or are Undetermined No Sapovirus, Rotavirus, Adenovirus, and Astrovirus was detected in the patient sample				
The sample has a Ct value > 35 cycles, but < 40 cycles the result is either false positive or weakly positive. Consult Supervisor. If instructed, repeat testing in duplicate				
Re-extraction would limit the possibility of it being a contaminated sample during the extraction/loading steps. If Then A Ct value is report as positive for 2/3 technical replicates A Ct value is not reproducible for 2/3 technical replicates A Ct value is not reproducible for 2/3 technical replicates				
If Then				
The amplification plot forms a normal curve.				
The amplification plot is NOT a normal curve. Consult Supervisor and troubleshoot.	k			
Export results into GI Panel Workflow Template. Print results and verify the CT values on the printout with the amplification plot and multicomponent plot.				
Enter control results into the QC Stat Tools and note on worksheet if QC has passed. Pass all worksheets to Supervisor for review.				
Proceed to PHGI_304_3600 Reporting GIDO Result Entry				

Procedure Notes:

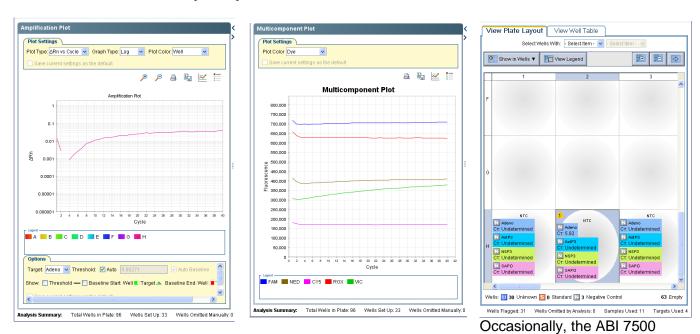
Method Limitations:

- A. This method has been only validated for fecal and vomitus samples extracted on the bioMérieux EasyMag.
- B. VIC-labelled probe and slope drift.



This multicomponent plot exhibits a "signal drift" for the VIC-labelled probe. The reasoning for this is unknown and will occur with a change of dye. Nevertheless, it has been validated against strong positives will not cause false positives. ABI has confirmed that this phenomenon is acceptable.

C. Poor Analysis by Software:



Occasionally, the software will calculate the math for some samples improperly. As you can see above, the background calculation has produced a positive detection at a Ct value of 5.82(automatic threshold). With a strong Ct value of 5.82 you would expect your multicomponent plot to produce both a strong fluorescence and a normal sigmoidal curve; however upon inspection we can see that there is no evidence of exponential amplification or any amplification for that matter.

There are two options here: you can disregard the amplification based on the multicomponent plot or you can alter the background calculations by manually setting the background cycle count cycles between 3 and 15. Cycle 3 is where actual target length template doubling is occurring and cycle 15 is an approximate value for a typically strong positive that will appear on a plate if the plate contains any positives.

References:

Pang, Xiaoli L., et al. *Journal of Medical Virology: Enhanced Enteric Virus Detection in Sporadic Gastroenteritis using a Multi-Target Real-Time PCR Panel: A One-Year Study.* 86 Vol. John Wiley & Sons Inc, 09/01/2014. Web. 2 Oct. 2015.

Zeng, S. -Q, et al. "One-Step Quantitative RT-PCR for the Detection of Rotavirus in Acute Gastroenteritis." Journal of Virological Methods 153.2 (2008): 238-40. Web.

Feeney, S. A., et al. *Journal of Medical Virology: Development and Clinical Validation of Multiplex TaqMan Assays for Rapid Diagnosis of Viral Gastroenteritis*. 83 Vol. John Wiley & Sons Inc, 09/01/2011. Web. 2 Oct. 2015.

Oka, Tomoichiro, et al. "Detection of Human Sapovirus by real-time Reverse transcription-polymerase Chain Reaction." Journal of Medical Virology 78.10 (2006): 1347-53. Web.

Final Validation Report, *Gastroenteritis Viral Panel Detection using qPCR;* Trevor Hird, MMGP and Environmental Microbiology Laboratory Affiliations, 2016.

Appendix A

Preparation of 20x Primer/Probe mix:

Prepare the GI Panel 20x primer-probe mix used in the qPCR cocktail according to the following recipe:

(Confirm the stock concentration in order to finalize the 20x Mix Recipe)

20x Mix Recipe						
Primer/Probe	[Stock] uM	[20x Mix] uM	[Final] nM	Volume(ul) of stock to add (500ul/rxn final vol.)		
Ast-F	200	9	450	22.5		
Ast-R	200	9	450	22.5		
Ast-P	100	3	150	15		
AdV-F	200	8	400	20		
Adv-R	200	8	400	20		
Adv-P	100	4	200	20		
Sav1F	200	4	200	10		
Sav5F	200	4	200	10		
Sav124F	200	4	200	10		
Sav124R	200	4	200	10		
Sav5TP	100	2	100	10		
Sav124TP	100	2	100	10		
NSP3-F	200	6	300	15		
NSP3-R	200	6	300	15		
NSP3-P	100	2	100	10		
	•		1x IDTE	280		
			Total Volume	500		

Note: Store the 20x mix at -20°C in the dark. Excessive exposure to light may affect the fluorescent probes. <u>Do not</u> perform more than 20 freeze-thaw cycles. **If you expect to freeze-thaw the 20x mix more than ten times, consider aliquoting to minimize the number of freeze-thaw cycles.**

Revision History

Version	Date	Writer	Description of Change
0.0	July 1, 2016	Trevor Hird	Training Draft
1.0	November 27, 2017	Angela Kong/Stephanie Man	Initial Release