



Noro-, Rotavirus, C. diff

PCR Test for Norovirus, Rotavirus, C. difficile (tcdA/tcdB)

Instructions for use Gebrauchsanweisung

Introduction

Clostridioides difficile (C. difficile) is a gram-positive, spore-forming anaerobic bacillus consisting of toxigenic and non-toxigenic strains. C. difficile is one of the most common pathogens of nosocomial diarrhea, especially in patients with antibiotic-associated diarrhea. In recent years, the number and severity of cases has increased in Europe and worldwide.1 The symptoms of C. difficile infection (CDI) ranges from mild diarrhea to severe life-threatening pseudomembranous colitis, although colonization of C. difficile does not necessarily lead to a symptomatic infection. The disruption of the balance of gut microorganisms e.g. due to antibiotically treatment may cause toxigenic C. difficile to establish, dominate colonization and might be the start of CDI.2

Factors involved in the pathogenesis of *C. difficile*, are the genes *tcdA* and *tcdB* that code for toxin A (enterotoxin) and toxin B (cytotoxin) which are located in a 19.6 kb chromosomal region called the pathogenicity locus (PaLoc). Most pathogenic strains are toxin A and B positive (A+B+), but some pathogenic variant isolates may be toxin A negative and B positive (A-B+). They are the cause of *C. difficile*-associated disasses.3 Some strains of toxigenic *C. difficile* also produce a toxin called *C. difficile* transferase (CDT), or binary toxin.4

Norovirus belongs to the virus class Caliciviridae and is known to be a seasonal disease. It is a highly contagious pathogen that is responsible for the majority of cases of acute gastroenteritis worldwide. Norovirus infections in humans are mainly caused by genogroup II and I, while genogroup II being the predominant cause of Norovirus outbreaks. Infections of the gastrointestinal tract by Norovirus are characterized by sudden onset symptoms such as nausea, vomiting, diarrhea, and abdominal pain. The virus is particularly prevalent in communal settings such as hospitals, schools, and nursing homes, where it can easily spread from person to person. The illness typically lasts only a few days but can cause more severe complications in immunocompromised individuals or the elderly,5,6,7

Rotavirus belongs to the virus class Reoviridae. It is a significant cause of acute gastroenteritis, particularly in infants and young children. Its viral infection of the gastrointestinal tract is characterized by symptoms such as severe diarrhea, vomiting, fever, and dehydration. Rotavirus-associated gastroenteritis can lead to severe complications, including hospitalization and even death, making it a major public health concern. Efforts to prevent and control rotavirus infections, such as through vaccination programs, have been crucial in reducing the burden of this disease. Acute gastroenteritis caused by both Norovirus and Rotavirus are self-limiting diseases although if untreated can cause a critical course of disease especially in the youngest or elderly. 7,8,9,10

Package Contents

15 Vivalytic Noro-, Rotavirus, C.diff test cartridges for the detection of *Clostridioides difficile* (tcdA/tcdB toxin genes), human Norovirus genogroup I/ II and Rotavirus type A.

Pathogen List		
Clostridioides difficile (tcdA/tcdB)	Norovirus genogroup I/II	Rotavirus Type A

Intended Use

The Vivalytic Noro-, Rotavirus, C.diff test is an automated qualitative in vitro diagnostic test based on real-time polymerase chain reaction (PCR) for the detection of nucleic acids from toxigenic Clostridioides difficile (toxin genes tcdA/tcdB), Human Norovirus genogroup I/ II and Human Rotavirus type A from swabs samples from liquid or soft human stool to aid in the diagnosis of acute gastrointestinal infections of symptomatic individuals.

Results should not be used as the sole basis for diagnosis, treatment or other patient management decisions. Positive results do not exclude co-infection with other pathogens. The agent(s) detected may not be the definite cause of disease. Negative results do not exclude a C. difficile, Norovirus, Rotavirus infection or another gastrointestinal infection. Results must be clinically correlated with patient history, clinical observations and epidemiological information. Other diagnostic information is necessary to determine patient infection status. Intended for use with a Vivalytic one analyser by healthcare professionals only in laboratory settings.

Safety Information

These Instructions for Use contain test-specific information only. For additional warnings and instructions refer to the Instructions for Use provided with your Vivalytic one analyser (chapter device safety information). Only use Vivalytic cartridges and accessories approved for the Vivalytic one analyser. Take care to avoid any contamination when handling patient samples and cartridges. When sample was spilled on the cartridge, do not use the cartridge, and dispose it.

For in vitro diagnostic use by trained healthcare professional.



WARNING

- Always follow good laboratory practice to ensure the proper performance of this test.
- · Make sure to wear appropriate personal protective equipment (PPE).
- Do not use a cartridge if the sealed pouch or the cartridge itself is visibly damaged.
- · Do not touch or scratch the detection area of the cartridge.
- · Do not reuse a cartridge.
- Do not use expired cartridges. The expiration date can be found on the packaging and the cartridge label.
- packaging and the cartridge label.
 Do not wait longer than 15 minutes after opening the cartridge pouch to begin the test.
 This maintains hygiene and avoids performance loss due to humidity. Prolonged exposure to humidity has a negative impact on test performance.
- · Do not shake a cartridge that contains a sample.
- · Do not turn the cartridge upside down.
- Place the cartridge on a clean and flat surface only.
- Do not use sample types, media and volumes that are not approved for the test.
- Always follow good laboratory practice to ensure the proper performance of this test.
- Biological specimens, transfer devices, and used cartridges should be considered capable of transmitting infectious agents requiring standard precautions. Handle potentially infectious patient samples and cartridges according to national laboratory standards and dispose samples and cartridges according to regional and laboratory standards.
- · Be compliant with the national safety regulations and practices.

Note: Further information can be found in the safety data sheet (SDS) of the product. Please contact the customer support of your local distributor.

Additional Equipment & Consumables Required but not Provided

- Bosch Vivalytic one analyser (reference number F09G300115)
- Pipettor (100–1000 μl)
- Sterile filter pipette tips 100 –1000 μl
- Swab collection kits
- _ Regular Flocked Swab FLOQSwabs® 552C (COPAN Italia s.p.a.)
- Transport medium
- _ eNAT® 606C (COPAN Italia s.p.a.)
- Suitable protective clothing

Test Principle

Vivalytic Noro-, Rotavirus, C. diff is a qualitative real-time PCR based test.

Storage und Usage Conditions

Product is stable until the expiry date if stored at +15 °C to +25 °C. Storage and usage conditions can be taken either from the cartridge, pouch, or box label. Cartridge has to be used at +15 °C to +25 °C, relative humidity <65 %, within 15 min upon pouch opening. This maintains hygiene and avoids performance loss due to humidity. Prolonged exposure to humidity has a negative impact on test performance.

Reagents All reagents necessary for the sample processing are integrated into the cartridge. The processing includes cell lysis, nucleic acid extraction, DNA amplification and detection.

Reagents are PCR bead, binding buffer, washing buffer, and elution buffer. The PCR bead contains the DNA polymerase, primers, and probes. Binding buffer facilitates binding of nucleic acids during the purification process. Washing buffer is a formulation of different salts and solvents to remove impurities e. g. proteins during the extraction process. Elution Buffer is a low-salt buffer and contains the purified nucleic acids at the end of the extraction process.

Sample Type/Medium

The test is intended for use with stool swab samples in eNAT® medium (Regular Flocked Swab FLOQSwabs® 552C, eNAT® transport medium Ref. 606C, COPAN Italia s.p.a.).

Collect and store samples as indicated in the manufacturer's instructions. In case the sample is not processed immediately after sample collection, nucleic acids will be preserved in eNAT° transport medium for up to 4 weeks at room temperature and at 4°C and up to 6 months at -20° C to -80° C (see Instructions for Use eNAT° transport medium11).

Sample Preparation

Use a Regular Flocked Swab FLOQSwabs® (COPAN Italia s.p.a.) to collect a small amount of stool by inserting the tip of the flocked swab into the stool sample and rotate it. Bloody, slimy, or watery area of stools should be selected and collected. After collection, examine the swab to make sure there is fecal material visible on the tip. In case it is not, again insert the flocked swab into the stool sample and rotate taking care that all the area of the swab tip is in contact with the sample. Make sure that the swab is just covered with stool and remove excess stool by gently rotating the swab against the rim of the sample. After collection transfer the swab into the 2 ml Copan eNAT® tube with eNAT® transport medium. Hold the swab shaft between thumb and finger, mash and mix the stool specimen against the side of the tube to evenly disperse and suspend the specimen in the preservation medium. Hold the swab shaft close to the rim of the tube, bend it at a 180 degrees angle to break it off at the marked breakpoint. Discard the broken upper part of the swab shaft and tighten the cap. Shake the sample tube containing the swab sample and eNAT® medium (COPAN Italia s.p.a.) thoroughly at least for 10 seconds for homogenization. Use a pipettor in order to fill 300 µl of homogenized patient sample in the sample input of the cartridge. Ensure to only pipet from the supernatant (top of the sample) to prevent the carryover of stool particles. In case of an excess amount of particles in the sample it is recommended to place the sample tube on a flat surface and let the particles sediment for 5 minutes. Do not use viscous samples that are difficult to pipette.

Test Result

After automatic processing of the sample with the Vivalytic *one* analyser the test result shown on the screen of the Vivalytic *one* analyser. The time to result is about 58 min.

For high titer specimens results are available after less than 41 min and the test run can be terminated earlier (see Chapter Test Termination).

The sample is classified either as *C. difficile* positive, Rotavirus and *C. difficile* positive, Norovirus and *C. difficile* positive, Rotavirus positive, Norovirus and Rotavirus positive, Norovirus and Rotavirus positive, Norovirus positive, negative or invalid. In case of a positive detection of a pathogen, the test is considered valid even if the Human Control is negative.

Detection of the human cell based whole process control (Human Control) in negative samples shows a successful extraction procedure and excludes an inhibition of the PCR reaction. Interpretation of results is shown in the table below.

Norovirus genogroup I/II	Rotavirus Type A	C. difficile (tcdA / tcdB)	Human control	Validity	Result
-	-	+	+/-	valid	Sample is considered <i>C.</i> difficile positive.
-	+	+	+/-	valid	Sample is considered Rotavirus and <i>C. difficile</i> positive.

+	-	+	+/-	valid	Sample is considered Norovirus and <i>C. difficile</i> positive.
+	+	+	+/-	valid	Sample is considered Norovirus, Rotavirus and <i>C.</i> difficile positive.
-	+	ı	+/-	valid	Sample is considered Rotavirus positive.
+	+	ı	+/-	valid	Sample is considered Norovirus and Rotavirus positive.
+	-	-	+/-	valid	Sample is considered Norovirus positive.
-	-	-	+	valid	Sample is considered negative.
-	-	-	-	invalid	Not evaluable. 1

¹ Retesting is recommended

PCR - Curve and Cq Value

Real-time PCR curves (software-modified) are shown and classified as positive or negative by the software. In case of positive curves, the respective Cq value is displayed. Inconclusive results are marked by the software (\(\triangle \) \). Retesting is advised.

Invalid or Failed Tests

A test is rated as invalid if neither target DNA nor Human Control is detected. Possible reasons for an invalid run might be poor sample quality due to a partial or complete absence of human cellular material in the sample. Results are displayed for an invalid test but are not allowed to be used for diagnostic interpretation. Pay attention to use the correct sample type, sample collection and storage of the sample and cartridges prior to the test run. If required, repeat the analysis with a new sample. In case of a failed test, first check for correct operating conditions of the Vivalytic one analyser (refer to Vivalytic one analyser. If the problem persists, contact the customer support of your local distributor.

Test Termination

As soon as a valid, positive result is shown for one of the three pathogens on the screen, the user has the option to finish the test.

Test Report

In the printed test report, pathogen, results, control and information on user, patient and Vivalytic *one* analyser are listed with a signature field.

Notice to Users in EU

Any serious incident that has occurred in relation to the device, should be reported to the manufacturer and the competent authority of the Member State in which the user and/or patient is established.

Limitations

The results of the Vivalytic Noro-, Rotavirus, C.diff test must be interpreted by a trained healthcare professional only. The results of the Vivalytic Noro-, Rotavirus, C.diff test must not be used as the sole parameter for diagnosis.

- A negative result does not exclude pathogens being present in the sample at a level below assay sensitivity or other pathogens being present not covered by this assay.
- There is a risk of false negative or false positive results due to improperly collected, transported, or handled samples.
 - In borderline cases atypical PCR characteristics (e. g. flat curve with low or high Cq-value) can occur. In case of atypical characteristics results are not allowed to be used for diagnostic interpretation. Inconclusive results are marked by the

software. Retesting is advised.

- Vivalytic Noro-, Rotavirus, C.diff is a qualitative real-time PCR test and does not provide a quantitative result.
- A positive result does not necessarily mean that viable pathogens are present.
- The test detects C. difficile carrying tcdA/tcdB genes, but does not differentiate between toxin producing and non-toxin producing strains
- A negative result does not preclude a Norovirus, Rotavirus, C. difficile or coinfection. It is recommended that negative tested samples are interpreted in the context with additional laboratory data
- An excess amount of stool may have inhibitory effects on the assay performance.
- A reduced performance was observed for tcdA+/tcdB- *C. difficile* strains
- An excess amount of Rotavirus A virus particles can lead to a reduced performance of the C. difficile PCR
- No exclusivity for Rotavirus Type C (porcine Rotavirus) could be achieved.
 Therefore a misidentification of Rotavirus Type C as Type A cannot be excluded.

Analytical Performance Evaluation

Analytical Sensitivity (Limit of Detection)

The limit of detection of the Vivalytic Noro-, Rotavirus, C. diff test was determined as the lowest concentration of analyte that can be consistently detected (295% of samples tested under routine laboratory conditions using a defined type of sample). (Table 1)

Inclusivity

To evaluate inclusivity, an *in silico* analysis (BLAST alignment) of the genomic sequence of various relevant Noroviruses, Rotaviruses and *Clostridioides difficile* strains against the sequence of the PCR primers and hydrolysis probe used in the Vivalytic Noro-, Rotavirus, C. diff test for amplification and detection of the respective pathogens was performed. Inclusivity could be shown for strains listed in Table 2.

Exclusivity/ Analytical Specificity

To exclude cross-reactivity (exclusivity), an *in silico* analysis (BLAST alignment) of the target region of Norovirus, Rotavirus and *Clostridioides difficile* against the genomic sequence of various other pathogens representing common gastrointestinal pathogens or closely related species was conducted. For the detection system of Rotavirus, sequence matches in the probe and primer area could be detected for porcine Rotavirus (group C), concluding a possible amplification. There was no evidence of an interference for the detection system of Norovirus and *Clostridioides difficile*. (Table 3).

Reproducibility

The reproducibility of the Vivalytic Noro-, Rotavirus, C. diff test was established using a panel with 3 different concentrations of Norovirus, Rotavirus and Clostridioides difficile. At 3 test sites, each mix was tested on the same set of Vivalytic instruments by the same operator with 3 LOTs in 4 replicates on 3 days, respectively. The obtained positivity rates for the different combinations were correlated to the expected positivity rate (Table 4a).

Repeatability

The repeatability of the Vivalytic Noro-, Rotavirus, C. diff test was established using a panel with 1 concentration (3x c95) of Norovirus, Rotavirus and Clostridioides difficile. At 1 test site, the mix was tested on the same set of Vivalytic instruments by the same operator with 3 LOTs in 20 replicates, respectively, yielding a total of 60 observations per target pathogen. The obtained positivity rates for the different combinations were correlated to the expected positivity rate (Table 4b).

Interferences

Interferences were evaluated for endogenous and exogenous substances, that are potentially present in the patient sample. Refer to Table 5 for substances that have the potential to interfere with the test.

Clinical Performance evaluation Sensitivity and Specificity

Sensitivity and specificity results derived from native liquid and soft human stool samples. Samples were collected in a clinical setting and compared with results of reference methods.

Samples for testing with Vivalytic Noro-, Rotavirus, C. diff cartridges were freshly used or frozen for storage and prepared as describe above in eNAT® (COPAN Italia

S.p.A.). In case of reference testing samples were prepared according to recommendations of used reference methods. In total, 159 samples were analysed. Sensitivity or Positive Percent Agreement (PPA) was calculated as 100 % x TP/ (TP+FN). Specificity or Negative Percent Agreement was calculated as 100 % x TN / (TN+FP). The results of the clinical performance evaluation are shown in **Table 6, Table 7 and Table 8.**

Technical Support

If you require any support, technical help or have additional questions, please contact your local distributor or visit the Bosch Vivalytic website at www.bosch-vivalytic.com.

References

Kuijper EJ, Coignard B, Tüll P; ESCMID Study Group for Clostridium difficile; EU Member States; European Centre for Disease Prevention and Control. Emergence of Clostridium difficile-associated disease in North America and Europe. Clin Microbiol Infect. 2006 Oct;12 Suppl 6:2-18.

Čzepiel J, Dróżdż M, Pituch H, Kuijper EJ, Perucki W, Mielimonka A, Goldman S, Wultańska D, Garlicki A, Biesiada G. Clostridium difficile infection: review. Eur J Clin Microbiol Infect Dis. 2019 Jul;38(7):1211-1221. doi: 10.1007/s10096-019-03539-6. Epub 2019 Apr 3. PMID: 30945014; PMCID: PMC6570665.

³Kordus SL, Thomas AK, Lacy DB. Clostridioides difficile toxins: mechanisms of action and antitoxin therapeutics. Nat Rev Microbiol. 2022 May;20(5):285-298.

⁴Gerding DN, Johnson S, Rupnik M, Aktories K. Clostridium difficile binary toxin CDT: mechanism, epidemiology, and potential clinical importance. Gut Microbes. 2014 Jan-Feb;5(1):15-27.

⁵Lopman B, Brown D, Koopmans M: Human caliciviruses in Europe. Journal of Clinical Virology 2002; 24:137-160

⁶Oh D, Gaedicke G, Schreier E: Viral agents of acute gastroenteritis in German children: Prevalence and molecular diversity. J Med Virol 2003; 71:82-93

⁷Schneider T, Mankertz J, Jansen A., Schreier E, Zeitz M: Norovirusinfektionen – häufigste Ursache akuter Gastroenteritiden in den Wintermonaten. Deutsches Ärzteblatt 2005; 38:2551-2556

⁸Parashar UD, Hummelmann EG, Bresee JS et al.: Global illness and deaths caused by rotavirus disease in children. Emerg Infect Dis 2003; 9:565–572

⁹Mas Marques A, Diedrich S, Schreier E: Group A Rotavirus Genotypes in Germany 2005/2006. Arch Virol 2007; 152:1743–1749

10 ACHERON ELL Weiber FOF USE BY COS AN ENTARMIC Affects A ance Placer Dis 2004 4:97 version EIFU002R01 Date 2021.06.

Symbols

Manufacturer SN Serial number Date of manufacture Temperature limit Expiry date Do not use if package is damaged Lot number For single use only Reference number Consult Instructions for Use REF Contains <n> tests in vitro diagnostic medical device IVD





Pipette the indicated sample volume in the sample input of the cartridge as marked by the black triangle.

WARNING



Hazard component in cartridge

(F09G300919; binding buffer RB-BB-34a)

Guanidinium chloride; guanidine hydrochloride

Hazard statements

H302 Harmful if swallowed. H315 Causes skin irritation.

H319 Causes serious eye irritation.

Precautionary statements

Wash hands thoroughly after handling. P264 Wear protective gloves and eye/face protection. P280

IF ON SKIN: Wash with plenty of water and soap. P302+P352

IF IN EYES: Rinse cautiously with water for several minutes. Remove P305+P351+P338

contact lenses, if present and easy to do. Continue rinsing.

If eye irritation persists: Get medical advice/attention. P337+P313 P501

Dispose of contents/container in accordance with local/regional/

national/international regulation

Table 1 Limit of Detection (LoD)	
Clostridioides difficile (tcdA/tcdB)*	10,000 copies/mL
Rotavirus group A**	10,000 copies/mL
Norovirus GII**	168 700 conies/ml

*Determined by using Vircell material (surrogate strain R20291 Ribotype 027 (toxin A + B positive) for a spiking approach in eNAT® (COPAN) supplemented with approximately 1000 Human Bronchial/Tracheal Epithelial Cells (hPBTEC) and 6% (v/v) Stool Matrix Negative Control (artificial simulated matrix). **Determined by using quantified clinical isolates (Rotavirus group A, Norovirus GII) for a

spiking approach in eNAT® (COPAN) supplemented with approximately 1000 Human Bronchial/Tracheal Epithelial Cells (hPBTEC) and 6% (v/v) Stool Matrix Negative Control (artificial simulated matrix). Norovirus GI was confirmed at the 3x LoD (as determined for Norovirus GII) using artificial nucleic acid constructs.

Table 2 Inclusivity
Clostridioides difficile toxin A gene (tcdA)
Clostridioides difficile toxin B gene (tcdB)
Rotavirus group A
Norovirus GI/GII

Strains validated for inclusivity via BLAST alignment

Table 3 – Exclusivity	
Abiotrophia defective	Enterococcus faecium Enterococcu
Acinetobacter baumannii	gallinarum Enterococcus hirae Enterococcu
Acinetobacter lwoffii	raffinosus Enterovirus Escherichia blatta
Adenovirus	Escherichia coli (Shiga toxin producing)
Aeromonas caviae	(STEC)
Aeromonas hydrophila	Escherichia coli (Enteroaggregative) (EAEC)
Alcaligenes faecalis	Escherichia coli (Enteroinvasive) (EIEC)
	Escherichia coli (Enteropathogenic) (EPEC)
Anaerococcus tetradius	Escherichia coli (Enterotoxigenic) (ETEC)
Arcobacter butzleri	Escherichia coli (non-pathogenic) (K-12
Arcobacter cryaerophilus	MG1655)
Ascaris lumbricoides	Escherichia coli (Enterohemorrhagic) (EHEC)
Astrovirus	Escherichia fergusonii
	Escherichia hermannii
Bacillus cereus	Escherichia vulneris Eubacterium rectale
Bacillus subtilis	Flavonifractor plautii
Bacteroides caccae	Fusobacterium varium
Bacteroides fragilis	Gardnerella vaainalis
Bacteroides merdae	Gemella morbillorum
Bacteroides stercoris	Giardia lamblia
Bacteroides thetaiotaomicron	Giardia muris
Bifidobacterium adolescentis	Grimontia hollisae
Bifidobacterium bifidum	Hafnia alvei
Bifidobacterium longum	Helicobacter fennelliae
Blastocystis hominis	Helicobacter pylori
Campylobacter coli	Helicobacter cinaedi
Campylobacter concisus	Helicobacter hepaticus
Campylobacter curvus	Hepatitis A virus
Campylobacter fetus	Herpes Simplex Virus 2
Campylobacter gracilis	Hymenolepis nana
Campylobacter helveticus	Klebsiella oxytoca
Campylobacter hominis	Klebsiella ozaenae
Campylobacter hyointestinalis	Klebsiella pneumoniae
Campylobacter jejuni	Lactobacillus acidophilus
Campylobacter lari	Lactobacillus reuteri
Campylobacter mucosalis	Lactococcus lactis
Campylobacter rectus	Leminorella grimontii
Campylobacter showae	Listeria grayi
Campylobacter sputorum	Listeria innocua
Campylobacter upsaliensis	Listeria monocytogenes
Candida albicans	Megamonas hypermegale
Candida catenulate	
Cedecea davisae	
Cedecea davisae Chilomastix mesnili	

Table 3 – Exclusivity Citrobacter amalonaticus	A4
Citrobacter amaionaticus Citrobacter freundii	Morganella morganii Norovirus GIV
Citrobacter freundii Citrobacter koseri	Pentatrichomonas hominis
Citrobacter koseri Citrobacter sedlakii	
Clostridium beijerinckii	Peptoniphilus asaccharolyticus Peptostreptococcus anaerobius
Clostridium bifermentans	·
Clostridium bijermentans Clostridium bolteae	Plesiomonas shigelloides
Clostridium bottelle Clostridium botulinum	Porphyromonas asaccharolytica
Clostridium butyricum	Prevotella melaninogenica
Clostridium chauvoei	Proteus mirabilis
Clostridium difficile (non-toxigenic)	Proteus penneri
Clostridium fallax	Proteus vulgaris
Clostridium haemolyticum	Providencia alcalifaciens
Clostridium histolyticum	Providencia rettgeri
Clostridium innocuum	Providencia stuartii
Clostridium methylpentosum	Pseudomonas aeruginosa
* *	Pseudomonas fluorescens
Clostridium nexile	Pseudomonas putida
Clostridium novyi	Rotavirus B
Clostridium orbiscindens	Rotavirus C
Clostridium paraputrificum	Rotavirus D
Clostridium perfringens	Rotavirus F
Clostridium ramosum	Rotavirus G
Clostridium scindens	Rotavirus H
Clostridium septicum	Rotavirus I
Clostridium sordellii	Ruminococcus bromii
Clostridium sphenoides	Saccharomyces boulardii
Clostridium spiroforme	Salmonella enterica
Clostridium sporogenes	Salmonella enterica spp. enterica serovar Typhimurium
	Salmonella enteritidis
Clostridium symbiosum	Salmonella subterranea
Clostridium tertium	Sapovirus
Clostridium tetani	Serratia liquefaciens
Collinsella aerofaciens	Serratia marcescens
Corynebacterium genitalium	Shigella boydii
Coxsackie virus	Shigella dysenteriae
Cryptosporidium canis	Shigella flexneri
Cryptosporidium cuniculus	Shigella sonnei
Cryptosporidium felis	Staphylococcus aureus
Cryptosporidium fetus	Staphylococcus aureus subsp. aureus
Cryptosporidium hominis	Staphylococcus epidermidis
Cryptosporidium meleagridis	Stenotrophomonas maltophilia
Cryptosporidium muris	Streptococcus agalactiae
Cryptosporidium parvum	Streptococcus dysgalactiae
Cytomegalovirus	Streptococcus dysgalactiae subsp.
Desulfovibrio piger	dysgalactiae
	Streptococcus intermedius
Diantamooka frasilis	Streptococcus pyogenes
Dientamoeba fragilis	Streptococcus salivarius
Diphyllobothrium latum	Streptococcus uberis
	Taenia saginata
Echovirus	Trabulsiella guamensis
Echovirus Edwardsiella tarda	-
Echovirus Edwardsiella tarda Eggerthella lenta	Trabulsiella guamensis
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis	Trabulsiella guamensis Veillonella parvula
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana	Trabulsiella guamensis Veillonella parvula Vibrio cholerae
Echovirus Edwardsiella tarda Eggerthella lenta Encephalliczoon intestinalis Endolimax nana Entamoeba coli	Trabulsiella guamensis Veillonella parvula Vibrio cholerae Vibrio mimicus
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni	Trabulsiella guamensis Veillonella parvula Vibrio cholerae Vibrio mimicus Vibrio parahaemolyticus
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni Entamoeba histolytica	Trabulsiella guamensis Veillonella parvula Vibrio cholerae Vibrio mimicus Vibrio parahaemolyticus Vibrio vulnificus
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni Entamoeba histolytica Entamoeba moshkovskii	Trabulsiella guamensis Veillonella parvula Vibrio cholerae Vibrio-parahaemolyticus Vibrio vulnificus Yersinia bercovieri Yersinia enterocolitica
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni Entamoeba histolytica Entamoeba moshkovskii Entamoeba polecki	Trabulsiella guamensis Veillonella parvula Vibrio eholerae Vibrio mimicus Vibrio parahaemolyticus Vibrio vulnificus Yersinia bercovieri Yersinia enterocolitica Yersinia rheterocolitica Yersinia frederiksenii
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni Entamoeba histolytica Entamoeba moshkovskii Entamoeba polecki Enterobacter aerogenes	Trabulsiella guamensis Veillonella parvula Vibrio-cholerae Vibrio-mimicus Vibrio parahaemolyticus Vibrio vulnificus Yersinia bercovieri Yersinia enterocolitica Yersinia enterocolitica Yersinia frederiksenii Yersinia intermedia
Dysgonomonas capnocytophagoides Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni Entamoeba histolytica Entamoeba moshkovskii Entamoeba polecki Enterobacter aerogenes Enterobacter cloacae	Trabulsiella guamensis Veillonella parvula Vibrio cholerae Vibrio parahaemolyticus Vibrio vulnificus Yersinia bercovieri Yersinia enterocolitica Yersinia enterocolitica subsp. enterocolitica Yersinia intermedia Yersinia mollaretii
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni Entamoeba histolytica Entamoeba moshkovskii Entamoeba polecki Enterobacter aerogenes Enterobacter cloacae Enterococcus casseliflavus	Trabulsiella guamensis Veillonella-parvula Vibrio-cholerae Vibrio-parahaemolyticus Vibrio parahaemolyticus Vibrio vulnificus Yersinia bercovieri Yersinia enterocolitica Yersinia enterocolitica subsp. enterocolitica Yersinia intermedia Yersinia mollaretii Yersinia pseudotuberculosis
Echovirus Edwardsiella tarda Eggerthella lenta Encephalitozoon intestinalis Endolimax nana Entamoeba coli Entamoeba hartmanni Entamoeba histolytica Entamoeba moshkovskii Entamoeba polecki Enterobacter cerogenes Enterobacter cloacae	Trabulsiella guamensis Veillonella parvula Vibrio cholerae Vibrio parahaemolyticus Vibrio vulnificus Yersinia bercovieri Yersinia enterocolitica Yersinia enterocolitica subsp. enterocolitica Yersinia intermedia Yersinia mollaretii

Strains validated for exclusivity via BLAST alignment

Table 4a Reproducibility							
Target	С	No of total tests	No of pos. tests	No of neg. tests	Proportion of positive/ negative result _s (%)	95 % Wilson- Score confidence interval (%)	95 % Pearson- Clopper confidence interval (%)
none	blank	108	0	108	100	96.6% - 100%	96.6% - 100%
Norovir us GII	3x c95	108	108	0	100	96.6% - 100%	96.6% - 100%
	<c100< td=""><td>108</td><td>108</td><td>0</td><td>100</td><td>96.6% - 100%</td><td>96.6% - 100%</td></c100<>	108	108	0	100	96.6% - 100%	96.6% - 100%
Rotavir us A	3x c95	108	108	0	100	96.6% - 100%	96.6% - 100%
	c100	108	108	0	100	96.6% - 100%	96.6% - 100%
Clostridi oides	3x c95	108	106	2	98.1	93.5% - 100%	93.5% - 99.8%
difficile	<c100< td=""><td>108</td><td>107</td><td>1</td><td>99.1</td><td>94.9% - 100%</td><td>95.0% - 100%</td></c100<>	108	107	1	99.1	94.9% - 100%	95.0% - 100%

³x c95 = 95 % predicted positive agreement

c200 = 100 % predicted positive agreement (Norovirus GII: verified with 3,33x106 copies/mL, c100 = 3,8x107 copies/mL; Clostridioides difficile: verified with 1,83x105 copies/ml; c100 = 5,48x105 copies/ml)

Table 4b	Table 4b Repeatability							
Target	LOT	No of total tests	No of pos. tests (3x <i>c95</i>)	Proportion of positive/ negative results (%)	95 % Wilson- Score confidence interval (%)	95 % Pearson- Clopper confidence interval (%)		
Norovir us GII	1	20	20	100	83.9% - 100%	83.2% - 100%		
	2	20	20	100	83.9% - 100%	83.2% - 100%		
	3	20	20	100	83.9% - 100%	83.2% - 100%		
	Total	60	60	100	94% – 100%	94% – 100%		
Rotavir us A	1	20	20	100	83.9% - 100%	83.2% - 100%		
	2	20	20	100	83.9% - 100%	83.2% - 100%		
	3	20	20	100	83.9% - 100%	83.2% - 100%		
	Total	60	60	100	94% – 100%	94% – 100%		
Clostrid ioides	1	20	20	100	83.9% - 100%	83.2% - 100%		
difficile	2	20	20	100	83.9% - 100%	83.2% - 100%		
	3	20	20	100	83.9% - 100%	83.2% - 100%		
211 205	Total	60	60 tive agreem	100	94% – 100%	94% – 100%		

³x c95 = 95 % predicted positive agreement

Table 5 – Tested Substances for Interference

No interference detected

c100 = 100 % predicted positive agreement

Cholesterol; 0.5 % (v/v)

Mucus; 3 mg/mL

Palmitic acid; 2mg/mL

Stearic Acid; 2mg/mL

Triglyceride; 0.8 % (w/v)

Whole blood; 3.0 (v/v)

Barium Sulfate; 1.3 % (w/v)

GLYICILAX for adults; 1.0 % (w/v)

GRÜNWALDER Sennalax film-coated tablet; 0.1 mg/mL

Hemorrhoid ointment with witch hazel; 1 % (v/v)

LOPERAMID-ratiopharm acute 2 mg film-coated tablet; 0.2 % (v/v)

MAALOXAN 25 mVal Liquid; 0.1 mg/mL

METRONIDAZOL Aristo 400 mg tablet; 0.5 % (w/v)

Matrix Negative Control (artificial simulated matrix).

Naproxen axicur® tablet; 0.5 % (w/v)

Nonoxynol-9; 1 % (w/v)

Nystatin STADA®; 1 % (w/v)

Postericort ointment; 0.5 % (v/v)

Claversal 4 g/60 ml clysms; 0.5 % (v/v) Vancomycin; 1.4 mg/mL

Vaseline; 1 % (w/v)

Interference was experimentally verified at 3xLoD for Norovirus GII, Rotavirus A and C. difficile using a spiking approach in eNAT® (COPAN) plus 1000 hPBTEC and 6% (y/y) Stool

Table 6 – Norovirus: Clinical Sensitivity (PPA) [1] and Specificity (NPA) [2] for

[1] 94.23% (84.05 – 98.79%)*

[2] 100.00% (96.48 - 100%)

Table 7 – **Rotavirus**: Clinical Sensitivity (PPA) [3] and Specificity (NPA) [4] for samples in eNAT® (95 % confidence interval clinical study)

[3] 92.59% (81.11 - 97.94%)*

[4] 97.03% (91.56 - 99.38%)**

 $\label{lem:conditional} Table 8 - \textbf{C. difficile} : Clinical Sensitivity (PPA) [5] and Specificity (NPA) [6] for samples in eNAT® (95 % confidence interval, clinical study)$

[5] 97.96% (89.15 - 99.95%)*

[6] 98.18% (93.59 - 99.78%)**

In total, 159 clinical samples were tested within the scope of the clinical study. In this data set, 5: samples were found positive for Rorovirus, 54 samples were found positive for Rotavirus and 49 samples were found negative for all pathogens. All samples were tested with the two reference tests from Seegene (Allplex™ GI-Bacteria(I) and Allplex™ GI-Virus Assay) and Vivalytic Noro-, Rotavirus, C. diff to determine the clinical performance for Vivalytic. Discrepant results were verified using RIDAGENE RIDA®GENE HSP (R-Biopharm) (12 samples).

^{*3} samples were detected as false negative for Norovirus.

^{*4} samples were detected as false negative for Rotavirus.

^{**3} samples were detected as false positive for Rotavirus.

^{*1} sample was detected as false negative for C. difficile.

^{**2} samples were detected as false positive for *C. difficile*.

Vivalytic Noro-, Rotavirus, C. diff - Annex

Revision 01	Initial document
Revision 02	Pre-release changes

For more information see www.bosch-vivalytic.com







