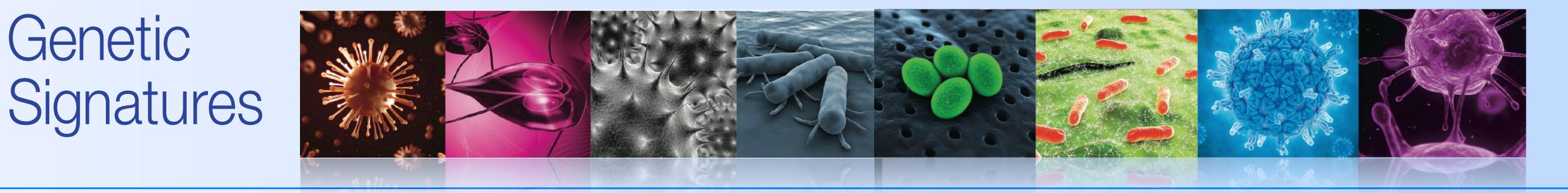
Multiplex PCR detection of all major gastrointestinal pathogens employing a novel universal extraction method J.R. Melki*, S.P. Siah, K. Kaur, J. Nair, N. Coulston, D.S. Millar (Sydney, AU)



Abstract

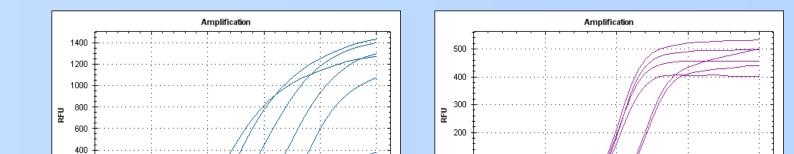
Materials & Methods

Results

Figure 3. Amplification plots obtained using the *EasyScreen*[™] Viral assay.

Genetic

Signatures



Gastrointestinal disease (GI) is a major

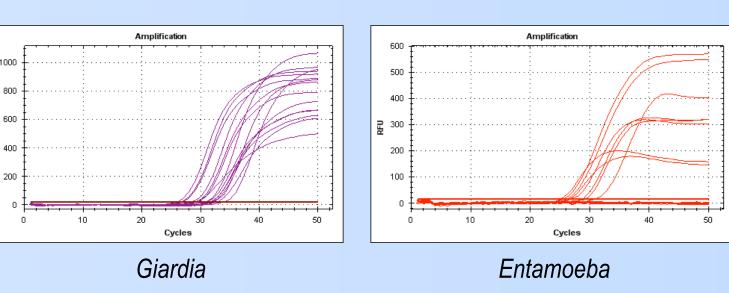
A universal sample processing method was devised that lysed and simultaneously converted the nucleic acids of bacteria, viruses and protozoan parasites from the primary patient sample. The sample buffer protects the labile RNA species from the harsh conditions required for complete lysis of tough organisms such as Giardia. The procedure comprises a single tube method during which the faecal sample is incubated in extraction buffer for 15 minutes. Samples were then purified using a column based method or automated platforms such as Roche, Qiagen and Thermo. PCR can then be carried out on most real-time instruments including those from Roche, Qiagen, Cepheid, ABI, Biorad and Stratagene. All reagents required from sample to result are included simplifying the method for the end user.

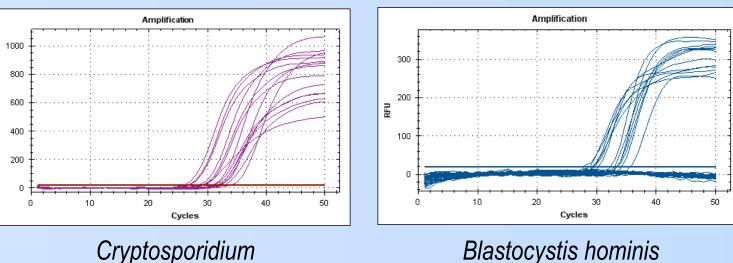
All assays were linear from 10⁶-10 copies and no cross reactivity was observed between individual primers and a larger number of bacterial and fungal non-target species. Over 500 clinical samples have been assessed and compared to conventional techniques such as culture, EIA and microscopy with excellent concordance. The method developed here is therefore suitable to the rapid and sensitive screening of primary patient material for a wide range of common Gl pathogens.

cause of morbidity and mortality worldwide. GI can be caused by a wide range of infectious agents including viral, bacterial and protozoa. Human viral gastroenteritis can be caused Noroviruses, Rotaviruses, Adenoviruses, Astroviruses and Sapovirus. Of these Norovirus is the most commonly isolated agent as the cause of acute viral gastroenteritis (1). According to the CDC 43% of bacterial GI infections are caused by Salmonella, followed by Campylobacter (33%), Shigella (17%), Escherichia coli (4.1%) and Yersinia (0.9%). Another cause of bacterial GI are hypervirulent strains of Clostridium difficile particularly PCR ribotype 027 (2). Among parasites Giardia intestinalis, Cryptosporidium spp and Entamoeba histolytica are considered the most common and important causes of diarrhea (3) although other species such as Dientamoeba fragilis and Blastocystis hominis have also been implicated in GI disease. Thus the diagnosis of GI can be

Figure 1. Amplification plots obtained using the *C. difficile EasyScreen*TM detection kit.

Figure 2. Amplification plots obtained using the *EasyScreen*[™] Parasite assay.





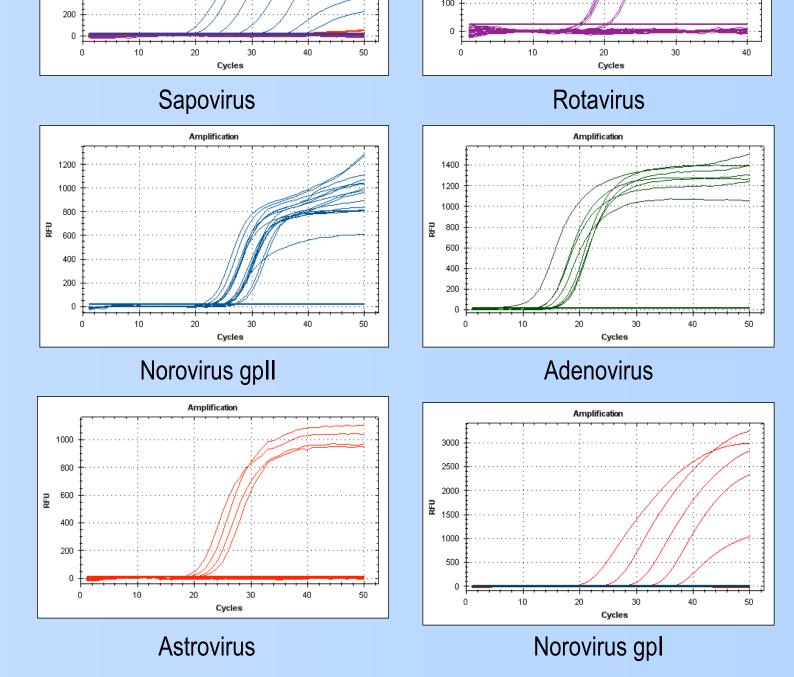


Table 4. Results obtained using the *EasyScreen*[™] Viral assay.

	EIA positive	EasyScreen Viral positive
Norovirus	81	81 (All group II)
Rotavirus	*21	17
Adenovirus	2	2
Astrovirus	5	6
Sapovirus	N/A	0

*The Rotavirus EIA cross reacts with the vaccine strain

Table 5. Results obtained using the *EasyScreen*[™] Bacterial assay.

challenging and involve specialists in microbiology, virology and parasitology.

Objective

In order to simplify the detection of causative agents of GI we have developed rapid real time multiplex PCR (mPCR) panels for all major GI pathogens (see Table 1). All assays share a universal sample processing method and incorporate our previously described 3base[™] technology (4). Furthermore, conventional GI diagnosis can in some instances take up to 5 days (5) to provide a definitive result. To reduce this time we aimed to produce assays with sample to result turnaround time in as little as 3 hours.

Table 2. Three independent Clinical studies using the *EasyScreen*TM *C. diffiicle* assay.

Cycles

	Study #1(n=70)
C. difficile Culture ⁺	70 (100%)
EIA [*]	43 (61.4%)
EasyScreen C. difficile*	53 (75.7%)

2

600

	(n=73)	
	Positive	%
C. difficile Culture ⁺	63/72	88
EIA [*]	31/72	43
Illumigene [*]	56/73	76.7
Quik Chek Complete [*]	35/52	67.3
EasyScreen C. difficile*	56/73	76.7

	Study #3 (n=74)	
Culture Toxin +ve	61/74 (82.4%)	
EasyScreen +ve ¹	65/74 (87.8%)	
Illumigene +ve ²	65/74 (87.8%)	
Ausdiagnostic +ve ³	36/69 (52.2%)	
Faecal Toxin +ve ⁴	33/74 (44.6%)	
¹ 61/61 CT positive samples (100%)		
² 59/61 CT positive samples (96.7%)		
³ 36/58 CT positive samples (62.1%)		
⁴ 33/74CT positive samples (45.6%)		

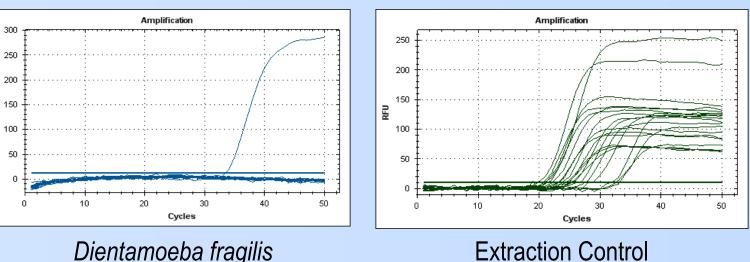


Table 3. Results obtained using the *EasyScreen*[™] Parasite assay.

	Faecal Samples (n=81)		Purified DNA (n=50)	
	Microscopy	EasyScreen	Previous PCR	EasyScreen
Giardia intestinalis	33	37	15	15
Cryptosporidium spp	15	15	15	15
Dientamoeba fragilis	12	13	12	9
Entamoeba complex	N/A	7	8	8
Entamoeba histolytica	0	0	2	2
Blastocystis hominis	15	20*	0	1
No Pathogen Identified	11	6	N/A	N/A
Mixed Infections	7	23	N/A	5

Using the *EasyScreen*TM GI panels we consistently achieved better results that conventional techniques such as culture, microscopy and EIA. The *EasyScreen*TM *C.difficile* detection kit demonstrated 100% concordance when tested against the "Gold-Standard" of Culture Toxin (see Table 2). In addition the method also showed improved performance when tested against two independent molecular assays. The *EasyScreen*TM Parasite assay also demonstrate improved detection when compared to Microscopy and EIA (see Table 3).

	Cartaro	24090010011
Salmonella	32	31
Campylobacter	40	41
Shigella	1	1
Clostridium difficile	4	4
Yersinia entercolitica	0	0
Listeria monocytogenes	0	0
No Pathogen Identified	3	2

Discussion

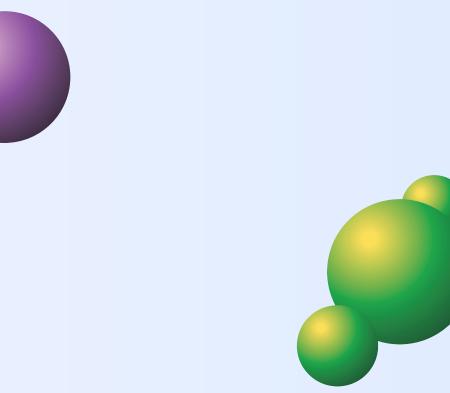
The assays developed here may be used as a complete screening system for the diagnosis of all major GI pathogens from primary clinical samples. The assays are simple and employ universal sample preparation conditions thereby streamlining the process of pathogen detection from faecal material. All assays have incorporated controls for sample processing and inhibition to ensure assay robustness and reliability. Assays can be run on virtually all purification and real time instruments found in major hospital and pathology laboratories.

Sample to results time is less than 3

hours, allowing for rapid diagnosis

References

facilitating optimal patient management.

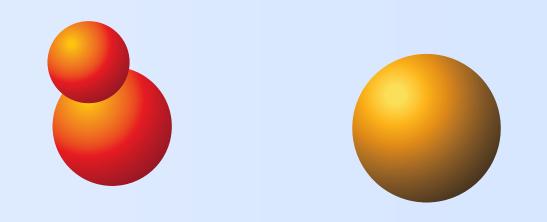


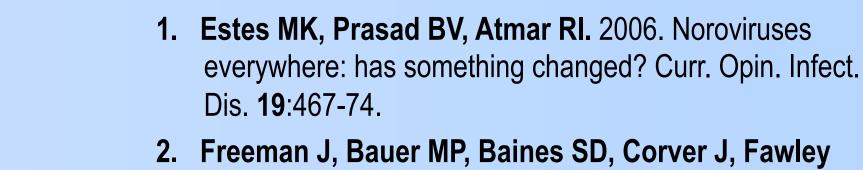
*Not toxigenic culture

Table 1. Targets detected by the EasyScreen[™] GI Panels.

Product #	Description	Microorganisms Detected
CDD001	EasyScreen™ C. difficile Detection Kit	Toxigeneic C. difficile (targets both tcdA and tcdB)
CDD002	<i>EasyScreen</i> ™ C. difficile Reflex Kit	Hypervirulent <i>C. difficile</i> incl. 027 & 078 (Targets (i) <i>tcdC</i> gene deletion at position 117 (ii) binary toxin gene (iii) <i>gyrA</i> gene mutation (fluroquinolone resistance))
EB001	EasyScreen™ Enteric Bacteria Detection Kit	(i) Salmonella spp. (ii) Campylobacter spp. (iii) Shigella spp. (iv) Yersinia sp., (v) toxigenic C. difficile (vi) Listeria monocytogenes
EP001	EasyScreen™ Enteric Parasite Detection Kit	(i) Cryptosporidium spp., (ii) Giardia intestinalis, (iii) Dientamoeba fragilis, (iv) Entamoeba Complex and (v) Blastocystis hominis
EV001	EasyScreen™ Enteric Viral Detection Kit	i) Norovirus I, (ii) Norovirus II, (iii) Astrovirus, (iv) Rotavirus, (v) Sapovirus and (vi) Adenovirus
EP002	EasyScreen™ Enteric Microsporidia Detection Kit*	(i) Enterocytozoon bieneusi (ii) Encephalitozoon intestinalis
EC001	EasyScreen [™] Pathogenic E. coli Detection Kit*	(i) Enterohaemorrhagic <i>E. coli</i> (including O104 strain) (ii) Enterotoxigenic <i>E. coli</i> (iii) Enteroinvasive <i>E. coli</i> (iv) Enteropathogenic <i>E. coli</i> (v) Enteroaggregative <i>E. coli</i>

*Coming soon





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