A novel method for rapidly diagnosing the causes of diarrhoea

C S J Probert, P R H Jones, N M Ratcliffe

Diarrhoea due to infection is a major cause of morbidity and mortality. In England and Wales, there are approximately 15 000 cases of Clostridium difficile infection per annum (www.PHLS.co.uk). It is estimated that in an average sized district general hospital, 100 cases of C difficile infection can be expected each year with an extra annual cost of £400 000 and 2100 lost bed days.1 This is equivalent to £60M for England and Wales as a whole. In the USA, rotavirus infection accounts for 48% of viral gastroenteritis and 56 000 hospitalisations per year,2 while in developing countries 600 000 children die from rotavirus each year.3

Despite the development of enzyme linked immunosorbent assays (ELISAs) and molecular techniques to determine the aetiology of infectious diarrhoea, in most cases there is a delay of several days between the collection of a stool sample and a microbiological diagnosis. Indeed, a delay of 8.2 days has been reported recently.4 The delay is, in part, due to shipping of the samples to an appropriate laboratory but mainly due to the time required to complete the diagnostic techniques in common use. The limited availability of expensive microbiological techniques in developing countries compounds the problem. In some epidemics, stool samples have been flown to the USA for microbiological processing. Depending on the patient’s circumstances, the results of a diagnostic delay is likely to range from lost productivity to prolonged hospitalisation or death. There is a pressing need to make a rapid accurate diagnosis, the volatile organic compounds associated with diarrhoea were analysed.

Methods: Stool samples were collected from 35 patients with infectious diarrhoea and from six healthy controls. Gaseous compounds were extracted from a headspace using solid phase microextraction and analysed using gas chromatography and mass spectroscopy.

Results: Characteristic patterns of volatile gases were found for the main causes of infectious diarrhoea in hospitals. Furan species without indoles indicated Clostridium difficile, ethyl dodecanoate indicated rotavirus, ammonia without ethyl dodecanoate suggested other enteric viruses, and the absence of hydrocarbons and terpenes indicated Campylobacter infection. Conclusion: These results could be the basis of rapid near patient diagnosis of infectious diarrhoea.

Materials and methods

Patients

Stool samples were obtained from 38 subjects with diarrhoea and from six healthy controls. For 35 subjects, the aetiology of the diarrhoea had been previously determined by routine microbiological assessment. The samples were a mixture of fresh stool and frozen archived samples. The cause of diarrhoea was known for each but no other clinical data were recorded. Of these samples, diarrhoea was due to C difficile (n = 6), Campylobacter (n = 5), rotavirus (n = 5), small rounded structured virus (SRSV, n = 9), adenovirus (n = 5), astrovirus (n = 2), and giardiasis (n = 3).

Three of the subjects had a clinically diagnosed exacerbation of inflammatory bowel disease involving the colon without microbiological assessment. No specimens had been sent for microbiological testing although none had received antibotics and each responded to treatment for the relapse.

The healthy controls (aged 40–65 years; three males) each provided a stool sample the form of which was normal for them; none had gastrointestinal symptoms and none had previously been diagnosed with a chronic gastrointestinal disorder.

In addition, samples were obtained from nine subjects (aged 0.5–49 years; two females) who developed diarrhoea following bone marrow transplantation (BMT). Each had received prophylactic broad spectrum antibiotics. Full microbiological assessment for common bacterial and viral causes for diarrhoea were performed as well as ELISA for C difficile toxin. No bacterial pathogens were identified. These samples were used as controls to compare with those from patients with C difficile. However, as the aetiology of post BMT diarrhoea is often unexplained, these samples were excluded from the remainder of the study and not used in the generation of summary statistics.

Analysis

Stool samples (approximately 0.75 g) were portioned into 10 ml sealed headspace vials (Supelco) designed for solid phase microextraction (SPME). Gas within the headspace

Abbreviations: ELISA, enzyme linked immunosorbent assay; VOCs, volatile organic compounds; SRSV, small rounded structured virus; BMT, bone marrow transplantation; SPME, solid phase microextraction; GC, gas chromatography; MS, mass spectroscopy; PPV, positive predictive value; NPV, negative predictive value

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was then sampled and analysed using gas chromatography (GC) and mass spectroscopy (MS). The technique used was based on that which we have employed to identify diseased vegetables.6,7

Samples were equilibrated in a water bath (37°C) and their volatile compounds adsorbed from the headspace onto the SPME fibre (Supelco, 85 µm carboxen/polydimethylsiloxane). Any samples not extracted within one hour of collection were stored in vials at –18°C. Sample volatile compounds were desorbed from the fibre thermally via splitless injection (HP 5890 II, 1.5 minutes, 250°C) onto a thick film column (Supelco, SPB-1 Sulphur, 30 m x 0.32 mm, β 20). Volatile compounds were eluted from the column (35°C (5 minutes) x 10°C/minute – 250°C (2 minutes), 1 ml/min) directly into a quadrupole mass spectrometer (HP 5971) full scan (positive mode, 15–60 amu. 2 minutes, 19–350 amu until 28.5 minutes). Compounds were identified by spectral analysis undertaken using proprietary Hewlett Packard software (G1034C, V1.05) running the NBS75K library, and also by their elution order.

Statistics
The proportion of samples, from each group of patients, that exhibited a particular characteristic were compared with the proportion of the remainder exhibiting the same characteristic using the χ2 test with Yates’ correction. Positive and negative predictive values, and sensitivity and specificity were also calculated based on these proportions.

RESULTS
VOCs were successfully adsorbed onto short 1 cm fibres using a recently developed SPME technique. Thermal desorption and analysis were undertaken via a small bench top GC-MS system. The composition of VOCs of stools from normal subjects were remarkably similar (fig 1A shows a typical example) but among patients with various types of diarrhoea, the distribution of VOCs was consistently different. Characteristic changes in the volatile pattern occurred with specific types of diarrhoea (fig 1B shows a typical example from a patient with C difficile).

Normal subjects
Analyses of the volatiles emitted from the stool of normal subjects (n = 6) showed that the predominant class of volatile was the phenols, which accounted for 20–50% of the total volatiles (fig 2), 4-methylphenol being by far the most prevalent. Indole and 3-methylindole were also abundant (5–25%) in normal subjects, as were a group of volatiles classified, without detailed differentiation, as terpenes and hydrocarbons (10–20%). A range of organic acids were also present but acids as a class of compounds were present in all the samples and therefore cannot be considered as marker compounds for diarrhoea. The principle volatile components of normal flatus were thus 4-methylphenol, indole, 3-methylindole, and limonene. Benzaldehyde was ubiquitous although less abundant.

Indoles were present in most subjects with diarrhoea (fig 2). Their occurrence is related to the metabolism of tryptophan by Escherichia coli. Indeed, a simple laboratory test for E coli based on the detection of indole has been in clinical use since 1963.7

Clostridium difficile
In flatus from patients (n = 6) with diarrhoea due to C difficile, furans (principally 2-furancarboxaldehyde and 5-methyl-2-furancarboxaldehyde) were particularly prevalent (25–55% of the measured volatiles) (fig 2). 3-Methylindole was present in a minority of patients with C difficile (1/6). Stool samples from patients with diarrhoea following BMT
were also unlikely to contain 3-methylindole (0/9). High levels of 5-methyl-2-furancarboxaldehyde were found in 5/6 *C difficile* patients but only in one of the remaining 38 samples of patients with infectious diarrhoea and exacerbation of inflammatory bowel disease. Used alone the presence of 5-methyl-2-furancarboxaldehyde was significantly associated with *C difficile* infection (\( \chi^2 = 22.2, p = 0.000002 \)). Furthermore, 5-methyl-2-furancarboxaldehyde without measurable 3-methylindole occurred exclusively among patients with *C difficile* (\( \chi^2 = 20.4, p = 0.000002 \)). This combination had a positive predictive value (PPV) of 100% but two cases of *C difficile* would have been overlooked, lowering the sensitivity to 66% and the negative predictive value (NPV) to 95%. In contrast, 2-furancarboxaldehyde without measurable 3-methylindole was found in 5/6 cases with two “false positive” cases among the remainder (\( \chi^2 = 22.2, p = 0.000003 \), sensitivity = 83%, specificity = 97%, PPV 83%, and NPV 97%). All of these associations and their respective sensitivity and specificities, and PPV and NPV are summarised in table 1.

None of the nine samples of diarrhoea following antibiotics associated with BMT contained furans however 3-methyl indole was suppressed. These samples were not used for statistical analysis although to have done so would have strengthened the argument regarding *C difficile*.

**Rotavirus**

It is a striking observation that ethyl dodecanoate was ubiquitous in the VOCs of stools from patients with rotavirus (n = 5). Only one other sample, from a patient with enteritis due to adenovirus infection, contained this and we cannot exclude the possibility of dual pathology. It is also of interest that dodecanoic acid was detected in four of five of the rotavirus samples and absent from all but one of the remaining 39 samples analysed. Propyl dodecanoate was not detected in any of the samples analysed except for one patient with rotavirus. Thus ethyl dodecanoate was a very sensitive (100%) and specific (97%) marker for rotavirus (table 1). Ethanol was also ubiquitous with rotavirus but it was also present in the flatus from two of the normal subjects and six patients with diarrhoea. The presence of ethanol in many beverages precludes its use as a marker compound, except in specific circumstances.

**Norwalk virus and other enteric infections**

Patients with diarrhoea due to SRSV2 (often called the Norwalk virus) (n = 9) were studied. Ammonia was found in seven of the samples but it is not known whether all nine subjects had the same strain of virus. Furthermore, one sample yielded very few volatiles and was the smallest volume of stool in the study. Ammonia was also found in both subjects with astrovirus, two of five with adenovirus, one of three with giardiasis, and one of five with rotavirus. Therefore, ammonia on its own cannot be used to diagnose the Norwalk virus although we have never detected it among patients with a bacterial infection and so it may have a role in the diagnosis of viral enteritis. Indeed, ammonia was found in 12/19 cases of non-rotavirus enteritis and giardiasis compared with just one, a rotavirus case, of the remaining 23 samples (\( \chi^2 = 13.4, p<0.0001 \)). Ammonia is a difficult gas to detect and volatility depends to a high degree on the pH of the sample. It may be that some sample preconditioning is required for an ammonia assay to be of significant diagnostic value.

Ammonia without ethyl dodecanoate was found in 11/19 cases of non-rotavirus enteritis but not in any other sample (\( \chi^2 = 16.3, p = 0.00005 \)). This combination has 100% specificity and 58% sensitivity for non-rotavirus viral enteritis.

**Campylobacter jejuni**

Among patients with *Campylobacter* (n = 5), phenols were abundant (10–40% of measured volatiles) as were indoles (20–30%). There were also more volatile organic acids (30–35% of measured volatiles) than in any other group of patients. No VOCs belonging to the terpenes/hydrocarbon groups were found. In contrast, all other samples except three contained terpenes/hydrocarbons (\( \chi^2 = 19.5, p = 0.00001 \)). This had 100% sensitivity and 92% specificity with an NPV of 100%.

**DISCUSSION**

We have combined the recently developed advances in solid phase microextraction with the novel approach of volatile profile analyses from stool samples in order to determine the cause of infectious diarrhoea. Despite the complexity of the VOCs in normal stool, we have found changes in specific diseases.

Furans were prevalent among the VOCs from patients with *C difficile* infection but furans were not found among the VOCs from normal controls nor in stool samples from the majority of subjects with other forms of infectious diarrhoea. Furans are likely to be related to bacterial metabolism of furanose from dietary fructose. Many *Clostridia* species are capable of fermenting sugars and amino acids. Efforts have been made to rapidly identify *C difficile* in cultures by the presence of p-cresol and caproic acid.** However, this

### Table 1 Diagnostic value of volatile organic compounds (VOCs) from stool samples from patients with infectious diarrhoea

<table>
<thead>
<tr>
<th>Marker</th>
<th>Present in specific disease</th>
<th>Present in remainder</th>
<th>( \chi^2 )</th>
<th>p Value</th>
<th>Sensitivity (%)</th>
<th>Specificity (%)</th>
<th>PPV (%)</th>
<th>NPV (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>C difficile</em> (i)</td>
<td>3MI negative</td>
<td>1/6</td>
<td>3.3</td>
<td>0.06</td>
<td>83</td>
<td>65</td>
<td>28</td>
<td>96</td>
</tr>
<tr>
<td><em>C difficile</em> (ii)</td>
<td>2FC positive</td>
<td>6/6</td>
<td>11.5</td>
<td>0.00007</td>
<td>100</td>
<td>80</td>
<td>40</td>
<td>100</td>
</tr>
<tr>
<td><em>C difficile</em> (iii)</td>
<td>3MI negative, 2FC positive</td>
<td>5/6</td>
<td>22.2</td>
<td>2.6E–6</td>
<td>83</td>
<td>97</td>
<td>83</td>
<td>97</td>
</tr>
<tr>
<td><em>C difficile</em> (iv)</td>
<td>3MI negative, 5M2FC negative</td>
<td>5/5</td>
<td>22.2</td>
<td>1.3</td>
<td>83</td>
<td>97</td>
<td>83</td>
<td>97</td>
</tr>
<tr>
<td><em>C difficile</em> (v)</td>
<td>3MI negative, 2FC negative</td>
<td>4/4</td>
<td>20.4</td>
<td>4.3E–6</td>
<td>83</td>
<td>97</td>
<td>83</td>
<td>97</td>
</tr>
<tr>
<td>Campylobacter</td>
<td>No terpenes/hydrocarbons</td>
<td>5/5</td>
<td>19.6</td>
<td>1E–5</td>
<td>100</td>
<td>92</td>
<td>63</td>
<td>100</td>
</tr>
<tr>
<td>Rotavirus</td>
<td>EDD positive</td>
<td>5/5</td>
<td>13.9</td>
<td>2E–7</td>
<td>100</td>
<td>97</td>
<td>83</td>
<td>100</td>
</tr>
<tr>
<td>Non-rotavirus</td>
<td>Ammonia positive, EDD negative</td>
<td>11/19</td>
<td>15.4</td>
<td>8.6E–5</td>
<td>63</td>
<td>96</td>
<td>92</td>
<td>77</td>
</tr>
</tbody>
</table>

The strength of association is shown using the \( \chi^2 \) test with Yates’ correction, sensitivity and specificity, and positive predictive (PPV) and negative predictive (NPV) values.

3MI, 3-methylindole; 2FC, 2-furancarboxaldehyde; 5M2FC, 5-methyl-2-furancarboxaldehyde; EDD, ethyl dodecanoate.
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C S J Probert, Department of Medicine, Bristol Royal Infirmary, Bristol, UK

P R H Jones, N M Ratcliffe, Faculty of Applied Sciences, Centre for Research in Analytical, Materials, and Sensor Sciences, University of the West of England, Coldharbour Lane, Bristol BS16 1QY, UK

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Authors’ affiliations

C S J Probert, Department of Medicine, Bristol Royal Infirmary, Bristol, UK

P R H Jones, N M Ratcliffe, Faculty of Applied Sciences, Centre for Research in Analytical, Materials, and Sensor Sciences, University of the West of England, Coldharbour Lane, Bristol BS16 1QY, UK

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