

Faecal sterols analysis for the identification of human faecal pollution in a non-sewered catchment

D. Sullivan*, P. Brooks*, N. Tindale*, S. Chapman**, W. Ahmed* ***

* Faculty of Science, Health and Education, University of the Sunshine Coast, Maroochydore DC 4558, Queensland, Australia. (E-mail: PBrooks@usc.edu.au)

** SEQ catchments and Faculty of Science, Health and Education, University of the Sunshine Coast, Maroochydore DC 4558, Queensland, Australia.

*** Department of Environment and Resource Management, Queensland 4068, Australia.

Abstract

In this study, faecal sterols were used to identify human faecal pollution in a non-sewered catchment in Southeast Queensland, Australia. In all, 36 water samples were collected from six sites on six occasions and the concentration of sterols were determined using gas chromatography and mass spectrometry. The stanols concentration in water samples generally increased with increased catchment runoff. After moderate rainfall, high coprostanols levels found in water samples indicated human faecal pollution via defective septic systems. In contrast, it appears that during dry weather human faecal pollution is not occurring in the study catchment. Sterol profiles also pointed to a cattle farm polluting during modest catchment runoff. The method used in this study was able to identify the sources of faecal pollution to the catchment due to rainfall.

Key words

Faecal pollution, faecal source tracking, faecal sterols, coprostanol, septic systems

Introduction

Septic systems are designed to accept domestic wastewater and prevent microbial and chemical pollutants from entering surface and ground waters. However, these systems may fail and could release nutrients and pathogenic microorganisms into the environment (Ahmed *et al.*, 2005, Geary and Gardner, 1998; Griffin *et al.*, 2001). There are approximately 102,000 septic systems in service in Southeast Queensland, of which 60-80% are thought to be failing (Ahmed *et al.*, 2005; Jelliffe, 1995). While detailed description of failed septic systems is not described in the literature, it appears that clogging of the absorption field is the leading cause of septic system failure (Moore, 1990).

Faecal pollution from point and non-point sources has traditionally been assessed by enumerating faecal indicator bacteria such as *Escherichia coli* and enterococci commonly found in the intestine of warm-blooded animals (Baudišová, 1997). However, while the presence of such indicator bacteria in surface waters can be seen as a measure of the quality of the water, it does not provide definitive information with respect to possible sources. In view of this, faecal source tracking (FST) methods have been developed to identify the most

51 likely source(s) of faecal pollution in surface waters (Field and Samadpour, 2007). These
52 methods could be broadly categorized as microbial methods and chemical methods. In recent
53 years, a number of microbial methods such as biochemical fingerprinting (Ahmed et al.,
54 2005), Antibiotic resistance analysis (Parveen *et al.*, 1997), Human- and cattle-specific
55 Bacteroides markers (Bernhard and Field, 2000), human- and bovine-specific viruses (Fong
56 *et al.*, 2005; Hundesa *et al.*, 2006) have been used to identify human and animal faecal
57 pollution in environmental waters.

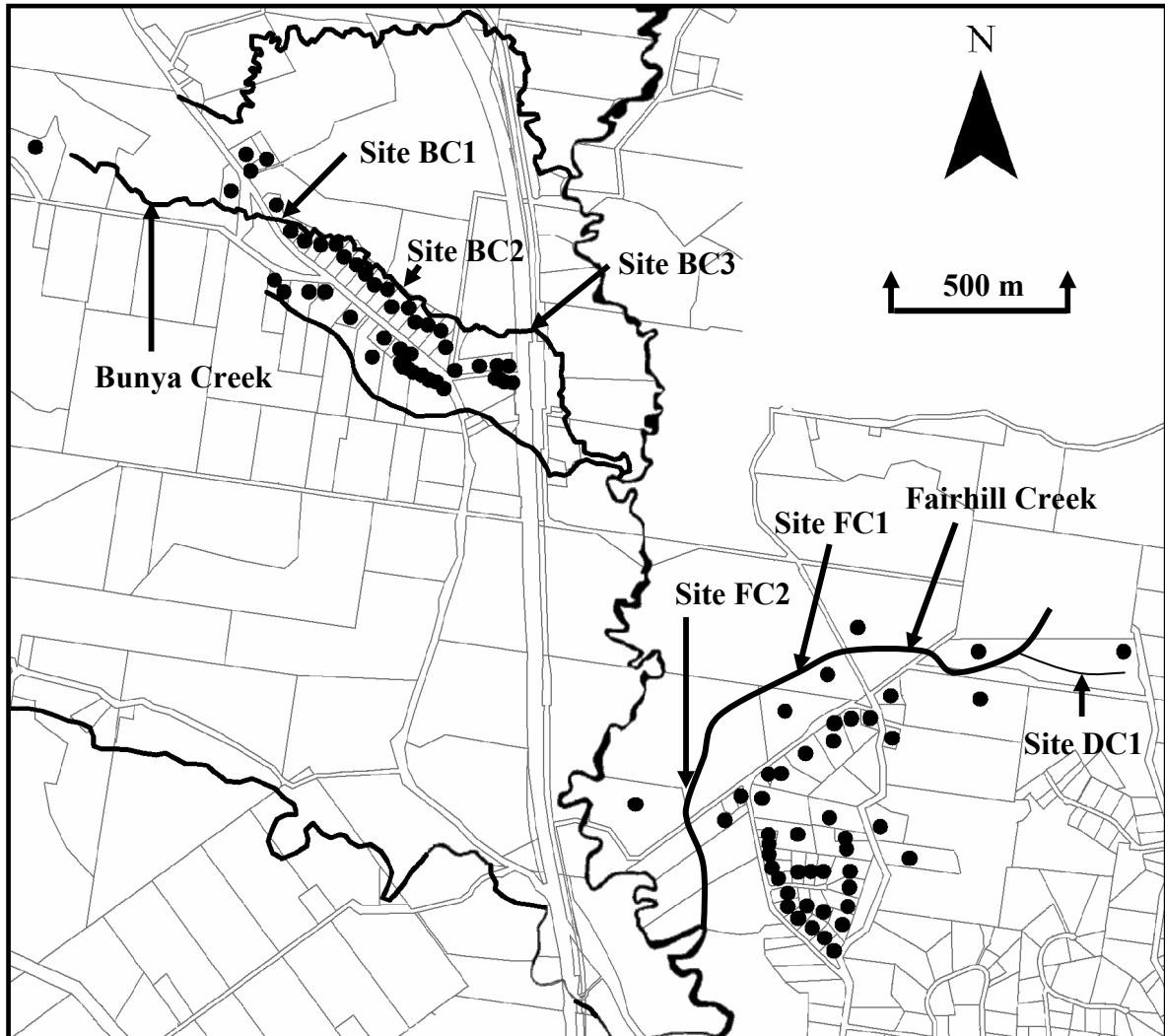
58
59 However, the performance of some microbial methods has not been fully evaluated, or are
60 still under evaluation and to date none of these methods are considered as ‘pioneer’ or ‘gold
61 standard’ in terms of identifying the sources of faecal pollution (Field and Samadpour, 2007).
62 On the other hand, chemical methods such as faecal sterols and stanols have also been used
63 extensively for FST (Leeming *et al.* 1996; Suprihatin *et al.*, 2003). Coprostanol is the major
64 sterol in human faeces, generally comprising about 40-60% of the total sterol content
65 (Leeming *et al.*, 1994). Coprostanol is considered a biomarker of human faecal pollution
66 (Leeming, 1997; Murtaugh and Bunch, 1967). However, the use of coprostanol alone as a
67 biomarker can lead to false indication of results as coprostanol is also present in the faeces of
68 some other animals. In addition, small amounts can be generated from cholesterol in
69 anaerobic sediments (Mudge and Gwyn Lintern, 1999). In addition to coprostanol, various
70 other sterols can be found in humans and animals as a result of different diets, variation in
71 digestive tracts and the diversity of gut microflora. The ratio of coprostanol with other faecal
72 sterols has been proposed as an improved chemical method to identify the sources of human
73 and herbivore faecal pollution (Bull *et al.*, 2003; Leeming, 1997). Animals such as dogs and
74 birds generally do not have faecal sterols in their faeces or it could be present in low
75 concentration (Leeming *et al.*, 1996). However, faecal matters from these animals can be
76 distinguished from humans and herbivores by comparing the concentration of *E. coli* and
77 enterococci commonly found in the faeces of these animals with the concentration of the
78 same indicators found in the faeces of human and herbivores based on the faecal sterol
79 concentrations. Such analysis may provide additional information to identify the sources of
80 dog and bird faecal pollution.

81
82 In this study, we used faecal sterol analysis to determine whether failing septic systems are
83 contributing faecal pollution into the environment. Surface water samples were collected
84 from a non-sewered catchment (i.e. North Maroochy Catchment, Southeast Queensland,
85 Australia) that contained a high density of septic systems and were tested for the presence of
86 faecal sterols. Sterol ratios were then used to provide further evidence, or to identify the most
87 likely sources of faecal pollution in the study catchment.

88 89 **Methods**

90 *Study catchment and sampling sites.* North Maroochy Catchment was chosen for this study
91 because the entire catchment is serviced by septic systems. These systems are not currently
92 being monitored, and they have the potential to fail and transport pollutants into the
93 environment. Three creeks (i.e. Bunya, Fairhill and Davidson) were chosen for water
94 sampling (Figure 1). Samples were collected from three sites (i.e. site BC1, BC2 and BC3) in
95 the Bunya Creek, two sites (i.e. FC1 and FC2) in the Fairhill Creek and one control site
96 (DC1) in the Davidson Creek. Bunya Creek is a small first order stream which receives water
97 mainly from a natural spring located upstream of the creek. Sample site BC1 was located
98 close to the spring and site BC2 was located 600 m downstream and is characterized by
99 nearby residential blocks with septic systems. The BC3 site is a further 800 m downstream
100 and below the residential development and is characterized by cattle grazing. Site FC1 was

101 located in close proximity to a cluster of older residential blocks in the Fairhill Creek. Site
102 FC2 was located 1.2 km downstream from the older residential blocks. The water flow at this
103 site is primarily associated with storm events. The control site was located in the upstream
104 region of the Davidson Creek, headwaters of the Fairhill Creek. The surrounding area of the
105 control site consisted mostly of rainforest with minimal anthropogenic impacts (< 3 septic
106 systems).



137 Figure 1. Map showing sampling sites and location of septic systems (●) in North Maroochy
138 Catchment.

140 *Analytical standards.* The sterol standards – coprostanol (5β -cholestan- 3β -ol), cholestane
141 (5α -cholestane), cholesterol (cholest- 5 -en- 3β -ol), stigmasterol (24-ethylcholesta- 5 , 22-E-
142 dien- 3β -ol), sitosterol (24-ethylcholest- 5 -en- 3β -ol) and sitostanol (24-ethyl- 5α -cholestan- 3β -
143 ol) were purchased from Sigma (Australia). Epicoprostanol was purchased from Steraloids
144 (USA). The derivatizing agent trimethylsilylimidazole (TMSI) was also purchased from
145 Sigma (Australia).

146 *Sample collection and preparation.* Water samples were collected in 10 L pre-cleaned
147 polyethylene containers from the six sites on six occasions using aseptic techniques. The
148 samples were then transported to the laboratory, kept at 4°C and processed within 24 h.
149 Appropriate volume (i.e. 5-10 L) of each water sample was filtered through 142 mm glass

150 fibre filters (Advantec, Tokyo, Japan). For lipid extraction of the particulate material, two-
151 phase extraction (utilizing methanol and hexane) was performed using a modification of the
152 one-phase CHCl_3 -MeOH- H_2O Bligh and Dyer method (1959). The filter discs (containing
153 particulates) were transferred into 250 mL bottles containing methanol (30 mL), hexane (30
154 mL), sodium carbonate (1 gm) and internal standard (10 μg cholestane in 10 μl
155 dichloromethane) and tumbled for 24 h. The samples were further filtered through 110 mm
156 filter papers (Whatman, Grade number 2) and transferred into liquid separating funnels. The
157 hexane phase was retained and washed with 15 mL Milli-Q water, dried with anhydrous
158 sodium sulphate and then concentrated using nitrogen. The residue was dried and
159 reconstituted in hexane (1 mL) and trimethylsilylimidazole (TMSI) (20 μl) in 1.5 mL vials.
160 The samples were then allowed to stand for 3 h at room temperature before being analysed.
161 Samples from the septic systems near the creeks, were collected directly from the outlet of
162 the septic system. The samples were weighed to determine the septic content. Extraction was
163 performed according to the methanol-hexane method described above.

164 *Analysis of sterol based lipid by gas chromatography and mass spectrometry.* Extracts were
165 analysed using a Varian 3900 Gas Chromatograph (GC) (Hansen Way, Palo Alto, USA). The
166 injector temperature was 320°C, and the split was shut for 0.5 min then opened to 50:1. The
167 oven temperature was programmed at 200°C on injection and increased at 20°C/min to 240°C,
168 then increased at 3°C/min to 320°C and held for 5 min. The GC was coupled (transfer line
169 280°C) to a Varian Saturn 2100T Mass Spectrometer (MS) with compound ionization by
170 electron impact energy at 70 eV. The positive fragment ions were analysed over a mass range
171 of 200-550 m/z. Sterols were quantified by reference to standard solutions and expressed as
172 ng/L of water. For interpretation of the faecal sterol profiles found in water samples, and to
173 identify possible human and animal faecal pollution, C27:C29 and 5 β :5 α ratio comparisons
174 were used (Leeming *et al.*, 1996; Leeming *et al.*, 1998).

175

176 **Results and discussions**

177 *Concentrations of sterols in environmental samples.* Low levels of coprostanol and
178 epicoprostanol (ranging from 0-4.0 X 10⁰ ng/L) were found in water samples collected on
179 occasion 1 except the sample from site FC1 which had a coprostanol level of 2.3 X 10¹ ng/L
180 (see Table 1). The water sample from this site also had high levels of cholesterol (6.5 X 10³
181 ng/L) and sitosterol (1.1 X 10⁴ ng/L). Increased levels of coprostanol and epicoprostanol
182 were found in water samples collected on occasion 2 with the highest found in site BC1 (1.2
183 X 10³ ng/L and 20 X 10¹ ng/L respectively). The high level of coprostanol suggesting human
184 sourced faecal pollution. The concentrations of other sterols were also high in this site. Prior
185 to this occasion, the catchment experienced moderate rainfall (36 mm) and as a result surface
186 and sub-surface runoff from agricultural areas and failing septic trenches may have increased
187 the levels of sterols in receiving waters. In contrast, the sample from site BC2 had lower
188 levels of coprostanol (2.2 X 10² ng/L) and epicoprostanol (3.7 X 10¹ ng/L), although, the
189 levels of cholesterol, sitosterol and sitostanol were high. Site BC3 showed similar patterns
190 except the coprostanol and epicoprostanol concentrations were higher than that of site BC2.
191 The lower levels of sterols in downstream sites (i.e. BC2 and BC3) could be due to the
192 dilution associated with more water flow, which may have masked certain sterols levels. It is
193 also possible that sterols bound particles may have settled in to the sediments in these sites.
194 The sterols are highly hydrophobic molecules which are found to be absent (insoluble) in the
195 clear water column. This is consistent with the research literature (Leeming *et al.*, 1996).
196 However, this could not be confirmed in this study as samples were not collected or tested
197 from the sediments.

198

199 The concentrations of stanols were relatively higher in site FC1 compared to site FC2 and the
200 upstream control site DC1. All sites had lower levels of coprostanol (ranging from 1.0×10^0 -
201 2.3×10^1 ng/L) and epicoprostanol (up to 1.0×10^0 ng/L) with the lowest found in site DC1
202 on occasion 3. The concentrations of other sterols were also lower on this occasion. This is
203 probably because during this occasion the catchment did not experience any significant
204 rainfall. Higher coprostanol level was again found in all samples on occasion 4 when the
205 catchment had just received 14 mm of rainfall. However, no epicoprostanol was detected in
206 any of these samples. On occasion 5, site BC1 had higher levels of coprostanol (2.0×10^2
207 ng/L) and epicoprostanol (3.4×10^1 ng/L) compared to sites BC2 and BC3. In contrast, sites
208 FC1, FC2 and the control had lower stanols level. On occasion 6, site BC1 had

Table 1: Sterol profile (ng/L) of water samples collected from six sites on six occasions.

Sampling sites	Sampling events (rainfall)*	Sterols (ng/L)						
		Coprostanol	Epicoprostanol	Cholesterol	5 α Cholestanol	24 Et-coprostanol	Sitosterol	
BC1	Event 1 (6 mm)	6.0 X 10 ⁰	2.0 X 10 ⁰	1.7 X 10 ³	1.4 X 10 ²	0.0 X 10 ⁰	3.8 X 10 ²	1.5 X 10 ²
	Event 2 (36 mm)	1.2 X 10 ³	2.1 X 10 ²	7.6 X 10 ²	3.7 X 10 ²	4.8 X 10 ²	3.8 X 10 ²	9.1 X 10 ¹
	Event 3 (1 mm)	2.3 X 10 ¹	0.0 X 10 ⁰	9.0 X 10 ²	2.3 X 10 ¹	3.0 X 10 ⁰	9.7 X 10 ²	4.5 X 10 ⁰
	Event 4 (14 mm)	2.0 X 10 ¹	0.0 X 10 ⁰	8.7 X 10 ²	6.9 X 10 ¹	0.0 X 10 ⁰	4.3 X 10 ²	1.1 X 10 ²
	Event 5 (28 mm)	2.0 X 10 ²	3.4 X 10 ¹	3.8 X 10 ³	4.2 X 10 ²	7.4 X 10 ¹	1.3 X 10 ³	2.0 X 10 ²
	Event 6 (30 mm)	3.0 X 10 ⁴	1.0 X 10 ³	3.6 X 10 ³	3.6 X 10 ³	1.2 X 10 ³	9.9 X 10 ²	1.1 X 10 ³
BC2	Event 1 (6 mm)	4.0 X 10 ⁰	0.0 X 10 ⁰	5.7 X 10 ²	4.6 X 10 ¹	1.2 X 10 ⁰	1.2 X 10 ²	4.5 X 10 ¹
	Event 2 (36 mm)	2.2 X 10 ²	3.7 X 10 ¹	1.2 X 10 ³	2.8 X 10 ²	2.2 X 10 ²	1.5 X 10 ³	3.4 X 10 ¹
	Event 3 (1 mm)	5.0 X 10 ⁰	0.0 X 10 ⁰	2.6 X 10 ²	2.9 X 10 ¹	8.0 X 10 ⁰	2.0 X 10 ²	6.9 X 10 ¹
	Event 4 (14 mm)	5.1 X 10 ¹	0.0 X 10 ⁰	1.1 X 10 ³	1.3 X 10 ²	0.0 X 10 ⁰	2.8 X 10 ²	1.1 X 10 ²
	Event 5 (28 mm)	9.2 X 10 ¹	1.2 X 10 ¹	1.2 X 10 ³	1.4 X 10 ²	8.0 X 10 ¹	7.7 X 10 ²	1.9 X 10 ²
	Event 6 (30 mm)	7.0 X 10 ¹	1.1 X 10 ¹	1.2 X 10 ³	1.3 X 10 ²	1.0 X 10 ²	3.2 X 10 ²	1.3 X 10 ¹
BC3	Event 1 (6 mm)	4.0 X 10 ⁰	0.0 X 10 ⁰	3.9 X 10 ²	3.4 X 10 ¹	2.3 X 10 ¹	8.0 X 10 ¹	2.9 X 10 ¹
	Event 2 (36 mm)	4.3 X 10 ²	9.3 X 10 ¹	1.5 X 10 ³	3.3 X 10 ¹	6.3 X 10 ²	1.5 X 10 ³	3.3 X 10 ²
	Event 3 (1 mm)	2.4 X 10 ¹	1.0 X 10 ⁰	5.2 X 10 ²	5.3 X 10 ¹	3.7 X 10 ¹	5.3 X 10 ²	9.2 X 10 ¹
	Event 4 (14 mm)	6.3 X 10 ¹	0.0 X 10 ⁰	8.3 X 10 ²	8.2 X 10 ¹	4.5 X 10 ¹	3.1 X 10 ²	5.4 X 10 ¹
	Event 5 (28 mm)	4.0 X 10 ¹	0.0 X 10 ⁰	6.1 X 10 ²	8.4 X 10 ¹	2.0 X 10 ¹	1.7 X 10 ²	4.4 X 10 ¹
	Event 6 (30 mm)	4.0 X 10 ¹	0.0 X 10 ⁰	5.1 X 10 ²	5.1 X 10 ¹	8.0 X 10 ¹	9.6 X 10 ¹	5.8 X 10 ¹
FC1	Event 1 (6 mm)	2.3 X 10 ²	0.0 X 10 ⁰	6.5 X 10 ³	3.5 X 10 ¹	4.3 X 10 ¹	1.1 X 10 ⁴	4.5 X 10 ²
	Event 2 (36 mm)	3.4 X 10 ²	1.7 X 10 ²	3.7 X 10 ³	9.3 X 10 ²	6.0 X 10 ²	2.2 X 10 ³	7.0 X 10 ²
	Event 3 (1 mm)	5.0 X 10 ⁰	0.0 X 10 ⁰	6.3 X 10 ²	4.1 X 10 ¹	7.0 X 10 ⁰	5.6 X 10 ²	5.3 X 10 ¹
	Event 4 (14 mm)	4.5 X 10 ¹	0.0 X 10 ⁰	1.9 X 10 ³	3.5 X 10 ¹	5.6 X 10 ¹	6.6 X 10 ²	1.5 X 10 ²
	Event 5 (28 mm)	8.0 X 10 ⁰	0.0 X 10 ⁰	1.1 X 10 ³	1.8 X 10 ¹	9.0 X 10 ⁰	1.7 X 10 ²	1.9 X 10 ¹
	Event 6 (30 mm)	3.4 X 10 ²	1.0 X 10 ¹	3.0 X 10 ³	2.0 X 10 ²	1.4 X 10 ²	1.0 X 10 ³	1.3 X 10 ²
FC2	Event 1 (6 mm)	0.0 X 10 ⁰	0.0 X 10 ⁰	7.9 X 10 ²	4.5 X 10 ¹	9.0 X 10 ⁰	2.8 X 10 ²	2.7 X 10 ¹
	Event 2 (36 mm)	2.9 X 10 ¹	0.0 X 10 ⁰	1.3 X 10 ³	9.1 X 10 ¹	7.1 X 10 ¹	1.0 X 10 ³	7.4 X 10 ¹
	Event 3 (1 mm)	1.6 X 10 ¹	0.0 X 10 ⁰	4.8 X 10 ²	2.6 X 10 ¹	1.0 X 10 ¹	8.2 X 10 ²	0.0 X 10 ⁰
	Event 4 (14 mm)	2.1 X 10 ¹	0.0 X 10 ⁰	1.1 X 10 ³	1.6 X 10 ¹	0.0 X 10 ⁰	7.8 X 10 ²	1.1 X 10 ¹
	Event 5 (28 mm)	4.7 X 10 ¹	1.7 X 10 ¹	2.6 X 10 ³	1.6 X 10 ²	1.3 X 10 ²	1.5 X 10 ³	5.6 X 10 ²
	Event 6 (30 mm)	6.5 X 10 ¹	9.0 X 10 ⁰	8.2 X 10 ²	6.5 X 10 ¹	8.0 X 10 ¹	5.3 X 10 ²	1.0 X 10 ²
DC1	Event 1 (6 mm)	1.0 X 10 ⁰	0.0 X 10 ⁰	4.5 X 10 ¹	3.0 X 10 ⁰	0.0 X 10 ⁰	1.7 X 10 ¹	3.0 X 10 ⁰
	Event 2 (36 mm)	8.3 X 10 ¹	2.8 X 10 ¹	1.6 X 10 ³	1.5 X 10 ²	6.0 X 10 ¹	2.7 X 10 ³	2.0 X 10 ²
	Event 3 (1 mm)	1.0 X 10 ⁰	0.0 X 10 ⁰	1.1 X 10 ²	1.0 X 10 ¹	0.0 X 10 ⁰	5.6 X 10 ¹	1.0 X 10 ⁰
	Event 4 (14 mm)	1.5 X 10 ¹	0.0 X 10 ⁰	6.6 X 10 ²	6.7 X 10 ¹	1.2 X 10 ¹	2.9 X 10 ²	1.6 X 10 ²
	Event 5 (28 mm)	0.0 X 10 ⁰	0.0 X 10 ⁰	3.6 X 10 ²	9.0 X 10 ⁰	0.0 X 10 ⁰	9.0 X 10 ¹	1.2 X 10 ¹
	Event 6 (30 mm)	2.2 X 10 ³	1.6 X 10 ²	5.5 X 10 ²	4.6 X 10 ²	8.0 X 10 ²	2.7 X 10 ²	1.3 X 10 ²

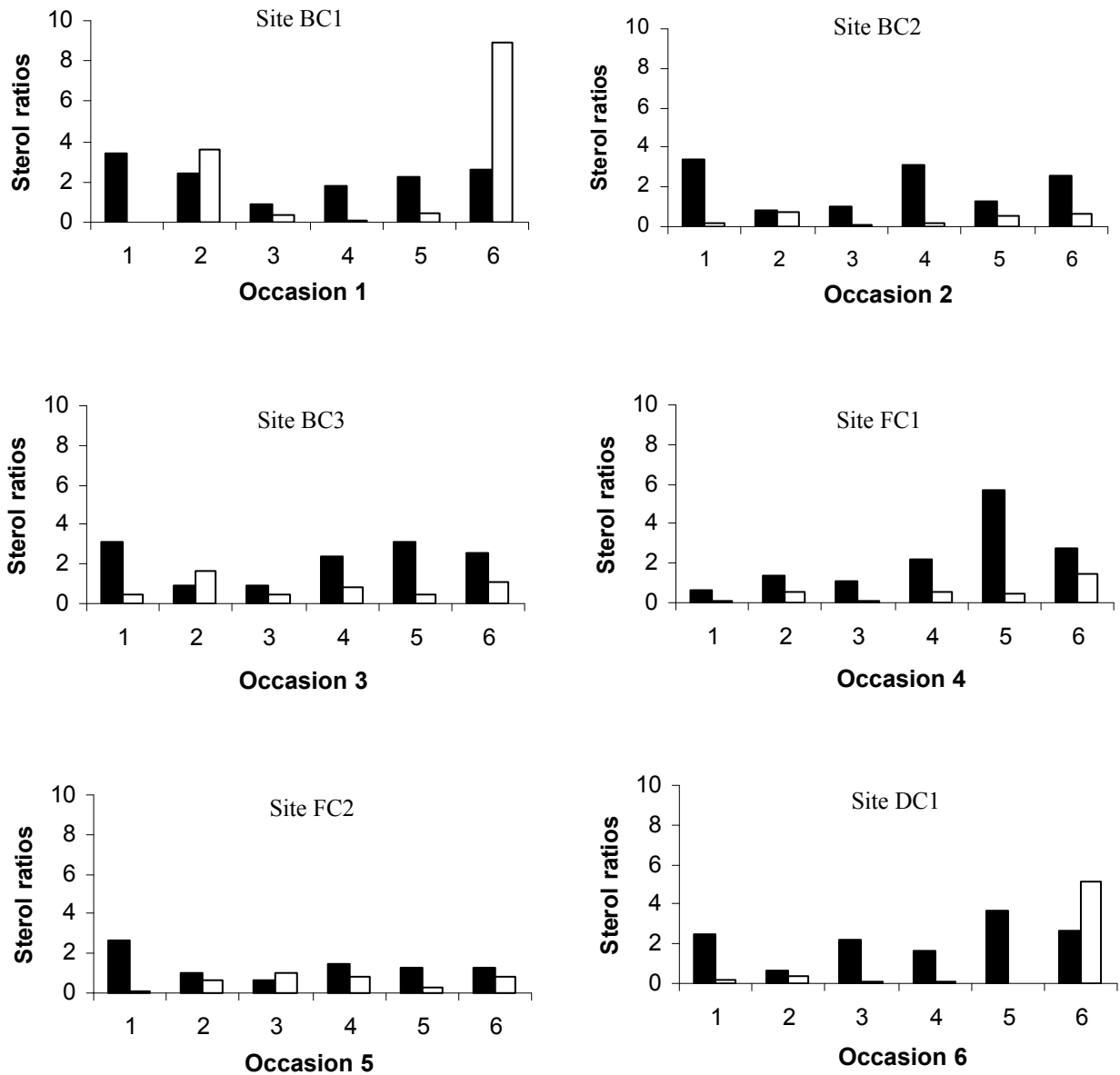
* Rainfall occurring two days prior to sampling

212 very high levels of coprostanol (approximately 3.0×10^4 ng/L) and epicoprostanol (1.0×10^3
213 ng/L) (this was following 30 mm of rainfall). The presence of high levels of stanols at this
214 site after rainfall events were followed by performing a sanitary inspection. A defective septic
215 system nearby was identified as the most likely source. The trench was located within 5 m
216 distance from the creek and was seeping. For confirmation, samples were collected from the
217 tank and the trench (i.e. soil sample). The sample from the tank had higher levels of
218 coprostanol (8.6×10^6 ng/L) while the sample from the trench also had coprostanol but the
219 level was 14 fold lower than that of the tank. This was expected given soil filtering and
220 catchment flushing during rainfall. The concentrations of other sterols were also high at site
221 BC1. However, lower levels of stanols were found in downstream sites (BC2 and BC3). Site
222 FC1 had higher coprostanol level than site FC2. Interestingly, site DC1 also had higher
223 coprostanol (2.2×10^3 ng/L) and epicoprostanol (1.6×10^2 ng/L). This was surprising as the
224 control site was located in a relatively pristine area with minimal human impacts (< 3 septic
225 tanks). It is possible that seepage from these septic systems could be considered as a
226 contributing factor. This could not be confirmed due to access restriction to these septic
227 systems.

228
229 It must be noted that, coprostanol can be found in other animal species such as pigs, cattle,
230 sheep and cats however the concentration is 10 times lower than human faeces (Leeming *et*
231 *al.*, 1996). Based on our data, it appears likely that the high levels of coprostanols in water
232 samples have originated from failing septic trenches. High levels of coprostanols were also
233 detected in Port Philip Bay, Australia near a sewage treatment plant indicating sewage
234 contamination (O'Leary *et al.*, 1999; Leeming *et al.*, 1998). A recent study also reported the
235 evidence of septic system failure by matching unique bacterial patterns from septic tanks with
236 those found in water samples collected from adjacent creeks (Ahmed *et al.*, 2005).

237
238 *Sterols ratios for source tracking.* The presence of coprostanol alone (lower concentration)
239 may not be sufficient enough to provide the evidence of human faecal pollution as other low
240 level inputs into the aquatic environment are possible. To overcome this, the knowledge of
241 the ratios of particular sterols and stanols in faeces has led to ratio analysis. In recent studies,
242 ratios across a range of C₂₇:C₂₉ sterols and 5 β :5 α stanols have given a more specific measure
243 of pollution (Bull *et al.*, 2003; Leeming *et al.*, 1996; Nash *et al.*, 2005). When C₂₇:C₂₉ and
244 5 β :5 α ratios are both greater than 1, the faecal source is likely to be of human origin. Ratios
245 (C₂₇:C₂₉ and 5 β :5 α) < 1, are indicative of mixed faecal pollution and C₂₇:C₂₉ < 1 and 5 β :5 α
246 > 1 ratios are indicative of herbivore faecal pollution. The ratio analysis was also used as
247 confirmation of the presence of human faecal pollution or others (i.e. herbivores and/or
248 mixed). The ratio of C₂₇:C₂₉ and 5 β :5 α in samples from site BC1 ranged between 0.93 to
249 3.41 and 0.02 to 8.91 respectively (see Figure 2). Both ratios were above 1 for two samples
250 (occasions 1 and 6) indicated human sourced faecal pollution at site BC1. The ratios of 0.93
251 and 0.38 (occasion 3) at this site suggesting mixed faecal pollution. The ratios in samples
252 from site BC2 ranged between 0.79 to 3.43 and 0.18 to 0.70 suggesting that human faecal
253 pollution is not the dominant source. The ratios of 0.79 and 0.70 (occasion 2) at site BC2
254 suggested mixed faecal pollution. The sterol ratios of 0.92 and 1.61 at site BC3 on occasion
255 2, strongly suggest herbivores as the major sources of faecal pollution. This is not surprising
256 as site BC3 is characterized by cattle farming. The ratios of 2.57 and 1.10 at this site on
257 occasion 6 are indicative of human faecal pollution. The sterol ratios of C₂₇:C₂₉ > 1 and 5 β :5 α
258 < 1 at site BC3 on occasions 4 and 5 suggesting humans (C₂₇:C₂₉ > 1, 5 β :5 α > 1) and cattle
259 (C₂₇:C₂₉ < 1, 5 β :5 α > 1) are not the dominant sources. Based on our data, it appears that during
260 dry events, septic systems may not contribute faecal pollution into the creeks. The sterol
261 ratios of C₂₇:C₂₉ > 1 and 5 β :5 α < 1 in sites FC1 on occasions 1 to 5 suggest humans (C₂₇:C₂₉

262 >1 , $5\beta:5\alpha >1$) and cattle ($C_{27}:C_{29} <1$, $5\beta:5\alpha >1$) are not the dominant sources of faecal
 263 pollution. However, the $C_{27}:C_{29}$ and $5\beta:5\alpha$ ratios of 2.74 and 1.45 on occasion 6 suggesting
 264 the presence of human faecal pollution. The sterol ratios of $C_{27}:C_{29} >1$ and $5\beta:5\alpha <1$ in site
 265 FC2 suggest again that humans and cattle are not the dominant sources of faecal pollution.
 266 The ratios at this site suggested mixed faecal pollution. Mixed faecal pollution was also
 267 observed in samples from the control site with the exception that a sample (occasion 6)
 268 indicated the presence of human faecal pollution.



306 Figure 2: $C_{27}:C_{29}$ (■) and $5\beta:5\alpha$ (□) ratios in water samples collected from six sites in North
 307 Maroochy Catchment. ($C_{27}:C_{29}$ and $5\beta:5\alpha$) >1 indicate human faecal pollution; ($C_{27}:C_{29}$
 308 and $5\beta:5\alpha$) <1 , indicate mixed faecal pollution; and ($C_{27}:C_{29} <1$ and $5\beta:5\alpha >1$) indicate
 309 herbivore faecal pollution.

312 **Conclusions**

313 In this study, faecal sterol analysis was used to identify the presence of human sourced faecal
314 pollution or others (non-point sources) in two adjacent creeks of North Maroochy Catchment.
315 It appears that stanols concentrations generally increased with increased catchment runoff.
316 After moderate rainfall, high coprostanols levels found in water samples indicated human
317 faecal pollution and defective septic systems are the most likely sources of pollution. The
318 human signal was traced on one occasion to a defective septic system. In contrast, it appears
319 that during dry weather human faecal pollution is not occurring in the study catchment. **The**
320 **advantages of faecal sterol analysis are that this method is cheaper and can be performed**
321 **rapidly compared to faecal source tracking methods that require the development of a**
322 **database of faecal indicator bacteria such as antibiotic resistance analysis.** One major
323 limitation of the method is that the sources could not always be identified due to dilution and
324 mixing of several sources. Another limitation is that the faecal sterols analysis does not
325 provide any information regarding public health risks associated with faecal pollution.
326 Nonetheless, the presence of human faecal pollution in the environment is alarming and such
327 data could be valuable to water quality managers who are charged with protecting water
328 quality and public health.

329

330 **Acknowledgements**

331 This study was supported by a grant from SEQ Catchments. Our thanks to Ms. Gloria
332 Macintosh for assisting in collecting water samples.

333

334 **References**

- 335 Ahmed, W., Neller, R. and Katouli, M. 2005. Evidence of septic system failure determined by a
336 bacterial biochemical fingerprinting method. *Journal of Applied Microbiology*, 98(4): 910-
337 920.
- 338 Baudišová, D. 1997. Evaluation of *Escherichia coli* as the main indicator of faecal pollution. *Water*
339 *Science and Technology*, 35(11-12): 333-336.
- 340 Bernhard, A.E. and Field, K.G. 2000. A PCR assay to discriminate human and ruminant feces on the
341 basis of host differences in *Bacteroides-Prevotella* genes encoding 16S rRNA. *Applied and*
342 *Environmental Microbiology*, 66(10): 4571-4574.
- 343 Bligh, E.G. and Dyer, W.J. 1959. A rapid method for total lipid extraction and purification. *Canadian*
344 *Journal of Biochemistry and Physiology*, 37(8): 911-917.
- 345 Bull, I.D., Elhmmali, M.M., Roberts, D.J. and Evershed, R.P. 2003. The application of steroidal
346 biomarkers to track the abandonment of a Roman wastewater course at the Agora (Athens,
347 Greece). *Archaeometry*, 45(1): 149-161.
- 348 Field, K.G. and Samadpour, M. 2007. Fecal source tracking, the indicator paradigm, and managing
349 water quality. *Water Research*, 41(16): 3517-3538.
- 350 Fong, T.T., Griffin, D.W. and Lipp, E.K. 2005. Molecular assays for targeting human and bovine
351 enteric viruses in coastal waters and their application for library-independent source tracking.
352 *Applied and Environmental Microbiology*, 71(4): 2070-2078.
- 353 Geary, P.M. and Gardner, E.A. 1998. Sustainable on-site treatment systems. ***In: Proceedings of***
354 ***the Eighth National Symposium on Individual and Small Community Sewage Systems,***
355 ***American Society for Agricultural Engineers, St. Joseph, MI. 12-20.***
- 356 Griffin, D., Lipp, E., McLaughlin, M. and Rose, J. 2001. Marine recreation and public health
357 microbiology: quest for the ideal indicator. *Bioscience*, 51(10): 817-825.
- 358 Hundesa, A., Maluquer de Motes, C., Bofill-Mas, S., Albinana-Gimenez, N. and Girones, R. 2006.
359 Identification of human and animal adenoviruses and polyomaviruses for determination of
360 sources of fecal contamination in the environment. *Applied and Environmental Microbiology*.
361 72(12): 7886-7893.
- 362 Jelliffe, P. A. 1995. Management of on-site effluent disposal: conclusions from a study on the
363 performance of 101 systems in Maroochy Shire, **February-March, 1994.**

- 364 Leeming, R., Ball, A., Ashbolt, N., Jones, G. and Nichols, P. 1994. Distinguishing between
365 human and animal sources of faecal pollution. *Chemistry in Australia*, 61, 434-435.
- 366 Leeming, R., Ball, A., Ashbolt, N. and Nichols, P. 1996. Using faecal sterols from humans and
367 animals to distinguish faecal pollution in receiving waters. *Water Research*, 30(12): 2893-
368 2900.
- 369 Leeming, R. 1997. Use of faecal sterols and bacterial indicators to discriminate sources of
370 faecal pollution entering Lake Macquarie, Newcastle, NSW, CSIRO Report 96-HWC1,
371 Report prepared for the Hunter Water Corporation.
- 372 Leeming, R., Nichols, P.D. and Ashbolt, N. J. 1998. Distinguishing sources of faecal pollution
373 in Australian Inland and Coastal waters using sterol biomarkers and microbial faecal
374 indicators. Report No. 204. Water Services Association of Australia, Melbourne, Australia.
- 375 Moore, J. A. 1990. Why do septic system fail? Corvallis, OR: Oregon State University
376 Extension Service. EC1340.
- 377 Mudge, S. M. and Gwyn Lintern, D. 1999. Comparison of sterol biomarkers for sewage with
378 other measures in Victoria Harbour, B.C., Canada. *Estuarine, Coastal and Shelf Science*,
379 48(1): 27-28.
- 380 Murtaugh, J.J. and Bunch, R.L. 1967. Sterols as a measure of faecal pollution. *Journal of Water*
381 *Pollution Control Federation*, 39(3): 404-409.
- 382 Nash, D., Leeming, R., Clemow, L., Hannah, M., Halliwell, D. and Allen, D. 2005.
383 Quantitative determination of sterols and other alcohols in overland flow from grazing land
384 and possible source materials. *Water Research*, 39(13): 2964-2978.
- 385 O'Leary, T., Leeming, R., Nichols, P. D. and Volkman, J. K. 1999. Assessment of the sources,
386 transport and fate of sewage derived organic matter in Port Phillip Bay, Australia using the
387 signature lipid coprostanol. *Marine and Freshwater Research*, 50(6): 547-556.
- 388 Parveen, S., Murphree, R.L., Edminston, L., Kaspar, C.W., Portier, K.M., Tamplin, M.L., 1997.
389 Association of multiple antibiotic resistance profiles with point and nonpoint sources of
390 *Escherichia coli* in Apalachicola Bay. *Applied and Environmental Microbiology*, 63(7): 2607-
391 2612.
- 392 Suprihatin, I., Fallowfield, H., Bentham, R. and Cromar, M. 2003. Determination of faecal
393 pollutants in Torrens and Patawolonga catchment waters in South Australia using faecal
394 sterols. *Water Science and Technology*, 47(7-8), 283-289.