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2 Title: Interpreting drinking water quality samples: understanding
3 contamination pathways at the point of collection

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22 Interpreting drinking water quality samples: understanding 23 contamination pathways at the point of collection

24 Abstract

25 2.2 billion people lack access to safely managed drinking water, with microbiological water
26 quality the major barrier to the achievement of SDG target 6.1. Microbiological water
27 quality is strongly influenced by environmental and hand hygiene. We demonstrate the
28 significant contribution of contamination from hygiene at water collection point localities
29 and at the household level to drinking water quality. Comparison of water quality samples
30 from 5,080 water systems at the ‘point of delivery’ (PoD) to those from the ‘point of
31 collection’ (PoC) demonstrated that 60% of the *E. coli* positive results at the PoC were
32 caused by contamination from local hygiene-related pathways, with seasonal variability.
33 Additionally, household hygiene contamination pathways contributed a further 5% to 70%
34 of *E. coli* positive detections in samples from the ‘point of use’ (PoU) based on analysis of 38
35 Multiple Indicator Cluster Survey (MICS) datasets. Better reporting of PoC and PoD methods
36 is required to recognise this contribution from poor hygiene and ensure data are
37 interpreted appropriately, thereby enabling more efficient targeting and prioritisation of
38 interventions. Further revisions to drinking water quality sampling protocols are required to
39 reflect the increase in (unchlorinated) piped water systems with potential for considerable
40 biofilm growth, with implications for release of *Pseudomonas* and other opportunistic
41 pathogens during sampling. There is immediate benefit to be gained from expanding
42 communication and hygiene education to encourage cleaning of taps and spouts for
43 drinking water.

44 Keywords

45 Drinking water, sampling methods, *E. coli*, faecal contamination, point of collection

46 1 Introduction

47 Faecal contamination is the primary limiting factor to get universal access to safely managed
48 drinking water (Greenwood et al., 2024). An estimated 2.2 billion people lack access
49 (UNICEF/WHO JMP, 2023), but adequate and available data on water quality is limited.
50 Recent approaches to address these data gaps through modelling have estimated the
51 number is closer to 4.4 billion people globally without access to safety managed drinking
52 water (Greenwood et al., 2024). This large estimate is realistic given the seasonal variability
53 in water quality (Bangladesh Bureau of Statistics and UNICEF Bangladesh, 2021; Charles et
54 al., 2022), however, to advance drinking water safety we need to understand the
55 implications of the data, and particularly, how it was sampled.

56
57 There are three common methods for sampling drinking water quality: sampling water from
58 taps that have been cleaned, taps that have not been cleaned, and from a vessel served at
59 the household level “as you would give a child to drink.” Here and throughout this paper we
60 use the term ‘tap’ to refer both to the devices that control outflow from reticulated systems
61 and the spouts of handpumps. We propose that water samples that are collected from
62 cleaned taps should be termed Point of Delivery (PoD) samples. International standards on
63 water quality sampling generally recommend that sampling be done at this point. To ensure
64 a clean tap, the tap should be disinfected by flaming (or appropriate disinfection method)

65 and flushed (APHA/AWWA/WEF, 2018; British Standards Institution, 2006; International
66 Organization for Standardization, 2006) or a clean tap should be chosen
67 (APHA/AWWA/WEF, 2018). These methods for *E. coli* and faecal indicator bacteria sampling
68 are commonly derived from chlorinated, piped systems (Charles et al., 2020).

69
70 Water that is sampled from taps that have not been cleaned can be termed Point of
71 Collection (PoC) samples. Samples collected in the household by asking for a cup of water
72 “as you would give a child to drink” are termed Point of Use (PoU) samples. These sampling
73 practices are common in international development, such as the Multiple Indicator Cluster
74 Survey (MICS) approach to nationally representative water sampling (JMP/UNICEF/WHO,
75 2016). PoC sampling is included in international standards to “know the quality of the water
76 as it is consumed” which is considered to be appropriate for “special situations, e.g.
77 outbreaks” (International Organization for Standardization, 2006). Water systems in low-
78 and middle- incomes countries typically include a higher proportion of small and non-piped
79 water systems. Private on-premises connections are less prevalent and this has implications
80 for primary contamination hazards.

81
82 Sampling drinking water from the PoC is becoming a de facto standard in international
83 development work. Early guidance developed for small community supplies (WHO, 1997)
84 differentiated approaches to sampling ‘water as consumed’, collected with flushing and
85 cleaning with a cloth, from ‘quality of water excluding contamination from the tap’ which
86 required flaming and flushing. Fifteen years later, guidance for the rapid assessment of
87 drinking-water quality was aiming to expand understanding of the extent to which “quality
88 of drinking-water from improved sources deviates from the assumption that it is safe” with
89 large scale testing programmes; here the methodology was refined to cleaning the “tap or
90 outlet with a clean, dry cloth” and then flush it before sampling (WHO and UNICEF, 2012).
91 As these approaches expanded into UNICEF’s MICS, the methodology for *E. coli* sampling
92 reduced cleaning further to what is now accepted as PoC samples, specifying only that
93 water be flushed for 30 seconds before sampling (JMP/UNICEF/WHO, 2016). Other
94 guidance does not specify how to sample (e.g. CDC, n.d.).

95
96 While sampling at the PoC is increasingly common, the details of the sampling method are
97 often not explicit in publications, reducing the comparability of results. A review of 171
98 articles that report faecal indicator bacteria results from PoC and PoU drinking water quality
99 sampling in low- and middle-income countries found that 44% did not reference a standard
100 method for sample collection and processing (Sikder et al., 2021). Sikder et al. (2021) did not
101 comment on PoC cleaning practices but they noted that 61% of studies did not report
102 anything about sampling sterility – here the focus was the sterility of sampling vessels and
103 whether the sampler sanitized their hands. This lack of standardisation in reporting presents
104 challenges in interpreting results, and limits the potential for aggregation of studies. A
105 previous systematic review (Bain et al., 2014) of faecal contamination of drinking water in
106 LMICs did not consider sampling methods, including papers using both PoD methods (e.g.
107 Luby et al., 2006) and PoC methods (Mertens et al., 1990; Rufener et al., 2010). In more
108 recent literature we find the same challenge, with recent literature on water quality in
109 Bangladesh exhibiting variable methodology reporting between studies, often providing no
110 data on sampling methods (Ercumen et al., 2017a; Escamilla et al., 2013; Ferguson et al.,

111 2011) , or reporting variable methods for either flushing (Ercumen et al., 2017b; Ferguson et
112 al., 2011) or full decontamination for PoD sampling (Doza et al., 2020).

113
114 PoD, PoC and PoU sampling methods support understanding of different hazards. Wright et
115 al. (2004) suggested that the PoD samples can lead to an understating of ‘source
116 contamination’, however, this misrepresents the utility of these different types of samples.
117 Mertens et al. (1990) demonstrated the difference in water quality from communal
118 tubewells between PoD and PoC: at the PoD only 5% of samples were contaminated with
119 faecal coliforms and contamination was at ‘low levels’ compared to at PoC, when around
120 50% of samples were contaminated, with average contamination similar to levels found in
121 unprotected shallow wells. In their work, over 30 years ago, Mertens et al. reflected that
122 communal water points are sites of activity for humans and animals, highlighting
123 contaminated hands as a potentially important source of contamination. Contamination at
124 the PoC is important for public health, but it should be recognised that the contamination is
125 coming from a wider range of pathways than just the water system, with local hygiene being
126 a primary consideration. At the PoU, still further pathways of contamination contribute to
127 the quality of water assessed, including potential sources of contamination in the household
128 management of drinking water, in transport and storage, as well as localised hygiene issues.
129 It is widely recognised that contamination commonly increases between the PoC and PoU
130 (Bain et al., 2021; Wright et al., 2004).

131
132 These three sampling methods evaluate the risks to the drinking water consumer in
133 different ways, capturing different contamination pathways. In this paper, we analyse how
134 different contamination pathways impact on water quality sample results, drawing on
135 secondary data from international and project level surveys to compare results from PoD,
136 PoC and PoU samples. We explore how interpretations of these different types of sample
137 results differ for drinking water management and the implications for advancing drinking
138 water safety. Lastly, we share results on the unintended results of cleaning taps for
139 sampling in unchlorinated piped systems. Overall, we conclude that the role of local hygiene
140 conditions in drinking water contamination at the point of collection are largely overlooked.

141 2 Materials and methods

142 To compare contamination of PoD, PoC and PoU drinking water quality samples with *E. coli*
143 we utilised multiple secondary data sets. *E. coli* is the most precise indicator of recent faecal
144 contamination in drinking water (Charles et al., 2020). To frame the analysis we broadly
145 characterise the contamination pathways in Figure 1, where PoD captures contamination
146 from the water catchment environment, such as groundwater contamination from latrines,
147 and from infrastructure, such as leaks in pipes. PoC additionally captures pathways
148 associated with ‘local hygiene conditions’; this potentially includes contamination from
149 users hands and animals that touch the tap. PoU capture pathways associated with
150 collection container contamination, storage behaviours, and ‘household hygiene’, the latter
151 is defined for the purposes of this paper as the contribution of hygiene to drinking water
152 contamination in the process of accessing the water and the hygiene of the drinking vessel.

153
154 *[insert here] Figure 1. Drinking water contamination pathways contributing to sample*
155 *results at PoD, PoC and PoU*

156

157 There are three components to the analysis. Firstly, we explore the relative contribution of
158 the ‘local hygiene’ contamination pathways to drinking water quality through comparison of
159 samples at the PoD and PoC to expand the empirical evidence for the difference in these
160 measures. Secondly, we focus on the contribution of the ‘household hygiene’ pathway at
161 the household level, addressing the contribution to water quality at the PoU through
162 comparison of with samples at PoC that indicated no contamination. Lastly, we recognise
163 the lack of reflective evidence that is available on these methods, and particularly on their
164 adaptation to the diverse range of water systems in use. For this we contribute an analysis
165 of PoD and PoC sampling method results in different types of water systems, to advance
166 discussion on considerations when sampling.

167

168 2.1 Contribution of contamination pathways from PoD and PoC

169 Datasets that included paired waterpoint samples for *E. coli* for PoD (sampled after
170 disinfection or decontamination and flushing) and PoC (sampled with flushing but no
171 cleaning) were used to compare the contamination pathways. This sampling approach is
172 applied by the Laboratory of Environmental Health, icddr,b across their research and quality
173 assurance programmes in Bangladesh (Table 1). ‘Tap’ is used throughout to identify the
174 point of delivery for the water for simplicity, but data is almost exclusively from handpumps
175 on tubewells (>99%) with the exception of Study 8. These data are largely from the
176 Rohingya refugee camps in Bangladesh’s Cox’s Bazar peninsula, thus they are not
177 representative of the national level of contamination (Bangladesh Bureau of Statistics and
178 UNICEF Bangladesh, 2021). Study 8 in hospitals is of water piped through the hospital for
179 use in healthcare provision; drinking water points are often separate from these larger
180 piped networks.

181

182 The sampling methodology instructs that waterpoints be flushed for approximately two
183 minutes before the PoC sample is taken. The waterpoint is then decontaminated, either
184 flamed with alcohol-soaked cotton wool for metal or cleaned with an alcohol wipe for
185 plastic outlets, in line with international standards (APHA/AWWA/WEF, 2018; International
186 Organization for Standardization, 2006). The waterpoint is then flushed and the PoD sample
187 is taken. All samples are stored in coolers with ice maintaining the temperature at 4 to 10°C
188 for transport and analysed within 24 hours of sampling in the Laboratory of Environmental
189 Health at icddr,b using membrane filtration (see individual studies for more details). For the
190 analysis in this paper, the data were pooled across the datasets with PoD-PoC pairs analysed
191 to compare the different levels of contamination using descriptive statistics in SPSS (version
192 28). Results were plotted in a Sankey plot using the networkD3 package with R version 4.1.1.
193 Results are presented using the WHO risk categories: <1 cfu/100 mL = low risk; 1 – 10
194 cfu/100 mL = intermediate risk; 11 – 100 cfu/100 mL = high risk; and >100 cfu/100 mL = very
195 high risk.

196

197 Table 1. Summary of Bangladesh water quality datasets used in the analysis. All data
198 collected and analysed by icddr,b.

	Site	System types	Sampling months	N of paired samples	Programme
1	Narail, Faridpur, Pirojpur	Household and community	02/2019	N=286	Baseline survey for ASWA-II (DFID, 2017)

	and Sylhet districts	tubewells with handpumps			
2	Rohingya refugee camp	Shallow and deep tubewells, taps on piped systems	01-02/2018, 10-11/2019, 01-02/2020, 01/02,2021	N=946	Quality assurance programme
3	Rohingya refugee camp	Shallow and deep tubewells with handpumps	03-12/2018	N=3161	Study of contamination pathways (Mahmud et al., 2019)
4	Matlab	Shallow and deep tubewells with handpumps	11/2019	N=125	School water quality assessment (Fischer et al., 2021)
5	Khulna	Shallow and deep tubewells, pond sand filter, taps on piped systems	01/2020, 06/2020, 03/2021	N=299	Seasonal assessment of community and school water points (Hoque et al., 2021)
6	Rohingya refugee camp	Tubewells with handpumps	01-02/2022	N=128	Bhasan Char site
7	Matlab	Tubewells with handpumps	04/2018	N=31	Water Quality Monitoring
8	7 hospitals across 7 districts	Deep tubewell supplying building piped water network	10-12/2022, 5-6/2023, 10-11/2023, 2/2024	N = 104	Hospital water quality

199

200 2.2 Contribution of household hygiene contamination pathways at the PoU

201 Data from MICS were accessed for 38 datasets that represent countries or areas that
 202 collected data on water quality for MICS round 6 (2021-23). Data were not population
 203 weighted so they represent the samples taken. MICS water quality sampling methodology
 204 uses PoC samples and PoU samples, for which household respondents are asked to provide
 205 a cup of water “as you would give a child to drink”. 100mL samples are analysed by
 206 membrane filtration using compact dry growth media plates and incubated with body heat
 207 (Bangladesh Bureau of Statistics and UNICEF Bangladesh, 2021; JMP/UNICEF/WHO, 2016).

208 For the analysis in this paper, two-tailed Pearson correlations on national rates of
209 contamination were performed in IBM SPSS v28. The increase in contamination between
210 PoC and PoU is widely reported, and from various contamination pathways. Here we
211 specifically focus on contamination pathways associated with household hygiene, excluding
212 water handling practices in the household such as storage. The analysis included only
213 households where a) water is not contaminated at the PoC (PoC = 0 *E. coli* per 100mL) and
214 b) PoU samples were collected from a cup that the respondent filled directly from the PoC
215 so that there is no household storage or treatment. Using this subset, the proportion
216 demonstrating contamination with *E. coli* at the PoU from these local hygiene-related
217 pathways was calculated.
218

219 2.3 Contribution of contamination pathways in unchlorinated piped water systems

220 Dataset 8 (Table 1) was collected at hospitals, sampling from various water access points on
221 the sites. All water originated from tubewells on site but were accessed in different ways
222 and for different purposes. These included: tubewell samples, including from handpumps (n
223 = 22), taps connected to piped systems for drinking water (n = 26), taps connected to piped
224 systems for healthcare use (n = 56). Piped systems used for healthcare use included storage
225 tanks. No disinfection was present in any of the systems. In addition to *E. coli* results, water
226 samples were analysed for bacterial indicators linked to the presence of biofilms (Makris et
227 al., 2014), including *Pseudomonas* are an opportunistic pathogen which is a particular
228 concern for immunocompromised patients which can form biofilms in piped systems and is
229 associated with hospital acquired infections (Loveday et al., 2014). Samples were taken as
230 for other studies. The analytical methods are briefly described. Total coliforms: 100mL
231 samples were filtered through a 0.22 μ m pore size membrane filter (Millipore Corp.), which
232 was then placed on membrane fecal coliform (MFC) agar plates. After incubation at 37°C for
233 18 to 24 hours, blue coloured colonies were counted (Islam et al., 2001). Total bacterial
234 count: 100 μ l samples were inoculated onto a nutrient agar plate using the drop plate
235 technique (Hoben and Somasegaran, 1982) and incubated at 37°C for 18 to 24 hours. The
236 total number of colonies were counted. *Pseudomonas aeruginosa*: 100 ml samples were
237 filtered using a 0.22 μ m pore size membrane filter (Millipore Corp.) and placed onto a
238 cetrimide agar (BD Difco) plate. The plate was incubated at 35 \pm 2°C for 18 to 48 hours. After
239 incubation, colonies surrounded by a blue-green pigment and fluoresced under short
240 wavelength (254 nm) ultraviolet light were presumptively identified and counted as
241 *Pseudomonas aeruginosa* (Crone et al., 2020; Zimbrow et al., 2009). This dataset was used to
242 explore the impact of decontaminating the tap in different systems including unchlorinated
243 piped systems. Descriptive statistics were used to assess the change in concentrations of
244 pathogens and indicators with decontamination.
245

246 3 Results

247 3.1 Contribution of contamination pathways from PoD and PoC

248 The analysis of secondary datasets at scale allowed assessment of the contribution of
249 contamination from (a) water catchment environment and infrastructure pathways as
250 detected at the PoD, and (b) local hygiene pathways as detected as additional
251 contamination at PoC. The eight studies provided 5,080 paired samples of water quality in
252 Bangladesh (Table 2). Contamination with *E. coli* from environment and infrastructure
253 pathways (PoD method) was detected in 7.3% of samples (n=370). Contamination with *E.*

254 *coli* in samples using PoC methods was present in 14% of samples (n=712), with local
 255 hygiene pathways associated with 48.2% of detection PoC. Only one PoD-PoC pair was *E.*
 256 *coli* detected in a PoD sample where it hadn't been detected in the PoC sample. Across the
 257 studies, 6.7% of taps were contaminated. Figure 2 demonstrates the change in risk class
 258 between PoD and PoC methods (for those taps where *E. coli* was detected at PoC): 60%
 259 classified as low risk (*E. coli* absent) at PoD, with an additional 16.0% having a reduced risk
 260 risk category after cleaning (lowered *E. coli* concentration).

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Table 2 Comparison of risk category from drinking water samples at PoD and PoC

	Risk class		Water as delivered (PoD)				Total
			Low	Intermediate	High	Very high	
Water as collected (PoC)	Low	N	4379	1	0	0	4380
		%	100.0	0.0	0.0	0.0	100
	Intermediate	N	297	123	2	0	422
		%	70.4	29.1	0.5	0.0	100
	High	N	40	73	58	0	171
		%	23.4	42.7	33.9	0.0	100
	Very high	N	6	9	32	72	119
		%	5.0	7.6	26.9	60.5	100
	Total	N	4722	4722	206	92	72
		%	92.7	92.7	4.1	1.8	1.4

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[insert here] Figure 2 Contribution of local hygiene contamination pathways at the tap: changes in risk category from drinking water samples at PoD and PoC, for the subset of samples with E. coli detected before decontamination.

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The majority of samples are from tubewells in the Rohingya refugee camps and are not representative of Bangladesh nationally, nor are they equally representative of all water system types. Across the eight studies, the proportion of samples with no *E. coli* detected at the PoD ranged from 40.5% to 97.9%, and for PoC ranged from 25.8% to 90.4% (Table 3), with variability by system type and season described below. The contribution of water catchment environment and infrastructure pathways varied, with high contamination in studies 6 and 8. High variability between the contribution of local hygiene pathways, as indicated by the difference in the risk level between PoD and PoC sampling, was also evident: ranging from 0% to 81.8%. More complex infrastructure was associated with a greater contribution of contamination from the environment and infrastructure pathway: water from water points connected to unchlorinated piped networks was more frequently and heavily contaminated compared to water from tubewells.

284 Table 3. Variation in contributions of contamination from different pathways

Study	n	PoD low risk	PoC low risk	Contamination from local hygiene pathways	Contribution of local hygiene pathways to overall contamination	Contamination increased by local hygiene pathways
		PoD = 0	PoC = 0	PoC-PoD	PoC >0, PoD = 0	PoC = > 0, PoD < PoC
		%	%	%	%	%
1	286	97.9	88.5	9.4	81.8	90.9
2	946	96.2	90.4	5.8	60.4	75.8
3	3161	95.5	89.6	5.9	56.8	72.0
4	125	94.4	88.8	5.6	50.0	85.7
5	299	89.0	78.3	10.7	49.2	49.2
6	128	43.0	25.8	17.2	23.2	54.7
7	31	87.1	87.1	0.0	0.0	0.0
8	116	40.5	30.2	10.3	16.0	30.9

285
 286 As seasonality has been found to significantly affect water quality (Charles et al., 2022),
 287 results were compared by sampling months for school, community and household settings
 288 (Table 4) excluding study #8. Across the studies, there were small seasonal variations in *E.*
 289 *coli*, with *E. coli* more commonly present in the colder, drier winter period of December to
 290 February. The contribution from water catchment environment and infrastructure pathways
 291 was greatest in the December to February period. The contribution from local hygiene
 292 pathways was greatest in the monsoon (June – August) and post-monsoon periods
 293 (September – November) periods. The contribution of local hygiene pathways was lowest in
 294 the pre-monsoon (March – May). This variability demonstrates the differing importance of
 295 these pathways seasonally.

296
 297 Table 4. Seasonal differences in contamination (excluding study 8)

Months	Average temperature (Climatic Research Unit (CRU), n.d.)		Average precipitation (Climatic Research Unit (CRU), n.d.)	PoD low risk		PoC low risk		Contamination from local hygiene pathways	Contribution of local hygiene pathways to overall contamination		Contamination increased by local hygiene pathways	
	Min	Max		PoD=0		PoC=0			PoD-PoC		PoC >0, PoD = 0	
	°C	°C	mm	n	%	n	%	%	n	%	n	%

DJF	13.4	26.6	30	1082	90.8	995	83.5	7.3	87	44.4	129	65.8
MAM	22.1	32.8	388	1859	94.9	1767	90.2	4.7	92	48.2	117	61.3
JJA	25.7	31.7	1226	1356	94.9	1233	86.3	8.6	123	62.8	148	75.5
SON	22.4	30.9	515	378	95.0	350	87.9	7.0	28	58.3	38	79.2
Total				4675	94.0	4345	87.3	6.6	330	52.3	432	68.5

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299 3.2 Contribution of household hygiene contamination pathways at the PoU

300 Data on PoC and PoU *E. coli* contamination was analysed based on MICS data from 38
 301 datasets (Table 5) to identify the contribution of household hygiene pathways, specifically
 302 hygiene of the user when filling a drinking water vessel and any contamination of the
 303 drinking water vessel. Across these countries and regions, samples from the PoC ranged
 304 from 89.6% without *E. coli* detected (i.e. low risk) in Turks and Caicos Islands to only 5.1% in
 305 Tuvalu. At the PoU, samples without *E. coli* detected ranged from 80.0% in Kosovo to only
 306 0.7% in Chad. To focus on the role of household hygiene pathways for contaminating water
 307 at the PoU, only households that collected the PoU water sample directly from the PoC
 308 were considered in further analysis. In the 38 datasets that were considered, the level of
 309 data that could therefore be included varied (Table 5). More than 70% of PoU samples were
 310 collected directly from the PoC in Georgia, Iraq, Kosovo, Samoa, Suriname and Tunisia. Less
 311 than a quarter of samples were collected directly from the source in 17 countries, with four
 312 countries excluded from the analysis due to insufficient data availability.

313

314 The proportion of households where no contamination was identified from household
 315 hygiene pathways (no detectable *E. coli* at the PoC that were also low risk for PoU samples
 316 collected from cups filled directly at the PoC) ranged from 29.5% in Nepal to 91.7% in Turks
 317 and Caicos Islands (Table 5). Figure 3 presents the variable impact of contamination from
 318 household hygiene pathways to drinking water at the PoU. In Nepal, 70.5% of households
 319 contaminated their drinking water through poor hygiene in collection of the water or poor
 320 hygiene of the cup. While there are correlations between the rate of contamination at PoC
 321 and PoU with national statistics on sanitation (open defecation and safely managed
 322 sanitation) and hygiene (basic access and no facility), there is no relationship between these
 323 sanitation and hygiene statistics and the number of households contaminating their water
 324 at the PoU (see supplementary material Table S2). PoU water quality was weakly correlated
 325 with the prevalence of collecting PoU samples from the source, supporting the argument
 326 that water sources on premises provide better water quality for users.

327

328 Table 5. Contamination of water quality at PoC & PoU based on MICS6 data

Country/area	Proportion of samples low risk (without <i>E. coli</i> detected)				PoU samples from cups filled directly at the PoC		No contamination identified from household hygiene pathways	
	PoC=0		PoU=0		n	%	n	%
	n	%	n	%				
Algeria	2641	84.3	2210	70.4	1518	48.4	1108	84
Bangladesh	3750	61.8	1101	18.1	345	5.7	75	33.9

Central African Republic*	323	30.8	131	12.5	64	6.1	8	61.5
Chad*	275	13.1	15	0.7	180	8.3	0	0
Democratic Republic of the Congo	1103	40.3	842	30.7	457	16.7	107	75.9
Dominican Republic	1126	44	581	22.7	507	19.8	137	55.7
Fiji	622	57.1	528	48.5	1035	95	483	79.7
Gambia	1021	57.9	489	27.7	64	3.6	35	71.4
Georgia	1590	65.5	1376	56.6	1804	74.3	1137	88.3
Ghana	1695	53.6	767	24.3	1011	32	527	72.9
Guinea-Bissau	657	36.8	333	18.7	102	5.7	18	47.4
Guyana	586	40.5	403	27.9	919	63.6	230	56.1
Honduras	2121	52.7	1672	41.5	2432	60.4	1196	73
Iraq	3866	57.8	3327	49.8	5365	80.2	2279	76.4
Kiribati	68	11.5	48	8.1	320	54.3	10	30.3
Kosovo	899	82.5	872	80	999	91.7	754	91.4
Laos	570	17.3	506	15.4	839	25.5	55	39.9
Lesotho	918	68.6	638	47.6	44	3.3	30	83.3
Madagascar	502	15.4	313	9.6	148	4.5	51	89.5
Malawi	1209	38.7	261	8.4	186	6	44	55.7
Mongolia	2106	81.1	2073	79.8	344	13.2	266	91.1
Nepal	524	21.4	310	12.7	585	24	64	29.5
Pakistan Baluchistan	379	14	201	704	734	27.2	40	52.6
Pakistan Khyber Pakhtunkhwa	537	16.3	176	5.3	508	15.4	26	39.4
Pakistan Punjab	4455	64.9	2859	41.7	3627	52.9	1446	61.4
Pakistan Sindh	979	41.2	405	17.1	597	25.1	114	40.1
Samoa	458	69.3	395	59.8	553	83.7	334	85
Sao Tome and Principe*	236	69	200	58.5	21	6.1	12	85.7
Sierra Leone	179	10.2	58	3.3	290	16.6	25	59.5

State of Palestine	1483	81.5	1265	69.5	1168	64.2	781	83.5
Suriname	888	54.8	563	34.8	1301	80.4	381	49.8
Togo	376	34.6	126	11.6	118	10.8	53	80.3
Tonga	153	28.2	115	21.2	361	66.5	54	57.4
Tunisia	2127	79.8	1916	71.9	2454	92.1	1700	85.3
Turks & Caicos Islands	232	89.6	205	79.2	137	52.9	111	91.7
Tuvalu*	8	5.1	17	10.8	78	49.7	0	0
Viet Nam	1670	50.5	1807	54.6	884	26.7	355	71.9
Zimbabwe	926	45.3	428	20.9	230	11.3	135	77.6
* Countries had fewer than 30 samples at PoU collected direct from a clean source. These results are not included in further analysis.								

329

330

331 *[insert here] Figure 3. Contribution of household hygiene contamination pathways:*

332 *proportion of households with water quality contaminated between the PoC and PoU*

333

334 3.3 Contribution of contamination pathways in unchlorinated piped water systems

335 The results above for PoD focused primarily on smaller infrastructure such as tubewells
 336 where results almost uniformly show reductions in contamination with *E. coli* after
 337 decontamination (PoC > PoD). However, in unchlorinated piped systems in hospitals,
 338 significant increases in contamination were observed after decontaminating taps, either by
 339 flaming or wiping with ethanol. In these systems, water quality from the source (or nearby
 340 tubewell) was typically low risk for *E. coli* at the point of delivery (81% for deep tubewell
 341 sources n=16; 77% for handpumps, n=9). Contamination at the taps used for healthcare was
 342 common with only 54% of taps free from *E. coli* contamination at the PoD and 50% at PoC.
 343 *Pseudomonas aeruginosa* were detected in water from 96% (n=54/56) healthcare use taps
 344 at PoC. Contamination was visible when wiping inside taps. Contamination in taps used for
 345 drinking water was slightly less common as these included taps on filtration systems. In
 346 these settings, contamination often increased after cleaning and flushing the taps (Figure 4),
 347 particularly for *Pseudomonas aeruginosa* which are associated with biofilm formation.

348

349

350 *[insert here] Figure 4. Change in concentration of organism in samples before and after*
 351 *cleaning and flushing taps for healthcare use in unchlorinated, piped hospital water systems*
 352 *(top), and in all taps and spouts sampled in hospitals (bottom)*

353

354 4 Discussion

355 The results demonstrate the influence of different contamination pathways on water quality
 356 sample results. Understanding the relative contributions of different contamination
 357 pathways influences the interpretation of different sample types and has important
 358 implications for drinking water safety management. Here, we focus on operational

359 understanding of water systems, however, the interpretation also relates to public health
360 and regulator understandings of health risks and appropriate interventions.

361
362 PoD samples characterise the water as delivered. It should not be interpreted as a 'source'
363 sample because it includes the impact of the engineered system (Bartram and Hunter,
364 2015), although it is often difficult to sample source water quality without using
365 infrastructure, especially in groundwater systems. For drinking water managers, PoD results
366 can be clearly interpreted as demonstrating the water quality that the water system
367 (source, treatment, distribution) is capable of providing. As such PoD samples can help
368 identify where interventions are required in the source, treatment, or distribution system,
369 including at the handpump or spring. PoD samples will not further differentiate between
370 source, treatment or distribution contamination pathways, so more distributed sampling is
371 recommended to identify vulnerabilities in complex systems such as piped networks (World
372 Health Organization, 2024).

373
374 PoC samples identify the user's risk of exposure to faecal contamination via drinking water
375 associated with where they collect their water from. Our results demonstrate that local
376 hygiene at the tap is a significant contamination pathway. This has implications for human
377 health. PoC samples are assumed to represent the quality of water that is available or is
378 collected (e.g. Rufener et al., 2010), however, there are limitations to this assumption.
379 Contamination on the tap will likely have high temporal variability due to both the nature of
380 the contamination source from people and animals accessing the tap, and the potential for
381 rapid decreases with die-off of bacteria due to heat and drying of the tap or washing of the
382 tap during use. The authors' fieldwork observations of user behaviour at the tap have
383 indicated that people (and animals) regularly touch water supply outlets, with different
384 collection vessels, priming methods, cleaning practices and mobility impacting the potential
385 for transference of contamination onto the outlet and therefore into the water collected.
386 For drinking water managers, results from PoC samples cannot differentiate between the
387 need to clean the tap, and the need for interventions to the water system.

388
389 PoU samples identify contamination in drinking water as consumed by a child, which
390 includes contamination originating from the water system and local hygiene, from
391 household water management, and from household hygiene including contamination on
392 the cup and on hand hygiene. Our analysis demonstrates that the contribution from
393 household hygiene varies, with household hygiene contributing contamination in up to 70%
394 of samples. For drinking water managers, understanding faecal contamination at the PoU
395 does not aid operational decisions due to the multiple contamination pathways involved,
396 however, in-household sampling approaches remain important to understand chlorine
397 residuals and contamination from the infrastructure, such as lead.

398
399 Consistent reporting of sampling methods is essential to enable appropriate interpretation
400 of results. In the literature, PoU sampling methods are generally clearly communicated.
401 However, there is often poor description of sample collection and processing (Sikder et al.,
402 2021), with sampling methodologies to differentiate PoD and PoC methods often unclear.
403 These two methods are not differentiated further in systematic reviews of drinking water
404 quality (e.g. Bain et al., 2014; Kostyla et al., 2015; Sikder et al., 2021; Wright et al., 2004),
405 with each using a combination of papers that use either PoD or PoC or do not report the

406 sampling method clearly. This poor reporting can be found in current papers, for example,
407 Murei et al. (2024) report on tap water quality samples and draw conclusions about
408 relationships with sanitary inspections with no details of the sampling method. Our results
409 (Figure 2) demonstrate that around 60% of sources that are contaminated with *E. coli* at the
410 PoC are free from contamination at the PoD. This large difference makes it imperative for
411 research and practice to explicitly communicate the sampling methods used to enable
412 useful interpretation and comparison of results. Sampling guidance should be updated to
413 clearly differentiate the types of samples, purposes and how to report them.

414
415 Cleaning the tap or spout is not just a sampling practice but should be a standard part of
416 hygiene education programmes. Contamination of communal water points from poor
417 hygiene presents a risk of disease transmission, between households and also between
418 vulnerable individuals at shared taps at schools and hospitals. However, there is a gap in the
419 guidance and programme documents on cleaning at the point of water distribution.
420 Literature addresses hygiene education, with a focus on the household, or infrastructure
421 operation and maintenance, but cleaning the tap is overlooked. For example, WaterAid's
422 Technology Notes (WaterAid, n.d.) provide guidance on hygiene education and water source
423 development, but do not mention cleaning handpumps. In guidance for small-community
424 supplies (WHO, 1985) it is advised that "improved water sources should be used
425 hygienically" but there is no reference to cleaning. Early drafts of current sanitary inspection
426 guidance (World Health Organization, 2024) propagated that approach, with hygiene only
427 focused on household practices. This was revised to include cleaning as a daily action for
428 management of water systems based on discussion of the preliminary results presented in
429 this paper.

430
431 Further guidance might consider recommending handwashing before water collection to
432 reduce contamination of water points. Our results demonstrate that flaming can remove
433 the contamination, but more research is needed to understand how frequently and what
434 materials are needed to reduce the risk of disease spread. Anecdotal evidence from
435 partners in Uganda suggests washing with just water can remove contamination, which is a
436 practice that is appropriate for the individual user. Understanding the drivers of
437 contamination, how quickly water collection outlets are contaminated after cleaning and
438 the efficacy of different cleaning methods could inform the design of interventions that can
439 improve water safety for users. Our analysis demonstrates that improving guidance on
440 cleaning taps could substantially increase access to safer drinking water: In Bangladesh, we
441 estimate an additional 35 million people would be considered to have access to safe
442 drinking water (i.e. without *E. coli*) if the water collection outlets were cleaned, with a
443 further 11 million people having access to cleaner drinking water with lower concentrations
444 of *E. coli*. This would lower the proportion of people nationally without safe water from 40%
445 to 19%.

446
447 Piped drinking water access has increased substantially in recent years, with an additional
448 1.8 billion people estimated to have access to piped water since 2000 (UNICEF/WHO JMP,
449 2023). However, the quality of that water is not always free from faecal contamination, with
450 chlorination not routinely practiced, creating new challenges in managing water safety. The
451 results from the unchlorinated piped systems in hospitals that are presented in this paper
452 show that tap cleaning methods such as flaming (for metal fittings) and using ethanol wipes

453 (for plastic fittings) increased the concentrations of bacteria in drinking water samples,
454 particularly for bacteria that are associated with biofilms, even after flushing (Figure 4). For
455 *Pseudomonas aeruginosa* over 70% of samples demonstrated increased concentrations
456 after cleaning the tap. It was not possible to do repeated samples to measure how
457 concentrations vary over time post-cleaning, and further PoD sampling was stopped to
458 avoid increasing risks for patients at the facilities. Similar results have been described by
459 Wang et al. (2012) with an increase in bacteria liberated into the water from disrupted
460 biofilm post-cleaning. Further research is needed to understand what sampling methods are
461 appropriate in these settings. And with limited knowledge of the impact of decontamination
462 on the scale and duration of release of such pathogens such as *Pseudomonas* and
463 *Legionella*, decontamination should be attempted with caution in distribution systems
464 serving vulnerable populations e.g. elderly, the immune-compromised patients in hospitals.
465

466 5 Conclusion

467 We demonstrated the substantial contamination of drinking water from local hygiene
468 conditions at the point of collection. Local hygiene sources, such as from people and animals
469 touching the tap, contributed to contamination of drinking water in 76% of water samples
470 where *E. coli* was detected, including being the sole source of contamination in an average
471 of 60% of samples. This impact of hygiene is also notable in samples from within the
472 household, at the PoU, where hand- and household-hygiene is the primary source of
473 contamination in some contexts, but with high variation by context. These pathways of
474 contamination related to hygiene have public health implications for drinking water users,
475 with a variety of interventions to address them. Recognising the contributions of these
476 pathways, and embedding this in sampling programmes, can improve understanding of the
477 contextual and seasonal drivers of contamination and support advances towards safer
478 drinking water.
479

480 One key intervention that could improve drinking water quality is cleaning the tap or spout,
481 especially before collecting drinking water. Behaviour interventions and hygiene have
482 focused on water management in the household and overlooked behavioural aspects at the
483 point of collection. It is important to note the limitations in the evidence, however, with the
484 potential for unintended consequences where cleaning may impact on biofilms resulting in
485 release of pathogens. There is increasing application of complex water supply infrastructure
486 in piped water systems without chlorination or other treatments to minimise biofilm
487 formation. More research is needed to refine guidance on the frequency and methods of
488 cleaning, and implications for systems with substantial biofilm growth. However, this does
489 not negate the immediate benefit to be gained from expanding communication and hygiene
490 education to encourage cleaning of the tap.
491

492 A key part of recognising the contribution of hygiene pathways is to improve report. To
493 date, there are inconsistencies in reporting drinking water sampling methods across the
494 literature. We call on authors to ensure transparency by reporting if they use
495 decontamination techniques as part of water sampling methods and urge caution in
496 drawing comparisons across studies using different or unclear methods.
497

498 Data statement

499 The MICS datasets analysed during the current study are available from the MICS website,
500 <https://mics.unicef.org/>

501

502 The icddr,b datasets in Table 1, used and analysed during the current study, are available
503 from the corresponding author on reasonable request.

504 Author Contribution Statement

505 K.C. and D.J. conceptualised the study. K.C., S.N. and L.O. contributed to the study design
506 and conceived the analysis. Z.M., L.O. and K.C. collected and curated the data. K.C.
507 performed the analyses. K.C. and S.N. undertook visualisation of the published work. K.C.
508 and D.J. supervised the study. K.C. wrote the initial draft of the manuscript. K.C., S.N., L.O,
509 Z.M. and D.J. contributed to reviewing and editing the manuscript.

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517

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- 659
660

List of Figures

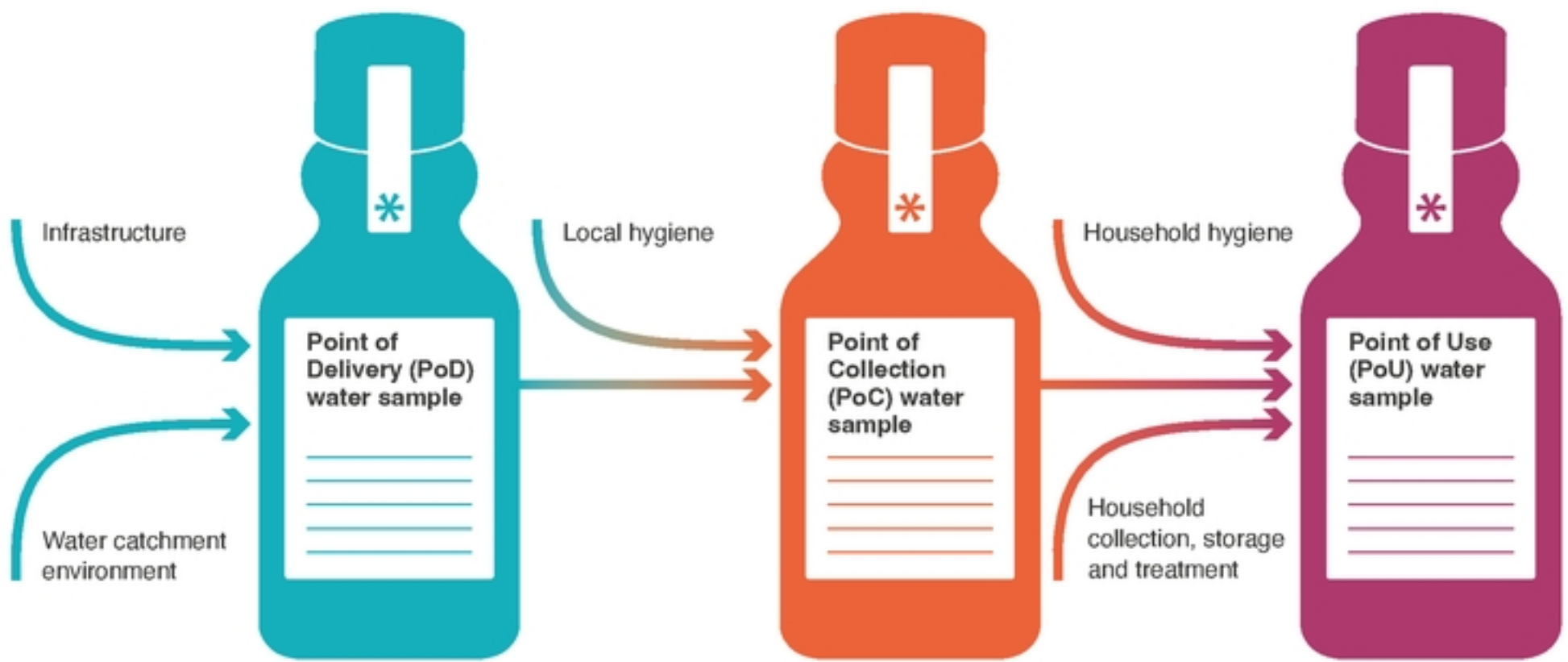
Figure 1. Drinking water contamination pathways contributing to sample results at PoD, PoC and PoU

Figure 2. Contribution of local hygiene contamination pathways at the tap: changes in risk category from drinking water samples at PoD and PoC, for the subset of samples with E. coli detected before decontamination.

Figure 3. Contribution of household hygiene contamination pathways: proportion of households with water quality contaminated between the PoC and PoU

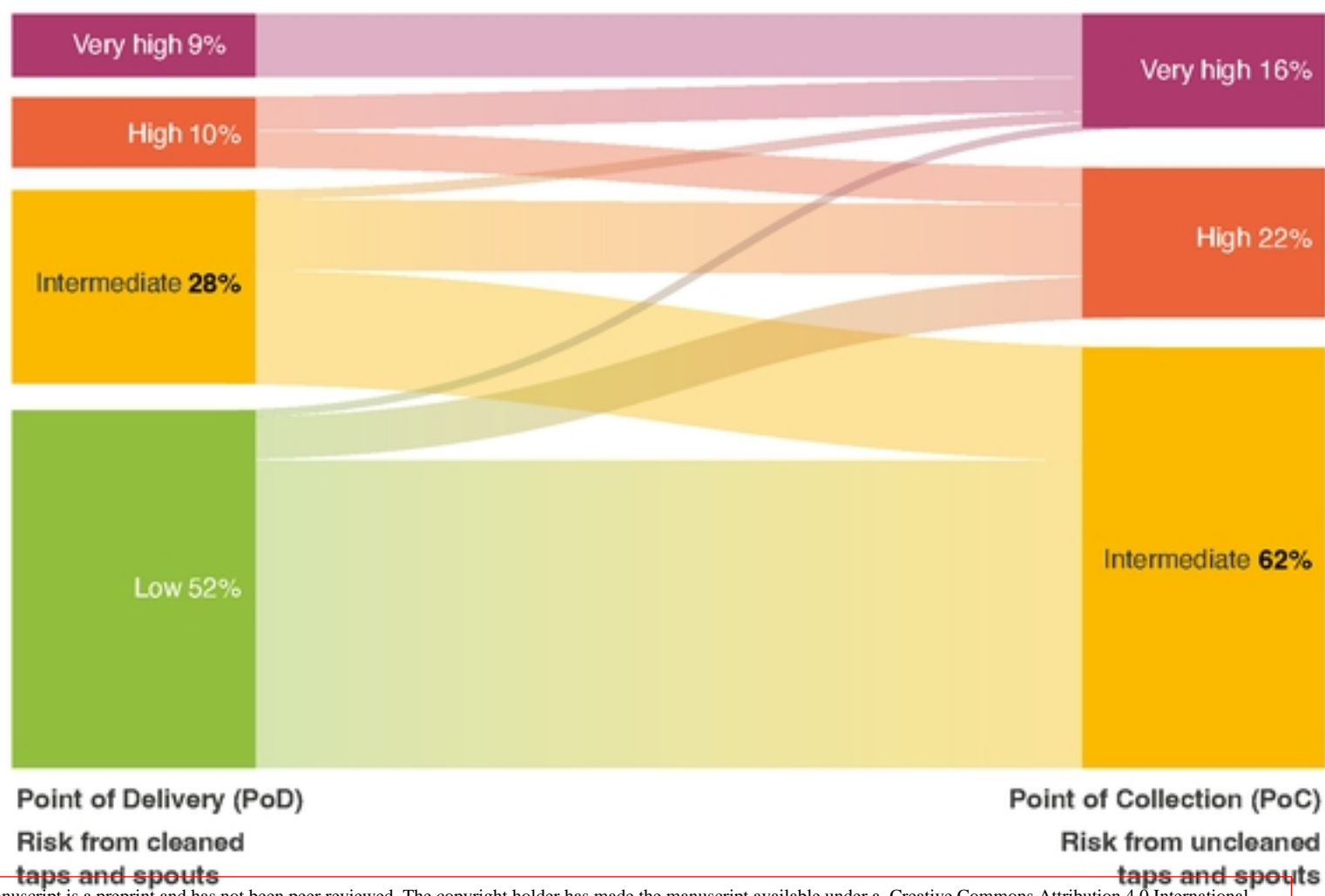
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Figure 4. Change in concentration of organism in samples before and after cleaning and flushing taps for healthcare use in unchlorinated, piped hospital water systems (top), and in all taps and spouts sampled in hospitals (bottom)



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Figure 1



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Figure 2

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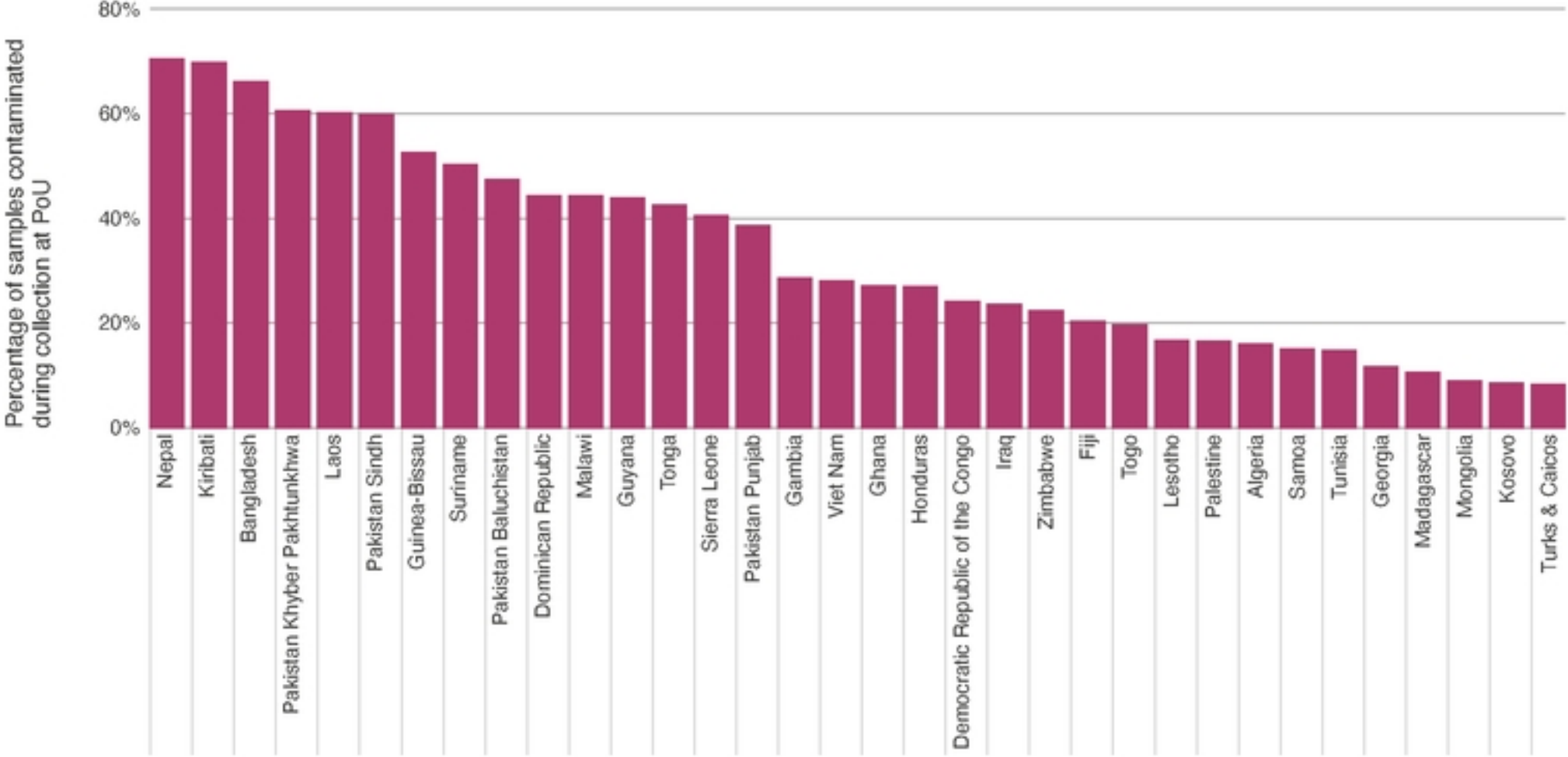


Figure 3

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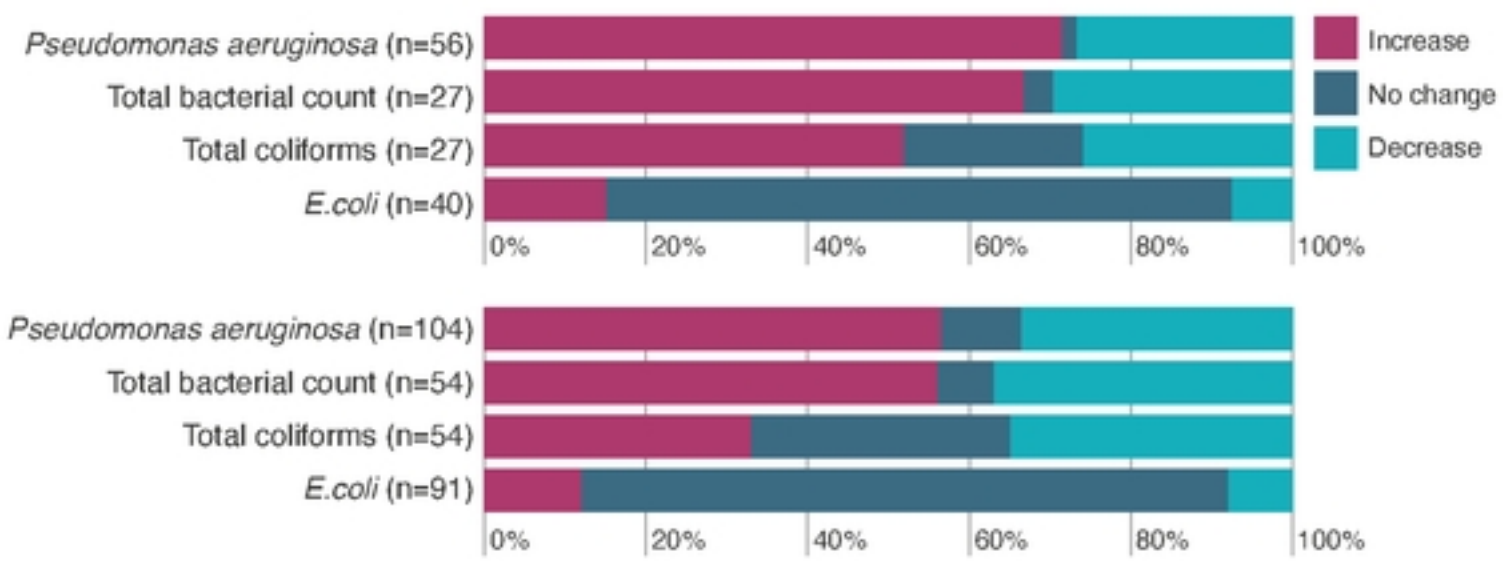


Figure 4