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A QUANTITATIVE MICROBIAL RISK ASSESSMENT MODEL FOR TOTAL COLIFORMS AND *E. COLI* IN SURFACE RUNOFF FOLLOWING APPLICATION OF BIOSOLIDS TO GRASSLAND

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27 **Abstract**

28 The land application of treated municipal sludge ('biosolids') may give rise to surface runoff
29 containing coliforms during episodic rainfall events, which may be potentially harmful to human
30 health if not fully treated in a water treatment plant (WTP). This study used surface runoff water
31 quality data generated from a field-scale study in which three types of biosolids (anaerobically
32 digested (AD), lime stabilised (LS), and thermally dried (TD)) were spread on micro-plots of
33 land and subjected to three rainfall events at time intervals of 24, 48 and 360 hr following
34 application. Under the assumption that this water directly entered abstraction waters for a WTP
35 without any grassed buffer zone being present, and accounting for stream dilution, die-off rate
36 and modelling various performance scenarios within the WTP, the aim of this research was to
37 conduct a human health risk assessment of coliforms (total and faecal), which may be present in
38 drinking water after the WTP. Two dose response models for probability of illness were
39 considered for total and faecal coliform exposure incorporating two different exposure scenarios
40 (healthy populations and immuno-compromised populations). The simulated annual risk of
41 illness for healthy populations was below the US EPA and World Health Organisation tolerable
42 level of risk (10^{-4} and 10^{-6} , respectively). However, immuno-compromised populations may still
43 be at risk as levels were greater than the tolerable level of risk for that subpopulation. The
44 sensitivity analysis highlighted the importance of residence time in a stream on the bacterial die-
45 off rate.

46

47 **Keywords:** biosolids, coliforms, risk, surface-runoff, water treatment

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55 **Introduction**

56 The application of treated municipal sewage sludge (“biosolids”) to agricultural land as a
57 fertiliser can offer an excellent source of nutrients (nitrogen, phosphorus and potassium),
58 increase organic matter and water absorbency, and reduce the possibility of soil erosion. It is also
59 a cost-effective way to dispose of municipal waste and reduce over-reliance on landfill whilst
60 cutting down on tipping fees. However, biosolids can also be non-point source contributors of
61 heavy metals, human pathogens and xenobiotics (Clarke and Cummins, 2014; McCall et al.,
62 2015; Peyton et al., 2016). Therefore, it is imperative that all biosolids are effectively treated to
63 remove pathogens and contaminants to a “safe level” prior to being used as a land conditioner or
64 fertiliser.

65 More than 10 million tonnes of sewage sludge was produced in the European Union (EU) in
66 2010 (Eurostat, 2014). Although EU policy favours the recycling of resources (COM, 2014),
67 including sludge, national sludge recycling policy varies throughout Europe. In some countries,
68 such as the Republic of Ireland, up to 80% of sludge is reused in agriculture (Eurostat, 2014),
69 whereas in other countries, such as Germany, the land application of sludge is prohibited. This is
70 due to the considerable public acceptance issues surrounding the reuse of treated sludge as a
71 fertiliser. The main fear is that the presence of organic and inorganic contaminants in biosolids
72 may accumulate in the food chain, or cause the contamination of soil and water (Clarke et al.,
73 2015). The US Environmental Protection Agency (USEPA) Part 503 regulations classify
74 biosolids according to Class A and Class B standard. Class A biosolids contain a faecal coliform
75 density below 1000 most probable number (MPN)/ g of total solids (dry matter, DM), whereas
76 Class B biosolids contain a geometric mean faecal coliform density of less than 2×10^6 MPN / g
77 of total solids (DM) (USEPA 2006). In the USA, the land application of certain types of
78 biosolids requires a class B designation, which must satisfy three different criteria, one of which
79 includes faecal coliforms whose level cannot exceed 2×10^6 MPN/g (Pascual-Benito et al.,
80 2015). In the EU, sewage sludge production is regulated by the Sewage Sludge Directive
81 86/287/EC. It does not specify limits for pathogens but instead specifies general land use,
82 harvesting and grazing limits to provide protection against the risk of infection (Sobrados-
83 Bernardos and Smith, 2012). A revision of the Sewage Sludge Directive (Working Document
84 3rd Draft) states that “the use of microbial indicators to evaluate the hygienisation of treated

85 sludge is based on fulfilling the limits of *E. coli* to achieve a 99.9% reduction and to less than $1 \times$
86 10^3 cfu/g dry weight, produce a sludge containing $< 3 \times 10^3$ spores of *Clostridium perfringens*/g
87 (DM) and absence of salmonella. spp in 50 g (DM)” (EC 2000). Furthermore, the Working
88 Document also states that sludge produced by conventional treatment shall at least achieve a 2
89 \log_{10} reduction of *E. coli* (Mininni et al., 2014). European countries are allowed to include their
90 own parameters in their national regulations (Pascual-Benito et al., 2015). For instance, in France
91 the standards for maximum concentrations of pathogens in biosolids cannot exceed 8 MPN/10 g⁻¹
92 DM for *salmonella*, whereas in Finland, the number of *E. coli* must be less than 1000 cfu and
93 *Salmonella* must not be detected in 25 g of biosolids (Mininni et al., 2014). Meanwhile, in
94 Ireland the standards for maximum concentrations must not exceed 1×10^3 MPN g⁻¹ which is
95 equivalent to Class B biosolids under the USEPA Part 503 regulation (Fehily Timoney and
96 Company 1999).

97 Following land-spreading of biosolids, there are two main scenarios which can lead to human
98 infection. First, pathogens may be transported *via* overland or sub-surface flow to surface and
99 ground waters, and infection may arise *via* ingestion of contaminated water or accidental
100 ingestion of contaminated recreational water (Jaimeson et al. 2002; Tyrrel and Quinton 2003).
101 Alternatively, it is possible that viable pathogens could be present on the crop surface following
102 biosolids application, or may become internalised within the crop tissue, where they are
103 protected from conventional sanitization (Itoh et al. 1998; Solomon et al. 2002). In this case, a
104 person may become infected if they consume the contaminated produce. Faecal coliform
105 numbers in the stabilised biosolids can be high, up to 10^5 g⁻¹ DM (Schwarz et al., 2014). Gerba
106 and Smith (2005) reported general survival times for bacteria in soil to be 2-12 months, whilst
107 Lang et al. (2007) reported survival times of enteric micro-organisms in sludge-amended soil
108 varying between 24 hours to 2 years. The disparities in survival rates are difficult to define due to
109 “knowledge gaps” with regards to decay mechanisms and the complex interactions between the
110 environment and soil-specific factors that result in the decay of enteric bacteria (Schwarz et al.,
111 2014). Therefore, it is critical to accurately determine the pathogen risk associated with land
112 application of sewage sludge to fully understand the potential for environmental loss and
113 consequently, human transmission.

114 Coliforms are bacteria that are always present in the digestive tract of animals including humans,
115 and are found in their waste. They are also found in soil and plant material. Total coliform (TC)
116 bacteria are common in the environment and, with a few exceptions, are generally harmless
117 (USEPA 2013). They are typically used as an indication of other pathogens in drinking water.
118 Faecal coliform bacteria are gram negative, non-spore forming rods that are found in the
119 intestines and faeces of humans and other warm blooded animals. In general, human faecal waste
120 gives rise to the highest risk of waterborne diseases (Odonkor and Ampofo, 2013). The
121 predominant faecal coliform is *Escherichia coli* (USEPA 2006). *E. coli* is currently recognised
122 by the World Health Organisation (WHO) as the best faecal indicator bacteria for monitoring
123 faecal contamination of drinking water and faecal coliforms are suggested as an acceptable
124 alternative (WHO 2011). *E. coli* is found in all mammal faeces at concentrations of 10^9 g^{-1} , but
125 does not multiply significantly in the environment (Edberg et al., 2000). High levels of these
126 bacteria indicate the presence of pathogens that cause waterborne diseases (Selvaratnam and
127 Kunberger, 2004). Most coliform bacteria do not cause disease; however, some rare strains of *E.*
128 *coli*, particularly O157:H7, can cause serious illness. As few as 10 cells can cause serious illness
129 or even death (Liu et al., 2008). Diseases and illness that can be contracted in water with high
130 faecal coliform counts include typhoid fever, hepatitis, ear infections (Oram, 2014),
131 gastroenteritis and, dysentery (Gruber et al., 2014).

132 The WHO recommends that either *E. coli* or faecal coliforms be used as indicators of faecal
133 contamination of water. The WHO guideline value for faecal coliforms (none detected in any
134 100 ml sample) is reflected in the standards of most OECD members and low-middle income
135 countries (Bain et al., 2014). In their Guidelines for Drinking Water Quality, the WHO have
136 developed a risk classification to prioritise interventions as higher levels of indicator organisms
137 are generally indicative of greater levels of faecal contamination. The risk classification is based
138 on the number of indicator organisms in a 100 ml sample which includes <1 'very low risk', 1-
139 10 'low risk', 10-100 'medium risk', > 100 'high risk' or 'very high risk' (WHO, 2011).

140 During wastewater treatment, the sludge component of the waste becomes separated from the
141 water component. As the survival of many microorganisms and viruses in wastewater is linked to
142 the solid fraction of the waste, the numbers of pathogens present in sludge may be much higher
143 than the water component (Straub et al. 1992). Although treatment of municipal sewage sludge

144 using lime, anaerobic digestion, or temperature, may substantially reduce pathogens, complete
145 sterilisation is difficult to achieve and some pathogens, particularly enteric viruses, may persist.
146 Persistence may be related to factors such as temperature, pH, water content (of treated sludge),
147 and sunlight (Sidhu and Toze, 2009). Similarly, there is often resurgence in pathogen numbers
148 post-treatment, known as the ‘regrowth’ phenomenon. Taskin et al. (2011) reported a sudden
149 increase in *E. coli* density in anaerobically digested (AD) biosolids immediately after high speed
150 centrifuge dewatering, a phenomena known as ‘reactivation’ and is separate from growth during
151 the storage of dewatered biosolids cake. There are also links to contamination within the
152 centrifuge, reactivation of viable, but non-cultural, organisms, storage conditions post-
153 centrifugation (Zaleski et al. 2005), and proliferation of a resistant sub-population due to newly
154 available niche space associated with reduction in biomass and microbial activity (McKinley and
155 Vestal 1985). Iranpour and Cox (2006) observed reoccurrence of faecal coliforms in post-
156 digested biosolids from thermophilic anaerobic digestion treatment. The explanations for
157 reoccurrence may be linked to 1) incomplete destruction of the faecal coliforms during treatment,
158 2) contamination from external sources during post-digestion, or 3) a large drop of the post-
159 digestion biosolids temperature to below the maximum for faecal coliform growth.

160 The European Drinking Water Directive 98/83/EC states that drinking water entering the
161 distribution system should contain zero coliforms and zero *E. coli* in 100 mls (EC 2000). Despite
162 advances in drinking water treatment, the WHO estimates that about 1.1 billion people globally
163 drink unsafe water and the vast majority of diarrhoeal disease (88%) stem from unsafe water,
164 lack of hygiene and sanitation (Ashbolt, 2004).

165 The objective of this work was to develop a quantitative microbial risk assessment (QMRA)
166 model for coliforms in drinking water assuming the application of biosolids to agricultural land
167 and resulting surface runoff entered abstraction waters for a water treatment plant (WTP).

168 **Materials and methods**

169 **Model development**

170 A quantitative drinking water treatment model was developed that was capable of predicting
171 likely human exposure and resulting risk from TC and *E. coli* present in the drinking water
172 without the possibility for attenuation to waters used for WTPs. This represents a pessimistic

173 scenario as, in reality, biosolids would not be spread to the edge of the field and that grassed
174 buffer zones would be in place. The model was created in Microsoft Excel 2010 with the add-on
175 package @Risk (version 6.0, Palisade Corporation, New York, USA). Uncertainty and
176 variability can be accounted for in the model by means of probability density distributions and
177 are represented in the model's equations by name (e.g. triangular, uniform). Data from peer
178 reviewed scientific literature were incorporated at various steps of the drinking water treatment
179 (i.e. coagulation and flocculation, sedimentation and disinfection). A process-based approach to
180 modelling TC and *E. coli* fate and human exposure considers total concentration in surface
181 runoff, dilution rate, bacteria die-off rate, drinking water treatment (primary, secondary and
182 tertiary) and human consumption (adult and child).

183 **Biosolid and soil characterisation**

184 Three types of biosolids were investigated in this study. They were: anaerobically digested
185 biosolids from the UK (AD-UK) and Ireland (AD-IRE), and lime stabilized (LS) and thermally
186 dried (TD) biosolids. With the exception of ADUK, all biosolids originated from the same
187 wastewater treatment plant (WWTP) in Ireland. The ADUK biosolids were sourced from the
188 UK, and were used as part of an EU-funded FP7 project (END-O-SLUDG, 2014). The sludge
189 was collected and land applied to small field plots at the maximum legal application rate in
190 Ireland (Fehily Timoney and Company, 2014) and subjected to three successive rainfall events,
191 applied using a rainfall simulator, at time intervals of 24, (RS1) 48 (RS2) and 360 (RS3) hr after
192 application (Peyton et al., 2016). A soil-only control was also included in the experimental
193 design.

194 Three different scenarios (worst case, xxx and yyy) were completed to account for the
195 differences in time and surface runoff volumes. The mean and standard deviation of the surface
196 runoff (C_{surface-runoff}) of TC and *E. coli*, as measured by Peyton et al. (2016), is shown in
197 Table 1. Runoff results indicated that the AD-UK biosolids had significantly higher
198 concentrations of *E. coli* in the RS1 and RS2 rainfall events, and exceeded the recommended
199 standards of $> 1 \times 10^3$ MPN g⁻¹ (Fehily Timoney and Company 2014). All of the reported Irish
200 biosolids were some 10-fold below the Class A Irish standard (Peyton et al., 2016).

201

202 **Table 1.** Mean and standard deviation for total and faecal coliform in surface water.

Total coliforms			
Mean and standard deviation (n=15) (MPN/ 100 mls)			
	RS1	RS2	RS3
AD-UK	171840 ± 158962	133516 ± 247832	134860.6 ± 119499
TD	299620 ± 511723.2	615760 ± 629487.1	980600 ± 822835.8
LS	15858 ± 27155.13	628400 ± 820378.8	492000 ± 614760.4
AD-IRE	155220 ± 163536.4	309934.4 ± 503104	197840 ± 190432.9
Control	158220 ± 121426	32850.4 ± 22214.2	470360 ± 506376
<i>E. coli</i>			
Mean and standard deviation (n = 15) (MPN/ 100 mls)			
	RS1	RS2	RS3
AD-UK	7055.4 ± 10283.15	4476 ± 5622	210.6 ± 419.6
TD	456 ± 804.3	114 ± 106	44.6 ± 94.23
LS	138.2 ± 21.5	358.2 ± 730.8	39 ± 61
AD-IRE	14.8 ± 21.4	271.6 ± 518.6	199.6 ± 440.7
Control	34.2 ± 47	30.4 ± 51.8	4 ± 8.9

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205 As a “worst case scenario” it was assumed that surface runoff following biosolid application
 206 entered an adjacent stream without any chance of attenuation along the transfer continuum before
 207 delivery to the surface water body. This is atypical in terms of grassland management. Schueler

208 et al. (2000) reported on the effectiveness of stream buffers and faecal coliform removal, and
209 found that grass filter strips were effective in removing up to 70% of faecal coliforms. Similarly,
210 Coyne et al. (1995) found that grass filter strips removed up to 74% of faecal coliforms from
211 surface water. However, concentrations of faecal coliforms in surface water still exceeded
212 minimum concentration standards for primary water.

213 It was assumed that the runoff effluent in surface-water was then abstracted to a nearby WTP. To
214 account for TC and *E. coli* concentrations in surface water being discharged into the stream, this
215 study used a dilution factor (DF), which is the ratio of concentration in the effluent to
216 concentration in the receiving water after mixing in the receiving water (Colman et al., 2011).
217 This assumes a homogenous distribution of the bacteria in the river and does not account for
218 dispersion or advection. Dilution factors can vary between 1 (dry river bed in summer) up to
219 100,000. The EU Technical Guidance Document on Risk Assessment (2003) states that where
220 there is a lack of specific data, a default dilution value of 10 is recommended for sewage from
221 municipal WTPs when predicting environmental concentrations of contaminants in receiving
222 waters (EC 2003). Therefore, a default dilution factor of 10 was applied to the data to calculate
223 the predicted environmental concentrations in surface water (Eq.1).

$$224 \quad \text{PEC}_{\text{surface-water}} \text{ (MPN/100 mls)} = C_{\text{surface-runoff}} / \text{DF} \quad \text{Eqn. 1}$$

225 Where $\text{PEC}_{\text{surface-water}}$ is the concentration of coliforms (TC and *E. coli*) in surface waters
226 receiving wastewater effluent, DF is the dilution factor, and $(C_{\text{surface-runoff}})$ (MPN/100 mls) is the
227 concentration in surface water

228 The first order decay equation often used to describe bacterial die-off is expressed as Chick's
229 Law, and is used to describe the survival (die-off rate) of TC and *E. coli* in soil, manure, streams
230 and groundwater over time (Benham et al., 2006). Die-off is a function of temperature, nutrient
231 levels, competing bacteria and solar radiation (Hrudey, 2004). The rate of bacterial "die-off" is
232 greater in summer than winter due to higher temperatures and increased UV light (Murphy et al.,
233 2015). Wilkinson et al. (1995) reported enhanced coliform concentrations in streams during high
234 and rising flows following storm events. The die-off rate in stream (D-off) was calculated
235 according to Eq. 2:

$$236 \quad N_t = N_0 e^{(-kt)} \quad \text{Eqn. 2}$$

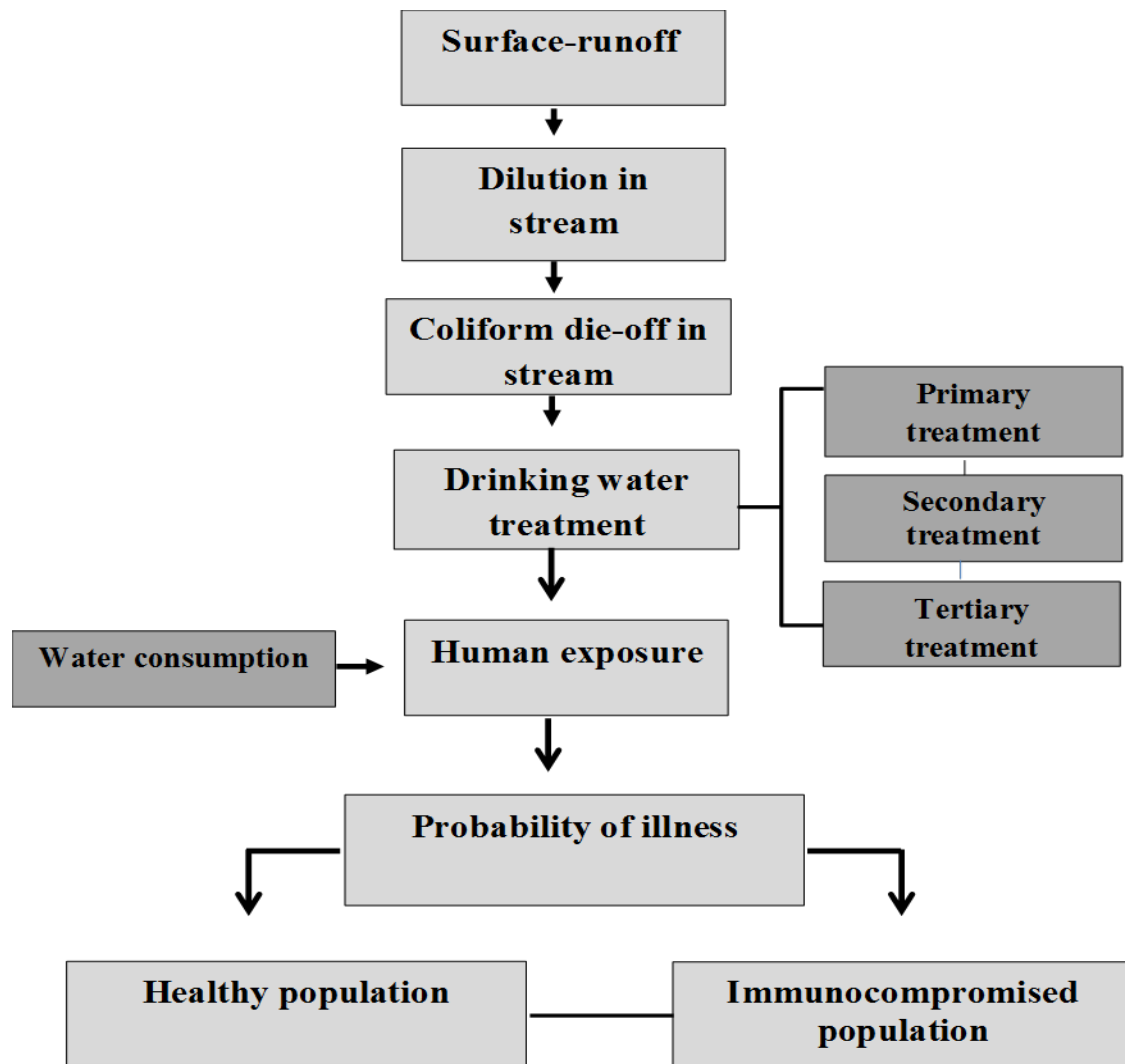
237 Where N_t is the number of coliforms at time t in surface-water (MPN/100 mls), N_0 is the original
238 number of coliforms following dilution in surface-water ($PEC_{\text{surface-water}}$) (MPN/100 mls), k is the
239 first order inactivation constant (d^{-1}), and t is the time in the stream (d^{-1}).

240 The k value was incorporated according to Schueler (2000), using a uniform distribution (values
241 min 0.7 and max 1.5 d^{-1}). k values in this range mean that about 90% of the bacteria present will
242 disappear from the water within 2 to 5 days. Therefore, it was assumed that water was abstracted
243 for drinking water treatment from the stream to a nearby WTP between 0 and 5 days. To account
244 for uncertainty, the time in stream “ t ” was fitted with a uniform distribution (min 0, max 5 day^{-1}).

245 **Drinking water treatment processes**

246 There are typically three stages to drinking water treatment (primary, secondary and tertiary)
247 (Figure 1). The three stages of drinking water treatment that were used were based on the Irish
248 Environmental Protection Agency’s (EPA) best practice guidelines for drinking water treatment
249 manuals (EPA 1995, 2002, 2011). It is assumed that operations within the drinking water
250 treatment process are running efficiently or stable (C-opt). However, to account for inefficiencies
251 in treatment operations, a sub-optimal (CS-opt) and failure (C-fail) option were incorporated into
252 the model. Poor operation of filters and inadequate disinfection may pose a risk to human health.
253 In recent times, many WTPs have become automated.

254



255

256 **Figure 1.** Flow diagram of the quantitative risk assessment drinking water model for coliforms
 257 in biosolids applied to grassland.

258

259 The first stage (primary treatment) considers the screening, storage, pre-conditioning and pre-
 260 chlorination of the water. In the current study, primary treatment was assumed to have a
 261 negligible impact on coliform removal. Secondary treatment involves the coagulation,
 262 flocculation, sedimentation and filtration of the influent. Coagulation/flocculation, sedimentation
 263 and filtration remove particles, including microorganisms (bacteria, viruses and protozoa) (WHO
 264 2011). The commonest types of coagulants used are aluminium-based (e.g. aluminium sulphate
 265 (alum) or polyaluminium chloride (PAC)). Both aluminium and ferric salts, either in monomer or
 266 polymeric forms, have been reported as effective coagulants in treating wastewater (Kang et al.,

267 2003; Pang et al., 2009). When properly performed, coagulation, flocculation and sedimentation
 268 can result in 1-2 log removal of bacteria, viruses and protozoa (WHO 2004). In accordance with
 269 the Irish EPA’s guidance manual (Ireland EPA 2002), the coagulant considered was aluminium
 270 sulphate ($Al_2(SO_4)_3$) (referred to as alum) for both TC and *E. coli*.

271 As faecal coliforms are the indicator organism for *E. coli*, reductions in *E. coli* counts were used
 272 to account for variability and uncertainty in the data. Pritchard et al. (2010) compared the
 273 efficacy of alum sulphate to more natural coagulants and reported *E. coli* reductions of 89%
 274 using 30-50 mg L⁻¹ of alum sulphate. Bulson et al. (1984) reported removal rates of *E. coli* of
 275 99.99% following a dose of 15 mg L⁻¹ of alum sulphate. A study conducted by Sarpong et al.
 276 (2010) showed that total coliform counts were reduced by 95% using a 5 ml dose of alum
 277 sulphate. Similarly, Bergamasco et al. (2011) reported a 99% reduction in total coliforms using a
 278 15 ml dose of alum sulphate. Thus, a uniform distribution was used to model coagulation,
 279 flocculation and sedimentation incorporating a decimal reduction to account for variability and
 280 uncertainty in the data (min 0.89, max 0.99).

281 As a “worst case scenario”, the model assumes a 90% probability of coagulation and flocculation
 282 occurring at an optimum stable run (Copt) and 5% probability for both sub-optimal (CS-opt) and
 283 failure (C-fail) (Table 2). When operating optimally, the model assumes a removal rate (uniform
 284 distribution min 0.89, max 0.99) (Table 3). When operating sub-optimally, the model assumes a
 285 removal of 50% of the optimal removal rate, and zero removal during failure events. It was
 286 assumed that aluminium sulphate was applied at an optimum dose of approximately 10 mg L⁻¹.

287

288 **Table 2.** Model inputs and distributions

Stage	Symbol	Description	Model /distribution	Units
Effluent (Surface-runoff)				
	$C_{\text{surface-runoff}}$	Initial concentration in surface runoff	Lognormal	MPN/ 100mls
Dilution	DF	Dilution in stream	(based on Table 1) Dilution factor (10)	-
	$PEC_{\text{surface-water}}$	Concentration of coliforms in surface-water following dilution	$C_{\text{surface-runoff}} / DF$	MPN/100 mls

Stage	Symbol	Description	Model /distribution	Units
Die-off	K	First order inactivation constant	Uniform	d ⁻¹
	t	Time in stream	Uniform (min 0, max 5)	d ⁻¹
	D-off	Die-off rate in stream	$N = N_0 \llbracket \exp (-kt) \rrbracket$	MPN/ 100mls
Secondary treatment				
	C-opt	Coagulation/Flocculation and sedimentation optimum	0.90	Probability
	CS-opt	Coagulation/Flocculation and sedimentation sub-optimum	0.05	Probability
	C-fail	Coagulation/Flocculation and sedimentation fail	0.05	Probability
	Cr	Coagulation/Flocculation and sedimentation reduction	Uniform (min 0.89, max 0.99)	Decimal reduction
	F-opt	Filter optimum run	0.9	Probability
	FS-opt	Filter sub-optimum run	0.1	Probability
	Frd	Filter reduction (rapid sand)	Uniform (min 0.74, max 0.99)	Decimal reduction
Tertiary treatment				
	D	Disinfection	Uniform (min 0.97, max 0.99)	Decimal reduction
Output	Pstt	Post-secondary and tertiary treatment	$Pstt = D\text{-off} \times (1-Cr) \times (1-Frd) \times (1-D)$	MPN/ 100mls
Human exposure				
Consumption	TWi	Tap water intake (adult)	Lognormal (mean 0.564,	L d ⁻¹
Output	Vcc	Viable coliforms/ <i>E. coli</i> consumed	$Pstt \times Twi$	MPN/ d ⁻¹
Dose response				
Output	I(H)	Probability of illness (healthy)	$1-EXP (- 0.0000005 \times Vcc)$	-
Output	I (Ic)	Probability of illness (immunocompromised)	$1-EXP (- 0.01 \times Vcc)$	-

290

291 The filtration process is the last treatment stages that can physically remove contaminants before
292 disinfection. One of the most popular filtration processes used in Ireland is the rapid gravity sand
293 process (Ireland EPA 1995). A study by Li et al. (2012) showed that direct rapid sand removal
294 can remove 0.6 - 1.5 log-units of total faecal coliform, depending on the loading rate and grain
295 size distribution. Mwabi et al. (2012) demonstrated that designing and building a bio-sand
296 filtration system was effective in removing 2 - 4 log₁₀ of coliform bacteria. Koivunen et al.
297 (2003) showed that tertiary treatment by the rapid sand filtration process found, on average, a
298 97% reduction of faecal coliforms and total coliforms in four conventional wastewater treatment
299 plants in Helsinki, Finland. In keeping with the Irish EPA's filtration manual guidelines, rapid
300 gravity filtration was considered in the model. Filtration can be stable or unstable due to
301 optimum, sub-optimum or failure of the coagulation/flocculation process. As a "worst case
302 scenario" the model assumes a 90% probability of filtration operating at an optimum stable run
303 (Fopt) and 10% probability for sub-optimal run (FS-opt). To model rapid sand filtration under
304 optimum conditions and to account for uncertainty and variability in the data, a decimal
305 reduction uniform distribution was assigned (min 0.74 max 0.99) (Table 2). When operating sub-
306 optimally, the model assumes a removal of 50% of the optimal removal rate.

307 **Disinfection**

308 Disinfection is the process by which an organism's viability/infectivity is destroyed with a
309 specific percentage of the population dying over some time frame defined as a rate (Betancourt
310 and Rose, 2004). Worldwide, chlorine is the most commonly used disinfection in drinking water
311 treatment, although other alternatives are being increasingly introduced such as ozonation,
312 ultraviolet irradiation, ultrasonic vibration, ultra-filtration, silver, bromide and iodine, membrane
313 filtration and granular activated carbon (GAC). Chlorine is added to provide a disinfectant
314 residual to preserve the water in distribution, where the chlorine is in contact with the water for a
315 longer period of time compared to the pre-chlorination process in primary treatment (Irish EPA,
316 2011). The principal factors that influence disinfection efficiency are the disinfection
317 concentration, contact time, temperature and pH (depending upon the disinfection) (Cotruvo et
318 al., 2013). Chlorination has been found to remove *E. coli* between 97-99% (O' Connor and O'
319 Connor 2001). However, a report by Igunnuugbemi et al. (2009) showed that water storage post

320 chlorination significantly reduced survival of *E. coli* and that the presence of *E. coli* following
321 chlorination could undermine the effectiveness of chlorination. To account for uncertainty in the
322 data, a uniform distribution (minimum 0.97, maximum 0.99) was assigned to model the
323 inactivation attributed to the disinfection process.

324 Removal of coliforms and bacteria (TC and *E. coli*) was quantified in terms of a decimal
325 reduction. The concentration of coliforms remaining after secondary and tertiary treatment in a
326 WTP was calculated by multiplying the level present post primary treatment by the decimal
327 reduction due to coagulation/ flocculation, sedimentation, filtration and disinfection. The
328 equation is:

$$329 \quad P_{\text{STT}} = D\text{-off} \times (1\text{-Cr}) \times (1\text{-Frd}) \times (1\text{-D}) \quad \text{Eqn.3}$$

330 Where: P_{STT} is the coliform concentration post-secondary and tertiary treatment (MPN/100 mls),
331 Cr is decimal reduction due to coagulation /flocculation and sedimentation, Frd is decimal
332 reduction due to filtration, and D is the decimal reduction due to disinfection.

333 **Human Exposure**

334 Water consumption in Ireland for adults was modelled using a lognormal distribution with a
335 mean and standard deviation value of $0.564 \pm 0.617 \text{ L d}^{-1}$ according to a survey on adult
336 consumption patterns conducted by the Irish Universities Nutrition Alliance (IUNA) which was
337 based on 1274 consumers. The same survey was used to model variation in adult body weight
338 (males and females) and a normal distribution with a mean and standard deviation value of $78 \pm$
339 16.5 kg was used (IUNA, 2011).

340 **Dose response model**

341 In order to assess the risk to human health from coliforms and *E. coli* associated with water
342 consumption, the potential exposure to the organism(s) in the daily drinking water intake was
343 estimated. Exponential models are widely used in microbial risk assessment (Teunis et al., 2004).
344 The exponential model assumes that pathogen-host interactions can describe the pathogen-host
345 survival probability by a discreet value (Haas et al., 2000). Two dose response models were
346 considered for TC and *E. coli* exposure incorporating two different exposure scenarios (healthy
347 populations and immuno-compromised populations). Immuno-compromised individuals include

348 patients on active anti-cancer drugs, HIV/AIDS and other chemotherapies. Allen et al. (2013)
349 defines an immuno-compromised individual as having a haematology profile showing abnormal
350 values for gamma globulins, white blood cells, red blood cells and liver function. The dose
351 response model estimated the probability of illness resulting from a certain level of exposure. An
352 exponential dose-response model was used for probability of illness, integrating an “r” value of
353 0.01 for immuno-compromised populations (I(Ic)) and an “r” value of 0.0000005 for healthy
354 population (I(H)) as proposed by Gale (2005). As a “worst case scenario”, the illness model was
355 parameterized with the assumption that the virulence of the pathogen is similar to *E. coli*
356 O157:H7. The *E. coli* O157:H7 strain is a particular serotype of the group referred to as
357 verocytotoxigenic *E. coli* (VTEC). VTECs produce verotoxins or shiga-like toxins that are
358 closely related to the toxin produced by *Shigella dysenteriae* (Cassin et al., 1998). The USEPA
359 have proposed an acceptable benchmark of 10^{-4} annual infection/illness probability per person per
360 year for *Shigella* (Grant et al., 2012). The WHO use the metric DALY (disability-adjusted life
361 year) to estimate severity and duration of a disease. The 10^{-6} DALY tolerable burden of disease
362 may be considered unrealistic and there have been proposals to introduce a less stringent burden
363 of risk such as the upper limit for excess lifetime risk of cancer of 10^{-5} or a 10^{-4} limit in line with
364 the USEPA limit (WHO 2011). Crockett et al. (1996) reported that ingestion of only 10-100
365 *Shigella* cells can lead to infection. The probability of illness per day can be expressed by:

366
$$P_i = 1 - \exp(-d \times r) \quad \text{Eqn. 4}$$

367 Where P_i is the probability of illness (d^{-1}), d is the dose and ‘r’ represents an exponential
368 parameter. The annual individual risk is calculated as:

369
$$P_{i(365)} = 1 - (1 - P)^{365} \quad \text{Eqn. 5}$$

370

371 **Sensitivity analysis**

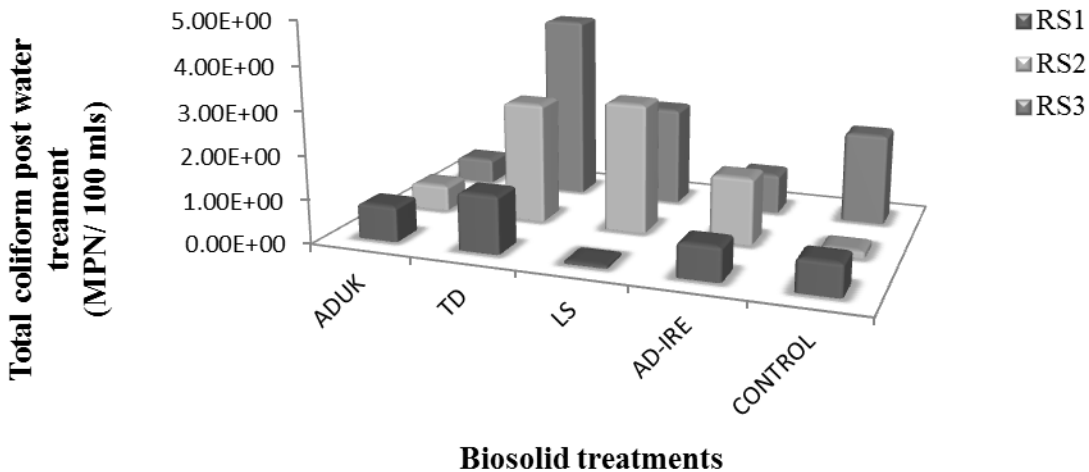
372 A sensitivity analysis, based on rank order correlation, was carried out to assess how the model’s
373 predictions are dependent on variability and uncertainty in the model input parameters.
374 Sensitivity analysis assesses how the model predictions are dependent on variability and
375 uncertainty in the model’s inputs. Monte Carlo simulation performs risk analysis by building

376 models of possible results by substituting a range of values—a probability distribution—for any
 377 factor that has inherent uncertainty or variability (Kavcar et al., 2009). It then iterates the results
 378 using a different set of random values from the probability functions. Ten thousand iterations
 379 were performed for each simulation.

380 **Results**

381 Figures 2 and 3 give the predicted results of TC and *E. coli* remaining following drinking water
 382 treatment under the scenarios considered. The model produced several output distributions (TC
 383 and *E. coli* concentration in effluent post WTP, viable coliforms consumed, and probability of
 384 illness) that can be used to compare the concentration of coliforms that were detected in surface
 385 runoff and their potential risk to human health. The model predicted that surface runoff arising
 386 from the land spreading of TD biosolids and ADUK biosolids produced the highest
 387 concentrations of TC and *E. coli*, respectively, in drinking water. The modelled mean TC and *E.*
 388 *coli* concentration in drinking water was highest when the surface runoff concentrations from the
 389 TD and ADUK, respectively, biosolids at each rainfall simulation time (24, 48 and 360 hr) were
 390 used as input into the model (mean concentration values 1.3, 2.7 and 4.2 MPN/100 mls for TC
 391 and TD biosolid treatment (Figure 2), and 7.3×10^{-2} , 4.7×10^{-2} and 2.4×10^{-3} for *E. coli* and
 392 ADUK biosolid treatment (Figure 3).

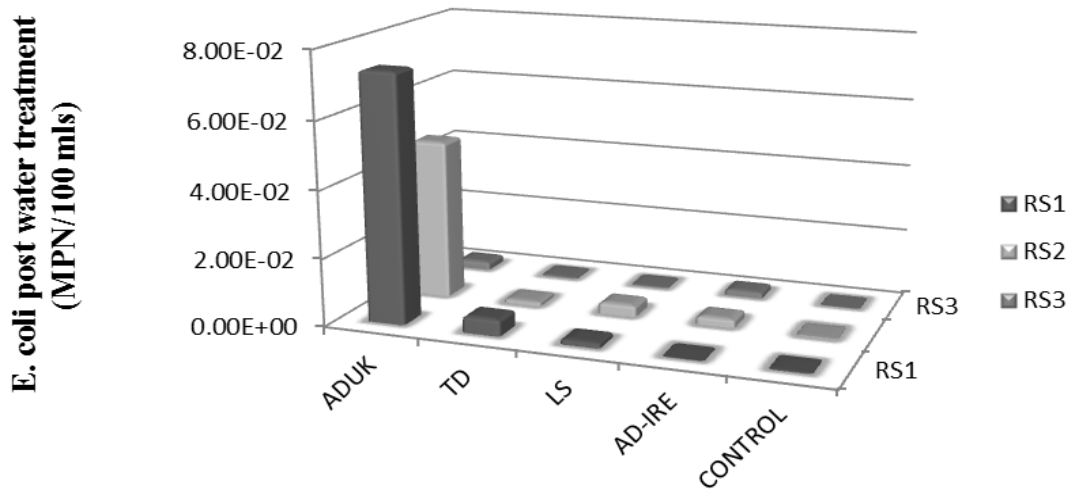
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394

395 **Figure 2.** Simulated mean total coliforms remaining following drinking water treatment

396



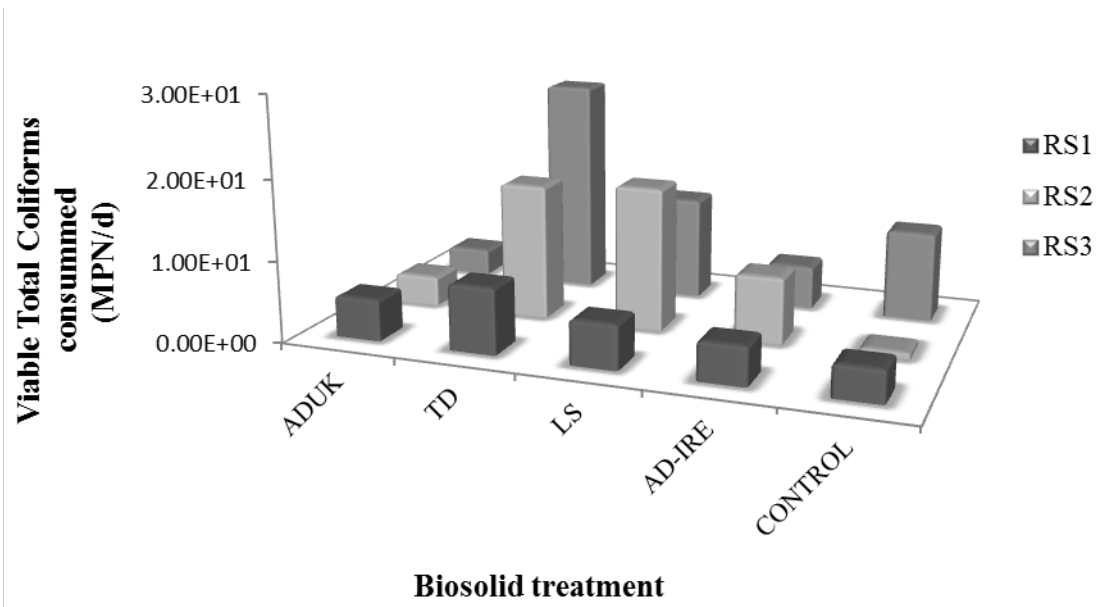
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398 **Figure 3.** Simulated mean *E. coli* remaining following drinking water treatment

399

400 The EU states that there should be 0 in 100 mls of coliform bacteria and *E. coli* following
401 drinking water treatment. The results for mean human exposure *via* drinking water consumption
402 show that for TC the greatest viable coliforms consumed was for the biosolid TD and LS
403 combining rainfall simulation times of 48 and 360 hr (RS2 and RS3), respectively (Figure 4)
404 (mean viable total coliform values 16.83 and 26.75 MPN d⁻¹, respectively) for TD (mean viable
405 total coliform values 17.74 and 12.82 MPN d⁻¹, respectively) for LS biosolids. The results for *E.*
406 *coli* show that the greatest viable coliforms consumed was for ADUK biosolids and rainfall
407 simulation times of 24 and 48hrs (RS1 and RS2) (Figure 5) mean viable faecal coliforms
408 consumed values 5.20×10^{-1} and 2.34×10^{-1} MPN d⁻¹, respectively.

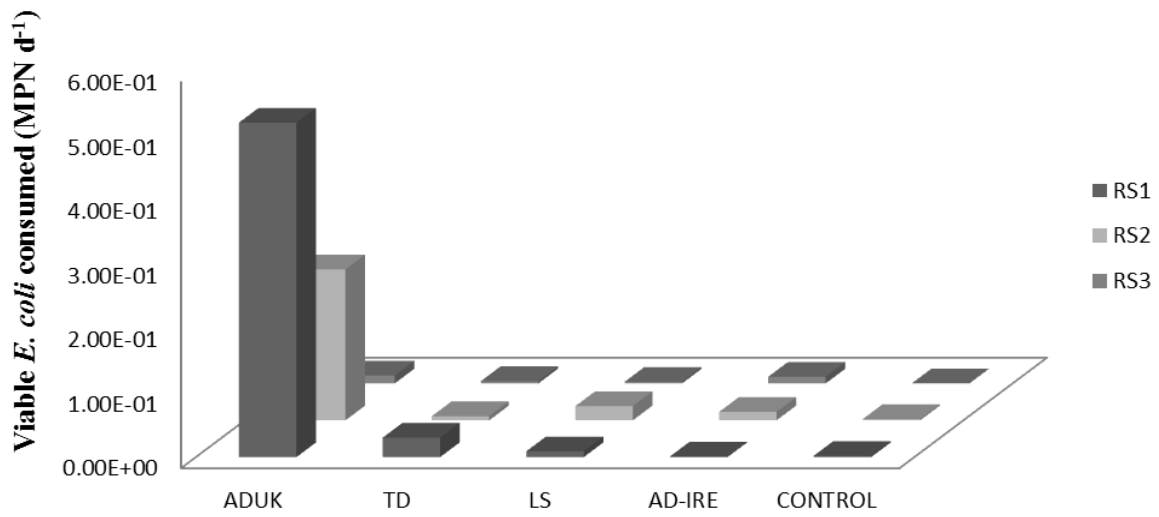
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410

411 **Figure 4.** Simulated mean viable total coliforms consumed

412



413

414 **Figure 5.** Simulated mean viable *E. coli* consumed

415

416 The results for probability of illness (healthy and immuno-compromised) are displayed in Table
417 3. For each scenario (healthy and immuno-compromised), the risk assessment model produced a
418 simulated probability of illness per day and per year. Compared to the healthy population, the
419 immuno-compromised population are more at risk of illness with mean annual values for TC and
420 immuno-compromised (9.92×10^{-1}) and LS biosolids treatment (RS1), (7.24×10^{-1} and $7.87 \times$
421 10^{-1}) for the TD biosolids treatment incorporating the RS2 and RS3 time frames, compared to
422 mean annual values for TC and healthy population for the same biosolid treatments and time
423 frames (1.01×10^{-3}) incorporating the LS biosolids treatment and RS1 time frame, mean annual
424 values for healthy population and TD biosolids treatment (2.77×10^{-3} and 4.27×10^{-3}
425 respectively), incorporating the RS2 and RS3 time frames. The mean annual values for *E. coli*
426 and immuno-compromised populations show that the ADUK biosolids and the RS1 and RS2
427 time frames had the greatest probability of risk (values 2.1×10^{-1} and 1.7×10^{-1} , respectively).
428 This is comparable to the healthy population for the same biosolid treatment and time frames
429 (mean annual values of 7.0×10^{-5} and 4.6×10^{-5} , respectively).

430

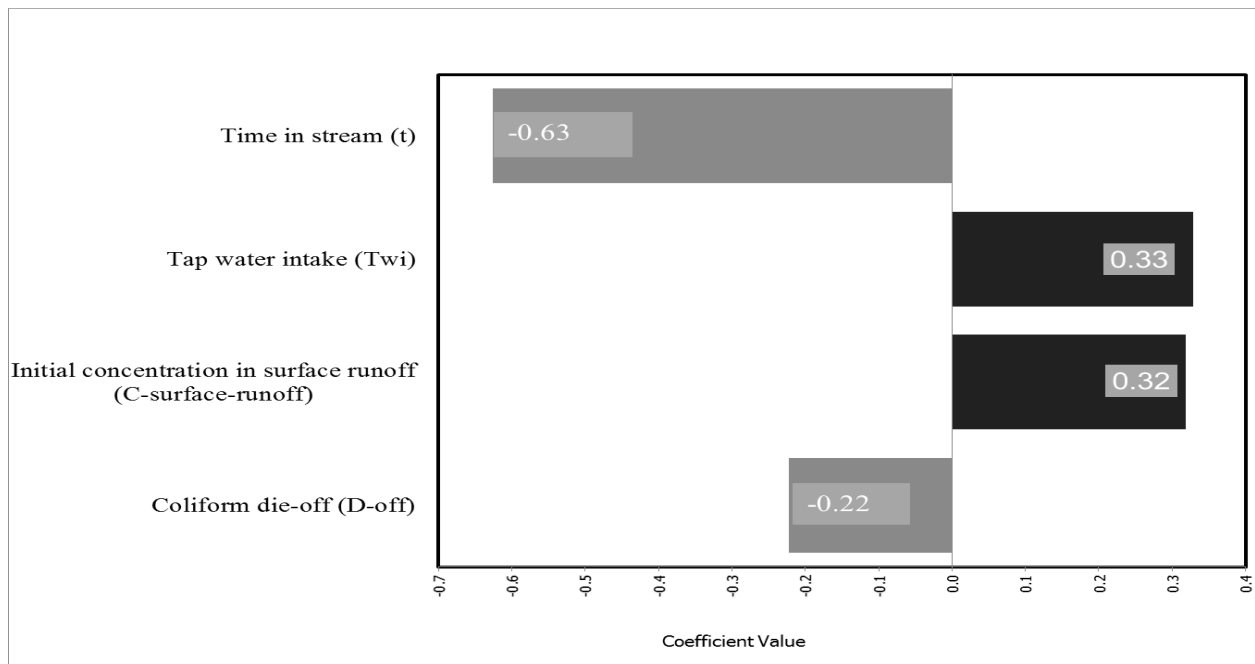
1 **Table 3.** Probability of illness for healthy and immuno-compromised populations

Probability of illness								
Biosolid treatment	Healthy population				Immuno-compromised population			
	(d ⁻¹)		(yr ⁻¹)		(d ⁻¹)		(yr ⁻¹)	
	TC	<i>E. coli</i>	TC	<i>E. coli</i>	TC	<i>E. coli</i>	TC	<i>E. coli</i>
RS1								
ADUK	2.57E-06	1.71E-07	8.90E-04	7.0E-05	2.81E-02	3.68E-03	5.62E-01	2.1E-01
TD	4.09E-06	1.46E-08	1.35E-03	4.2E-06	3.67E-02	2.86E-04	5.85E-01	4.1E-02
LS	2.76E-06	5.29E-09	1.01E-03	1.3E-06	5.22E-02	1.03E-04	9.92E-01	1.9E-02
AD-IRE	2.38E-06	3.94E-10	7.89E-04	1.4E-07	2.44E-02	7.87E-06	5.38E-01	2.5E-03
CONTROL	2.00E-06	9.51E-10	7.18E-04	3.9E-07	2.60E-02	1.90E-05	5.62E-01	6.2E-03
RS2								
ADUK	1.86E-06	1.23E-07	6.34E-04	4.6E-05	2.05E-02	2.1E-03	4.65E-01	1.7E-01
TD	8.41E-06	3.46E-09	2.77E-03	1.0E-06	6.73E-02	6.9E-05	7.24E-01	1.5E-02
LS	8.86E-06	9.26E-09	2.71E-03	3.4E-06	6.32E-02	1.8E-04	7.10E-01	3.5E-02
AD-IRE	4.11E-06	6.35E-09	1.41E-03	3.0E-06	3.87E-02	1.3E-04	5.94E-01	2.9E-02
CONTROL	3.91E-07	8.69E-10	1.42E-04	3.2E-07	6.83E-03	1.7E-05	3.43E-01	5.2E-03
RS3								
ADUK	1.66E-06	6.1E-09	5.99E-04	2.6E-06	2.30E-02	1.2E-04	5.31E-01	2.3E-02
TD	1.34E-05	1.2E-09	4.27E-03	1.2E-06	9.31E-02	2.4E-05	7.87E-01	6.7E-03
LS	6.41E-06	1.2E-09	2.20E-03	3.9E-07	5.47E-02	2.3E-05	6.82E-01	6.4E-03
AD-IRE	2.70E-06	5.1E-09	9.38E-04	2.3E-06	2.97E-02	1.0E-04	5.80E-01	2.3E-02
CONTROL	5.56E-06	1.1E-10	1.93E-03	4.6E-08	5.35E-02	2.2E-06	6.89E-01	8.6E-04

1 Sensitivity analysis was performed to investigate how variability of the outputs can be
 2 apportioned quantitatively to different sources of variability in the inputs. The analysis indicated
 3 that the LS and TD biosolids produced the highest concentration post WTP of TC, and ADUK
 4 produced the highest concentration of *E. coli*, in drinking water, therefore, a sensitivity analysis
 5 was conducted for the annual probability of illness for both biosolid treatments. Results for TC
 6 and *E. coli* show that the parameter of importance that affected the variance in model predictions
 7 was time in the stream (correlation coefficient -0.63 and -0.57, respectively) (Figures 6 and 7).
 8 This highlights the importance of residence time of bacteria in stream. The longer the bacteria
 9 are in the stream, the more likely the bacteria are subject to factors such as temperature, pH and
 10 photolysis, which may in-turn influence the growth or die-off rate of bacteria in a stream. The
 11 other parameters of importance were the tap water intake and initial concentrations in surface
 12 runoff (correlation coefficients 0.33 and 0.31, respectively, for Twi and 0.32 and 0.33,
 13 respectively, for C-surface-runoff). The die-off rate in the stream (-0.22 for TC and -0.20 for *E.*
 14 *coli*) was also of importance. The die-off rate is related to the residence time in the stream and is
 15 associated with sub-optimum conditions in the stream that influence bacterial growth.

16

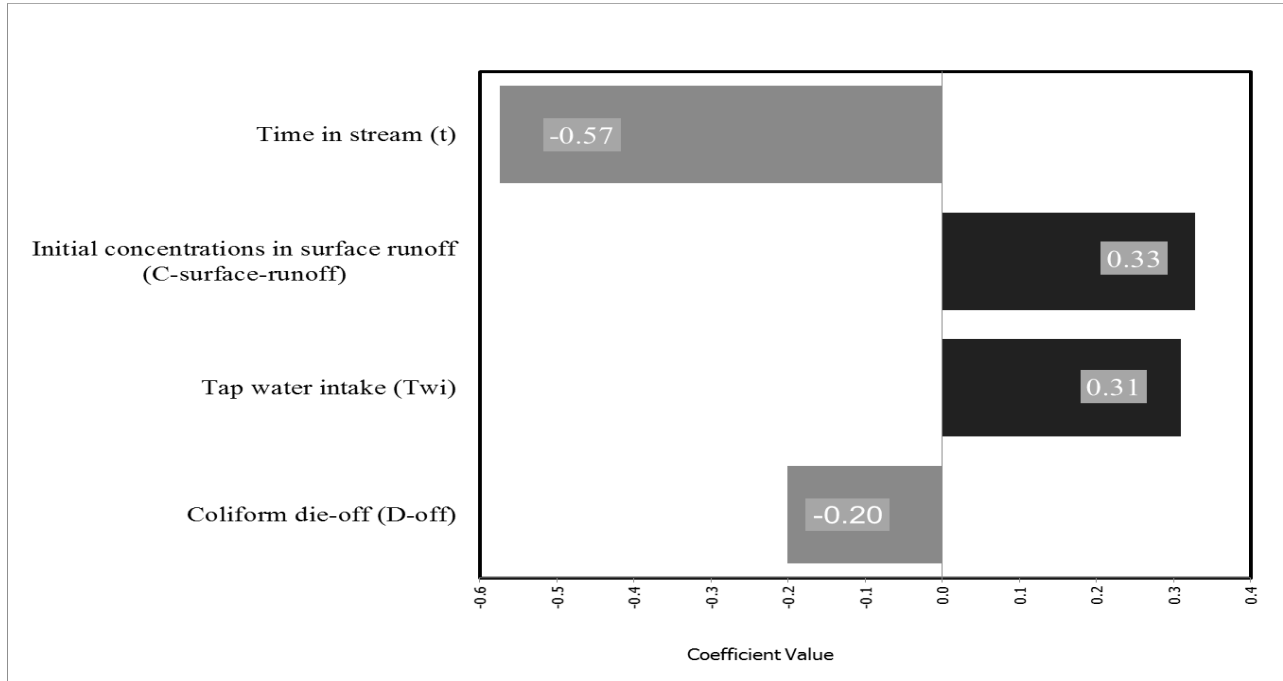
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18

1 **Figure 6.** Sensitivity analysis for TC annual probability of illness and TD biosolid treatment

2
3



4
5 **Figure 7.** Sensitivity analysis for *E. coli* annual probability of illness and ADUK biosolid
6 treatment

7 **Discussion**

8 Concentrations of TC and *E. coli* in surface runoff following the spreading of biosolids on
9 grassland were quantitatively assessed to study their fate in drinking water treatment and
10 subsequent consumption and human health effects. Initial concentrations of *E. coli* in surface
11 runoff were above the recommended standards of $> 1 \times 10^3$ MPN g^{-1} and were equivalent to class
12 B microbial matter under the USEPA Part 503 regulations. Surface runoff is distinguished from
13 other types of runoff in that it does not pass through the soil. Therefore, typical soil-pathogen
14 reactions (desiccation, photolysis, temperature and nutrients) may be by-passed depending on the
15 rate off rainfall. Concentrations of TC and *E. coli* in surface runoff in this study are comparable
16 to concentrations reported by Le Chevalier et al. (1991) and Schreiber et al. (2015). All TC and

1 *E. coli* concentrations had decreased by the third rainfall event (RS3; 360 hr) due to desiccation
2 of the pathogens in soil following the application of the biosolids.

3 The mean concentration of TCs after the WTP showed that the TD biosolids (RS2 and RS3) and
4 LS biosolids (RS2 and RS3) with the highest concentration of TC. This was attributed to initial
5 concentrations of TCs in the influent and the time in stream combined with the removal rates
6 associated with secondary treatment (e.g. coagulation/flocculation and sedimentation and
7 filtration). Thermal drying is recognised as more effective in pathogen removal than mesophilic
8 digestion and can achieve the time-temperature requirement for Class A biosolids (Iranpour and
9 Cox, 2006). However, regrowth of pathogens can occur in thermally dried biosolids (Zaleski et
10 al. 2005). Lloret et al. (2013) showed that the reduction in sludge retention time may be
11 responsible for presence of coliforms post treatment. Lloret et al. (2013) reported that a
12 minimum time of more than 10 days under thermophilic conditions is required to achieve
13 appropriate sanitation of sludge. Similarly, Iranpour and Cox (2006) reported the presence of
14 faecal coliforms after thermal drying, and attributed the reason to be the relatively short sludge
15 retention time of about 10 days.

16 The mean concentration of *E. coli* post drinking water show that the ADUK biosolids had the
17 greatest concentration of *E. coli* for RS1 and RS2 only. This was also attributed to the initial
18 concentration and the time in the stream of *E. coli* in the influent and associated drinking water
19 treatment removal rates. Although initial concentrations of TC and *E. coli* in surface water were
20 high, the effect of drinking water treatment significantly reduced overall TC and *E. coli*
21 concentrations with a 99.9% reduction across all treatments and time frames.

22 The mean viable consumption of TC and *E. coli* in drinking water showed the same trends as
23 mean TC and *E. coli* concentrations post drinking water treatment. Safe drinking water is a
24 human right and in developed countries it has become an “entitlement”. Water consumers rely on
25 the efficacy of drinking water treatment to produce a product that is pathogen free, odourless and
26 clear. However, indicator bacteria are known to regrow in finished drinking water. This was
27 highlighted in a report by Le Chevalier et al. (1991). The authors reported various factors
28 attributed to the occurrence of coliforms in drinking water including disinfectant residuals,
29 filtration and temperature. Bacterial growth can occur on any surface that is constantly wet, so
30 the internal surface of water distribution pipes is normally coated with a biofilm (Gray, 2010).

1 Although the concentrations of coliforms post drinking water treatment in this study were
2 significantly reduced, inefficiencies in drinking water treatment due to operational defects that
3 promote the regrowth of coliforms and other pathogens can be a cause of concern for drinking
4 water management.

5 Ideally water intended for human consumption should be pathogen free. However, in practice,
6 this is an unachievable goal. A consequence of variable human susceptibility to pathogens is that
7 exposure to drinking water of a particular quality may lead to health problems in different
8 populations (WHO 2011), particularly the very young and immuno-compromised. Enteric
9 pathogens are among the many agents that take advantage of the impaired or destroyed immune
10 system; therefore, sensitive populations are considerably more vulnerable and may need special
11 protection from waterborne microorganisms (Gerba et al., 1996). As *E. coli* is used as an
12 indicator that faecal matter is present, it may indicate the presence of pathogens that cause
13 waterborne diseases. The risk of illness for healthy populations was deemed negligible based on
14 the tolerable risk guidelines set by the USEPA and the WHO for *Shigella*. However, based on the
15 same guidelines, immuno-compromised populations may be at risk. Individuals who are truly
16 immuno-compromised would follow medical advice regarding food and water intake, thus
17 reducing the risk of illness.

18
19 The Sewage Directive has yet to address the bacteriological quality of treated water. The current
20 European legislation requires that the sludge be subjected to a process of stabilisation before land
21 application. With future demography increases and growing demand for water, the use of
22 reclaimed water will rise; therefore efforts to assess the treatment efficacy are vital.

23

24 **Conclusions**

25 Application of biosolids on grassland and subsequent simulated rainfall over three time frames
26 resulted in TC and *E. coli* counts in surface runoff. The concentrations of *E. coli* exceeded the
27 recommended standards being some 10-fold below the Class A Irish standard. This prompted the
28 need to investigate human exposure. Further analysis which included simulated dilution and die-
29 off rate in a stream, drinking water treatment, and human exposure following consumption of the
30 treated water resulted in a very low probability of illness based on the USEPA and the WHO

1 threshold of acceptable risk (10^{-4} and 10^{-6} , respectively) for healthy populations. However, the
2 risk of illness for immuno-compromised populations exceeded the thresholds of acceptable risk
3 by a factor of 3 for TC and a factor between 1-3 for *E. coli*. It is noted in such cases, susceptible
4 populations would be subject to medical advice regarding food and water intake, thus reducing
5 the risk of illness. The sensitivity analysis identified that the time in stream is an important
6 parameter as the longer the bacteria are in the water and being exposed to ultraviolet light,
7 varying temperature and pH, the greater the influence on bacterial growth. The risk assessment
8 model developed in this study may be of importance to local authorities or regulatory agencies to
9 evaluate the likely risk of *E. coli* entering potable water following biosolid application on
10 agricultural land. As this study only focused on coliforms, future studies are needed in order to
11 assess other compounds of concern e.g. pharmaceutical contaminants that may be present in
12 biosolids.

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3 Programme (2007-2013).

4

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29 [er%20Policy/Comments/Sewage%20Sludge/EWA_WD_sludge_en.pdf](http://www.ewa-online.eu/comments.html?file=tl_files/media/content/documents_pdf/European%20Water%20Policy/Comments/Sewage%20Sludge/EWA_WD_sludge_en.pdf)

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