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Validation of hydrodynamic and microbial inactivation models for UV-C treatment of milk in a swirl-tube 'SurePure Turbulator™'

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For consideration in Journal of Food Engineering

1	Validation of hydrodynamic and microbial inactivation models for UV-C
2	treatment of milk in a swirl-tube 'SurePure Turbulator [™]
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10	Abstract
12	The Positron Emission Particle Tracking (PEPT) flow visualisation technique has been applied to
13	determine the hydrodynamic performance of a full-scale transparent model of a SurePure
14	Turbulator [™] used for the microbial treatment of turbid dairy fluids using UV-C radiation. The effect of
15	flow rate upon the refreshment of fluid at the surface of the UV source has been investigated using
16	two model fluids each possessing the same viscosities as milk and cream respectively. The amount
17	of surface refreshment is modelled as a time density function close the surface of UV-C source and
18	incorporated into an existing first order microbial inactivation model and a Weibull distribution model.
19	Fitting to experimental data obtained for the inactivation of selected milk pathogens using the
20	Turbulator TM have demonstrated the superiority of the Weibull model. These models enable a more
21	precise estimation of UV-C energy requirement for the inactivation of the milk borne pathogenic
22	organisms to be made since the amount of surface refreshment affords a significant performance
23	enhancement.
24	Keywords
25	UV-C inactivation of milk pathogens, Positron Emission Particle Tracking, hydrodynamic model,
26	Weibull inactivation model
27	1. Introduction
28	Heat pasteurization is the most commonly used treatment for the microbial inactivation in milk and

30 causes thermal denaturation of proteins (Dietz *and* Erdman, 1989), a decrease in nutritional value, an

other dairy fluids. However, the process is known to have adverse effects on the final product as it

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31 increase in heat-generated aroma compounds and deterioration in sensory attributes. As a 32 consequence, emerging non-thermal technologies have been evaluated as possible alternatives 33 (Engin and Karagul Yuceer, 2012). The application of UV light for the treatment of turbid liquids has 34 been demonstrated and has growing industrial interest, despite the limited penetration depths 35 observed due to attenuation of UV light through the media (Bintsis et al., 2000). Despite this issue, 36 the processing advantages of UV light treatment have been demonstrated not only in terms of cost 37 (Koutchma, 2009), but also due to better preservation of critical food guality and health attributes after 38 treatment (Cohen and Birk, 1998; Orlowska et al., 2013). As reported by Choi and Nielsen (2005), 39 thermally pasteurized samples were different in colour and less preferred in all areas of consumer 40 acceptability compared to UV-irradiated samples. In their work ozone-treated cider had greater 41 sedimentation, lower sucrose content and a decrease in soluble solids using UV irradiation. Besides, 42 UV irradiation has been considered a more cost-effective method to produce safe apple cider with 43 minimal quality and consumer acceptability differences. Light wavelengths used in UV processing are 44 usually in the range of 100 to 400 nm, which is subdivided into UV-A (315 to 400 nm), UV-B (280 to 315 nm) and UV-C (200 to 280 nm). Long wavelength UV-A light has limited microbiocidal effects, 45 46 and for practical applications in foods its effectiveness has to be enhanced by the presence of 47 photosensitive compounds which diffuse into a microbial cell prior to irradiation. These photosensitive 48 compounds are expensive and their addition to foodstuffs is highly questionable on safety and toxicity 49 grounds. Therefore, recent studies have been conducted using UV-C light characterized by sufficient 50 energy of the photons to cause microbiocidal action by destruction of nucleic acids within 51 microorganisms. Several studies can be found in the literature where UV-C treatment of turbid and 52 opaque fluids has been used instead of a thermal process. Different types of liquid food products are 53 processed using UV-C light such as juice products (Koutchma and Parisi, 2004; Koutchma et al., 54 2007; Freitas, et al. 2015), cider (Unluturk et al., 2004), milk (Krishnamurthy et al., 2007; Matak et al., 55 2007), liquid egg (Kuda et al. 2012; Mendes de Souza et al. 2013) and white and red wine (Rizzotti et 56 al. 2015). In the work of Gayan et al. (2014) an overview of continuous flow UV liquid food 57 pasteurization is introduced. In this paper the main engineering aspects required for understanding 58 the current work are presented such as exposure time, UV radiation dose, absorption coefficient (α), and the corresponding penetration depth (λ) of food products. 59

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60 The treatment of low UV transmission (UVT) fluids can be considerably enhanced by manipulation of 61 the hydrodynamic environment within UV-C systems. Two different strategies are generally employed 62 to reduce the impact of low penetration depth in turbid fluids. The first strategy is the use of extremely 63 thin liquid films, in the range between 0.9 and 1.6 mm, to decrease the path length of the UV light 64 photons, thus avoiding problems associated with lack of penetration (Koutchma et al. 2007). A second strategy is to increase surface refreshment of the fluid in close proximity to the UV source; 65 such secondary flows may be induced by a swirling flow, as in the case in this work, or by exploitation 66 67 of Dean vortices in coiled tubes. In the latter case, the lamps and reflectors are placed both inside and outside the coiled tube, increasing not only the UV irradiance of the flowing liquid, but also its 68 69 uniformity (Koutchma, 2008).

The "SurePure TurbulatorTM" UV-C device used in this work exploits swirling turbulent flows and is 70 71 designed for continuous flow inactivation of turbid fluids such as milk. Fluid enters through a 72 tangential inlet to promote a swirling flow and then passes through an annular gap between the UV 73 lamp-containing quartz sleeve and the outer turbulator tube. The outer tube employs a wavy inner 74 wall designed to produce additional turbulence to further promote surface refreshment of fluid flow. 75 The outer wall has a spiral channel cut into it with a pitch of 5mm the grooves are sinusoidal in shape with an amplitude of 0.35 mm, so the gap varies from 0.9 mm to 1.6 mm. Previous work carried out by 76 77 Simmons et al. (2012) has involved fluid mechanical measurements to determine the degree of swirl, 78 and thus the degree of surface refreshment, which the fluid experiences on average. The fluid motion 79 through the device was determined using Positron Emission Particle Tracking (PEPT) which tracks 80 the motion of a radioactive tracer particle placed into the fluid within an exact model of the SurePure TurbulatorTM. The technique can detect the tracer position within < 0.5 mm. The experiments were 81 carried out for water at the design flow rate (4500 L hr⁻¹) and at 3380 L hr-1 and 2250 L hr⁻¹ 75% and 82 83 50% turndown respectively; the intensity of the swirl was found to decrease with flow rate. The work 84 highlighted the importance of various aspects of the design, with the inlet configuration found to be 85 critical to the degree of swirl and thus the rate of surface refreshment at the UV source.

In this previous work, the developed surface refreshment (hydrodynamic) model based upon the PEPT data was used to determine the fraction of the total residence time spent by the particle (and thus the fluid) as a function of distance from the UV-C light source. The model was developed on the basis of the Lambert-Beer law and first order lethality kinetics shown in equations (1) and (2) below:

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90

$$\frac{I}{I_0} = e^{-\alpha x} \tag{1}$$

(2)

91
$$\frac{N}{N} = \exp(-k_1 I_x t_x)$$

92 where x is the distance from the UV source (cm), I_0 is the fluence rate at the surface of the UV source, N is the concentration of survived microorganisms (CFU cm⁻³), N_0 is the initial concentration of 93 microrganisms, k_1 is the first order rate constant (m² J⁻¹), α is the absorption coefficient (cm⁻¹) and I_x is 94 the fluence rate at distance x from the UV-C source (W m⁻²). t_x is the residence time spent by the 95 particle, on average, at distance x from the source, as determined from PEPT experiments. For 96 highly opaque fluids with $\alpha > 200$ cm⁻¹, such as milk, the UV light attenuation is so excessive that only 97 98 fluid reaching the surface of the UV-C source receives sufficient treatment (Koutchma, 2009), i.e. as x 99 \rightarrow 0.

In this paper, this surface refreshment modelling approach is further developed and used to estimate 100 the required UV-C dose and time to achieve specific microbial reduction in milk and milk cream 101 products using the Turbulator[™] as a function of flow rate. PEPT experiments have been carried out 102 103 over a broader range of flow conditions using two model fluids with viscosities representative of milk products and cream respectively. Thus, PEPT has been used to obtain the time, t_{0.5}, spent by the fluid 104 at a distance of less than 0.5 mm from the surface of the UV-C source as a function of flow rate and 105 106 fluid viscosity. Comparison of the first order inactivation kinetics data obtained for selected milk pathogens as Serratia marcescens, Aeromonas hydrophila, Escherichia coli and Listeria 107 108 Monocytogenes (Crook et al. 2014) have identified inadequacies in the first order assumption used by 109 Simmons et al. (2012), therefore the model is extended to include a Weibull frequency distribution 110 model (Albert and Mafart, 2005).

111

112 2. Materials and Methods

113 2.1 Flow conditions

114 Over the range of conditions used, both milk and cream can be considered as Newtonian fluids.

115 Aqueous solutions of glycerol at concentrations of 40 % and 50% by weight at 20°C were used to

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116 match the viscosities of milk and cream respectively at commercial processing temperatures. The

117 physical properties of the fluids at 20°C are presented in Table 1.

- 118 The SurePure Turbulator[™] rig was the same as used in the previous work and comprises two model
- 119 turbulator sections manufactured from poly(methyl methacrylate) (PMMA), one mounted on top of the
- 120 other (Figure 1). The liquid is pumped from a supply tank through the bottom turbulator section and
- 121 exits from the top section back to the tank, forming a closed circuit flow loop.
- 122 The flow rates chosen were 7.5% above and below the design rate of 4000 L hr⁻¹ for the milk mimic,
- reflecting commercial operating conditions. Experiments with the cream mimic were carried out at
- 4000 L hr⁻¹ only. Flow properties are shown in Table 2, together with values of superficial velocities
- and their corresponding values of Reynolds number. The Reynolds number is calculated from
- 126

$$\operatorname{Re} = \frac{d_e U \rho}{\mu}, \qquad (3)$$

where d_e is the hydraulic diameter of the flow conduit (four times the flow cross-sectional area divided by the wetted perimeter), *U* is the superficial velocity, ρ is the fluid density and μ is the fluid dynamic viscosity.

130 2.2 PEPT experiments

131 The details of the experimental protocol carried out in this study are described in Simmons et al. 132 (2012). The experiments carried out in this paper however utilised an improved radiotracer particle of a smaller size, 250 µm, compared with the 500 µm resin bead used previously which as a density 133 lower than water (0.98 kg m⁻³). The maximum value of Stokes number, St, reached in the previous 134 135 experiments was 0.024 (500 µm particles) whereas for the experiments carried out in this work using mimic milk and cream fluids the maximum value of St is 0.002. Although these values are larger than 136 137 those for optical diagnostic methods such as Particle Image Velocimetry (for which St < 0.0001 using 138 10 µm tracers), the values are still sufficiently low that the mean flow will be tracked with a high degree of accuracy (Adrian, 1991). Beside, the size reduction had the advantage that the particle was 139 140 more resistant to damage through the pump and flow loop and thus acquisition times improved with a 141 consequent increase in the measured number of passes per experiment.

- 142 PEPT is performed by mapping of the position the particle in both space and time using the
- 143 Birmingham ADAC Forte positron camera. To reconstruct the particle position, triangulation of the

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144 gamma ray pairs is performed which allows the location of the particle to be detected with a reasonably high spatial resolution several hundred times per second, e.g. a tracer travelling at 1 m s⁻¹ 145 146 (which is of the order of the flow velocity within the turbulator) can be located to within < 0.5 mm for a 147 250 µm particle at an acquisition rate of 250 Hz. The range of the PEPT experiments was limited by the 400 mm width of the detector heads and therefore two detector positions were used to record the 148 149 motion of the PEPT particle within the UV section of the turbulator: regions A and D for the first position, and regions B and C for the second position, as shown in Figure 1. The total number of 150 151 particle passes recorded through the top and bottom turbulators are shown in Table 3, where a pass 152 is defined as a series of contiguous particle locations where the PEPT particle motion is observed in 153 either the top or the bottom turbulator UV section.

154 **2.3 Microbiological data**

The microbiological data were obtained from Crook et al. (2014). In their work, Serratia marcescens 155 156 ((SM), source ATCC and strain 13880), Aeromonas hydrophila ((AH), source ATCC and strain 7966) 157 Listeria monocytogenes ((EC), source ATCC and strain 4388), and Escherichia coli ((LM), source 158 ATCC and strain 43256) were selected as model milk pathogens. This group of vegetative pathogens 159 of concern is primary associated with milk and dairy products and milk related illnesses. The 160 microorganisms were prepared and added to UHT treated milk; such an approach allows the 161 determination of the initial bacteria population before and after treatment that is fundamental for the 162 following analysis (Rossito et al., 2012). UV-C treatment of the inoculated milk was then performed in a pilot-scale SurePure Turbulator[™] consisting of four turbulators connected in series. Each 163 164 turbulator contains an 11.8 W low pressure mercury UV bulb emitting at 254 nm installed in optically 165 pure quartz sleeve to separate the milk fluid from the UV bulb. The annulus volume of the turbulator is 0.675 L which at the design flow rate of 4000 L hr⁻¹ results in a mean residence time of 0.6 s for 166 167 each turbulator. The fluid is pumped in a 0.9 to 1.6 mm channel over the quartz sleeve at 4300 L hr⁻¹. 168 The milk was processed in a closed loop at a fixed temperature of 280 K until the required inactivation 169 of bacteria was achieved. The resulting sets of data were used to validate the developed surface 170 refreshment models and to enable calculation of the required UV-C exposure time.

171 2.4 Analysis of PEPT data

The PEPT data is analysed using the previous procedure in Simmons et al. (2012) and is outlined
here. A summary of the principal steps is presented below: a time density function, *f*, is calculated

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which is the fraction of the total time spent by all passes at a given radial distance from the lamp. Let the total number of measured passes be N_P , the time for each pass be τ_P , the number of measurement points in each pass be *J*, and the number of measurement points located in an annular volume of $(\pi/4)L(r_{m+1}^2 - r_m^2)$ be *j*. In addition, let *M* be the total number of annular shells between R_I and R_2 chosen to evaluate the function. Since the PEPT has a resolution of 0.5 mm, the radial gap between shells is set to a multiple close to this value i.e. 0.46 mm \approx 0.5 mm. Then, for a single pass

 $\tau_{r_{m+1},r_m} = \frac{J}{J}\tau_P$

181 and over all passes the time density function

182

 $f_{r_{m+1},r_m} = \frac{\sum_{i=1}^{N_P} \frac{j}{J} \tau_P}{\sum_{i=1}^{N_P} \tau_P}$

(5)

(4)

183 To ensure that the volume is conserved

184

 $\sum_{m=1}^{M} f_{r_{m+1},r_m} = 1$ (6)

Making the assumption that this fraction is representative of the history of the entire fluid volume passing through the turbulator, this allows a model for microbial reduction to be developed. This assumption is only strictly true for an infinite number of passes; however at least ~1000 passes are measured per flow rate as shown in Table 3.

Since the food fluids modelled are highly opaque, the expected fluence rate drops extremely rapidly with radial distance from the UV lamp surface. Therefore the time density function, *f*, can be used to build a simplified kill model based upon the average residence time spent by the fluid in the annular shell closest to the lamp surface; microbial kill in all other shells being assumed as negligible. The model for microbial inactivation used previously based upon all M shells is

194
$$\frac{N}{N_o} = \prod_{m=1}^{M} e^{-k_1 \bar{I}_{f_{r_{m+1},r_m}}\bar{\tau}}$$
(7)

195
$$\frac{N_4}{N_0} = \left[\frac{N}{N_0}\right]_{BOTTOM} \times \left[\frac{N}{N_0}\right]_{TOP}^3$$
(8)

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196 where the arithmetic mean fluence \overline{I} , in each shell with radii between r_m and r_{m+1} , can be calculated

198
$$\bar{I} = \frac{I_0}{(r_{m+1} - r_m)} \int_{r=r_m}^{r=r_{m+1}} e^{-\alpha r} dr = \frac{I_0 (e^{-\alpha r_m} - e^{-\alpha r_{m+1}})}{\alpha (r_{m+1} - r_m)}$$

199 and $\overline{\tau}$ is the mean residence time:

$$\overline{\tau} = \frac{L}{U_0}$$

Since the shell closest to the UV-C source only receives significant fluence, (7-9) above can be simplified to consider microbial kill which occurs for a time t_x , where $x \rightarrow 0$. Within the measurement capability of the PEPT technique, this occurs for x < 0.5 mm after which the fluence rate will have decayed almost completely ($||I_o| < 10^{-5}$). Thus assuming only the shell closest to the lamp (r < 0.5 mm) is active the model reduces to

$$\left[\frac{N}{N_0}\right]_{r_m,r_{m+1}} = e^{-k_1 \bar{l} f_{0.5} \bar{\tau}}$$
(11)

where $f_{0.5}$ is the time density function for r < 0.5 mm which is different between bottom ($f_{0.5BOT}$) and top ($f_{0.5TOP}$) turbulators. Simmons *et al.* (2012) demonstrated that the values for the top turbulator are representative of all subsequent turbulators thus the density time distribution for uses $f_{0.5BOT}$ for the first and $f_{0.5TOP}$ for the remainder. This replaces equation (7) and the log inactivation over 4 turbulators can be obtained using (11) in (8) with \overline{I} calculated using (9) for the first shell only.

The model was further modified to adapt the non-linear kinetics described by the Weibull distribution model. The conventional way of calculating the efficiency of any treatment in food preservation is based on the assumption that survival curves of microorganisms are governed by first-order kinetics (2). A linear relationship between the number of surviving microorganisms and time is used. In many real cases the survival curves are not linear and present concavity which is well described by the Weibull model:

218
$$\frac{N}{N_o} = \exp\left(-k_1 I_x t_x^{\ n}\right)$$
(12)

(10)

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where n is the shape factor of the curve. Thus when n<1, when the distribution has a strong right skew, the semi logarithmic survival curve has a noticeable upward concavity. When n>1, the semi logarithmic survival curve has a pronounced downward concavity (Peleg *et al.* 1997). Modified Weibull models (Albert *and* Mafart, 2005) can better fit microbial inactivation data. The microbial model for 4 turbulators using the Weibull distribution equation (13) has been derived using equations (8) and (12). Parameters such as \overline{I} (17.67 W m⁻²), calculated using equation (10), $\overline{\tau}$ (0.565 s) and f_{0.5} (TOP and BOTTOM, see Table 4) are set constant in (11) and (13).

226

227
$$\frac{N_4}{N_0} = e^{-k_1 \bar{I} (f_{0.5_{BOT}} + 3f_{0.5TOP})\bar{\tau}}$$

(13)

Ģ

228 2.5 Statistical Analysis

All microbiological data published by Crook et al. (2014) were obtained in triplicate from which the corresponding standard deviations were calculated. The accuracy of the PEPT data is established by examination of the stability of the mean and standard deviation of the velocity; the fluctuation of these properties is less than 2.5% after the acquisition of ~1000 passes which is twice as accurate as obtained previously. Corresponding errors on both cumulative fraction and time fraction calculations are thus also of the order of 2.5%.

235

236 3. Results and Discussion

237 3.1 Flow behaviour of milk model fluids

Due to process conditions, only turbulent and transitional flow regimes have been considered in this
 work. For the milk mimic, Table 2 shows that the Reynolds numbers are significantly above the critical

240 values of 2300 in all parts of the equipment, indicating fully developed turbulent flow conditions

241 throughout. In contrast, the Reynolds number for the more viscous cream mimic fluid has a value

242 which indicates transitional flow (between laminar and fully developed turbulent flow).

- 243 The cumulative fraction of passes as a function of non-dimensional radius is shown for milk and
- cream model fluids in Figure 2. Corresponding data for water presented in Simmons *et al.* (2012)
- showed a similar general trend. The cumulative fraction at the highest flow rate is shifted slightly to

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the left for the top turbulator in Figure 2a, otherwise the data appear to be only weakly dependent onflow rate and viscosity.

The time density functions (refer to paragraph 2.4 equation 5), *f*, for both fluids in the top and bottom turbulators are shown in Figure 3. The data show that the time density function exhibits a positive skew for all flow rates. Table 4 shows that the values of $f_{0.5}$ are ~0.1 for the cream and ~0.2 for the milk for the top turbulator, the values for the bottom turbulator are reversed. The reason for the reversal is unclear; however the entry boundary conditions for the fluid entering the bottom turbulator (from the pump) would be expected to be different from the top turbulator, which is fed by the one beneath it, and all subsequent turbulators.

255 For both top and bottom turbulators the trends indicate a strong effect of viscosity but only a weak 256 effect of flow rate over the narrow range of +/- 300 L/hr measured, each increment of 7.5% in flow 257 rate gives a change of less than 5% in the value of f_{0.5}. The value of f_{0.5} for water, obtained at the 258 same value of non-dimensional radius of 0.08 in Simmons et al. (2012), ranges from 0.06-0.08, which 259 indicates that the increased viscosity, in turbulent regime, actually leads to a slightly larger proportion 260 of the fluid being refreshed at the inner surface. The value of $f_{0.5}$ for water, obtained at the same value of non-dimensional radius of 0.08 in Simmons et al. (2012), ranges from 0.06-0.08, which indicates 261 that the increased viscosity, in turbulent regime, actually leads to a slightly larger proportion of the 262 263 fluid being refreshed at the inner surface. The value for the more viscous fluid is 0.2 which is larger 264 than the values of 0.06-0.08 found for water. One could postulate that the turbulence within the flow, which dissipates energy, will act to reduce the angular momentum of the flow due to the stochastic 265 266 nature of the random velocity fluctuations. As the turbulence level decreases (due to decreased 267 Reynolds number, Table 2), the advective swirls generated by the inlet chambers and the sinusoidal 268 wall shape of the Surepure Turbulator within the flow which is introduced, may persist longer. The 269 swirl generated by turbulence may disrupt the advective swirls thus it can affect negatively the amount 270 of surface refreshment. However the resolution of these measurements does not allow this 271 hypothesis to be tested. A full computational fluid dynamics would be required to confirm if this is 272 indeed the case which is beyond the scope of this paper. However, these results give some basis to 273 identify a benefit when using the turbulator to process higher viscosity fluids in turbulent regime.

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274 3.2 Hydrodynamic based models for microbial reduction: First order kinetic versus Weibull 275 distribution model 276 The data from the microbial inactivation study of Crook et al. (2014) have been used to compare and 277 contrast the first order and the Weibull inactivation models and generate microbial inactivation parameters as k1, N0 and n. Two approaches are used to account for the concavity in the data 278 279 observed for Escherichia coli O157:H7 inactivation in Figure 4. Firstly, the first order model is fitted over two linear ranges in the data, leading to two values of k_1 for each range (first order – double), 280 281 with the intercept between the two regions selected at the 5 log reduction point. Secondly, the Weibull model (13) is applied. Fitting parameters and R² values are shown in Table 5. As expected, 282 283 the single first order model does not fit the data and the double part first order shows improvement. 284 Finally, the Weibull model enables an improved fit to the whole data set with fitting parameters k_1 and 285 n.

The first order -double and Weibull models are fitted to the experimental microbial inactivation data for all 4 microorganisms in Figure 5. The microbial survival curves clearly show non-linearity. The fitted kinetic parameters (inactivation rate k_1 and n-values) are listed in Table 5. The values of the inactivation rate constant, k_1 are set as identical in both models for each specific microorganism to enable a direct comparison between first order and Weibull models. The range of fitted k_1 values for the tested microorganisms in milk are close to those of 0.03 to 0.06 m² J⁻¹ reported by Koutchma (2009).

It is important to point out that comparison of the values of UV inactivation rates k₁ among these four most common milk pathogenic organisms indicates that *Listeria monocytogenes* has the lowest UV inactivation rate and correspondingly the highest UV resistance in milk. The least UV resistant pathogenic organism in milk was *Escherichia coli* O157:H7 (Table 5).

For the Weibull distribution model, the changes in n-value can serve as a function of the UV resistance of the microorganisms, where the higher the n-value equates to a higher UV resistance such as n of 0.67 for *Listeria monocytogenes*. This is also shown by the concavity of the curve in Figure 5 where all the microbial inactivation data are plotted versus the UV-C exposure time. The time is related directly to the average residence time of the fluid in a single turbulator. The UV dose received is directly proportional to the exposure time near the UV source and the energy emitted by a single UV-C bulb. All the continuous lines represent the fittings of the Weibull distribution model

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which show good agreement with the experimental data. Comparing the coefficients of determination (R²) (see Table 5) for the Weibull model, the fits for *Aeromonas hydrophila* and *Escherichia coli* O157:H7 bacteria are the most accurate. *Serratia marcescens* and *Listeria monocytogenes* have the worst fit due to the double concavity in the range from 0 to 50 s. The first order model gives the poorest fits, as expected, with the exception of *Serratia marcescens*, however they allow the extrapolation of consistent results in a range of ~15% from the measured values.

Referring to Table 6 the experimental and the estimated data of UV energy to achieve 5-log reduction 310 in milk are presented. A maximum UV energy of 1100 -1120 kJ m⁻³ will be required for the process to 311 312 deliver a 5 log reduction of the most resistant pathogen, Listeria monocytogenes, in milk. Only in the 313 case of Serratia marcescens is there an underestimation of the UV energy required for the treatment 314 of milk, which is due to the double concavity of the trend. For the other microorganisms the models 315 overestimated the UV energy required with a maximum error of 16.6% for the first order model. As 316 expected the Weibull model gives better estimation with a maximum error of 12.2%. However, the 317 better fitting of experimental data is due to the additional parameter (n) in the Weibull model.

318 In the literature there are no studies that investigate the nature of n, apart from the mathematical aspects related to the concavity of the curve (Peleg et al. 1997). This is a limitation of the Weibull 319 320 model which can be used only as experimental data fitting model. While the linear model can be 321 employed for a simulation since the parameters of the model are better known in the literature. In 322 comparison, the linear model can be always used for an estimation of a working range of exposure 323 time or UV energy required for the inactivation of microorganisms using the cited k1 range. For all 324 presented data the 5 Log reduction always occurs in the initial (steep) slope. This means that the first 325 part of trend is key and thus only the first part of the first order - double fit (Figure 5) has to be 326 considered if the 5 Log reduction is the sole concern.

Both hydrodynamic models show lower values of the energy required for the UV-C treatment of different microorganisms compared to the values determined by Crook *et al.* (2014). This phenomenon can be explained by the overestimation of milk residence time in the turbulator for 5-log inactivation in their work. The use of the hydrodynamic models developed here allow for more accurate energy estimation because they take into account the effect of viscosity upon the fluid flow, thus the fluid refreshment at the lamp surface. The residence time at the surface of the lamp is

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affected by the viscosity, which is highlighted in this work by the difference between the values of $f_{0.5}$ for water, milk and cream.

335 4. Conclusions

The PEPT technique has been applied to determine the hydrodynamic performance of a full-scale transparent model of a SurePure TurbulatorTM used for pathogens inactivation in milk and cream using UV-C radiation. This study shows that the surface refreshment is enhanced when the fluid viscosity is increased at constant flow rate, which affects the residence time of the fluid on the surface of the UV-C lamp; a large difference between the values of $f_{0.5}$ obtained for water, milk and cream are observed. Conversely, the effect of flow rate on the values of $f_{0.5}$ is rather weak.

- 342 These results have been used to calculate "corrected" residence times for each fluid in the
- 343 Turbulator[™] and develop both first order and Weibull distribution inactivation models. Fitting these 344 models to microbial inactivation data obtained by Crook et al. (2014) has shown excellent agreement 345 with the latter Weibull model. The models thus enable a more accurate estimation of the required UV 346 energy for the inactivation of the microorganisms from the effective residence time. The results show 347 that the use of UV-C radiation combined with the surface refreshment flow principle in the turbulators
- 348 is effective for the inactivation of bacteria in low UV transmittance dairy fluids. The proposed models
- 349 allow precise calculation of the required UV-C exposure dose. The maximum UV energy required for
- 350 the process to deliver a 5 log reduction of the most resistant pathogen has been found for Listeria
- 351 monocytogenes. Generally, the predicted values of energy required are always overestimated within
- 352 ~16% using the double first order and ~12% using the Weibull model. Serratia marcescens is the only
- 353 case where there is an underestimation, which is due to the double concavity of the trend.

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Tables

Table 1: Experimental conditions

Fluid type	Flow rates	Fluid viscosity	Fluid viscosity	Density
	Q	μ	Standard	ρ
(-)	(L hr ⁻¹)	(Pa s)	Deviation	(kg m ⁻³)
Milk mimic: 40% Glycerol-water solution	3700; 4000; 4300	0.0036	0.0002	1104
Cream mimic: 50% Glycerol-water solution	4000	0.0123	0.0003	1130

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Flow rate	Flow rate	superficial	Reynolds	Reynolds	
		velocity	number	number	
			milk	cream	X
Q	Q×10 ⁴	Uc	Re _M	Re _c	2
(L hr ⁻¹)	(m ³ s ⁻¹)	(m s ⁻¹)	(-)	(-)	6
3700	10.28	1.88	5980		
4000	11.11	2.02	6430	2250	
4300	11.94	2.18	6940	2 -	
			2		
0					

Table 2: Reynolds numbers in the UV section for the flow conditions and fluids used

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Flow rate 'fluid'	Number of	Number of		
0	passes	recorded	Average number of	
Q	N _P	points, N	points per pass	
(L hr ⁻¹)	(-)	(-)	0	
'milk'				•
3700	922	6304	6.84	
4000	1284	9035	7.04	
4300	1080	7027	6.51	
'cream'				
4000	1134	7645	6.74	

Table 3: Summary of PEPT experiments and number of measured passes.

Table 4: Values of f0.5 at each flow condition.

 Table 5: List of the inactivation rate parameters of milk borne bacteria for the fitting by first

 order-Double and Weibull distribution models in milk.

-	Microorganisms	k₁ [m² J⁻¹]	Log(N₀) [-]	n [-]	R ² First order- Double	R ² Weibull
-	SM	0.035	7.34	0.58	0.89	0.79
-	AH	0.049	6.00	0.57	0.93	0.97
-	EC	0.079	7.13	0.45	0.86	0.92
-	LM	0.025	7.08	0.67	0.73	0.81

 Table 6: Estimation of UV energy (KJ m-3) required for 5 Log reduction of pathogen organisms

 in milk using linear and Weibull distribution models.

Microorganisms	Experimental	First	Error %	Weibull	Error %
	Data	Order-D	(FOD)	Distribution	(WD)
		(FOD)		(WD)	
SM	860	750	-12.7	800	-6.7
AH	645	750	+16.2	650	+0.7
EC	300	350	+16.6	330	+10.0
LM	980	1120	+14.2	1100	+12.2

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Figure 3: Plot of time density function, f, as a function of non-dimensional radius at each flow rate in (a) top TurbulatorTM; (b) bottom TurbulatorTM. Glycerol-water mixtures representing milk and cream are the working fluids.

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Figure 1: Schematic of rig and PEPT imaging (Simmons et al. 2012)

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Figure 2: Plot of cumulative fraction of passes as a function of non-dimensional radius at each flow rate for (a) top turbulatorTM; (b) bottom turbulatorTM. Glycerol-water mixtures representing milk and cream are the working fluids.

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MAS

C

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Highlights

- The PEPT technique has been applied to determine the hydrodynamic performance
- This study shows that the surface refreshment is enhanced when the fluid viscosity is increased at constant flow rate
- These results have been used to calculate "corrected" residence times for each fluid in the Turbulator[™]
- Both first order and Weibull distribution inactivation models have been developed

The models enable a more accurate estimation of the required UV energy for the inactivation of the microorganisms