

Reaction pathways of the chlorination of amino acids at Cl:AA \geq 2

1

2 **Graphical Abstract**

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4 **CHLORINATION OF AMINO ACIDS: REACTION PATHWAYS AND REACTION**
5 **RATES**

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16 compounds

17
18 **Abstract**

19 Chlorination of amino acids can result in the formation of organic monochloramines or organic
20 dichloramines, depending on the chlorine to amino acid ratio (Cl:AA). After formation, organic
21 chloramines degrade into aldehydes, nitriles and *N*-chloraldimines. In this paper, the formation
22 of organic chloramines from chlorination of lysine, tyrosine and valine were investigated.
23 Chlorination of tyrosine and lysine demonstrated that the presence of a reactive secondary
24 group can increase the Cl:AA ratio required for the formation of *N,N*-dichloramines, and
25 potentially alter the reaction pathways between chlorine and amino acids, resulting in the
26 formation of unexpected by-products. In a detailed investigation, we report rate constants for
27 all reactions in the chlorination of valine, for the first time, using experimental results and
28 modelling. At Cl:AA = 2.8, the chlorine was found to first react quickly with valine (5.4×10^4
29 $\text{M}^{-1} \text{s}^{-1}$) to form *N*-monochlorovaline, with a slower subsequent reaction with *N*-
30 monochlorovaline to form *N,N*-dichlorovaline ($4.9 \times 10^2 \text{M}^{-1} \text{s}^{-1}$), although some *N*-
31 monochlorovaline degraded into isobutyraldehyde ($1.0 \times 10^{-4} \text{s}^{-1}$). The *N,N*-dichlorovaline then
32 competitively degraded into isobutyronitrile ($1.3 \times 10^{-4} \text{s}^{-1}$) and *N*-chloroisobutyraldimine
33 ($1.2 \times 10^{-4} \text{s}^{-1}$). In conventional drinking water disinfection, *N*-chloroisobutyraldimine can
34 potentially be formed in concentrations higher than its odour threshold concentration, resulting
35 in aesthetic challenges and an unknown health risk.

36

37 **Introduction**

38 Organic chloramines, a subclass of nitrogenous disinfection by-products (N-DBPs), can form
39 in water when the organic nitrogen components of organic matter react with free chlorine¹ or
40 inorganic chloramines.^{2,3} It has been reported that N-DBPs have significantly higher toxicity
41 than the carbonaceous DBPs,^{4,5} and organic chloramines were identified by Bull et al.⁶ as a N-
42 DBP subclass of high interest based on their potential carcinogenic hazards. An *in vitro* study
43 by Laingam et al.⁷ also found that *N*-chloroglycine, *N*-chlorolysine, *N*-chloroethanolamine and
44 *N*-chlorohistamine were potentially genotoxic and cytotoxic to humans at parts-per-billion
45 concentrations.

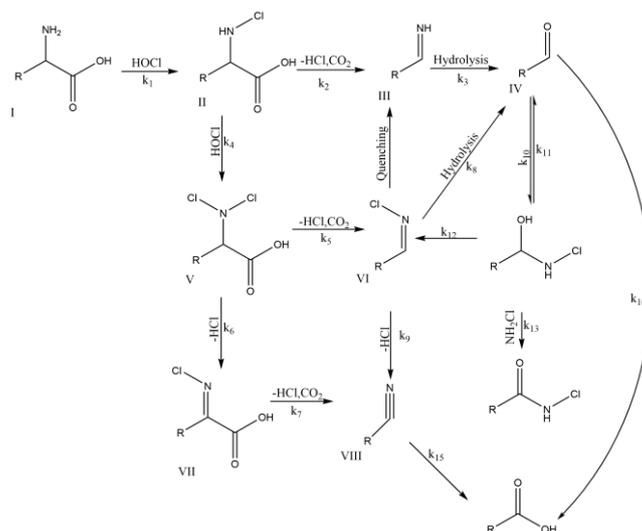
46 Amino acid concentrations in natural waters can range from 20 to 10000 $\mu\text{g L}^{-1}$,⁸
47 accounting for between 2 and 13% of dissolved organic carbon⁸ and up to 75% of dissolved
48 organic nitrogen⁹. In general, the concentration of combined amino acids such as proteins and
49 peptides are four to five times higher than free amino acids.^{8, 10} Although, free amino acids
50 only contribute a small fraction of the total amino acids,^{8, 11, 12} free amino acids are poorly
51 removed during biological filtration,¹³ and concentrations of free amino acids can even increase
52 after sand filtration.¹⁴ Hence, it is likely that free amino acids will be present in waters during
53 drinking water disinfection. Amino acids are known to form organic chloramines (*N*-
54 chloramino acids) when chlorinated,^{15, 16} and are believed to be the main contributors to the
55 formation of organic chloramines from organic matter.^{17, 18} Amino acids can also form organic
56 bromamines if brominated, or if bromine is present in the water.¹⁹ However, organic
57 chloramines formed from amino acids have been found to be unstable.^{20, 21} In our study of 18
58 organic chloramines formed from amino acids, we found that 12 had a half-life of less than 90
59 minutes.²⁰ Consequently, as the time between chlorination of drinking water and final
60 distribution to the end consumer increases, it becomes more likely for consumers to be exposed
61 to the degradation products of organic chloramines, rather than the organic chloramines
62 themselves. Therefore, a full assessment of the health impact of organic chloramine formation
63 in drinking water must include the identification of stable by-products from the reaction of
64 amino acids and chlorine.

65 Previous studies of organic chloramines formed from isoleucine, phenylalanine and valine
66 have shown that many organic chloramine degradation by-products retain structural
67 components from the original amino acid precursor,²²⁻²⁴ but that DBPs like the
68 trihalomethanes,²⁵ haloacetic acids²⁵ and haloacetonitriles²⁶ can also form. The main
69 degradation product identified from organic monochloramines has been the corresponding

70 aldehyde, while aldehyde, nitrile and *N*-chloraldimine degradation products have all been
71 detected from organic dichloramines.²³ A generalised formation and degradation pathway has
72 previously been proposed to fit these experimental results (Figure 1),^{23, 27, 28} however not all
73 the species proposed have been confirmed experimentally, in part due to the lack of suitable
74 analytical methods. In addition, rate constants for the reactions in this generalised formation
75 and degradation pathway have not yet been determined.

76 In this paper, we have further investigated the chlorination of amino acids using valine as
77 a model compound. To facilitate improved understanding of the reaction pathways of the
78 chlorination of valine, liquid chromatography- and gas chromatography-based mass
79 spectrometric methods were developed for the identification of organic monochloramines, and
80 aldehyde, nitrile and *N*-chloraldimine by-products. The effect of secondary functional groups
81 (apart from the amino acid group) on the formation of organic chloramines was also studied
82 using lysine, tyrosine, and valine. Finally, the rates of formation of the aldehyde, nitrile and *N*-
83 chloraldimine were also investigated from chlorination of valine using both modelling and
84 experimental results. This is the first study to investigate the rate of reaction for every step in
85 the reaction pathway for the chlorination of valine.

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Figure 1. Reaction pathways for the chlorination of amino acids. (Adapted from Conyers and Scully, 1993²³; Kimura, 2015²⁷; Yu and Reckhow, 2015²⁸)

91 **Materials and Methods**

92 **Formation of organic chloramines.** Chemicals and materials used are detailed in the
93 Supporting Information (SI) Text S1. The amino acids used in this study (Table S1) were
94 chosen because of data existing in previous studies (valine), or because of the presence of
95 specific secondary functional groups (lysine and tyrosine). In addition, the organic
96 monochloramines that form from these amino acids have half-lives of 45 min or longer,²⁰ which
97 was the time required for each chromatographic separation. Organic chloramines (0.03 mM or
98 1.6 mM) were formed by chlorination of individual amino acids (0.04 mM or 2 mM) at chlorine
99 to amino acid molar ratios (Cl:AA) of 0.8 and 2.8. The lower concentration (0.03 mM) was
100 used for gas chromatography-mass spectrometric (GC-MS) experiments, while the higher
101 concentration (1.6 mM) was used for liquid chromatography coupled with ultraviolet and high
102 resolution mass spectrometric (LC-UV-HRMS) experiments. The Cl:AA ratios were chosen
103 because previous studies have shown that only organic monochloramines form at Cl:AA = 0.8
104 and only organic dichloramines form at Cl:AA = 2.8.^{20, 22} The conditions used for the reaction
105 were: pH 7.5 at room temperature (20-25 °C). Reaction mixtures were reacted in the dark in air
106 tight vials.

107

108 **Analytical methods.** Amino acids and organic chloramines were measured using LC-UV-
109 HRMS analysis. The chromatographic conditions were as described in our previous work.²⁹
110 Analytes were detected first by UV ($\lambda=255$ nm) using an Accela photodiode array detector
111 (ThermoScientific, Waltham, USA), then by high resolution mass spectrometry (HRMS) using

112 a LTQ Orbitrap XL (Thermo Scientific) fitted with electrospray ionization operated in either
113 positive or negative ionization mode, as previously described.²⁰ HRMS screening used a full-
114 MS scan in the range 70-300 m/z with a mass resolution of 15000 at 400 m/z . Organic
115 chloramines were identified by comparing the measured mass and isotope pattern against the
116 theoretical mass (< 5 ppm, relative error) and the predicted isotope pattern, while amino acids
117 were confirmed by comparison with their respective analytical standards. Data was processed
118 using X-calibur QualBrowser 2.0.7 SP1 (Thermo Scientific). The HRMS instrumental
119 conditions are provided in the Supporting Information (Table S2).

120 Volatile degradation by-products, namely the aldehyde, nitrile and *N*-chloraldimine, were
121 extracted using an automated headspace solid-phase microextraction (HS SPME) GC-MS
122 method previously developed by our group,³⁰ using a 75 μm CAR/PDMS fibre. Sodium
123 sulphate (3 g) and the internal standard solution (1,2-dibromopropane- d_6 in methanol, 100 μg
124 L^{-1} , 2 μL) were added to each sample (10 mL in a 20 mL glass vial). Samples were agitated at
125 500 rpm at 60 $^\circ\text{C}$ for 10 minutes before the SPME fibre was inserted into the sample headspace
126 for 15 minutes and then transferred to the GC injector port for thermal desorption at 300 $^\circ\text{C}$ for
127 3 minutes. GC-MS analysis used an Agilent 6890N gas chromatograph coupled with an Agilent
128 5975 mass selective detector using conditions reported in SI Table S3. The samples were
129 analysed in both MS scan mode (20 – 320 m/z) to screen for the degradation products and single
130 ion monitoring mode to quantify selected aldehydes, nitriles and *N*-chloraldimines (quantifying
131 and qualifying ions are listed in SI Table S4). Liquid-liquid extraction (LLE) was used for the
132 extraction of by-products from the chlorination of lysine (organic solvent used was
133 dichloromethane) and also from the chloramination of isobutyraldehyde in anhydrous
134 conditions (organic solvent used was diethyl ether). The GC-MS program used for the LLE
135 was identical to that of the HS SPME-GC-MS method (Table S3), except that the injector
136 temperature was 250 $^\circ\text{C}$ and initial holding time was 3 min instead of 2 min.

137 Proton nuclear magnetic resonance (^1H NMR) spectroscopy was used to determine the
138 positions of substitution of the chlorine atoms on the *N*-chloramino acids formed. All NMR
139 spectra were obtained using a Bruker AVANCE III 400 MHz spectrometer (Bruker, Australia),
140 by dissolution of amino acids or the *N*-chloramino acids in 90% water and 10% D_2O , with
141 water signal suppression.

142

143 **Reaction rates for the reaction pathway of the chlorination of valine at Cl:AA = 2.8.** To
144 measure the reaction rates of the formation of *N*-chlorovaline, chlorinated valine solutions were
145 analysed by LC-UV-HRMS. Solutions were injected 5 min after chlorination, without

146 quenching. Both the amino acid and the organic chloramine were measured at 45 min intervals
147 until 185 min after chlorination. To measure the rate of formation and degradation of
148 isobutyraldehyde, isobutyronitrile and *N*-chloroisobutyraldimine, chlorinated valine solutions
149 were analysed by HS SPME-GC-MS, without quenching, at $t = 15$ min after chlorination, and
150 then at one hour intervals until $t = 7$ hours, and 12 hour intervals from $t = 12$ to 72 hours. As
151 the samples could not be quenched for the determination of organic chloramines, each data
152 point was determined from the average of two repeated experiments. The concentration of *N*-
153 chloroisobutyraldimine was estimated as the difference in concentration of isobutyraldehyde
154 between quenched and unquenched experiments at 7 hours. The measurements were then used
155 to plot the first-order linear kinetic plot, $\ln (A_t/A_0)$ vs time in seconds, where A_t was the mass
156 abundance at a specific time and A_0 was the initial mass abundance.

157 Reaction rates that could not be determined experimentally were predicted by iteration
158 through modelling using Kintecus,³¹ as described in the SI text S2. The accuracy of the
159 predicted rate constants was assessed by a sensitivity analysis to investigate the effect of
160 varying modelled rate constants from their chosen value.

161

162 **Results and Discussion**

163 **Effect of a reactive secondary functional group on the formation of organic**
164 **monochloramines and dichloramines.** The formation of organic monochloramines and/or
165 organic dichloramines (to be referred to as monochloramines and dichloramines) from the
166 chlorination of amino acids is controlled by the Cl:AA ratio and by the presence of secondary
167 functional groups that may also react with chlorine.^{20, 32} Formation experiments using a Cl:AA
168 = 0.8 were used in this study to produce monochloramines, while avoiding dichloramine
169 formation. In our previous work, we used LC-UV-HRMS to confirm formation of both *N*-
170 chloroisoleucine and *N*-chlorovaline after the chlorination of isoleucine and valine,
171 respectively.²⁰ The same technique was used to detect and identify monochloramines formed
172 from the chlorination of the other amino acids tested in this current study: lysine and tyrosine,
173 which have not previously been reported to be detected by MS methods. For all amino acids,
174 monochloramine peaks were detected in both UV spectra and MS spectra, and the mass to
175 charge ratios of these suspected monochloramine peaks were within 5 ppm error of their
176 theoretical values (Table 1) and the measured isotopic patterns were similar to the predicted
177 isotopic patterns. In addition, comparison of LC-UV-HRMS chromatograms of unquenched
178 samples and samples quenched with ascorbic acid showed that the suspected organic

179 monochloramine peaks were absent after quenching, indicating that the suspected peaks were
180 the organic monochloramines (Figure S1).

181

182 **Table 1. Comparison of the measured mass of suspected organic monochloramines and**
183 **their theoretical mass. The difference (ppm) in the measured and theoretical mass was**
184 **always less than 5 ppm.**

Compounds	Measured mass	Theoretical mass	Difference (ppm)
Positive [M+H] ⁺			
<i>N</i> -chlorolysine	181.07369	181.07383	-0.784
<i>N</i> -chlorotyrosine	216.04179	216.04220	-1.886
<i>N</i> -chlorovaline	152.04660	152.04728	-4.491

185

186 Formation experiments using Cl:AA = 2.8 were designed to preferentially form
187 dichloramines from the amino acid precursors,^{20, 22} except for lysine where Cl:AA = 4.8 (Cl:N
188 = 2.4) was used, as discussed in more detail below. As dichloramines are, in general, not
189 detectable by UV,^{20, 22, 23, 33} suspected dichloramine peaks were confirmed by LC-HRMS,
190 including the exact mass and isotopic ratio. In addition, proton nuclear magnetic resonance (¹H
191 NMR) spectroscopy was used to determine the positions of substitution of the chlorine atoms
192 on the organic chloramines that formed, and LLE or HS SPME GC-MS were used to identify
193 by-products. *N,N*-Dichloramine species were not detected by LC-HRMS for lysine or valine.
194 This might be due to the reported instability of the *N,N*-dichloramine species formed from α -
195 amino acids.³⁴ However, other dichloramine species were detected for lysine and tyrosine,
196 which was unexpected. The exact mass to charge ratios of the detected dichloramines were
197 within 5 ppm of the theoretical values (-2.742 ppm error for lysine and -1.886 ppm error for
198 tyrosine). Lysine has two amine nitrogens in its structure and, when chlorinated, can produce
199 a chloramine functional group at both of the nitrogens. At Cl:N = 0.4, two monochloramine
200 peaks were detected from lysine (Figure S2a), suggesting that two species of monochloramine
201 were formed. At Cl:N = 1.2, only one dichloramine peak was detected (Figure S2b), most likely
202 the *N,N'*-dichlorolysine, given the subsequent detection of the corresponding aldehyde, 5-
203 chloraminopentanal, by LLE and analysis with GC-MS. When the Cl:N ratio was increased to
204 2.4, no peaks corresponding to any dichlorolysines were detected by LC-HRMS, and the
205 corresponding aldehyde of *N,N',N'*-trichlorolysine, 5-dichloraminopentanal, was detected by
206 LLE GC-MS. The formation of *N,N'*-dichlorolysine at a Cl:AA ratio of 2 and *N,N',N'*-
207 trichlorolysine at a Cl:AA ratio of 4 was also inferred by Conyers and Scully,³⁵ through the
208 detection of the corresponding aldehydes.

209 Although accurate chemical formula of the chlorinated products formed from tyrosine and
210 detected by LC-HRMS could be determined from the measured accurate mass to charge ratio,
211 LC-HRMS could not definitively indicate the position of the chlorine atoms, as chlorine was
212 the first atom to be removed during fragmentation. Therefore, samples suspected to contain *N*-
213 monochlorotyrosine and *N,N*-dichlorotyrosine were analysed using ¹H NMR (Figure S3) and
214 ¹³C NMR (Figure S4) spectroscopy. Complete details of the NMR results are discussed in SI
215 Text S3, with a summary provided here. Tyrosine has two reactive functional groups (phenol
216 and amino acid), and the NMR results confirmed that *N*-monochlorotyrosine was formed at
217 Cl:AA = 0.8. However, at Cl:AA = 2.8, new aromatic signals consistent with chlorine
218 substitution on the aromatic ring were observed, and a mixture of *N*-monochloro-3-
219 chlorotyrosine and *N,N*-dichlorotyrosine in a slightly lower than 1:1 ratio was formed. Finally,
220 when the Cl:AA ratio was increased to 12, degradation of the aromatic ring of tyrosine was
221 observed.

222 Thus, while all amino acids tested demonstrated similar reaction pathways, with formation
223 of *N*-monochloroamino acids at Cl:N ≤ 1 and formation of dichloroamino acids at Cl:N ≥ 2, results
224 from the chlorination of lysine and tyrosine at Cl:AA = 2.8 demonstrate that the presence of
225 other reactive groups in the amino acids can delay formation of the *N,N*-dichloramine to higher
226 Cl:AA ratios, or lead to formation of dichloramine isomers other than the *N,N*-dichloramine
227 species.

228
229 **Investigation of the reaction pathways of degradation of *N*-chloroamino acids using**
230 **valine as a model compound.** In order to better understand the generalised formation and
231 degradation pathways for *N*-chloroamino acids, the rate of reaction for each pathway was
232 investigated using experimental and modelling methods. Valine was chosen as a model amino
233 acid for this investigation due to the detection of its *N*-monochloroamino acid and the detection
234 of all three degradation products and valine is reported to be one of the most abundant amino
235 acids in surface waters.^{36, 37} The reaction rates for each reaction step in the reaction of valine
236 with chlorine were determined experimentally or predicted by iteration through modelling
237 using Kintecus.³¹

238 The investigation of reaction rates for each reaction pathway in the chlorination of valine
239 as a model amino acid (Table 2) showed that the *N,N*-dichloramine was formed from a stepwise
240 reaction through the formation of the *N*-monochloramine, and that the initial formation (first
241 15 min) of the aldehyde at Cl:AA > 2 was from the degradation of the *N*-monochloramine,
242 rather than from the *N*-chloraldimine. The investigation also confirmed that the nitrile and the

243 *N*-chloraldimine both formed from the *N,N*-dichloramine *via* competitive reaction pathways.
 244 After formation, the *N*-chloraldimine continued to degrade slowly to either the aldehyde or the
 245 nitrile. The nitrile and aldehyde also degraded slowly to presumably the corresponding
 246 carboxylic acid; in the case of valine, isobutyric acid. The formation of isobutyric acid was
 247 confirmed from the GC-MS detection and confirmation of isobutyric acid in the reaction
 248 mixture from the chlorination of valine. Further discussion of these pathways is presented
 249 below.

250

251 **Table 2: Proposed reaction rates for the chlorination of valine.**

Reaction (number)	Rate	Units
Valine + HOCl → <i>N</i> -monochlorovaline (k_1)	5.4×10^4	$M^{-1} s^{-1}$
<i>N</i> -Monochlorovaline → isobutyrimine (k_2)	5.0×10^{-4}	s^{-1}
Isobutyrimine → isobutyraldehyde (k_3)	5.0×10^4	s^{-1}
<i>N</i> -Monochlorovaline + HOCl → <i>N,N</i> -dichlorovaline (k_4)	4.9×10^2	$M^{-1} s^{-1}$
<i>N,N</i> -Dichlorovaline → <i>N</i> -chloroisobutyraldimine (k_5)	7.0×10^{-5}	s^{-1}
<i>N,N</i> -Dichlorovaline → 2-(chlorimino)-3-methylbutanoic acid (k_6)	1.0×10^{-5}	s^{-1}
2-(Chlorimino)-3-methylbutanoic acid → isobutyronitrile (k_7)	1.0×10^{-2}	s^{-1}
<i>N</i> -Chloroisobutyraldimine → isobutyraldehyde (k_8)	4.0×10^{-6}	s^{-1}
<i>N</i> -Chloroisobutyraldimine → isobutyronitrile (k_9)	1.0×10^{-6}	s^{-1}
Isobutyraldehyde + NH_2Cl → 1-(Chloroamino)-2-methylpropan-1-ol (k_{10})	24.3	$M^{-1} s^{-1}$
1-(Chloroamino)-2-methylpropan-1-ol → isobutyraldehyde (k_{11})	0.247	s^{-1}
1-(Chloroamino)-2-methylpropan-1-ol → <i>N</i> -chloroisobutyraldimine (k_{12})	9.75	s^{-1}
1-(Chloroamino)-2-methylpropan-1-ol + NH_2Cl → <i>N</i> -Chlorisobutyramide (k_{13})	2.7×10^{-4}	$M^{-1} s^{-1}$
Isobutyraldehyde → isobutyric acid (k_{14})	3.0×10^{-6}	s^{-1}
Isobutyronitrile → isobutyric acid (k_{15})	3.0×10^{-5}	s^{-1}

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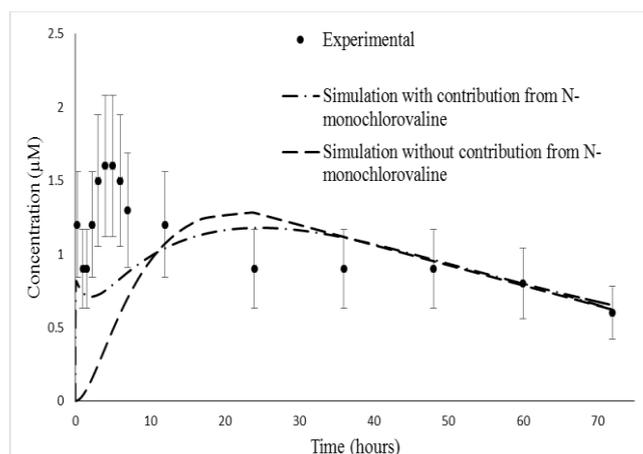
253 When the Cl:AA ratio is less than 1, the aldehyde is reported to be formed from the
 254 monochloramine through an imine intermediate (structure III in Figure 1).²³ Although the
 255 detection of this imine was not possible by LC-MS or GC-MS, the rate of formation of *N*-
 256 monochlorovaline ($k_1 = 5.4 \times 10^4 M^{-1} s^{-1}$ or $2.4 \times 10^2 s^{-1}$ for a HOCl concentration of 4.48 mM
 257 as Cl_2) and the rate of formation of isobutyraldehyde ($1.5 \times 10^{-4} s^{-1}$) were both determined
 258 experimentally. Comparison of these rates showed that the formation of *N*-monochlorovaline
 259 from the chlorination of valine is much faster than the rate of isobutyraldehyde formation.
 260 Therefore, the rate limiting step for the formation of isobutyraldehyde is either the degradation
 261 of *N*-chlorovaline into isobutyrimine (k_2) or the degradation of isobutyrimine into
 262 isobutyraldehyde (k_3). We previously found the rate of degradation of *N*-chlorovaline to
 263 isobutyrimine, k_2 , to be $1.0 \times 10^{-4} s^{-1}$,²⁰ which is similar to the rate of formation of
 264 isobutyraldehyde determined in the current study. This suggests that the rate of degradation of

265 isobutyrimine into isobutyraldehyde, k_3 , is faster than the degradation of *N*-monochlorovaline
266 into isobutyrimine (k_2), and that formation of isobutyrimine is the rate determining step for
267 formation of isobutyraldehyde from valine. Modelling using the Kintecus software indicated
268 that $k_3 = 5.0 \times 10^4 \text{ s}^{-1}$, and this suggests the imine very quickly underwent hydrolysis to the
269 aldehyde after formation. All reaction rates related to the formation of isobutyraldehyde were
270 investigated using a sensitivity analysis (SI Figure S7), and results of this were used to optimise
271 the final reaction rates, reported in Table 2, and SI Table S6.

272 It has previously been proposed that, when the Cl:AA ratio is > 2 , aldehydes can form
273 through hydrolysis of *N*-chloraldimine, a by-product of dichloramine degradation,²² rather than
274 from degradation of the monochloramine (Figure 1). However, the same study²² and
275 subsequent studies^{23, 33} also suggested the *N*-chloraldimines are stable. To assess the
276 importance of *N*-chloroisobutyraldimine in the formation of isobutyraldehyde, Kintecus was
277 used to predict isobutyraldehyde concentrations in both the presence and absence of
278 monochloramine degradation into isobutyraldehyde (Figure 2). Experimentally,
279 isobutyraldehyde showed an instantaneous formation, followed a gradual degradation. In
280 comparison, modelling without a contribution from monochloramine degradation showed
281 delayed formation of isobutyraldehyde compared to the experiment. This suggested that
282 isobutyraldehyde was not formed solely from the hydrolysis of *N*-chloroisobutyraldimine
283 (VI→IV) in experimental studies, but also from the degradation of the monochloramine
284 (II→III→IV). Thus, at Cl:AA > 1 , *N*-monochlorovaline may either react with HOCl to form
285 *N,N*-dichlorovaline (II→V) or be degraded into isobutyraldehyde (II→III→IV). We believe
286 this difference might be due to the hydrolysis of *N*-chloroisobutyraldimine to isobutyraldehyde
287 being catalysed by a reagent present in the system (likely to be HOCl), but not included in the
288 modelling.

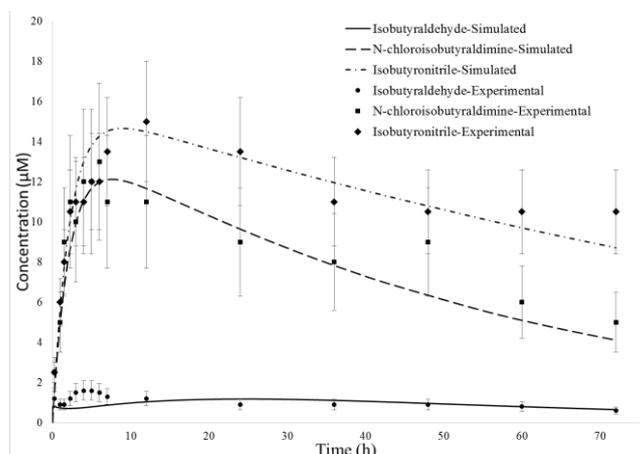
289 Figure 3 shows that, experimentally, the concentrations of isobutyronitrile and *N*-
290 chloroisobutyraldimine, both by-products of *N,N*-dichlorovaline, were much higher than the
291 concentration of isobutyraldehyde. Thus, most *N*-monochlorovaline must react with chlorine
292 to form *N,N*-dichlorovaline (k_4) rather than degrading into isobutyraldehyde. Therefore, the
293 reaction rate k_4 must be faster than the degradation of *N*-monochlorovaline. Again, a sensitivity
294 analysis of the reaction rates related to the formation of isobutyronitrile (SI Figure S8) and *N*-
295 chloroisobutyraldimine (SI Figure S9) were undertaken to optimise the final reaction rates,
296 reported in Table 2, and SI Table S6. When the final proposed reaction rates presented in Table
297 2 were used to simulate the trends of formation of isobutyraldehyde, isobutyronitrile and *N*-
298 chloroisobutyraldimine, results were very similar to the experimental trends of formation

299 (Figure 3), indicating that the proposed rate constants were reasonable estimations for the
300 reaction steps.



301
302 **Figure 2.** Concentrations of isobutyraldehyde from experimental chlorination of valine (30
303 μM), and from simulation of isobutyraldehyde formation with and without a contribution
304 from the degradation of *N*-monochlorovaline (Cl-Val).

305



306
307 **Figure 3.** Concentrations of isobutyraldehyde, isobutyronitrile and *N*-chloroisobutyraldimine
308 formed over 72 hours from both experimental and simulated chlorination of valine (30 μM).
309 The experimental concentration of *N*-chloroisobutyraldimine was estimated from the
310 difference in the isobutyraldehyde concentrations in quenched and unquenched samples.
311

312 Both isobutyronitrile and *N*-chloroisobutyraldimine are degradation products of *N,N*-
313 dichlorovaline (Figure 1). The rates of formation of isobutyronitrile and *N*-
314 chloroisobutyraldimine were experimentally determined from the increase in their
315 concentrations over seven hours (Figure 3), assuming a pseudo first order reaction, and were
316 very similar, at $1.0 \times 10^{-4} \text{ s}^{-1}$ and $9.2 \times 10^{-5} \text{ s}^{-1}$, respectively, suggesting that both of these
317 products are formed from *N,N*-dichlorovaline in competitive pathways. Previous work^{23, 33} with
318 other amino acids has also shown nitriles and *N*-chloraldehydes both formed from *N,N*-
319 dichloramines *via* competitive pathways. While it is proposed that isobutyronitrile can also

320 arise from the degradation of *N*-chloroisobutyraldimine,²³ the hydrolysis of *N*-
321 chloroisobutyraldimine into isobutyraldehyde was found to be faster than the dechlorination
322 into isobutyronitrile (discussed in detail later in this section).

323 Finally, *N*-chloroisobutyraldimine degradation into isobutyraldehyde and isobutyronitrile
324 was also investigated. Accurate measurements of the rate of degradation of *N*-
325 chloroisobutyraldimine would require a pure *N*-chloroisobutyraldimine solution. However, the
326 synthesis of pure *N*-chloroisobutyraldimine proved challenging. While *N*-
327 chloroisobutyraldimine (V) can theoretically be formed through chloramination of
328 isobutyraldehyde (IV) (Figure 1), there was no reaction observed between isobutyraldehyde
329 and inorganic monochloramine in an aqueous solution, even when inorganic monochloramine
330 was present in 20 times molar excess. This result is consistent with previous reports that the
331 formation of *N*-chloraldehydes from the chloramination of aldehydes in aqueous solutions is
332 very slow.^{27, 38} Therefore, *N*-chloroisobutyraldimine was produced in anhydrous conditions;
333 inorganic monochloramine was first produced in water and then extracted using diethyl ether.
334 The concentration of inorganic monochloramine in the diethyl ether was assumed to be the
335 difference in inorganic monochloramine concentration in the aqueous layer before and after
336 the extraction. The isobutyraldehyde standard in methanol was then added to the inorganic
337 monochloramine/diethyl ether extract at a monochloramine:isobutyraldehyde molar ratio of
338 10. The solution was then analysed by GC-MS for the formation of *N*-chloroisobutyraldimine
339 every 60 minutes (time for each chromatographic separation) for 12 hours using liquid
340 injection. Both *N*-chloroisobutyraldimine and isobutyronitrile were detected in the reaction
341 mixture, with the concentration of isobutyronitrile increasing over time (Figure S5). In this
342 experiment, the formation of isobutyronitrile must be from the degradation of *N*-
343 chloroisobutyraldimine as no other pathway exists. This result confirmed that *N*-
344 chloroisobutyraldimine can degrade into isobutyronitrile. However, further investigation into
345 the degradation of *N*-chloroisobutyraldimine using Kintecus modelling (Table 2) indicated that
346 the hydrolysis of *N*-chloroisobutyraldimine into isobutyraldehyde ($k_8 = 4.0 \times 10^{-6} \text{ s}^{-1}$) was faster
347 than the dechlorination into isobutyronitrile ($k_9 = 1.0 \times 10^{-6} \text{ s}^{-1}$). Both experimental and
348 modelled results showed a decrease in the concentration of isobutyraldehyde and
349 isobutyronitrile after formation. This is likely due to the degradation of isobutyraldehyde and
350 isobutyronitrile into isobutyric acid. Finally, calculation using the rates of formation of *N*-
351 chloroisobutyraldimine and isobutyronitrile indicated that the half-life of *N,N*-dichlorovaline
352 was around 100 min. This suggests that *N,N*-dichlorovaline was not detected because of the

353 absence of a suitable analytical technique for its detection, rather than *N,N*-dichlorovaline being
354 inherently too unstable.

355

356 **Implications of this work for drinking water disinfection.** This study of the formation and
357 degradation pathways of the chlorination of valine indicates that, when Cl:AA > 2, chlorine
358 will react quickly with valine to form *N*-monochlorovaline ($5.4 \times 10^4 \text{ M}^{-1} \text{ s}^{-1}$), with a slower
359 formation of *N,N*-dichlorovaline from *N*-monochlorovaline ($4.9 \times 10^2 \text{ M}^{-1} \text{ s}^{-1}$). Some *N*-
360 monochlorovaline will degrade to isobutyraldehyde ($5.0 \times 10^4 \text{ s}^{-1}$), while *N,N*-dichlorovaline
361 will competitively degrade to either isobutyronitrile ($1.3 \times 10^4 \text{ s}^{-1}$) or *N*-chloroisobutyraldimine
362 ($1.2 \times 10^4 \text{ s}^{-1}$). As the reaction pathways of most amino acids with chlorine are similar, the
363 speciation of degradation products (aldehydes, nitriles and *N*-chloraldimines) from other amino
364 acids is expected to be similar to that of valine, although the reaction rates of the various reactions
365 will be dependent on each individual amino acid. Amino acids with a reactive side chain, like
366 tyrosine and lysine, may not favour the formation of the *N,N*-dichloramine, depending on the
367 reactivity of the side chain functional group, and may potentially produce different degradation
368 products.

369 While organic chloramine formation is of interest due to the potential toxicity of these
370 compounds, this study has shown that the aldehyde, nitrile and *N*-chloraldimine by-products
371 are more likely to be found in finished drinking water, with potential issues for odours in
372 drinking water and as yet unknown health implications. The formation of stable aldehydes and
373 *N*-chloraldimines from the chlorination of amino acids could result in odour problems in
374 chlorinated waters.^{37, 39} For example, the odour threshold concentration of isobutyraldehyde is
375 reported to range from 0.9 to 4 $\mu\text{g L}^{-1}$,^{39, 40} while the odour threshold concentration of *N*-
376 chloroisobutyraldimine is reported to be 0.20 $\mu\text{g L}^{-1}$.⁴⁰ The maximum molar concentration of
377 valine that has been found in a drinking source water is 63 nM.^{11, 37} If this raw water was
378 chlorinated under a conventional drinking water disinfection regime, with a minimum chlorine
379 residual of 0.5 mg L^{-1} (Cl:AA ≥ 2), and without treatment to remove free amino acids, 0.2 μg
380 L^{-1} (3 nM) of isobutyraldehyde and 2.0 $\mu\text{g L}^{-1}$ (19 nM) of *N*-chloroisobutyraldimine would be
381 formed, based on the respective molar conversions of 5% and 30% found in the current study.
382 While the potential concentration of isobutyraldehyde is below the previously reported odour
383 thresholds, the potential concentration of *N*-chloroisobutyraldimine of 2.0 $\mu\text{g L}^{-1}$ is 10 times
384 higher than its odour threshold concentration (0.2 $\mu\text{g L}^{-1}$), suggesting that formation of *N*-
385 chloroisobutyraldimine during chlorination of waters containing valine could result in an odour
386 in the drinking water, especially in distribution systems with contact times less than 24 hours,

387 but also potentially result in odour problems in longer distribution systems (contact times
388 longer than 24 hours) as *N*-chloroisobutyraldimine was found to be stable for more than 8 days.
389 Additionally, no toxicity information is available for aldehydes, nitriles and *N*-chloraldimines
390 in drinking water, and thus their presence would result in an unknown health risk. Therefore,
391 improved knowledge of the occurrence of free amino acids and the odorous DBPs formed from
392 the chlorination of amino acids would be beneficial for the water industry. For drinking water
393 systems where long retention times in the distribution system (contact times longer than 24
394 hours) are expected, the degradation pathways of aldehydes and nitriles (e.g. to the
395 corresponding carboxylic acids) should also be considered.

396

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405

406 **Supporting Information Available**

407 The Supporting Information includes: Chemicals and materials (Text S1); structures for amino
408 acids used (Table S1); LC-HRMS and GC-MS conditions (Table S2-S4); modelling parameters
409 and sensitivity test data (Text S2, Tables S5-S6 and Figures S6-S8); tyrosine NMR results
410 (Text S3 and Figure S3-S4); LC chromatograms for chlorination of valine and lysine (Figure
411 S1-S2); concentrations of isobutyraldehyde, *N*-chloroisobutyraldimine and isobutyronitrile
412 over time (Figure S5). This information is available free of charge via the Internet at
413 <http://pubs.acs.org>.

414

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