



Floating modular drug delivery systems with buoyancy independent of release mechanisms to sustain amoxicillin and clarithromycin intra-gastric concentrations

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3 1 **Floating modular drug delivery systems with buoyancy independent of release**
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5 2 **mechanisms to sustain amoxicillin and clarithromycin intra-gastric**
6
7 3 **concentrations**
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51 23
52 24 Keywords: oral delivery, release module, assembled system, drug combination,
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3 26 **Abstract**
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5 27 Release modules of amoxicillin and clarithromycin combined in a single dosage form
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7 28 designed to float in the gastric content and to sustain the intra-gastric concentrations
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9 29 of these two antibiotics used for the eradication of *Helicobacter pylori* have been
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11 30 studied.

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14 31 The modules having a disc shape with curved bases were formulated as hydrophilic
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16 32 matrices. Two modules of clarithromycin were assembled by sticking the concave
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18 33 base of one module to the concave base of the other, creating an internal void
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20 34 chamber. The final dosage form was a floating assembly of three modules of
21
22 35 clarithromycin and two of amoxicillin in which the drug release mechanism did not
23
24 36 interfere with the floatation mechanism.

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26
27 37 The assembled system showed immediate *in vitro* floatation at pH 1.2, lasting 5
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29 38 hours. The *in vitro* antibiotics release profiles from individual modules and
30
31 39 assembled systems exhibited linear release rate during buoyancy for at least eight
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33 40 hours. The predicted antibiotic concentrations in the stomach maintained for a long
34
35 41 time levels significantly higher than the respective Minimum Inhibitory
36
37 42 Concentrations. In addition, an *in vivo* absorption study performed on beagle dogs
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39 43 confirmed the slow release of clarithromycin and amoxicillin from the assembled
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41 44 system during the assembly's permanence in the stomach for at least 4 hours.
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45 Introduction

46 *Helicobacter pylori* (*H. pylori*) infection is an important factor in the development of
47 gastritis, gastric ulcer and gastric carcinoma¹. More than 50% of people worldwide²
48 are infected with this pathogen. Factors such as age and socio-economic conditions
49 affect its prevalence³. *H. pylori*~~*Helicobacter pylori*~~ mainly resides in the gastric
50 mucosa or at the interface between the mucous layer and the epithelial cells in the
51 antral region of the stomach. The first-line pharmacological treatment for *H. pylori*
52 ~~*Helicobacter pylori*~~ eradication is one week of oral therapy with high doses of two
53 antibiotics (clarithromycin and amoxicillin or metronidazole) combined with a proton-
54 pump inhibitor⁴⁻⁵, often exposing the patient to adverse effects. The complexity of an
55 administration schedule, combining different dosage forms taken together, may lead
56 to non-adherence to the therapeutic regimen, giving rise to the appearance of
57 resistant strains. Therefore, the systemic treatment failure is dependent on the
58 difficulty to reach and maintain an effective antimicrobial concentration on the gastric
59 epithelial surface⁶.

60 Because of these problems, medicines should be provided that are based on drug
61 release control enabling to sustain effective local drug concentration, thus simplifying
62 the regimen. Given that the bacterium lives in the gastric mucosa, a logical way to
63 improve the therapy would be to administer a gastro-retentive dosage form capable
64 of releasing the antibiotics for as long as possible in the bacterium niche. This
65 approach could provide effective and prolonged local levels of antibiotic even by
66 administering lower doses of drug.

67 Several described gastro-retentive dosage forms are floating preparations⁷⁻¹¹ but
68 most of them did not show a prompt floatation. Moreover, the release of the drug is
69 often disturbed by the floatation mechanism adopted. For example, frequently

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3 70 described floating drug delivery systems are swellable matrices, in which the
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5 71 floatation is determined by the bubbles, produced by a gas-generating agent,
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7 72 entrapped in the gel layer¹²⁻¹³. Since the bubbles develop in the gel layer, their
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9 73 presence interferes with the release of drug due to the fact that the gel layer controls
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11 74 the drug release. A drug dosage form based on module assembly technology could
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13 75 provide a modular delivery system for prolonged drug release in gastro-retentive
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15 76 conditions in which the adopted floatation mechanism does not interfere with the
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17 77 release control mechanism¹⁴⁻¹⁶. The release module having a disc shape with a
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19 78 convex and a concave base, has been formulated as a tableted hydrophilic matrix for
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21 79 floatation and drug release control. Two of these release modules assembled by
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23 80 sticking the concave base of one to the concave base of the other form an assembly
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25 81 with an internal void chamber that showed immediate floatation and gastro-retention
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27 82 in humans¹⁷. To the floating assembly, additional modules can be stacked to obtain
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29 83 multi-kinetics and multi-drug delivery systems¹⁸. In summary, module assemblage
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31 84 could allow for the delivery of more than one drug in a single dosage form at a
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33 85 specific time, proper rate and duration of release. Moreover, according to the number
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35 86 of modules assembled, the dose administered could be adjusted.

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37 87 Therefore, the aim of this work was thus to study the drug release mechanism and
38
39 88 kinetics of a floating system designed to sustain the intra-gastric concentrations of
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41 89 amoxicillin and clarithromycin in combination. By applying the module assembly
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43 90 technology¹⁴ for the construction of a floating prolonged release dosage form, it was
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45 91 postulated that the drug delivery system could float on gastric fluid without disturbing
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47 92 the drug release control. The objective would be to keep the concentration of the two
48
49 93 antibiotics above the respective minimum inhibitory concentration (MIC) in the
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51 94 stomach for 4-5 hours. Different modules of clarithromycin and amoxicillin prepared
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3 95 as controlled release formulations were assembled in one single drug delivery
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5 96 system (Figure 1). The *in vitro* antibiotic release rate and mechanisms from
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7 97 individual modules and assembled systems, together with the buoyancy
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10 98 performance of the assembly, were investigated. The drug concentration attainable
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12 99 in the stomach was predicted applying a pharmacokinetic model. Furthermore, to
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14 100 assess clarithromycin and amoxicillin release rate and site from the assembled
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16 101 system, a pilot *in vivo* study using dogs was conducted.
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20 103 **Materials and methods**

21 104 **Materials**

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25 105 Clarithromycin was supplied by Special Product's Lines (Rome, Italy) and amoxicillin
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27 106 purchased from Sandoz GmbH (Kundl, Austria). Hydroxypropylmethylcellulose
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29 107 polymers (HPMC K15M; HPMC K100M) were obtained from Colorcon Limited
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31 108 (Orpington, U.K.); polyvinylpyrrolidone (PVP K30) was purchased from BASF SE
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33 109 (Ludwigshafen, Germany) and polyethylene glycol (PEG 6000) was obtained from
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35 110 Lisapharma S.p.A. (Erba (CO), Italy). All other chemicals were standard
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37 111 pharmaceutical grade.
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42 113 **Methods**

43 114 *Preparation of prolonged release clarithromycin modules*

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47 115 Clarithromycin modules were prepared by compression of clarithromycin granules
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49 116 prepared by kneading 100 g of clarithromycin and 10 g of HPMC K15M using 40 ml
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51 117 of a 50:50 hydroalcoholic solution (v/v) containing 5 % w/v of PEG 6000 and 5 % w/v
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53 118 of PVP K30. Granules were obtained by using an oscillating arm granulator (Erweka
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55 119 AR400, Düsseldorf, Germany), equipped with a 0.8 mm mesh. Granules were dried
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3 120 in an oven for about 5 hours at 45 °C. Then, granules were blended with a 3 % w/w
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5 121 of talc and 1 % w/w of magnesium stearate in a Turbula® blender for 25 minutes.
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7 122 The blend was compressed at a tablet weight of 120 mg using a single-punch
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9 123 eccentric tableting machine (EKO Korsch, Berlin, Germany) equipped with 7.4 mm
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11 124 diameter cylindrical punches having a tip surface suitable for manufacturing convex
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13 125 or concave bases. Two different module shapes were compressed (see Figure 1).
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15 126 The male modules had an average thickness of 4.8 ± 0.2 mm and the female ones of
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17 127 4.4 ± 0.2 mm.
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23 129 *Preparation of prolonged release amoxicillin modules*

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25 130 Amoxicillin female modules were prepared by direct compression of a mixture of 100
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27 131 g of amoxicillin, 46.2 g of HPMC K100M, 6.2 g of talc and 1.6 g of magnesium
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29 132 stearate powders, using the same single-punch eccentric tableting machine
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31 133 equipped with 7.4 mm cylindrical punches. The female modules (see Figure 1d)
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33 134 were tableted at 154 mg. The average thickness was 5.8 ± 0.2 mm.
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38 136 *Assembly of Dome Matrix modules*

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41 137 The assembled systems were obtained by "clicking" together the modules by hand in
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43 138 the following way: one male (see Figure 1a) and one female module of
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45 139 clarithromycin (see Figure 1b) were interlocked concave to concave face. This
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47 140 assembly, named void configuration, ensured the floatation of the release system to
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49 141 be exploited *in vivo* for localizing and maintaining the drug release in the stomach.
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51 142 Additional female modules of clarithromycin or amoxicillin were stacked on the
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53 143 convex base of the female module of the void configuration assembly (see Figure 1).
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3 145 *Assay for clarithromycin content*

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5 146 The content of clarithromycin was determined following the method reported in the
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7 147 USP 34 ed. monograph "Clarithromycin Extended-Release Tablets". The HPLC
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9 148 apparatus and conditions were the following: Shimadzu Liquid Chromatograph LC-
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11 149 10AT (Shimadzu Europe GmbH, Duisburg, Germany); UV-VIS detector SPD-10A at
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13 150 210 nm; Luna C18 column 4.6 x 150 mm, 5 μ m (Phenomenex, Torrance, CA, USA)
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15 151 maintained at 50 °C; mobile phase: methanol:phosphate buffer pH 4.0 (65:35 (v/v));
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17 152 flow rate: 1 ml/min; injection volume: 50 μ l (Autosampler Model 542, ESA Inc.,
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19 153 Chelmsford, USA). The system suitability gave the following results: theoretical
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21 154 plates 2184; peak symmetry 1.01; RSD 1.7%.
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27 156 *Assay for amoxicillin content*

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29 157 The content of amoxicillin was determined following the method reported in USP 34
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31 158 ed. monograph "Amoxicillin Tablets". The HPLC apparatus and conditions were the
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33 159 following: Shimadzu Liquid Chromatograph LC-10AT (Shimadzu Europe GmbH,
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35 160 Duisburg, Germany); UV-VIS detector SPD-10A at 230 nm; μ Bondapak C18
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37 161 column 3.9 x 300 mm, 5 μ m (Waters Corp., Milford, MA, USA) maintained at 40 °C;
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39 162 mobile phase: acetonitrile:phosphate buffer pH 5.0 (2.5:97.5 (v/v)); flow rate: 0.7
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41 163 ml/min; injection volume: 10 μ l (Autosampler Model 542, ESA Inc., Chelmsford,
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43 164 USA). The system suitability gave the following results: theoretical plates 2253; peak
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45 165 symmetry 2.25; RSD 0.5%.
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52 167 *In vitro drug release of individual modules and assembled system*

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54 168 *Clarithromycin*

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56 169 The USP dissolution apparatus II with paddle rotating at 75 rpm was employed. The
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3 170 dissolution medium was Britton-Robinson buffer solution pH 3.0¹⁹ at 37 °C. 500 ml
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5 171 and 900 ml were used for dissolving the individual modules and the assembled
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7 172 systems respectively. Samples collected at fixed time points were filtered through a
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10 173 0.45 µm membrane (CA 0.45 µm, LabService Analytica, Bologna, Italy) and
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12 174 analyzed using the HPLC assay method.

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15 16 176 *Amoxicillin*

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18 177 The USP dissolution apparatus IV, equipped with a 12 mm diameter cell, was
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20 178 employed. Simulated gastric fluid without enzymes pH 1.2 at 37 °C was used as
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22 179 dissolution medium. The flow rate was set at 8 ml/min. Samples of 8 ml collected at
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24 180 fixed times were filtered through a 0.45 µm membrane and analyzed using the HPLC
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26 181 assay method.

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31 32 183 *Determination of floatation characteristics*

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34 184 As previously described²⁰⁻²¹, the buoyancy of the assembly was determined by
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36 185 measuring the resultant weight of the system made of three modules of
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38 186 clarithromycin and two modules of amoxicillin, submerged in simulated gastric fluid
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40 187 without enzymes at 37 ± 0.5 °C.

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44 45 189 *Simulation of clarithromycin and amoxicillin concentrations in the stomach*

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47 190 The intra-gastric concentrations of clarithromycin and amoxicillin were simulated and
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49 191 predicted using STELLA[®] software (isee systems, Lebanon, NH, USA). The slope of
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51 192 the amount released vs time linear profiles for the assembled system was elected as
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53 193 the intra-gastric release rate. The initial volume of fluid in the stomach in fed state
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55 194 was set at 1.5 L²²; the emptying rate followed a first order kinetic with t_{50%} of 0.20 h²³.

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3 195 The gastric residence of the dosage form was fixed at 5 h¹⁷.
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7 197 *Oral absorption study in dogs*
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9 198 An *in vivo* study to measure drug absorption upon oral administration was carried out
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11 199 using six male beagle dogs, weighing 15-20 kg. The study was performed at the
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13 200 Faculty of Health Sciences, Fernando Pessoa University (Portugal), approved by the
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15 201 competent ethics and scientific committee and conducted in accordance with EU
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17 202 Legislation²⁴. The dogs were fed before drug administration with a standard meal
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19 203 and water was left *ad libitum*. A hard gelatin capsule (Coni-Snap[®] size 00, Capsugel,
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21 204 Colmar, France) containing the assembly of Figure 1 (three modules of
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23 205 clarithromycin and two of amoxicillin) was administered. In order to check by X-ray
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25 206 the location in the stomach of the assembled system during *in vivo* drug release, 10
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27 207 mg of barium sulfate powder were introduced in the empty chamber of the
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29 208 clarithromycin void configuration assembly.
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31 209 Blood samples were collected immediately before and after 60, 120, 240, 480, 720
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33 210 min from drug administration. 200 µl of plasma was added to 200 µl of chilled
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35 211 acetonitrile containing the internal standards (oxacillin and erythromycin for
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37 212 amoxicillin and clarithromycin, respectively; 50 µl of each at concentration of 10
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39 213 µg/ml). The sample was mixed for at least 5 min to allow complete protein
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41 214 precipitation. After centrifugation at 16,000 X g for 10 min, 200 µl of the supernatant
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43 215 was transferred to a clean vial. The solvent was evaporated to dryness under
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45 216 nitrogen gas flow. The residue was reconstituted in 100 ml of 0.1% (v/v) aqueous
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47 217 formic acid and left to mix for 5 min. Samples were centrifuged again at 16,000 X g
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49 218 for 10 min. The supernatant was transferred to a polypropylene auto-sampler vial
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51 219 and stored at 5 °C until analysis by UPLC-MS/MS. The analytical method consisted
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3 220 of a Waters Acquity UPLC instrument coupled to a Quattro Premier XE tandem-
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5 221 quadrupole mass spectrometer (Waters Corp., Milford, MA). The analytical column
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7 222 was an Acquity UPLC BEH C18 (2.1 mm x 50 mm, 1.7 μm ; Waters Ltd., Dublin,
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9 223 Ireland), to which a 0.2 μm pre-column filter unit was added. The mobile phase was
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11 224 a gradient of solution A (0.1% FA in water) and solution B (0.1% FA in methanol),
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13 225 with an initial composition of 10% solution B. The mobile phase composition
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15 226 changed linearly from 10% B at 1 min to 80% B at 2 min and onward to 100% B at 4
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17 227 min. The composition was switched back to 10% B at 5 min and maintained until 6
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19 228 min. The flow rate was 0.3 ml/min, with a column temperature of 40 $^{\circ}\text{C}$. From each
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21 229 sample, 10 μl was injected in duplicate onto the column. The analysed compounds
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23 230 were detected via MS with an electrospray ionization interface in positive multiple
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25 231 reaction monitoring mode. Optimized multiple reaction monitoring settings for the
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27 232 individual drugs, including cone voltage and collision energy are shown in Table 1.
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29 233 The acquisition settings were as follows: capillary voltage, 3.4 kV; source
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31 234 temperature, 120 $^{\circ}\text{C}$; desolvation temperature, 300 $^{\circ}\text{C}$; desolvation gas flow, 600
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33 235 L/h; cone gas flow, 50 L/h; and dwell time, 80 ms.
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35 236 Data were acquired using Masslynx V4.1 software. For all compounds, calibration
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37 237 curves were obtained by plotting the peak area ratios of drug versus internal
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39 238 standard against the theoretical concentration. The LOQ values were 0.015 $\mu\text{g/ml}$ for
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41 239 amoxicillin and 0.024 $\mu\text{g/ml}$ for clarithromycin and the LOD values were 0.004 $\mu\text{g/ml}$
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43 240 and 0.006 $\mu\text{g/ml}$, respectively. The precision in terms of relative standard deviation
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45 241 was 4.87 % for amoxicillin and 3.11 % for clarithromycin.
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243 *Statistics*

244 Graphing was performed by means of KaleidaGraph software (version 4.5.2,

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3 245 Synergy Software, Reading, PA, USA). Confidence limits of the power equation
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5 246 exponent were determined from the slope of the log/log plot of data at 95%
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7 247 probability level.
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11 249 **Results and discussion**

12 250 *Formulation and manufacturing of modules and assembled systems*

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16 251 Clarithromycin and amoxicillin modules were formulated taking into account the
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18 252 biopharmaceutical characteristics of the active substances and the therapeutic
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20 253 program for ~~*H. pylori*~~*Helicobacter pylori* eradication. The high doses of these
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22 254 antibiotics, required in therapy, restricted the formulation design space, but at the
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24 255 same time simplified the module composition. Clarithromycin and amoxicillin oral
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26 256 recommended doses for *Helicobacter* eradication are 500 mg and 1000 mg
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28 257 respectively, twice a day for 7 days of treatment³⁻⁶. Given such high drug amounts
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30 258 and the need to construct a dosage form in a size convenient for swallowing, the
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32 259 modules used in this study had a diameter of 7.4 mm and a thickness between 4.4
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34 260 and 5.8 mm. With this size, each module could accommodate between 100 and 200
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36 261 mg of powder formulation and, for administration purposes, up to six assembled
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38 262 modules could be introduced into a Coni-Snap[®] size 00 capsule. The modules were
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40 263 prepared as hydrophilic matrices for prolonged release. Amoxicillin modules
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42 264 contained 100 mg of drug and HPMC K100M. The clarithromycin modules included
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44 265 100 mg of clarithromycin, HPMC K15M, PVP K30 and PEG 6000. The choice of
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46 266 excipients was mainly guided by the solubility of drugs at the relevant pH values. For
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48 267 instance, the modules of clarithromycin, which was the less soluble drug substance
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50 268 (6.9 g/L at pH 3.0²⁵), contained more soluble polymers compared to HPMC used in
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52 269 amoxicillin modules. Both module formulation exhibited compressibility
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3 270 characteristics favorable to module tableting. Their friability was lower than 1% and
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5 271 the diametric crushing strength was between 3.9 and 10.7 kg (Monsanto hardness
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7 272 tester).
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10 273 The modules of amoxicillin and clarithromycin were then assembled in different
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12 274 numbers and configurations. A total of five assembled modules formed the
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14 275 amoxicillin-clarithromycin site-specific delivery system used for floatation and release
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16 276 studies, as illustrated in Figure 1.
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21 278 *In vitro drug release rate of single modules and assembled systems*

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23 279 The individual modules and various assembled configurations, all substantially
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25 280 hydrophilic matrices, were studied with respect to drug release rate and kinetics. The
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27 281 release rate of the individual modules was tested in a flow-through apparatus at pH
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29 282 1.2 for amoxicillin²⁶ and in a paddle apparatus at pH 3.0 for clarithromycin¹⁹. The
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31 283 choice of different conditions was made in order to overcome the solubility and
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33 284 instability problems of clarithromycin and amoxicillin at acidic pH values^{19, 26-27}. The
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35 285 obtained release profiles (Figure 2) were analyzed using the power equation in which
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37 286 the value of time exponent gives indication on the mechanism of drug release²⁸. The
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39 287 analysis was extended up to 80% values of fraction released in consideration of the
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41 288 quasi-linear profile in the measured range. In the case of the clarithromycin modules
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43 289 (male and female), a quasi-constant release rate up to 80% of drug released was
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45 290 observed, with a non-significant difference between male and female in terms of
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47 291 rate. However, the exponent n value was significantly higher for the female module,
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49 292 indicating an effect of the module geometry on the release kinetics (see n values and
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51 293 95% confidence limits in Figure 2 caption). The quasi-linearity of the curve supported
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53 294 an erosive release mechanism of these matrices. In fact, in 4-5 hours clarithromycin
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3 295 modules were completely dissolved. Thus, clarithromycin modules in the medium at
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5 296 pH 3.0 exhibited a behavior more erosive than diffusive. While assuring the
6
7 297 maximum stability for clarithromycin, a value of 3.0 for the pH is in line with the pH
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9 298 values found in humans in fed conditions²⁹.

10
11 299 The release rate of amoxicillin from the female module was slower than that of
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13 300 clarithromycin female module, likely owing to the presence (30% w/w) of the high
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15 301 molecular weight hydrophilic HPMC polymer in the formulation. Moreover, *in vitro*
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17 302 release was measured at pH 1.2 using the flow through apparatus instead of the
18
19 303 paddle employed for clarithromycin. This apparatus was used to limit the amoxicillin
20
21 304 *in vitro* degradation at pH 1.2 medium²⁶. With this matrix the kinetics appeared linear
22
23 305 up to 80% of amoxicillin released, as reflected by the value of exponent n ($0.93 \pm$
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25 306 0.11). Amoxicillin was totally released within 6 hours from a gelled system that
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27 307 completely dissolved in 24 hours. Tested at pH 3.0, where amoxicillin is less
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29 308 soluble²⁷, the module showed a lower drug release (40% in 8 hours).

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34 309 The release rate and duration for both drugs from individual modules was judged
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36 310 adequate relatively to the assumed residence time in the stomach of the modules
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38 311 when assembled. In fact, in a previous *in vivo* study¹⁷ a period of gastro-retention for
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40 312 an assembled system in void configuration was determined between 4 and 5 hours.

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43 313 Two clarithromycin modules having the same formulation and mass, but male and
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45 314 female geometry, were then assembled in void configuration by interlocking their
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47 315 concave bases. To increase the total unit dose, additional clarithromycin female
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49 316 modules were stacked onto the convex female face of this void assemblage, giving
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51 317 rise to void/stacked assembled systems made of two, three or five clarithromycin
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53 318 modules. Thus, these assemblies were investigated for *in vitro* drug release at pH
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55 319 3.0 in paddle apparatus during floatation. Clarithromycin release profiles are

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3 320 reproduced in Figure 3 as amount released versus time.
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5 321 The profiles of amount released versus time show that the clarithromycin release
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7 322 rate increased with the number of modules from two to five, owing to the increased
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9 323 area exposed to dissolution medium. The release area exposed was 183.7 mm²,
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11 324 247.9 mm² and 376.4 mm² for the two modules, three modules and five modules,
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13 325 respectively.
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16 326 In the case of the two-module void assembly, the clarithromycin release rate was
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18 327 slowed down compared to the cumulative drug released from the male and female
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20 328 modules individually. This was attributed to the fact that, upon assembly by "clicking"
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22 329 together the male and female module, the internal surface was no more accessible
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24 330 to the solvent. After a small initial burst, clarithromycin was released at a quasi-
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26 331 constant rate as the system visibly underwent dissolution/erosion.
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29 332 The release profiles up to 80% of drug released were also fitted to the power
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31 333 equation and the release kinetics were identified from the values of the exponent n .
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34 334 The n values obtained with these assembled hydrophilic matrices (see caption to
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36 335 Figure 4) were significantly different from 0.5. This supported the dissolution/erosion
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38 336 mechanism of the drug/polymer matrix, as observed with several other hydrophilic
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40 337 matrices containing small amounts of polymer or soluble drugs and polymers³⁰.
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43 338 The addition of clarithromycin modules on the basic void assembly, pushed the
44
45 339 release kinetics towards higher linearity (see the exponent n values in Figure 4
46
47 340 caption), but the significance of the increase remained at $p \geq 0.2$. Since there was no
48
49 341 change in module composition, this kinetics shift has to be assigned to the
50
51 342 modification of the aspect ratio of the delivery system. In fact, the release from the
52
53 343 lateral surface of the assembled pile augmented with additional modules, whereas
54
55 344 the base contribution in the pile was not changed. It was previously observed with a
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3 345 soluble drug that increasing the length of the assembly by module addition, shifted
4
5 346 the release kinetics towards higher linearity³¹. The release from the lateral side of the
6
7 347 compressed disc was found to be kinetically different from that of the axial side³² and
8
9 348 the lateral side contribution to clarithromycin release in the assembly multiplied with
10
11 349 the number of stacked modules.

12
13
14 350 In the delivery system prepared for the *in vivo* study, two female amoxicillin modules
15
16 351 were stacked on a three-clarithromycin module assembled system, as indicated in
17
18 352 Figure 1. The *in vitro* amoxicillin and clarithromycin release profiles of the combined
19
20 353 assembled systems were determined by dissolution at pH 1.2 in flow-through
21
22 354 apparatus for amoxicillin and at pH 3.0 in paddle for clarithromycin. The release
23
24 355 profiles are illustrated in Figure 5.

25
26
27 356 Amoxicillin and clarithromycin release profiles from the combined system remained
28
29 357 quasi-linear. Individually, two assembled modules of amoxicillin (Figure 5) or three
30
31 358 assembled modules of clarithromycin (see Figure 3) showed a release rate higher
32
33 359 than that of amoxicillin and clarithromycin from the assembly. The assembly of three
34
35 360 modules of clarithromycin and two of amoxicillin reduced the exposed release area
36
37 361 compared to the independent drug assemblies. However, the release of the two
38
39 362 combined drugs in the assembled system at steady state remained very close to
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41 363 linearity.

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45 46 47 365 *Simulation of intra-gastric drug concentration*

48
49 366 In order to estimate the maintenance of clarithromycin concentration in the stomach
50
51 367 above the ~~MIC~~Minimum Inhibitory Concentration of *Helicobacter* (2 µg/mL)³³⁻³⁴, the
52
53 368 release rates of clarithromycin assembled systems made of three (300 mg) and five
54
55 369 modules (500 mg) (see Figure 3) were used for predicting intra-gastric
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3 370 concentrations by the STELLA software³⁵.
4
5 371 The simulated intra-gastric concentration of clarithromycin (Figure 6) from the
6
7 372 assembled systems containing three or five modules remained higher than the
8
9 373 clarithromycin MIC. The simulation supports the possibility of a steady drug
10
11 374 concentration maintained for the expected residence time of the system within the
12
13 375 stomach. The five-module system of clarithromycin could maintain the drug
14
15 376 concentration largely above the MIC for the time of residence considered. In the
16
17 377 case of the system with three clarithromycin modules, containing a dose of only 300
18
19 378 mg, the steady gastric drug concentration was threefold higher than the MIC.
20
21 379 Literature pharmacokinetics data of clarithromycin show that 2 hours after
22
23 380 administration of 500 mg of clarithromycin, the stomach tissue concentration was
24
25 381 10.5 µg/g in the antrum, 20.8 µg/g in the fundus and 4.2 µg/g in the mucus³⁶.
26
27
28
29 382 In humans, the gastro-retention could allow for a longer effective concentration of
30
31 383 clarithromycin directly in contact with *H. pylori*~~*Helicobacter pylori*~~ at a lower dose
32
33 384 than the usual systemic administration. The number of clarithromycin modules could
34
35 385 then be reduced from five to three, and the two eliminated clarithromycin modules
36
37 386 could be replaced by two amoxicillin modules, each containing 100 mg of drug.
38
39 387 From the combination of two amoxicillin modules with three clarithromycin modules,
40
41 388 the simulated concentration profile of clarithromycin was slightly reduced compared
42
43 389 to the corresponding three-module system in Figure 6. The concentration profile of
44
45 390 amoxicillin maintained for 5 hours steady values sixfold higher than the MIC that is
46
47 391 0.5 µg/mL³⁷. Although the amount of amoxicillin in the assembled system (200 mg)
48
49 392 was significantly lower than the usually prescribed dose (1000 mg), the simulation
50
51 393 suggested that this dose could maintain amoxicillin concentration in the stomach
52
53 394 higher than the MIC value for a prolonged time.
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3 395 *Floatation assessment of assembled system*

4
5 396 The male and female modules interlocked concave to concave base (void
6
7 397 assembly), in simulated gastric fluid displayed an immediate floating lasting more
8
9
10 398 than 5 hours. Increasing the number of modules added on the void assembly, a non-
11
12 399 immediate floatation was expected. However, the system with five assembled
13
14 400 modules, i.e., three clarithromycin and two amoxicillin, immediately floated on the
15
16 401 fluid as well, with an initial resultant weight value of 80 mg. The buoyancy force of
17
18 402 the assembled system measured over time increased to 100 mg, maintaining this
19
20 403 value for more than 5 hours, i.e., the time designed for completing drug release. This
21
22 404 indicates that the floating property of the delivery system is strengthened by the
23
24 405 presence of the swollen polymer. ~~As shown in Figure 7, showing the system during~~
25
26 ~~the floatation measurement,~~ the left side of the swollen system ~~during the floatation~~
27
28 ~~measurement~~ corresponds to amoxicillin modules and the right to clarithromycin
29
30 407 modules. ~~The picture of Figure 7 shows that~~ The system has two portions with
31
32 408 different physical behavior during release: amoxicillin is released from a more highly
33
34 409 swollen gel, whereas clarithromycin from a portion of the system that jellified less
35
36 410 and eroded more. Moreover, it can be observed that the external gel layer of the
37
38 411 matrix was not disturbed by the presence of air bubbles as expected from the
39
40 412 mechanism adopted for floatation.
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47 415 *Oral absorption in dogs (bioavailability pilot study)*

48
49 416 The drug release rate and the system gastro-retention were investigated by an oral
50
51 417 absorption test in dogs. Dogs were used primarily for assessing the drug release
52
53 418 kinetics, concomitantly to the system localization in the gastrointestinal (GI) tract.
54
55 419 The administered product consisted of three clarithromycin modules (two in void
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3 420 configuration) and two amoxicillin modules stacked on them. The clarithromycin
4
5 421 maximum plasma concentration (C_{max} , $10.56 \pm 3.66 \mu\text{g/mL}$) was achieved after 8
6
7 422 hours, with a profile indicating a prolonged absorption of drug (Figure 8). The
8
9 423 maximum amoxicillin plasma concentration (C_{max} , $1.53 \pm 0.44 \mu\text{g/mL}$) was achieved
10
11 424 after two hours. Then, the plasma levels were steadily maintained between 1.25 and
12
13 425 $1.15 \mu\text{g/mL}$ until twelve hours. ~~The plasma levels were lower than other values in~~
14
15 ~~dog reported in the literature³⁸⁻³⁹ for immediate release tablets of the same drugs.~~
16
17 426
18 427 The shape of the curves suggested that both drugs were slowly released over a long
19
20 428 period of time. ~~The plasma levels were lower than other values in dog reported in the~~
21
22 ~~literature³⁸⁻³⁹ for immediate release tablets of the same drugs.~~
23
24 429
25 430 During the blood sampling, the dog was examined by X-rays in order to
26
27 431 concomitantly localize the position of the system in the animal GI. The X-ray pictures
28
29 432 revealed that the assembled system was still in the dog stomach at 4 hours (Figure
30
31 433 9). The in vitro release ~~profiles behavior reflected in correlated with~~ the measured
32
33 434 plasma levels and the permanence of the delivery system inside the stomach.
34
35 435 ~~demonstrated that t~~ The goal of sustaining the antibiotic concentration in the
36
37 436 stomach using swellable release matrices assembled to form a floating delivery
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39 437 system ~~was realistic is affordable.~~
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439 Conclusion

440 A modular single unit dosage form containing clarithromycin and amoxicillin for *H.*
441 ~~pylori~~ *Helicobacter pylori* therapy was constructed by interlocking five drug modules.
442 The final dosage form included three assembled clarithromycin modules and two
443 amoxicillin modules stacked on them. The assembled system floated *in vitro* for
444 more than 5 hours on the surface of a dissolution medium at pH 1.2. The simulated

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3 445 drug intra-gastric concentrations over time were significantly higher than the MIC
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5 446 values of both antibiotics. An oral absorption experiment in dogs showed prolonged
6
7 447 plasma concentrations of clarithromycin and amoxicillin after administration of the
8
9 448 combined system. The X-ray examinations confirmed that the system was still in the
10
11 449 stomach 4 hours after administration.

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13
14 450 The ability of the assembled system to float in the stomach content does not interfere
15
16 451 with the drug release control mechanism since the floating mechanism is determined
17
18 452 by the internal void of the assembly and the drug release control by the walls of the
19
20 453 system. This is different and new compared to other technologies in which floatation
21
22 454 depends on the same mechanism that controls drug release (e.g. hydrophilic
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24 455 matrices entrapping air/gas bubbles in the swollen polymer). By maintaining a
25
26 456 prolonged direct contact between *H. pylori*~~Helicobacter pylori~~ and the antibiotic
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28 457 molecules, the assembled system has the potential to reduce the drug doses and the
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30 458 number of dosage forms administered daily.

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34 459 The assembly of modules having different drug content and/or different kinetics in a
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36 460 single dosage unit leads to the assumption of an increase in patient compliance and
37
38 461 adherence to the therapeutic regimen. Indeed, the actual size of the present unit
39
40 462 dose (5-module assembly inside a Coni-Snap 00 type capsule, comprising the two
41
42 463 drugs) is not significantly different from the average size (22 mm) of commercially
43
44 464 available capsules and tablets containing only amoxicillin (i.e., not combined with
45
46 465 clarithromycin) at dosages of 500 mg or higher⁴⁰.

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50 51 52 467 **Declaration of interest**

53
54 468 The authors report no declaration of interest.

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Table 1. Parameters of UPLC-MS/MS system: Q1, parent ion mass; Q3, daughter ion mass; CV, cone voltage; CE, collision energy; Rt, retention time. Oxacillin is the internal standard for amoxicillin, while erythromycin is the internal standard for clarithromycin.

Antibiotic	Q1 (m/z)	Q3 (m/z)	CV (V)	CE (eV)	Rt (min)
Amoxicillin	366.6	349.5	18	10	2.9
Oxacillin	402.6	243.4	20	14	2.6
Clarithromycin	748.9	558.8	30	30	2.7
Erythromycin	734.9	576.6	30	25	2.6

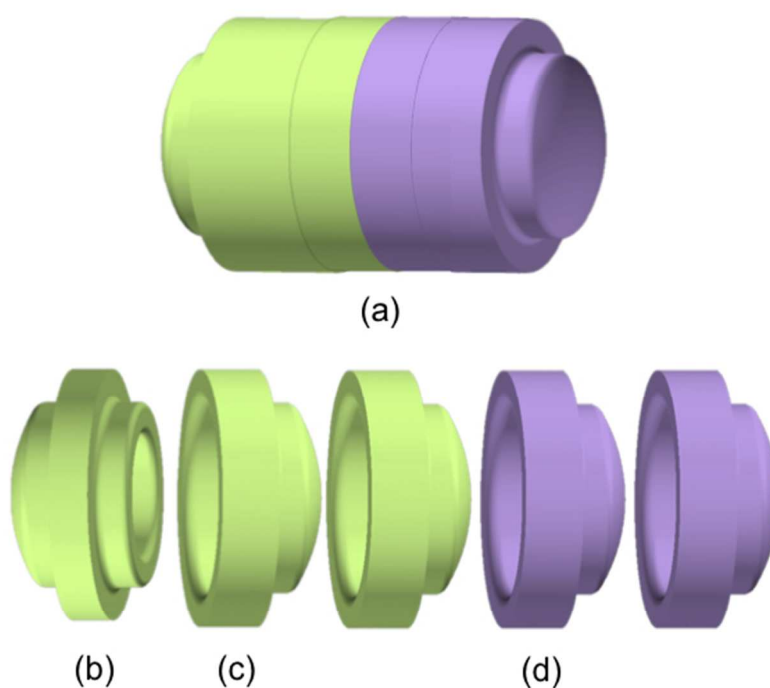


Figure 1. Modular assembled system for HP treatment: (a) clarithromycin controlled release modules (yellow: male (b) and female (c)) and amoxicillin controlled release female modules (purple, (d)). 60x45mm (300 x 300 DPI)

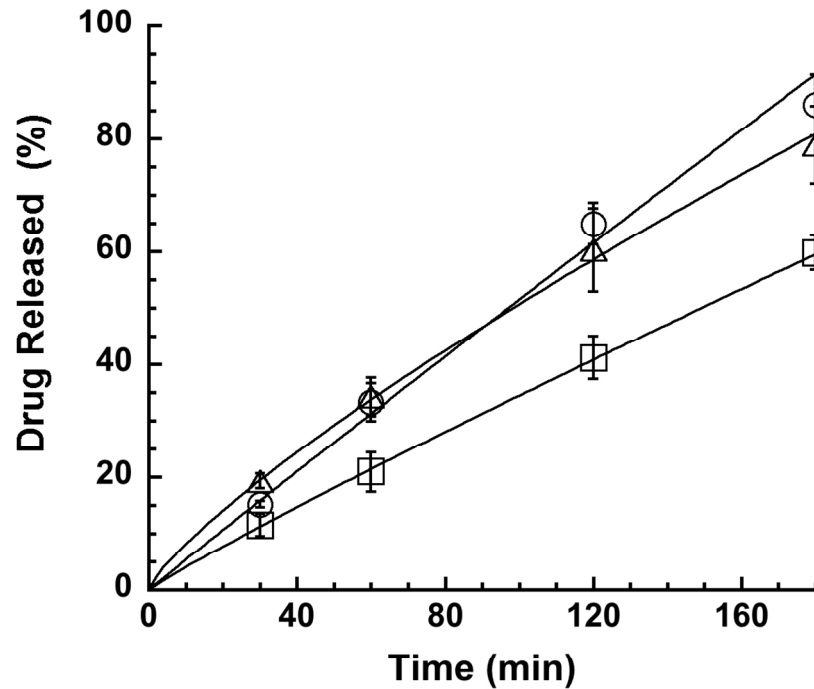


Figure 2. Release profiles of clarithromycin and amoxicillin individual modules (mean \pm S.D.; $n=3$). Lines represent the power equation fitting: (\circ) clarithromycin female module, drug released (%) = $0.573 \cdot \text{min}^{0.977 \pm 0.116}$; (Δ) clarithromycin male module, drug released (%) = $1.309 \cdot \text{min}^{0.794 \pm 0.063}$; (\square) amoxicillin female module, drug released (%) = $0.478 \cdot \text{min}^{0.929 \pm 0.105}$.

120x101mm (300 x 300 DPI)

Only

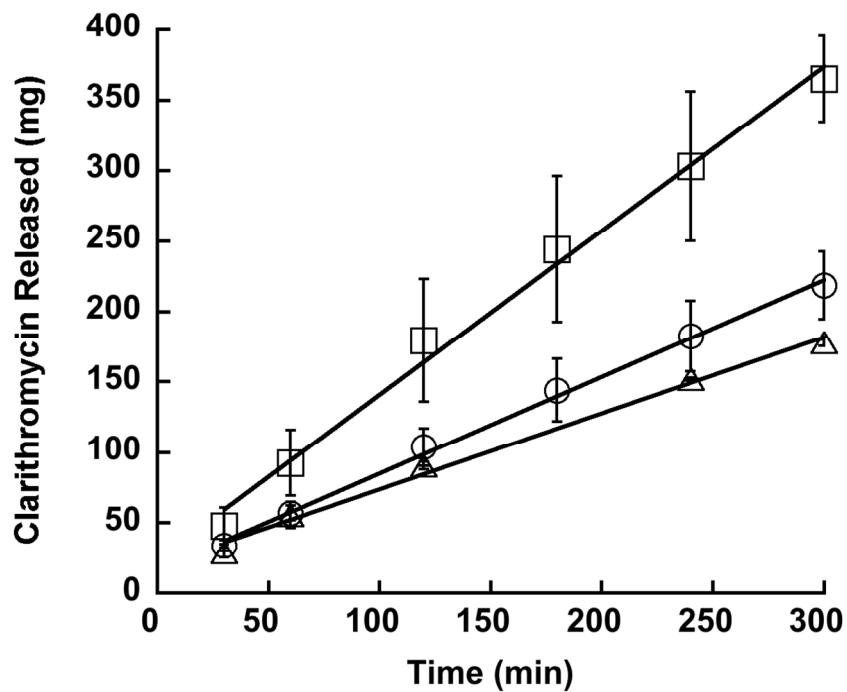


Figure 3. Amount of clarithromycin released from the assembled system (mean \pm S.D.; $n=3$). Lines represent the fitting to linear equation: (Δ) two-clarithromycin modules in void configuration, amount (mg) = $19.94 + 0.539 \text{ min}$; (\circ) three-clarithromycin modules, amount (mg) = $16.99 + 0.685 \text{ min}$; (\square) five-clarithromycin modules, amount (mg) = $24.83 + 1.164 \text{ min}$.

120x101mm (300 x 300 DPI)

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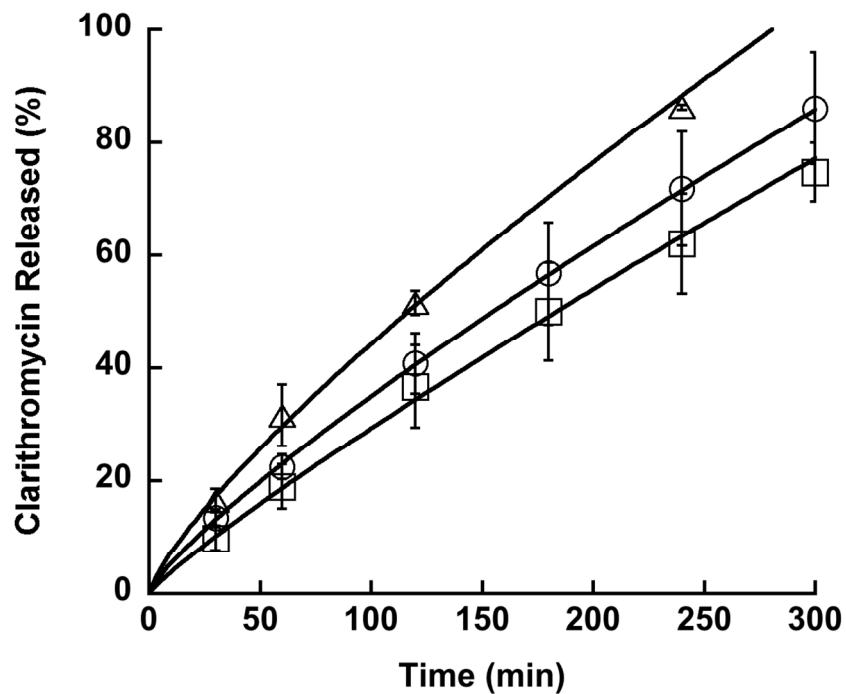


Figure 4. Percent release profiles of clarithromycin (mean \pm S.D.; $n=3$) from assembled system. Lines represent the fitting to the power equation: (Δ) two-clarithromycin modules in void configuration, drug released (%) = $1.174 \text{ min}^{0.788 \pm 0.085}$; (\circ) three-clarithromycin modules, drug released (%) = $0.812 \text{ min}^{0.817 \pm 0.073}$; (\square) five-clarithromycin modules, drug released (%) = $0.506 \text{ min}^{0.881 \pm 0.128}$.
120x101mm (300 x 300 DPI)

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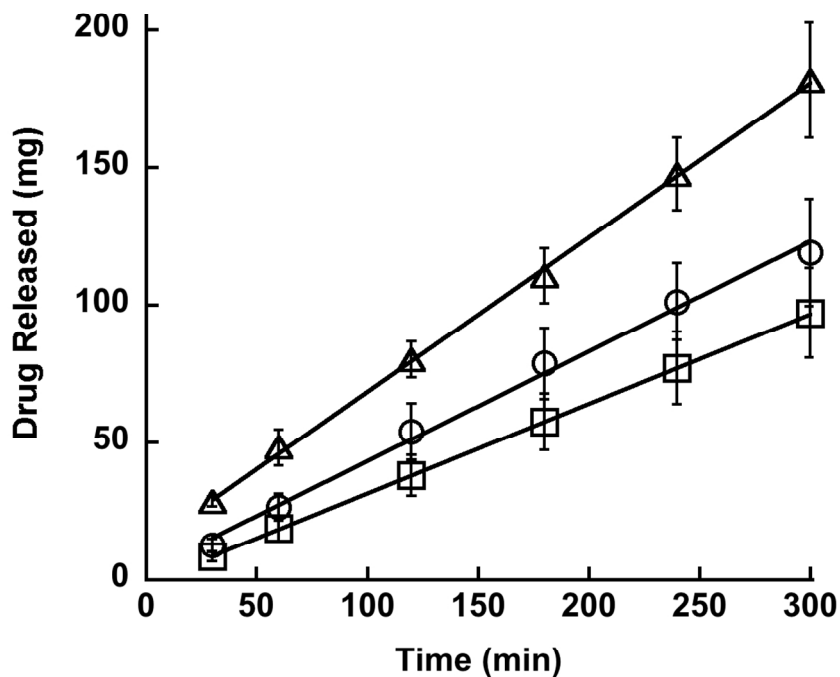


Figure 5. Amount released profiles (mean \pm S.D.; $n=3$). Lines represent the fitting to linear equation: (○) two-amoxicillin assembled modules, amount (mg) = $3.346 + 0.399 \text{ min}$; (□) amoxicillin from three-clarithromycin and two- amoxicillin module assembled system, amount (mg) = $-1.149 + 0.326 \text{ min}$; (Δ) clarithromycin from three-clarithromycin and two-amoxicillin module assembled system, amount (mg) = $12.447 + 0.561 \text{ min}$.
120x101mm (300 x 300 DPI)

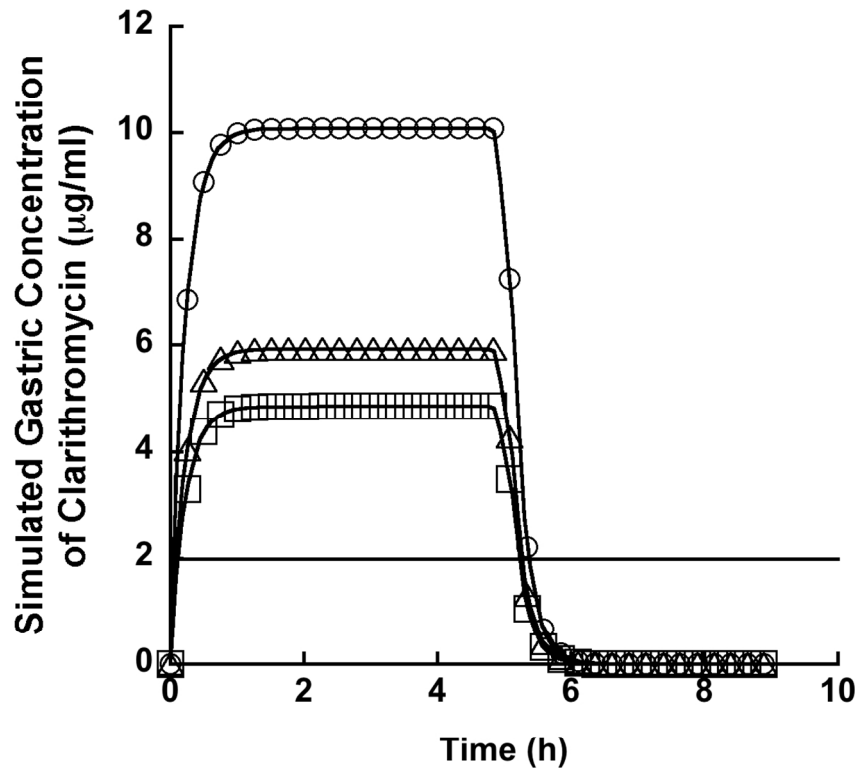


Figure 6. Simulated gastric concentration of clarithromycin from assembled system: (○) five-clarithromycin modules; (△) three-clarithromycin modules; (□) three-clarithromycin modules and two-amoxicillin modules. Straight line indicates clarithromycin MIC (2 µg/ml).

120x114mm (300 x 300 DPI)

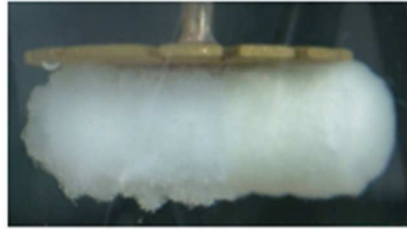


Figure 7. Five-module assembled system of clarithromycin and amoxicillin after 60 minutes of floatation.
18x11mm (300 x 300 DPI)

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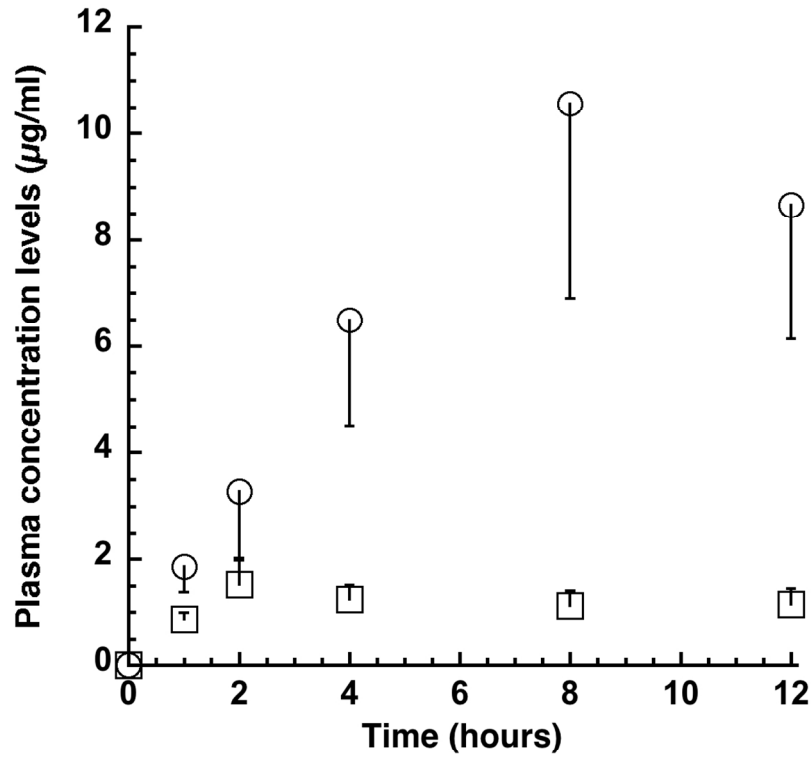


Figure 8. Plasma concentration levels after oral administration of the assembled system to dogs: (○) clarithromycin and (□) amoxicillin.
120x114mm (300 x 300 DPI)

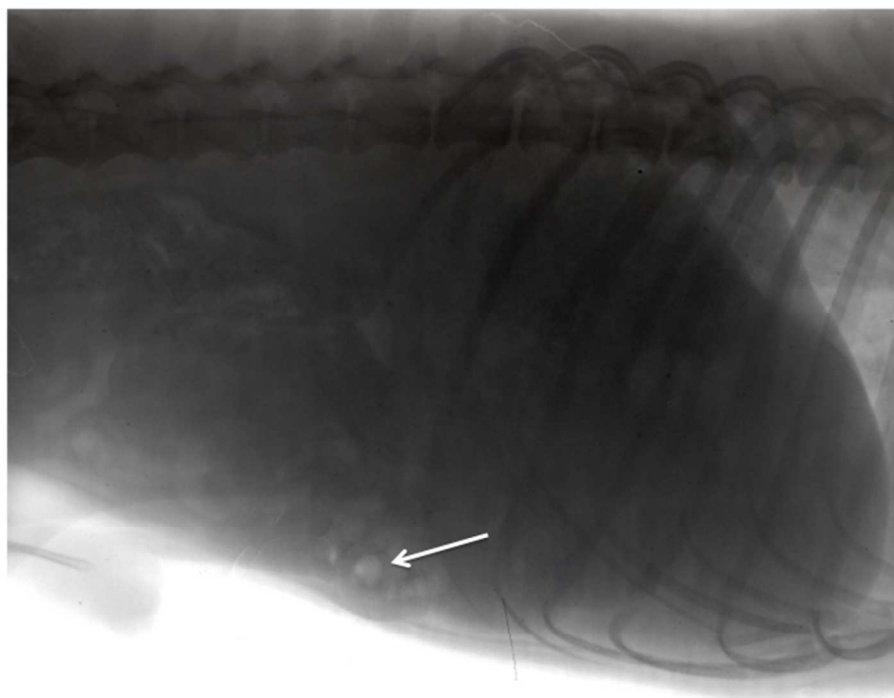


Figure 9. X-ray picture of beagle stomach 4 h from the administration.
60x45mm (300 x 300 DPI)

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