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1 **Analytical method for the simultaneous determination of a broad range of**
2 **opioids in influent wastewater: optimization, validation and applicability to**
3 **monitor consumption patterns**

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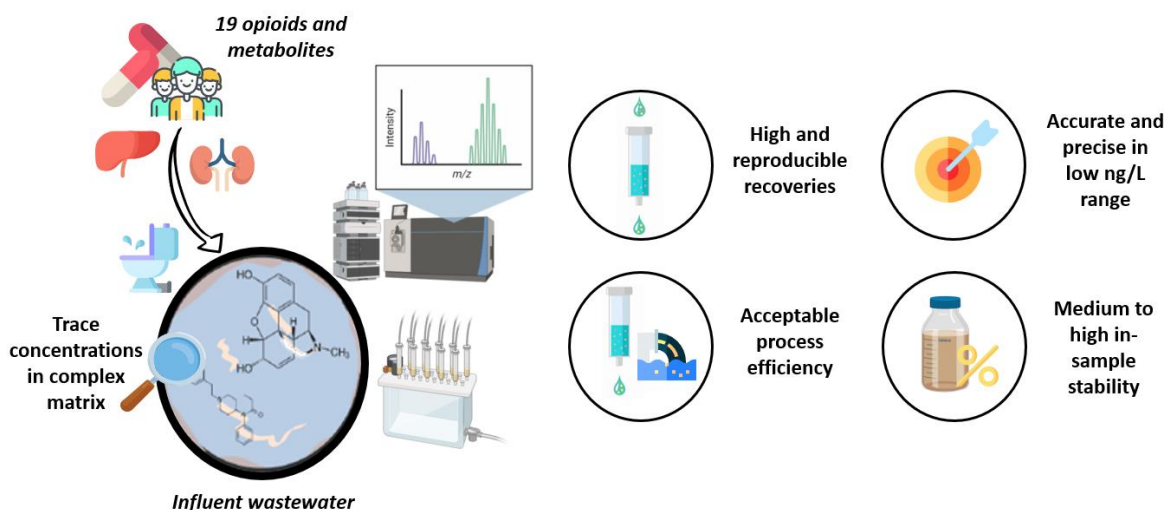
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10
11 **Abstract**

12 Wastewater-based epidemiology (WBE) employs the analysis of human metabolic biomarkers in influent
13 wastewater (IWW) to estimate community-wide exposure to xenobiotics (e.g. prescription opioids). The
14 low ng/L range of concentrations of these biomarkers and the complex matrix composition pose
15 bioanalytical challenges related to sample preparation and detection/quantification. Therefore, a sensitive
16 analytical method for the detection and analysis of 19 opioid biomarkers was optimized and validated
17 according to the European Medicines Agency guidelines. Oasis HLB cartridges were used for sample
18 concentration and the Atlantis T3 column with gradient elution resulted in sufficient separation of the
19 analytes. Absolute recoveries (RE) were highly reproducible and ranged between 50-93% with the exception
20 of 2-Ethylidene-1,5-dimethyl-3,3-diphenylpyrrolidine (EDDP). The lower limit of quantification (LLOQ)
21 ranged between 1 and 100 ng/L and was based on the analyte's concentrations found in IWW. Process
22 efficiency was acceptable for all biomarkers for which an isotope-labelled deuterated analogue was
23 available. All biomarkers showed high benchtop stability with the exception of buprenorphine, EDDP,
24 fentanyl and normorphine. Apart from buprenorphine and hydrocodone, all analytes under investigation
25 were detected at least once above LLOQ levels in five locations in Belgium, including Antwerp, Boom,
26 Brussels, Ostend and Koksijde. The highest population-normalized mass loads were found for tramadol, O-
27 desmethyltramadol and codeine. The proposed methodology was able to evaluate spatial differences in
28 opioid use.

29

30 **Graphical abstract**



31

32 **Credit author statement**

33

34 **Tim Boogaerts** – Conceptualization, data curation, formal analysis, investigation, methodology, validation, visualization, roles/writing – original draft, writing -review and editing

35

36 **Maarten Quireyns** – Data curation, formal analysis, investigation, methodology, validation, visualization, roles/writing – original draft, writing -review and editing

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38 **Adrian Covaci** – Conceptualization, writing – review and editing, funding acquisition, supervision

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40 **Hans De Loof** – Writing – review and editing, supervision

41 **Alexander L.N. van Nuijs** – Conceptualization, methodology, writing – original draft, writing – review and editing, funding acquisition, supervision

42

43 **Keywords:**

- 44 • *Wastewater-based epidemiology*
- 45 • *Opioids*
- 46 • *Mass-spectrometric analysis*
- 47 • *Validation*
- 48 • *Stability*
- 49 • *Process efficiency*

50

51 **Highlights**

- 52 • Validation of an analytical method for quantification of 19 opioids in wastewater
- 53 • Process efficiency acceptable for all analytes when deuterated analogue was available
- 54 • High in-sample stability for 15 opioid biomarkers
- 55 • Spatial differences in opioid use between five Belgian catchment areas

56

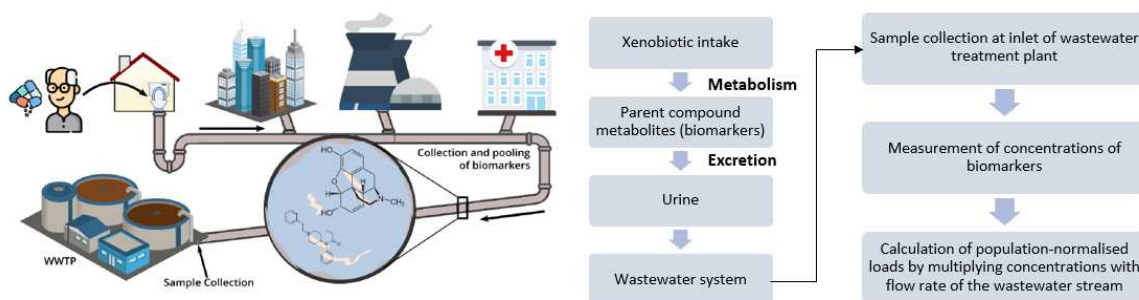
60 **Introduction**

61 Chronic pain affects about 20% of the adult population and has substantial financial costs [1]. Although
62 opioids (e.g. tramadol, oxycodone, fentanyl, etc) are widely considered as the most effective
63 pharmaceuticals for the treatment of acute pain, the risk of addiction exists to the current day in spite of a
64 continued and still promising search for safer opioids [2]. In the United States, the drastically increased use
65 of prescription opioids has led to a public health crisis of epidemic proportions in the last decades, as
66 reflected by the four-fold increase in admissions, morbidity and mortality for substance use disorder
67 treatment [3]. In Belgium, the use of prescription opioids has increased with 35% with the highest increase
68 reported for oxycodone [4]. For policy makers to monitor the burden of (prescription) opioids and react
69 swiftly and decisively, objective and timely data on the use and misuse of opioids in high spatio-temporal
70 resolution is necessary [5–8]. New patterns can become established in a short time, and could develop into
71 major problems before they have even been identified. This suggests that alternative and complementary
72 approaches are needed in order to rapidly provide an objective and clear picture on opioid consumption
73 within different communities.

74 Wastewater-based epidemiology (WBE) is an analytical approach based on the chemical analysis of specific
75 human metabolic excretion products (biomarkers) in influent wastewater. These biomarkers are collected,
76 transported and pooled by the sewage system. Therefore, wastewater contains a wealth of information
77 about the catchment population connected to a specific wastewater treatment plant (WWTP), as illustrated
78 in Figure 1 [9]. For biomarkers stable in wastewater and efficiently conveyed to a wastewater treatment
79 plant (WWTP), it is reasonable to assume that the collective amount excreted by the target population
80 within a given period (day, week, month, season, year, etc) is reflected by the mass load reaching the WWTP
81 in the corresponding interval. Concentrations of biomarkers in influent wastewater need to be analyzed
82 using sensitive and selective analytical techniques (e.g. chromatography coupled with mass spectrometry)
83 and these concentrations can then be multiplied with daily flow rates to obtain the mass loads of
84 biomarkers (in mg/day). The mass loads can be divided by the size of the population present in the
85 catchment area of a WWTP, resulting in population-normalized loads (in mg/day/1000 individuals). This
86 allows results to be compared between different WWTP and/or different time points. The WBE approach
87 can be applied to estimate different health-related aspects of populations. It has been used successfully in
88 measuring use of alcohol, tobacco, illicit drugs and other pharmaceutical classes, dietary habits, and to
89 investigate population exposure to pollutants such as pesticides, flame retardants, plasticizers, etc [10–17].
90 WBE could especially be useful to monitor quickly developing changes in the consumption patterns of
91 pharmaceuticals in near real time and to measure the extent of illegal consumption [9,18,19].

92 In contrast to the monitoring of amphetamine-type drugs using WBE, opioids have been only recently
93 received more attention [17,19–22]. Additionally, most WBE applications only include a minor selection of
94 opioid biomarkers (mostly morphine, codeine, heroine and methadone). However, only a few of these

95 publications quantitatively measure a broad range of biomarkers simultaneously [17,19–23] and others
96 only provide semi-quantitative measurements [24,25].



97

98 Figure 1 Schematic overview of the WBE approach

99 The aim of this study was to develop, validate and apply a analytical method capable of measuring a broad
100 range of opioids and metabolites simultaneously at low levels (ng/L) in influent wastewater (IWW). The
101 validated method will be applied to different catchment areas in Belgium to evaluate opioid consumption
102 at a population level.

103 **Materials and methods**

104 *Reagents and materials*

105 Reference standards and deuterated internal standards (IS) for investigated compounds were purchased
106 from Cerilliant Corporation (Texas, US), Chiron AS (Trondheim, NO), Grünenthal GmbH (Aachen, DE), LGC
107 Limited (Teddington, UK), National Measurement Institute (Canberra, AU) and Toronto Research Chemicals
108 (Ontario, CA). Reference and deuterated standards were of analytical grade and purchased as neat powders
109 or as solutions at concentrations of 1 mg/mL or 100 µg/mL in acetonitrile (AcN) and methanol (MeOH). AcN
110 and MeOH were purchased from Fischer Scientific (Loughborough, UK). Formic acid (analytical grade) was
111 obtained from Merck (Darmstadt, DE). Ultrapure water was prepared using an Elga LabWater Purelab Flex
112 system (Veolia Water Solutions & Technologies, Tienen, BE). Oasis® MCX and HLB solid-phase extraction
113 (SPE) cartridges (60 mg, 3 mL) were purchased from Waters (Massachusetts, USA). SPE was performed on
114 a Supelco 12- or 24-port VISIPREP™ vacuum manifold with a Welch 2023 self-cleaning dry vacuum system.
115 Samples were dried in a REACTI-TERM III #TS-18824 evaporator from Thermo Fischer Scientific
116 (Massachusetts, United States). Centrifugal filters (modified nylon 0.2 µm, 500 µL) were acquired from
117 Avantor (Pennsylvania, United States). An overview of all investigated analytes and the corresponding IS is
118 given in Table 1.

119

120 *Samples and sample treatment*

121 Influent wastewater (IWW) samples were collected from five different Belgian cities covering approximately
122 12% of the Belgian population (Table 2). IWW samples (500 mL) were collected over a 24h period during a
123 normal week (i.e. with no special events occurring), in a time- or volume-proportional manner, in order to
124 be representative for an entire day. During sample collection, the sample collection bottle was cooled (<4
125 °C) and after transport samples were stored at -20 °C until analysis. Daily flow rates (m³/day) were obtained
126 from the Flanders Environment Agency [26].

127 *Method development*

128 Liquid chromatography tandem mass spectrometry

129 The liquid chromatography (LC) system consisted of an Agilent® 1290 Infinity II Ultra High Performance LC
130 (UHPLC), with a degasser, a column thermostat, a binary pump, and an autosampler. In the final protocol,
131 chromatographic separation was carried out with an Atlantis® T3 column (150 mm x 2.1 mm, 3 µm)
132 maintained at 30 °C. The mobile phase consisted of (A) ultrapure water with 0.1% v/v formic acid, and (B)
133 methanol with 0.1% v/v formic acid. A gradient was optimised as follows: 0-0.5 min: 5% B; 0.5-5 min:
134 increase to 25% B; 5-16 min: increase to 30% B; 16-21 min: increase to 95% B; 21-21.1 min decrease to 5%
135 B and equilibration at 5% B up to 25 min. The flow rate was 0.275 mL/min and the injection volume was 4
136 µL.

137 The mass spectrometry (MS) system consisted of an Agilent® 6460 triple quadrupole mass spectrometer
138 equipped with an electrospray ionization source (ESI), operated in positive ionization mode. The source
139 parameters were optimised as follows: gas temperature 300 °C; gas flow 5 L/min; nebulizer pressure 32 psi;
140 sheath gas temperature 350 °C; sheath gas flow 11 L/min; and capillary voltage 3500 V. Dynamic multiple
141 reaction monitoring (dMRM) was used and at least two MRM transitions were used for each analyte: one
142 transition, usually the most abundant, was used as quantifier (Q) and the other was used as qualifier (q).
143 Identification of the analytes of interest in the extracted samples was based on the quantifier/qualifier
144 (Q/q) ratio and the relative retention time (RRT) (i.e. the ratio of the Rt of the compound to that of the IS).
145 A tolerance level for the Q/q ratio of 30% relative standard deviation (RSD) was set for the identification of
146 the compounds of interest within each batch. In addition, the RRT must be within ±2.5% RSD of that of the
147 calibration standard [27,28].

148 Sample preparation

149 An aliquot of 100 mL IWW was spiked with 100 µL of internal standard mixture resulting in a concentration
150 of 100 ng/L (except for TRA-D₆ with a final concentration of 500 ng/L). The aliquot was centrifuged for 30
151 min at 2465g prior to SPE to remove solid particles.

152 In order to obtain the highest and most reproducible absolute recoveries (RE), different SPE cartridges
153 (Oasis HLB (60 mg, 3 mL) and Oasis MCX (60 mg, 3 mL)), washing, and elution conditions were tested. The
154 RE was determined for each analyte by comparing the peak areas of the analyte spiked pre- and post-
155 extraction at 200 ng/L in tap water, following this equation [29]:

$$156 \quad RE (\%) = \frac{\text{peak areas of the standards spiked pre extraction}}{\text{peak areas of the standards spiked post extraction}}$$

157 *Equation 1 Absolute recovery (RE) calculations used to evaluate the different SPE conditions*

158 In the finalized protocol, Oasis HLB cartridges were gravitationally conditioned with 4 mL MeOH and
159 subsequently 4 mL ultrapure water at pH 7. The supernatant from the centrifugal tubes was loaded under
160 vacuum on the SPE cartridges. After loading of the samples, washing of the cartridges was done with 3 mL
161 20% (v/v) methanol in ultrapure water, under vacuum. The SPE cartridges were afterwards vacuum-dried
162 for at least 30 minutes. Elution of the analytes from the SPE cartridges was performed with 8 mL of 2% v/v
163 formic acid in methanol. Subsequently, the eluent was evaporated at 37 °C under a gentle nitrogen stream
164 and reconstituted in 150 µL of 5% methanol/95% ultrapure water + 0.1% v/v formic acid, which was the
165 same as the starting mobile phase conditions. The reconstituted extract was vortexed for 2.5 minutes,
166 centrifuged on a centrifugal filter (0.2 µm) at 10 000g, transferred to an autosampler vial with glass insert
167 and finally analyzed with liquid chromatography coupled to tandem mass spectrometry (LC-MS/MS).

168

169 *Method validation*

170 The analytical method was validated according to the guidelines for bioanalytical method validation by the
171 European Medicines Agency (EMA) with minor adjustments [30]. Tap water was used as an alternative
172 matrix for method validation because it is not possible to obtain blank influent wastewater matrix. Yet, tap
173 water is not the most ideal alternative because of lower matrix effects compared to real influent
174 wastewater. For this reason, standard addition experiments were applied to investigate whether the IS
175 robustly compensates for matrix interferences (see below).. Among performance features, linearity,
176 precision, accuracy, lower limit of quantification (LLOQ), calibration range and specificity/selectivity were
177 assessed during method validation using tap water.

178 Selectivity was evaluated by analyzing three zero blank samples (i.e containing no analytes or IS) for
179 interferences. The response of the quantifier transition in the zero blank should be <20% of the response
180 of the LLOQ for each biomarker and <5% for the response of the IS. The confirmation criteria based on RRT
181 and Q/q ratios (described earlier) were applied to distinguish matrix interferences from the analytes of
182 interest in IWW. Carry-over was assessed for each biomarker by injecting a zero blank sample after the
183 highest calibration level and comparing the response of the quantifier transition in this zero blank to the
184 response of the quantifier transition of the LLOQ sample. Limits were set at <20% for the analyte and <5%
185 for the IS.

186 For each biomarker, a seven-level calibration curve was prepared in tap water; linearity and curve fitting
187 were evaluated by testing weighting factors $1/x$ or $1/x^2$. Back-calculated concentrations of the seven
188 calibration standard levels should be within 15% of the nominal value, or within 20% at the LLOQ.

189 Within- and between-run accuracy and precision were determined using four quality control (QC) levels.
190 The LLOQ was equal to the lowest calibration level, the QC low (QCL) within three times the LLOQ, the QC
191 medium (QCM) between 30-50% of the calibration curve range, and the QC high (QCH) was between 75-
192 100% of the calibration curve range. For within-run and between-run accuracy and precision four and
193 twelve replicates were considered respectively for each QC level. For accuracy, mean concentration levels
194 should be 15% of the nominal values for the QC level; except for the LLOQ which should be within 20%. For
195 precision, the coefficient of variation (CV) should not exceed 15%; except for the LLOQ where the CV should
196 not exceed 20%.

197 The recommendations from Matuszewski et al. concerning process efficiency (PE) were modified and
198 applied [12,29]. PE is the product of matrix effects (ME) and RE during sample preparation, as defined by
199 Matuszewski et al [29]:

$$200 \quad PE = \frac{ME \times RE}{100} = \frac{\text{peak areas for standards spiked before extraction in blank matrix}}{\text{peak areas obtained in neat solution standards}} \times 100\%$$

201 *Equation 2 Calculation of the process efficiency according to Matuszewski et al [29].*

202 Standard addition was used to evaluate PE given the impossibility of obtaining blank influent wastewater
203 matrix:

$$204 \quad PE = \frac{(\text{concentration in spiked sample before extraction (C2)} - \text{concentration in a corresponding native sample (C1)})}{\text{concentration obtained in neat solution standard (A)}} \times 100\%$$

205 *Equation 3 Calculation of the process efficiency through standard addition experiments.*

206 Thus, the measured concentration in the C2 sample equals the sum of spiked and native concentration
207 (Figure S1). The proposed set-up was employed to test whether the method is sensitive enough to
208 distinguish the compounds under investigation from the matrix interferences, and to assess if the IS corrects
209 for potential ME and/or losses during sample preparation.

210 Six samples of pooled IWW originating from different sources were spiked to evaluate PE (Figure S1).
211 Multiple sources of IWW were used to mimic different IWW matrices. Each IWW pool (N = 6) was further
212 subdivided in 100 mL aliquots of non-spiked 'control' samples (C1 in Equation 3), and spiked samples (C2 in
213 Equation 3), as illustrated by Figure S1. The C2 samples were spiked with 100 μ L of a spiking mixture with
214 biomarker concentrations ranging from 250 to 1000 ng/L to ensure that spiked concentrations were
215 substantially higher (up to 10 times) compared to native concentrations. Native concentrations measured
216 in the control sample were subtracted from the concentrations found in the spiked IWW samples (if >LLOQ),
217 as illustrated in Eq 2. It was accepted that the IS compensates for possible loss during sample preparation
218 and matrix effects when PE was within 80-120% bias [12,29].

219 *Stability experiments*

220 The experimental set-up to assess in-sample stability was done accordingly to McCall et al [31]. In short, a
221 large wastewater pool was made by subsampling different IWW sources. This IWW pool was subsequently
222 divided in 3 aliquots of 1000 mL, including a non-spiked 'control' IWW sample and two spiked IWW samples
223 (Figure S2). Metabolites (M) and parent (P) compounds were spiked separately in different pools to insure
224 there was no interference from overlapping metabolization pathways. The M and P pools were spiked with
225 high concentrations of compounds (250-1000 ng/L) to ensure that spiked biomarkers were present in
226 substantially higher amounts compared to control samples. During the experiment, the aliquots were
227 placed at room temperature, and on a magnetic stirrer (300 rpm) to simulate sewer currents. The time
228 point the aliquots were spiked was considered the as time point 0 h. At specific time intervals (2 h, 6 h, 10
229 h, and 24 h) two aliquots of 100 mL were taken from each sample and prepared according to the sample
230 protocol outlined earlier, including the SPE drying step. The cartridges were stored at -20 °C until the
231 moment of elution, which was the next day. The aliquots taken at time interval 24 h were eluted after the
232 drying step, along with the other stored SPE cartridges. The time interval was set to 24 h since this
233 encompasses the residence times in most sewer systems in Belgium.

234 Native biomarker concentrations measured in the control non-spiked samples, when above LLOQ, were
235 subtracted from the concentrations in the spiked samples. Mean concentrations at time intervals 2 h, 6 h,
236 10 h and 24 h were normalized against time point 0 h to evaluate the stability.

237 Each biomarker was categorized as either high (<20% transformation), medium (20-60%), low (60-100%) or
238 variable (i.e. varying results found in different studies) stability over a 24-h period [31]. Biomarkers with
239 medium to low (and variable) stability are unreliable to perform further back-calculations and should, if
240 transformation is not corrected for, be excluded in WBE studies.

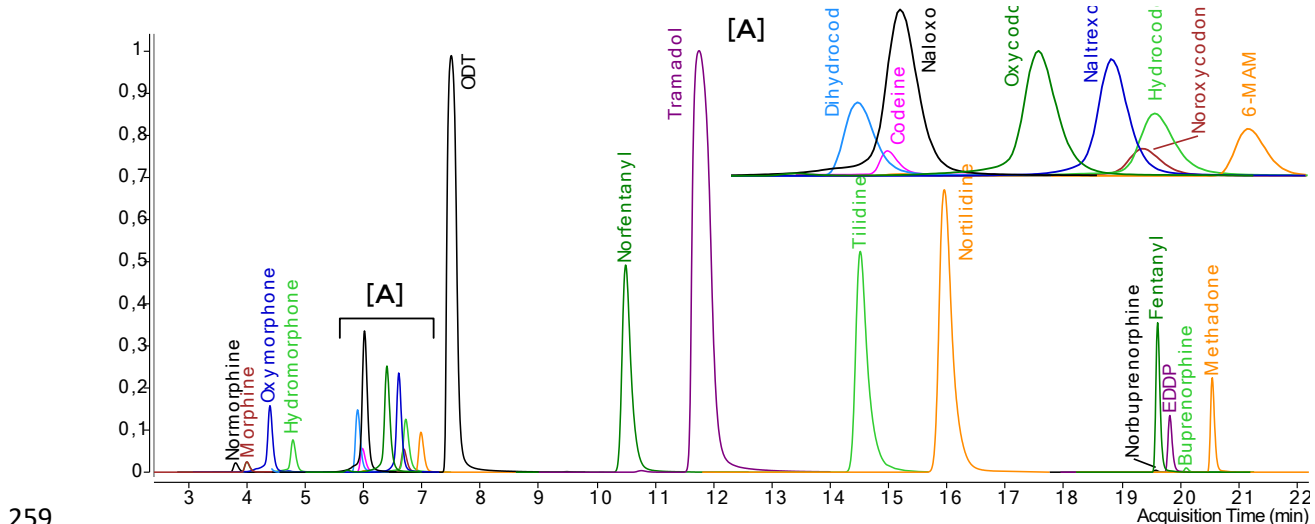
241 **Results and discussion**

242 *Method development & optimization*

243 *Liquid chromatography-tandem mass spectrometry*

244 For LC optimization, a standard mixture with all analytes of interest was injected on different reversed-
245 phase LC (RPLC) columns: a Kinetex® EVO C18 (100 mm x 2.1 mm, 2.6 µm) column, a Kinetex® Biphenyl
246 (100 mm x 2.1 mm, 2.6 µm) column and an Atlantis T3 (150 mm x 2.1 mm, 3 µm) LC column. This was done
247 in combination with the use of different organic phases (MeOH or AcN) and buffer conditions (0.1% v/v
248 HCOOH or 0.04% v/v HCOOH). Most of the analytical methods found in literature applied RPLC with the use
249 of modified C₁₈ LC columns [17,20,21]. The Atlantis® T3 column resulted in the best retention for the most
250 polar compounds, such as MOR and norMOR, and was therefore chosen for chromatographic separation
251 (Figure 2). norMOR eluted within the first 2 minutes when using the Kinetex® EVO C18 and Kinetex®

252 Biphenyl LC column (Figure S3). Initially, mobile phase A and B consisted respectively of ultrapure water
 253 and acetonitrile both with 0.1% v/v formic acid using a standard gradient of 0-0.5 min: 5% B; 0.5-15 min:
 254 increase to 95% B; 15-17 min: 95% B; 17-17.1 min: decrease to 5% B. Lowering the percentage of aqueous
 255 mobile phase at the beginning of the run did not result in sufficient retention of norMOR on these columns.
 256 For the organic mobile phase, MeOH with 0.1% v/v formic acid was chosen because this resulted in better
 257 retention and sensitivity of the analytes of interest. All compounds eluted within 21 minutes as illustrated
 258 by Figure 2.



260 Figure 2 Chromatographic overview of all MRM quantifier transitions in tap water spiked at the highest calibration level.
 261 The y-axis scale was represented as the abundance relative the most abundant peak (in this case tramadol)

262 Compound-dependent MS parameters, including fragmentor voltage, collision energy and ionization mode,
 263 were determined by injecting 1 ppm standard solutions without LC column. For each analyte, the transition
 264 with the highest signal-to-noise ratio (S/N) and absolute abundance was used as quantifier transition and
 265 the two remaining abundant transitions were used as qualifier transition (Table S1). Biomarkers which
 266 share the same transitions were separated sufficiently with the Atlantis T3 column. The effect of co-eluting
 267 matrix interferences on peak intensity was also considered when choosing quantifier and qualifier
 268 transitions since matrix-induced suppression or enhancement could result in poor or improved peak
 269 intensities.

270 TRA shared its transitions with a coeluting interferent (Figure S4). A reference standard was used to confirm
 271 the identity of the interference as O-desmethylvenlafaxine (ODV), a metabolite of the antidepressant
 272 venlafaxine and previously quantified in wastewater [12,32]. As a result, chromatographic separation was
 273 further optimized to separate these compounds. Baseline separation was achieved by increasing the
 274 aqueous gradient time at the moment of elution.

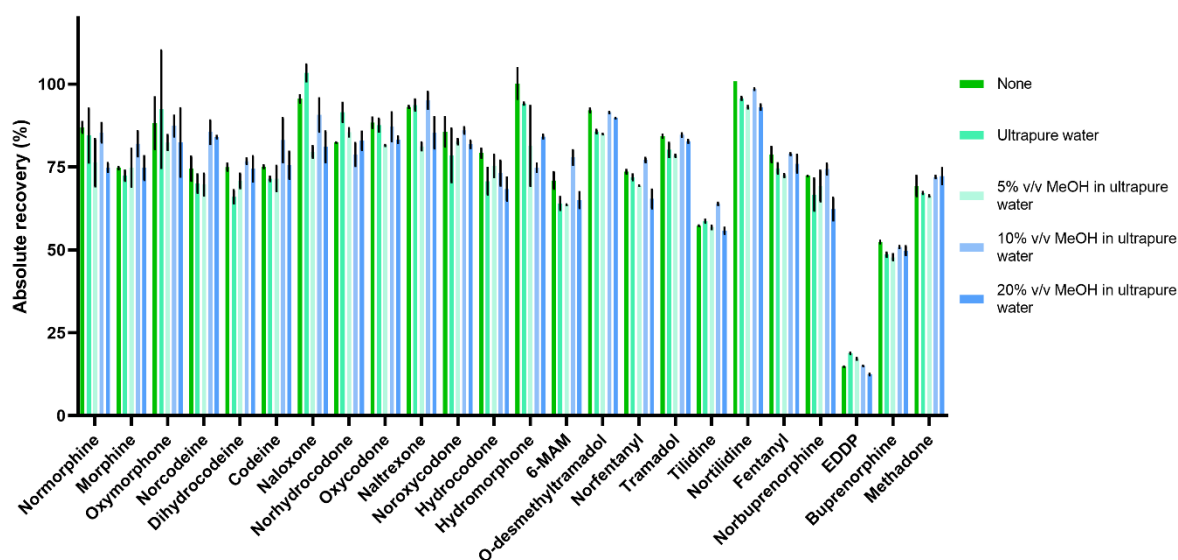
275 norHCD, norCOD and 6-MAM did not go through method validation due to sensitivity issues. This is less of
 276 a concern for norHCD as the parent drug hydrocodone is, since 2008, no longer available on the Belgian

277 market [33]. This is more important for norCOD since it is a distinct metabolite of COD, especially because
278 of the overlap in metabolic pathways between heroin (HER), MOR and COD (Figure S5).

279 Furthermore, PIR was excluded from the final method due to poor reproducibility of the reference standard
280 peaks in tap water. In some European countries, PIR is considered the first-choice opioid analgesic for pre-
281 and post-operative pain [34]. It should be noted that the global use of PIR is rather limited [35].
282 Consumption in Belgium is also low [4], making this exclusion of less concern. In 2017, piritramide was only
283 dispensed in Austria, Belgium, Curaçao, Czechia, Germany, Italy, Luxembourg, the Netherlands, Slovakia,
284 and Slovenia [35].

285 Sample preparation

286 Most of the multi-analyte sample preparation methods used for the extraction of opioids in IWW applied
287 SPE with Oasis MCX (or Strata-X-C) and Oasis HLB cartridges with loading volumes between 50-500 mL [20–
288 25,36] and used (acidified) ultrapure water as a washing solvent. Methanol (with or without pH modifier)
289 and dichloromethane/isopropanol mixtures were used most frequently as elution solvents. RE was assessed
290 to select the most optimal SPE conditions, including elution solvents, washing solvents, pH modifiers and
291 solvent volumes. Initially, extraction with both Oasis HLB and MCX cartridges resulted in high RE (Table S2),
292 which was in line with the findings of others [17,21]. However, the MCX procedure was discarded in an
293 early stage because of the high variability in RE among the different SPE experiments compared to the HLB
294 procedure. It should be noted that the RE of EDDP was low, yet highly reproducible with the HLB extraction.
295 However, the combination of a highly sensitive detection and considerable concentrations of EDDP in
296 influent wastewater allow the use of the HLB protocol. Table S2 indicates that an elution volume of 8 mL
297 resulted in the highest RE with the HLB procedure. Higher elution volumes did not yield in further
298 improvements. Overall, the use of MeOH and MeOH with acidic modifier (i.e. 2% v/v FA) as an elution
299 volume was similar. However, MeOH with 2% v/v FA was selected as elution solvent for sample preparation
300 since it resulted in slight improvements for a minor selection of biomarkers including norOXY and OMP.



301

302 Figure 3 Absolute recoveries (RE) (in % \pm SD, n = 2) in tap water for all compounds per protocol. Colours represent the
 303 different washing solvents tested.

304 In the analytical method proposed by Krizman-Matic et al., co-eluting matrix interferences were reported
 305 with the use of Oasis HLB [21], emphasizing the need for further optimization of the washing solvent. Figure
 306 3 illustrates the RE (in % \pm SD) using the HLB procedure with 8 mL of 2% v/v FA in MeOH as elution solvent
 307 and varying washing solvents. Washing of the HLB cartridges with 3 mL of 20% v/v MeOH proved to have
 308 a positive effect on the peak shape and signal intensity of the investigated compounds, potentially by
 309 washing away matrix interferences. No substantial loss in absolute recoveries (<20%) was observed with
 310 this washing solvent. RE were highly reproducible and ranged between 50-93% with the exception of EDDP.

311 *Method validation*

312 Performance criteria

313 For 19 compounds, the performance criteria met the requirements for method validation provided by the
 314 EMA guidelines, as illustrated in Table S3.

315 The method proved to be selective as analysis of three zero blank samples did not result in any interference.
 316 No significant carry-over occurred for most biomarkers (< 3%), however, carry-over was higher for
 317 methadone (17%) and fentanyl (10%) but still less than 20% of the LLOQ as recommended by EMA.
 318 Additionally, injection of IS did not interfere with the analytes and vice versa.

319 A linear calibration curve ranging from the low ng/L to low μ g/L range was obtained for 21 compounds in
 320 tap water. The LLOQ of the analytes of interest was between 1 ng/L and 100 ng/L, based on their
 321 concentrations found in IWW. A weighting of $1/x^2$ was considered more appropriate for biomarkers with
 322 measured concentrations at the lower end of the calibration curve, whereas $1/x$ was used for higher
 323 concentrations. Most biomarkers favored a curve weight of $1/x^2$ based on their low concentrations in IWW,
 324 as illustrated in Table S3.

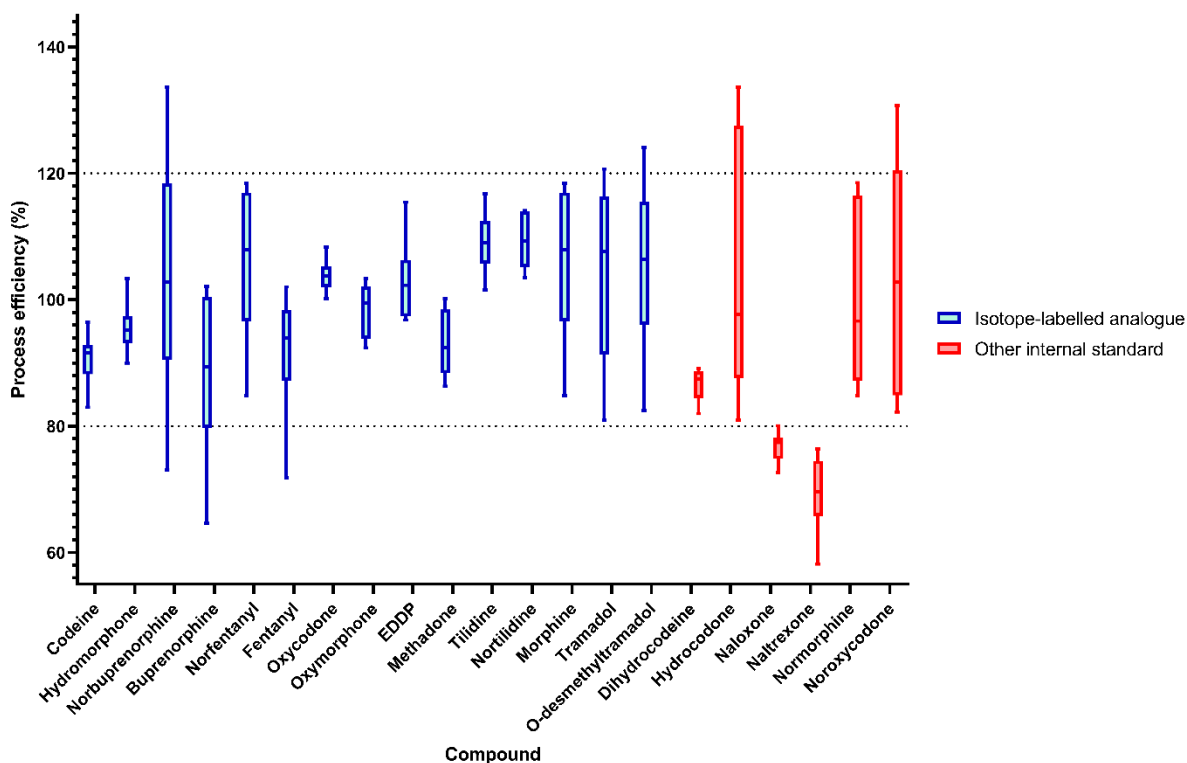
325 The within-run and between-run accuracy and precision measured at the QC levels met the acceptance
326 criteria. Within-run and between-run accuracy and precision results (Table S3) at four different spiking
327 levels (i.e. LLOQ, QC_{low}, QC_{mid}, QC_{high}) were respectively within the range of <15% bias and <15%CV.

328 The LLOQs in the present study range between 1-15 ng/L for all biomarkers of investigation with the
329 exception of tramadol which has a LLOQ of 100 ng/L. From an analytical perspective, it would be probably
330 possible to achieve lower LLOQs, however, the relevance of an LLOQ needs to be considered from a WBE
331 perspective. From a WBE point-of-view, the analytical approach needs to be sensitive enough to pick up trends
332 in consumption patterns of prescription drugs. Even though it is theoretically possible to obtain lower
333 detection levels, this might be less of interest because these concentrations correspond with very low levels
334 of consumption. For this reason, the proposed LLOQs of this analytical methodology are appropriate to
335 capture relevant consumption patterns in the use of prescription opioids.

336 Process efficiency

337 The PE experiment was designed to investigate whether the IS was able to correct for ME and/or loss of
338 analyte during extraction (i.e. RE). Deviations in PE originate from differences between the analyte and the
339 corresponding IS in wastewater due to different signal suppression or enhancement. This is especially
340 important for biomarkers for which no deuterated analogue was available as IS (Table 1). Figure 4
341 summarizes the results of the PE standard addition experiment. PE were within $\pm 20\%$ bias for all compounds
342 for which an isotope-labelled analogue was available. For DHC, HCD, norMOR and norOXY, the following IS
343 COD-D₆, HMP-D₃, MOR-D₃ and OXY-D₆ respectively were able to compensate for potential losses due to
344 matrix interferences or during SPE. For NLX and NTX, no isotope-labelled deuterated analogues were
345 available during method development. No alternative IS was found for NLX and NTX and therefore these
346 compounds were excluded from the final method. The high reproducibility of the PE of NLX would
347 potentially allow the use of a correction factor (CF) to accurately measure concentrations of NLX in
348 wastewater, but more research is needed to establish such a CF.

Evaluation of process efficiency (% \pm SD) ($N \geq 6$) through standard addition experiments



349

350 Figure 4 Estimation of process efficiency through standard addition experiments in influent wastewater.

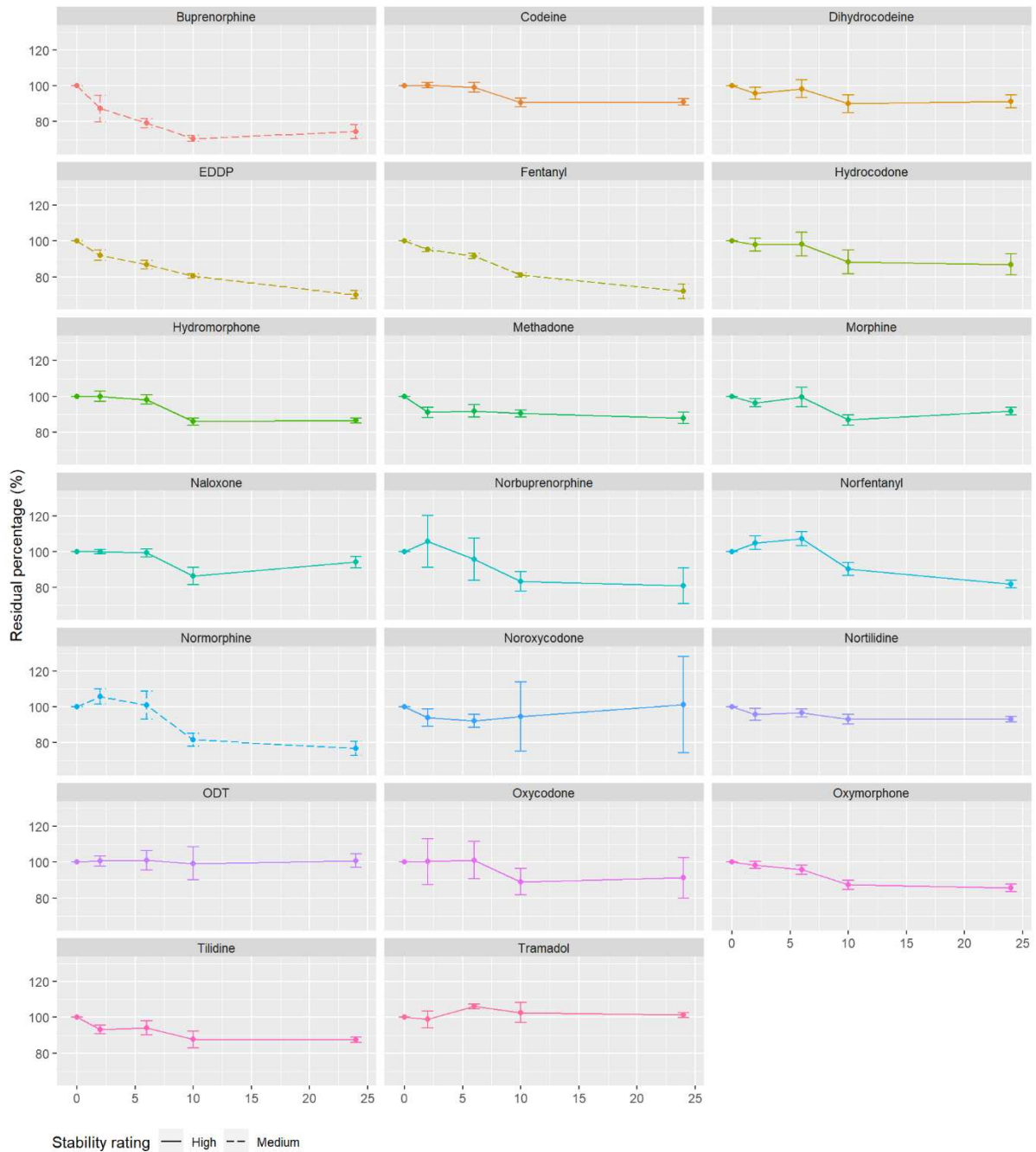
351 Stability

352 Benchtop stability of the analytes under investigation was investigated as in-sample biomarker
 353 transformation could result in over- or underestimation of population-normalized mass loads. This study
 354 did not evaluate in-sewer degradation in presence of a biofilm under gravity or rising main sewer conditions
 355 or in a pilot sewer study. The in-sewer transformation of the targeted biomarkers could lead to additional
 356 uncertainty. Additionally, this study did not investigate in-sample stability at -20 °C. However, McCall et al.
 357 indicate that most opioid compounds (e.g. morphine, oxycodone, fentanyl,...) are sufficiently stable during
 358 sample storage at -20 °C, with the exception of some compounds such as 6-MAM and heroin [31].
 359 Furthermore, we did not assess potential loss due to sorption to solid particulate matter, which would
 360 potentially lead to uncertainty in back-estimating biomarker population-normalised mass loads. Baker et
 361 al. found that the average proportion of solid particulate matter was >10% with regard to MTD, EDDP and
 362 FEN, but was acceptable for most opioid biomarkers (e.g. norCOD, DCD, TRA,...) [37]. It should also be noted
 363 that PE also evaluates if any potential loss of analyte due to sorption to particulates is corrected.

364 Figure 5 shows the residual percentages of the analytes of interest at five different time points (0 h, 2 h, 6
 365 h, 10 h and 24 h). 16 compounds showed high stability (<20% transformation) over 24 h. Stability of BUP,
 366 norMOR, FEN and EDDP was only medium, with more than 20% but less than 40% in-sample transformation.
 367 However, the use of BUP and FEN could be easily monitored through the use of their metabolites norBUP
 368 and norFEN. Similarly, the parent compound MTD could be used in the case of EDDP, which proves to be

369 stable in IWW (with 12% of the initial MTD load transformed). As illustrated by Figure S5, there is overlap
370 between the metabolic pathways of MOR, COD and HER and norMOR, therefore it is more appropriate to
371 measure MOR use. However, the benchtop stability of norMOR should be taken into account to accurately
372 measure its concentrations in influent wastewater. To our knowledge, it was the first time in-sample
373 stability was determined for HMP, norTIL, ODT and TIL. Even though the residual percentage of BUP and
374 FEN decreased in the IWW pools spiked with parent compound (P1 and P2, see Figure S2), we did not
375 observe a parallel increase in norBUP and norFEN in these pooled IWW samples. This means that BUP and
376 FEN are not degraded in norBUP and norFEN, further proving the suitability of these metabolite as
377 alternative biomarkers.

378 The results found in this study were comparable with the results found by Baker et al for most compounds
379 under investigation (Table S4) [38]. However, for BUP, MOR, OMP and norMOR, the data did not match as
380 Baker et al did report in-sample formation of MOR and OMP and no more than 20% transformation was
381 observed for BUP and norMOR in this study. According to McCall et al, these biomarkers should strictly be
382 excluded from the analytical method due to variable stability. Therefore, OXY use should be monitored
383 through its metabolite norOXY and for MOR a more specific and stable metabolite should be further
384 explored.



385

386 Figure 5 Benchtop stability in wastewater of opioids, transformation of biomarker at each time point. Mean residual percentages
 387 of four spiked samples, normalised against time of spiking are reported for time points 2 h, 6 h, 10 h and 24 h. The error bars
 388 represent the relative standard deviation (%RSD) between the concentrations measured in the replicate wastewater pools.

389 *Method application*

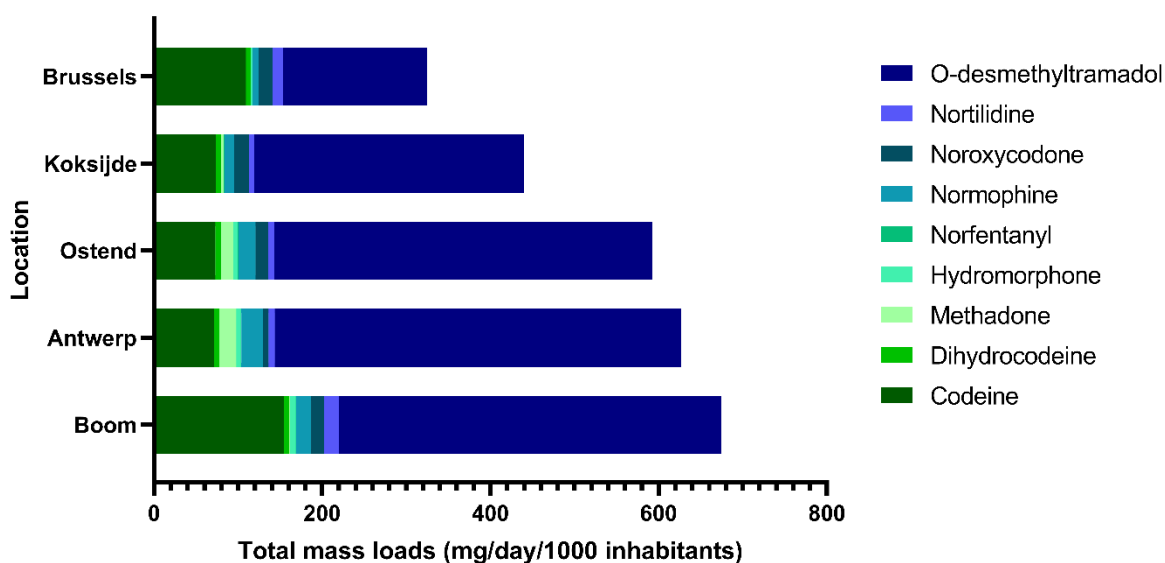
390 Spatial comparison of prescription opioid use

391 The applicability of the validated method was evaluated on 35 different IWW samples from 5 different
 392 WWTPs (Table 2). Apart from BUP and HCD, every validated biomarker was detected (>LLOQ) at least once.
 393 Calculated population-normalised mass loads of the investigated analytes are shown in Table S5. Highest
 394 population-normalised mass loads were reported for TRA, ODT and COD which is in line with sales and

395 dispensing data [39,40]. Although fentanyl is prescribed in a relatively high number of patients in Belgium
396 [4], population-normalised mass loads of norFEN and FEN are close to the LLOQ level because of its low
397 dose. The magnitude of the population-normalized mass loads found in this study were also comparable
398 with the results of other WBE applications on prescription opioids [17,20,22,23,41].

399 Figure 6 summarizes the total opioid population-normalized mass loads in all locations under investigations.
400 Note that only one biomarker (preferably the metabolite as it represents actual consumption) was included
401 for each opioid prescribed in Belgium. A limitation of this study is that only one week of sampling within
402 the same time period (i.e. Sep 2019) was included for the spatial comparison, and consumption rates might
403 be different in the sampling period compared to the rest of the year. Spatial differences were found for the
404 majority of prescription opioids among the different locations.

Spatial comparison of the total opioid population-normalized mass loads



405
406 Figure 6 Spatial analysis of total opioid consumption among different catchment areas.

407 At this moment HER use can only be measured based on morphine concentrations, after the correction for
408 morphine originating from consumed morphine and codeine. In the future, a specific biomarker for HER
409 needs to be found in order to investigate consumption patterns of this compound with less uncertainty.
410 The in-sewer transformation of 6-monoacetylmorphine (6-MAM) is substantial and paralleled with an
411 increase of morphine, as reported by Senta et al. [42]. This implies that at this stage it is difficult to estimate
412 HER consumption from 6-MAM.

413 Investigation of metabolite/parent compound ratios

414 For 6 compounds, the parent compound and metabolite were included in the analytical method. However,
415 BUP was never detected and FEN was only detected occasionally. Table 3 illustrates the metabolite to
416 parent compound ratios (M/PC). M/PC are expected to be relatively constant for each location. It should

417 be noted that (accidental) discharge of the parent drug can as also result in a decrease of the
418 metabolite/parent ratio.

419 The mean M/PC of EDDP/MTD is 1.33 ± 0.19 , which was comparable with the average EDDP/MTD ratio of
420 1.97 found in 44 other wastewater studies [43]. In all locations the population-normalised mass loads of
421 EDDP were higher than MTD. The norMOR/MOR ratio was relatively constant within each location. Based
422 on the mean influent concentrations of only 7 IWW samples, Boleda et al. reported a M/PC ratio of 7.9%,
423 which was lower compared to this study [44]. However, deviations in the norMOR/MOR ratio could be the
424 result of overlapping metabolic pathways for MOR. MOR could also be metabolized from COD and HER and
425 therefore COD and HER use could potentially influence the observed norMOR/MOR ratios. Baker et al found
426 a OMP/OXY ratio of 1.66 which is in line with the results found in this study [17]. No relevant literature
427 could be found for ratios norOXY/OXY and norTIL/TIL. As illustrated in Table 3, the metabolite was always
428 measured in higher concentrations than the parent drug. Variations in these ratios appear to be relatively
429 high, which could be related to their lower concentrations found in IWW compared to MTD, MOR and their
430 metabolites. It should be noted that this variability is not related to in-sample stability since OXY, TIL and
431 their metabolites showed <20% transformation for 24h.

432 Note that if the population (or rather the number of people using a drug) attached to a WWTP is low, the
433 variance associated with ratios will increase. This is because interindividual metabolization variability (poor
434 metabolizer, high metabolizer, ...) will be more predominant.

435 **Conclusions**

436 A sensitive analytical method based on SPE and LC-MS/MS was developed and validated for the
437 simultaneous measurement of 19 prescription opioids and their metabolites at trace concentrations in
438 influent wastewater. The PE was acceptable for all compounds for which an isotope-labelled deuterated
439 analogue was available. However, for NLX and NTX, no IS was found able to compensate for ME and
440 potential losses during extraction and therefore these compounds were excluded from the final method.
441 All compounds with the exception of BUP, EDDP, FEN and norMOR showed high benchtop stability.

442 Apart from BUP and HCD, all biomarkers were detected at least once in Belgian IWW. Wherever possible
443 metabolite/parent compound ratios were determined but, apart from EDDP/methadone, literature data
444 for comparison is scarce. The M/PC ratios were in line with the results found in other WBE studies on
445 opioids. Overall, metabolites were found in higher levels compared to parent compounds among all
446 location.

447 **Acknowledgements**

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450 **Figures**

451 *Figure 1 Schematic overview of the WBE approach*

452 *Figure 2 Chromatographic overview of all MRM quantifier transitions in tap water spiked at the highest calibration level. The*
453 *y-axis scale was represented as the abundance relative the most abundant peak (in this case tramadol)*

454 *Figure 3 Absolute recoveries (RE) (in % \pm SD, n = 2) in tap water for all compounds per protocol. Colours represent the different*
455 *washing solvents tested.*

456 *Figure 4 Estimation of process efficiency through standard addition experiments in influent wastewater.*

457 *Figure 5 Benchtop stability in wastewater of opioids, transformation of biomarker at each time point. Mean residual*
458 *percentages of four spiked samples, normalised against time of spiking are reported for time points 2 h, 6 h, 10 h and 24 h.*
459 *The error bars represent the relative standard deviation (%RSD) between the concentrations measured in the replicate*
460 *wastewater pools.*

461 *Figure 6 Spatial analysis of total opioid consumption among different catchment areas.*

462

463 **Tables**

464 *Table 1. Target biomarkers and corresponding internal standard for quantification purposes. (*) = compounds for which no*
 465 *isotope-labelled analogue was used; (**) = not transferred to method validation*

Pharmaceutical	Biomarker monitored in wastewater	Abbreviation	Internal standard
Buprenorphine	Buprenorphine	BUP	Buprenorphine-D ₄
	Norbuprenorphine	norBUP	Norbuprenorphine-D ₃
Codeine	Codeine	COD	Codeine-D ₆
	Norcodeine*, **	norCOD	Codeine-D ₆
	Morphine and metabolites		
Dihydrocodeine	Dihydrocodeine*	DCD	Codeine-D ₆
Fentanyl	Fentanyl	FEN	Fentanyl-D ₅
	Norfentanyl	norFEN	Norfentanyl-D ₅
Heroin	6-Monoacetylmorphine**	6-MAM	6-MAM-D ₃
	Morphine	MOR	Morphine-D ₃
Hydrocodone	Hydrocodone*	HCD	Hydromorphone-D ₃
	Norhydrocodone*, **	norHCD	
	Dihydrocodeine		
Hydromorphone	Hydromorphone	HMP	Hydromorphone-D ₃
Methadone	Methadone	MTD	Methadone-D ₉
	2-ethylidene-1,5-dimethyl-3,3-diphenylpyrrolidine	EDDP	EDDP-D ₃
Morphine	Morphine		Morphine-D ₃
	Normorphine*	norMOR	Morphine-D ₃
Naloxone	Naloxone*	NLX	Codeine-D ₆
Naltrexone	Naltrexone*	NTX	Norfentanyl-D ₅
Oxycodone	Oxycodone	OXY	Oxycodone-D ₆
	Noroxycodone*	norOXY	Oxycodone-D ₆
	Oxymorphone		
Oxymorphone	Oxymorphone	OMP	Oxymorphone-D ₃
Piritramide	Piritramide*, **	PIR	*
Tilidine	Tilidine	TIL	Tilidine-D ₆
	Nortilidine	norTIL	Nortilidine-D ₃
Tramadol	Tramadol	TRA	Tramadol-D ₆
	O-desmethyltramadol	ODT	O-desmethyltramadol-D ₆

466

467 *Table 2 Overview of the sampling locations and periods*

WWTP (City)	Period	Population serviced by WWTP	Sampling mode
Antwerp-South (Antwerp)	23/09/2019-29/09/2019	130,218	Time-proportional
Boom (Boom)	23/09/2019-29/09/2019	30,600	Time-proportional
Brussels-North (Brussels)	23/09/2019-29/09/2019	953,987	Volume-proportional
Ostend (Ostend)	23/09/2019-29/09/2019	159,000	Time-proportional
Wulpen (Koksijde)	23/09/2019-29/09/2019	78,441	Time-proportional

468

469

470 Table 3. Percentual mass load ratios of metabolite/parent drug per location. N.d.: The M/PC ratio could not be determined
 471 because either the parent or metabolite could not be detected in IWW.

	EDDP/MTD (±SD)	OMP/OXY (±SD)	NorMOR/MOR (±SD)	norOXY/OXY (±SD)	norTIL/TIL (±SD)
AZ	1.11 ± 0.12	n.d.	0.26 ± 0.02	n.d.	1.30 ± 0.28
BOO	n.d.	1.06 ± 0.78	0.38 ± 0.03	1.61 ± 0.85	1.76 ± 0.63
BRU	n.d.	1.57 ± 0.66	0.12 ± 0.01	4.80 ± 1.29	1.71 ± 0.64
OOS	1.41 ± 0.14	2.26 ± 0.70	0.29 ± 0.01	2.73 ± 0.37	1.87 ± 0.55
WUL	1.46 ± 0.12	2.57 ± 0.40	0.28 ± 0.03	3.25 ± 0.45	n.d.
Mean	1.33 ± 0.19	1.87 ± 0.68	0.27 ± 0.09	3.17 ± 1.34	1.66 ± 0.25

472

473

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