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3 Full title:

4 **Quantification of acetaminophen and its metabolites in plasma using**
5 **UPLC-MS: doors open to therapeutic drug monitoring in special patient**
6 **populations**

7

8 Short title:

9 **Quantification of acetaminophen and its metabolites**

10

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32 **Conflicts of Interest and Source of Funding**

33 The authors declare that they have no conflict of interest nor have they
34 received funding.

35

36 **ABSTRACT**

37 Background: Acetaminophen (APAP, paracetamol) is the most commonly
38 used drug for pain and fever in both the United States and Europe and
39 considered safe when used at registered dosages. Nevertheless, differences
40 between specific populations lead to remarkable changes in exposure to
41 potentially toxic metabolites. Furthermore, extended knowledge is required
42 on metabolite formation following intoxication, to optimize antidote
43 treatment. Therefore, the authors aimed to develop and validate a quick
44 and easy analytical method for simultaneous quantification of APAP, APAP-
45 glucuronide, APAP-sulfate, APAP-cysteine, APAP-glutathione, APAP-

46 mercapturate, and protein-derived APAP-cysteine in human plasma by ultra-
47 performance liquid chromatography-electrospray ionization-tandem mass
48 spectrometry (UPLC-MS/MS).

49 Methods: The internal standard was APAP-D4 for all analytes.
50 Chromatographic separation was achieved with a reversed-phase Acquity
51 UPLC HSS T3 column with a runtime of only 4.5 minutes per injected sample.
52 Gradient elution was performed with a mobile phase consisting of
53 ammonium acetate, formic acid in Milli-Q ultrapure water or in methanol at
54 flow rate of 0.4 mL/min.

55 Results: A plasma volume of only 10 μ L was required to achieve both
56 adequate accuracy and precision. Calibration curves of all six analytes were
57 linear. All analytes were stable for at least 48 hours in the autosampler; the
58 high quality control of APAP-glutathione was stable for 24 hours. The
59 method was validated according to the US Food and Drug Administration
60 guidelines.

61 Conclusions: This method allows quantification of acetaminophen and six
62 metabolites, which serves purposes for research, as well as therapeutic drug
63 monitoring (TDM). The advantage of this method is the combination of
64 minimal injection volume, a short runtime, an easy sample preparation
65 method, and the ability to quantify acetaminophen and all six metabolites.

66

67 **Keywords:**

68 Acetaminophen, metabolites, UPLC-MS/MS, therapeutic drug monitoring,
69 pharmacokinetics

70

71 **INTRODUCTION**

72 Acetaminophen (APAP, N-Acetyl-p-Aminophenol, or paracetamol) is
73 the most commonly used drug for pain and fever in both the United States
74 and Europe [1]. Acetaminophen is generally safe when used at registered
75 dosages, thereby titrated upon effect, with a maximum of 4 g/day in four
76 doses for adults. In children, dosage depends on age and weight as follows:
77 with <1 month—30-60 mg/kg/day in three doses; with age >1 month—up
78 to 90 mg/kg/day in four doses [2]. On the other hand, administration of
79 supratherapeutic doses of acetaminophen is the leading cause for liver
80 failure in the United States [3], mainly influenced by its drug metabolism.
81 This metabolism has been reported to deviate in (premature) neonates [4],
82 obese patients [5], and following supratherapeutic doses [6]. Such variability
83 in exposure to potentially toxic metabolites can be expected in other
84 specific populations as well, e.g., anorexic patients, patients from different
85 ethnical backgrounds, extreme elderly, pregnant women and their fetuses
86 [7], and in patients with drug- or genetically driven changes in CYP1A2
87 activity, e.g., omeprazole induction.

88 Acetaminophen is largely metabolized in the liver, predominantly by
89 sulfation and glucuronidation (Figure 1). In adults, sulfation encompasses
90 about 30% and glucuronidation about 55% [8-10]; only 2%-5% is excreted

91 unchanged in the urine [9, 10]. Approximately 5%-10% of acetaminophen is
92 metabolized by cytochrome P450 (CYP), primarily by CYP2E1 [11-13], to the
93 toxic metabolite N-acetyl-p-benzoquinone imine (NAPQI) [9, 14-16]. At
94 therapeutic doses, NAPQI is immediately inactivated by conjugation with
95 glutathione. However, without this detoxification route, NAPQI binds
96 covalently to cellular proteins and forms toxic protein adducts, such as
97 protein-derived acetaminophen-cysteine (APAP-cysteine). These protein
98 adducts may cause mitochondrial dysfunction and early oxidant stress [17-
99 19]. Consequently, hepatotoxicity can be caused by liver cell necrosis [3].
100 Although it was thought that depletion of 70% of total liver glutathione
101 would be necessary for NAPQI to begin with protein binding [20], protein-
102 derived APAP-cysteine was detected in serum from human after therapeutic
103 doses [21]. It is likely that either a threshold of protein-derived APAP-
104 cysteine needs to be exceeded for the development of toxicity or that
105 specific binding targets are spared at therapeutic doses [6].

106 Currently, acetaminophen concentrations have only been considered
107 important to measure for patients who are suspected of intake of a toxic
108 amount and for patients who show a decreased hepatic function. In that
109 case, acetaminophen is mostly analyzed with an immunoassay, not
110 measuring metabolites, despite their key role in acetaminophen
111 hepatotoxicity. Considering acetaminophen-metabolic routes, further
112 investigation for associations between exposure to acetaminophen
113 metabolites and toxicity is warranted, as well as exposure in specific

114 populations. In the case of acetaminophen intoxication, extended
115 knowledge of metabolite formation will assist in optimizing (antidote)
116 treatment. This also applies to intoxication upon chronic use of high
117 acetaminophen dosages. Currently, there remains a knowledge gap
118 regarding the optimal treatment with N-acetylcysteine infusion to prevent
119 or treat hepatotoxicity. These new insights suggest Therapeutic Drug
120 Monitoring (TDM) of metabolites in case of toxicity, or as part of standard
121 clinical care in certain populations. Ultimately, monitoring of APAP-
122 metabolite concentrations may prevent or reduce toxicity and optimize
123 therapy.

124 We developed and validated an analytical method for simultaneous
125 quantification of APAP, APAP-glucuronide, APAP-sulfate, APAP-cysteine,
126 APAP-glutathione, APAP-mercapturate, and protein-derived APAP-cysteine
127 in a minimal volume of human plasma by ultra-performance liquid
128 chromatography-electrospray ionization-tandem mass spectrometry (UPLC-
129 MS/MS), preceded by an easy sample preparation. We aimed to optimize
130 the sensitivity of the assay to minimize the required sample volume, which
131 allows measurement of the smallest volume samples from preterm infants.

132

133 **MATERIALS AND METHODS**

134 **Chemicals and reagents**

135 APAP, APAP-sulfate, APAP-mercapturate, APAP-glucuronide, and
136 APAP-cysteine solution were purchased from Santa Cruz Biotechnology

137 (Heidelberg, Germany). APAP-glutathione was obtained from Toronto
138 Research Chemicals (Eching, Germany) and APAP-D4 solution from Sigma
139 Aldrich Cerilliant (Zwijndrecht, The Netherlands). Methanol absolute LC-MS
140 grade and formic acid 99% ULC/MS grade were purchased from Biosolve BV
141 (Valkenswaard, The Netherlands). Water was purified by using a MilliPore
142 Advantage A10 system. External quality control samples for acetaminophen
143 were purchased from *Stichting Kwaliteitsbewaking Klinische*
144 *Geneesmiddelanalyse en Toxicologie* (KKG, The Hague, The Netherlands)
145 and Santa Cruz Biotechnology (Heidelberg, Germany).

146

147 **Stock solutions, calibrators, quality control samples, and internal standard**

148 Stock solutions of APAP, APAP-sulfate, APAP-glucuronide, and APAP-
149 cysteine were prepared at a concentration of 500 mg/L using methanol,
150 while stock solutions of APAP-mercapturate and APAP-glutathione were
151 prepared at a concentration of 100 mg/L using methanol. For each analyte,
152 two separate stock solutions were prepared with the same concentration,
153 for both calibration of standard samples and for QC samples. Stock solutions
154 were stored at -20°C .

155 The working solution, calibrator 8 (50 mg/L), was prepared by drying
156 500 μL of APAP, APAP-sulfate, APAP-glucuronide, and APAP-cysteine, and
157 2500 μL of APAP-mercapturate and APAP-glutathione in one glass tube at 40
158 $^{\circ}\text{C}$ under nitrogen flow until all methanol was evaporated. Subsequently, all
159 analytes were reconstituted in 5-mL human plasma and mixed for 30

160 seconds. Calibrators 1 through 7 (0.05–25 mg/L) and the lower limit of
161 quantification (LLOQ) standard (0.01 mg/L) were prepared by diluting
162 calibrator 8 with human plasma. Quality control (QC) samples were
163 prepared the same way, using the other stock solution. The working solution
164 was diluted with human plasma to get three concentrations: QC Low (0.20
165 mg/L), QC Medium (1.5 mg/L), and QC High (15 mg/L). Then, calibrators and
166 QC samples were transferred in 10- μ L portions to 1.5-mL tubes (Eppendorf)
167 and stored at -80 °C awaiting analysis.

168 The internal standard (IS) was APAP-D4. A working solution of the
169 internal standard was prepared in methanol at a concentration of 100 μ g/L
170 APAP-D4.

171

172 **Specimens**

173 Human blank plasma was obtained from the blood transfusion
174 laboratory of the Erasmus Medical Center Rotterdam. Because
175 acetaminophen is a regularly used drug, acetaminophen-free blood was
176 collected from volunteers. Blood was centrifuged to separate plasma from
177 the red blood cells. Plasma was pooled and collected in smaller tubes. These
178 tubes were stored at -20 °C awaiting analysis.

179

180 **Sample preparation**

181 All calibrators, QC samples, blank and patient samples were thawed at
182 least half an hour prior to preparation. Then, to 10 μ L of each standard and

183 sample, 40 μL of internal standard solution was added for protein
184 precipitation. The samples were mixed for 15 seconds and then centrifuged
185 for 5 minutes at 16000 $\times g$. Of about 30 μL of supernatant was taken from
186 each sample and transferred to amber auto sampler insert vials (VWR).
187 Next, 140 μL of 0.1% aqueous formic acid was added and the samples were
188 mixed for 15 seconds. The ratio of the aqueous and organic solvent in the
189 sample matched the ratio in the mobile phase at start of the gradient. The
190 blank sample, without internal standard, was prepared by adding 40 μL of
191 methanol instead of internal standard solution. For acetaminophen,
192 acetaminophen-D4, and APAP-cysteine, 4 μL of sample was injected into the
193 UPLC-MS/MS apparatus. For all the other analytes, 10 μL was injected
194 because of the lower sensitivity for these analytes.

195

196 *Protein-derived APAP-cysteine*

197 For quantification of protein-derived APAP-cysteine in patient
198 samples, the sample preparation was preceded with one extra step after
199 thawing the sample: The protein-bound fraction was removed by filtration
200 of an extra 130- μL plasma through an Amicon Ultra-0.5 Centrifugal Filter
201 Unit with Ultracel-10 membrane (Merck Chemicals, Amsterdam, The
202 Netherlands) and discarded afterward, in order to collect 10- μL plasma with
203 unbound APAP-cysteine. The concentration of protein-derived APAP-
204 cysteine is determined by calculating the difference in APAP-cysteine
205 concentrations before and after filtration. The sample was further prepared

206 as described for all other analytes, continuing with the addition of 40 μ L of
207 internal standard.

208

209 **Instrumentation**

210 The equipment used was a Dionex Ultimate UPLC system consisting of
211 an Ultimate 3000 RS UPLC pump, an Ultimate 3000 RS autosampler and an
212 Ultimate 3000 RS Column Compartment. The UPLC was connected to a triple
213 quadrupole Thermo TSQ Vantage MS with HESI probe (Thermo Scientific).
214 The software programs Chromeleon (version 6.8, Dionex, Thermo Scientific),
215 Xcalibur (version 2.1, Thermo Scientific), and LCquan (version 2.6, Thermo
216 Scientific) were used to control the system and analyze the data.

217

218 **UPLC conditions**

219 Chromatographic separation, based on affinity of the analytes with the
220 nonpolar stationary phase, was achieved with a reversed-phase Acquity
221 UPLC HSS T3 column (1.8 μ m, 2.1 \times 100 mm; High Strength Silica with a
222 bound trifunctional C18 alkyl phase). Gradient elution was performed with a
223 mobile phase consisting of 1 mL of a 154 mg/L solution of ammonium
224 acetate in formic acid (99%) in 1 L of Milli-Q ultrapure water (eluent A) and 1
225 mL of the same solution in 1 L of methanol (eluent B). Prior to the analysis,
226 the system was equilibrated at the starting conditions of 86% eluent A and
227 14% eluent B until pressure was stable. The multistep gradient was as
228 follows: from 0 to 0.8 minutes, eluent B was increased from 14% to 28%;

229 from 0.8 to 1.0 minute, eluent B was increased to 95%; from 1.0 to 2.0
230 minutes, eluent B was kept stable at 95%; from 2.0 to 2.2 minutes, eluent B
231 was decreased to 14%; from 2.2 to 5.3 minutes, eluent B was kept stable at
232 14%. The run ended at 5.3 minutes at starting conditions. The flow was kept
233 at 0.400 mL/minute during the entire runtime. The temperature for the
234 column oven was set at 40 °C and for the autosampler at 15 °C. In order to
235 quantify all analytes, a volume of 4 µL as well as 10 µL is injected, which
236 requires two runs per sample and therefore doubles the runtime to 10.6
237 minutes. For the quantification of protein-derived APAP-cysteine, a third run
238 is required.

239

240 **MS/MS conditions**

241 For the detection and quantification of acetaminophen and
242 metabolites, settings of the MS/MS were as follows: MS runtime of 4.5
243 minutes, experiment type was Selected Reaction Monitoring (SRM),
244 ionization at ESI+, spray voltage of 4000 V, vaporizer temperature at 375 °C,
245 sheath gas pressure with nitrogen at 50 psi, auxiliary gas pressure with
246 nitrogen at 20 psi, capillary temperature at 250 °C, and collision pressure at
247 1.5 mTorr. All other settings were specific for each analyte and were
248 determined by infusion experiments with academic solutions of each
249 analyte of 1 mg/L. The chosen transitions and settings are shown in Table 1.

250

251 **Assay validation**

252 Validation of the method was performed according to the US Food
253 and Drug Administration (2001) guidelines for bioanalytical methods [22].
254 The following validation parameters were investigated.

255

256 *Linearity*

257 The relation between the concentration of the calibrators and
258 response (ratio of peak areas of the analytes and the internal standard) was
259 tested with a calibration curve. This curve should be linear across the range
260 from 0.05 up to 50 mg/L. To make the calibration curve, eight calibrators
261 were prepared and analyzed. Linear least square regression was used to
262 analyze the data. It was decided to apply weighting $1/x$, which means that
263 calibrators with the lowest concentrations are more important for the
264 calibration line than calibrators with highest concentrations [23]. The
265 relative standard deviation (RSD) was required to be lower than 15%, and
266 the correlation coefficient (r) together with the determination coefficient
267 (r^2) were required to be at least 0.9950 and 0.9900, respectively.

268

269 *LLOQ and ULOQ*

270 The LLOQ was measured by analyzing the LLOQ calibrator (0.01 or 0.05
271 mg/L) six times in a row. Mean and standard deviation of the response
272 ratios of the six samples were measured. Imprecision and accuracy were

273 calculated and should be $\leq 20\%$ and between 80% and 120%, respectively.
274 The highest calibrator of the calibration curve was used as upper limit of
275 quantification (ULOQ).

276

277 *Accuracy*

278 Accuracy was measured by analyzing three QC concentrations ($n = 6$
279 for each concentration). The percentage deviation between measured
280 concentration and theoretical concentration was calculated, and should be
281 lower than 15%.

282

283 *Repeatability and reproducibility*

284 Repeatability was tested by analyzing three QC concentrations in six-
285 fold on the same day. For each concentration, mean and RSD were
286 calculated. Reproducibility was tested by analyzing each QC concentration in
287 duplicate on six different days. The mean response of the 12 concentrations
288 for each sample with their RSD was calculated. For both tests, RSD was
289 required to be lower than 15%.

290

291 *Stability*

292 In-process stability was determined by storing QC samples of three
293 concentrations ($n = 2$ per concentration) at 6 °C prior to preparation for 24
294 and 48 hours. Autosampler stability was determined by storing QC samples
295 ($n = 2$ per concentration) after sample preparation in the autosampler for

296 24, 48, and 72 hours. Response ratios were measured and compared with
297 response ratios of samples kept at -80°C prior to preparation. After sample
298 preparation, samples were directly analyzed. Recovery was required to be
299 between 90% and 110%.

300

301 *Matrix effect and recovery*

302 It is important to measure matrix effects and absolute recoveries in
303 the development of an LC-MS/MS method since ion suppression and ion
304 enhancement effects can be expected owing to interferences by matrix
305 compounds, stable-isotope-labeled internal standards and co-eluting
306 compounds [24]. In order to check whether the precision, the
307 reproducibility, and the stability of the concentration-signal ratio are
308 affected by interference of the matrix analytes, the method described by
309 Matuszewski et al. (2003) was used [25]. Five different lots of human plasma
310 were used. To two QC concentration levels (QC low and QC high) and a
311 blank sample (all three in duplicate), analytes were added before extraction.
312 The same set of QCs and blanks was prepared with the analytes added after
313 extraction. Also, a set of six samples was evaluated with only Milli-Q
314 ultrapure water instead of plasma. Matrix effects were calculated as follows:
315 $(\text{peak area of analyte spiked after extraction}) / (\text{peak area of analyte}$
316 $\text{prepared in Milli-Q ultrapure water}) \times 100\%$.

317 The recovery was calculated as the percentage ratio of the area of the
318 analytes spiked before extraction and the ones prepared in Milli-Q ultrapure

319 water. The mean and RSD were calculated of both matrix effects and
320 recovery. In the ideal situation, the mean matrix effects and recovery are
321 between 80% and 120% and the RSD of both parameters is $\leq 15\%$.
322 Furthermore, for each analyte, the IS-normalized matrix effect should also
323 be calculated by dividing the matrix effect of the analyte by the matrix effect
324 of the IS. The RSD of the IS-normalized matrix effect calculated from the
325 different lots of matrix should not be greater than 15%.

326

327 **Application to pediatric pharmacokinetic samples**

328 For the validation of the assay for clinical practice and research
329 purposes, the method has been applied to quantify acetaminophen and its
330 metabolites in plasma of children participating in a pediatric clinical study.
331 This observational prospective study was performed at the Department of
332 Anaesthesia and Intensive Care Medicine of Our Lady's Children's Hospital,
333 Dublin, Ireland, between January and November 2012. Children (with and
334 without Down's Syndrome) routinely received acetaminophen post-cardiac
335 surgery in a dose of 7.7 mg/kg for children below 10 kg bodyweight, and 15
336 mg/kg for children above 10 kg bodyweight. The study protocol was
337 approved by the local ethics committee.

338

339

340

341

342 **RESULTS**

343 **Linearity**

344 Linearity was achieved for each analyte in the range between the
345 LLOQ and the ULOQ (Table 2), with all RSDs to be lower than 15% and the
346 determination coefficient (r^2) to be 0.998 at the lowest. APAP was linear
347 from calibrator 1 up to and including 7; APAP-mercapturate and APAP-
348 cysteine from calibrator 1 up to and including 6; APAP-sulfate from
349 calibrator 2 up to and including 8; APAP-glucuronide, APAP-glutathione from
350 calibrator 1 up to and including 8. Quantification performance of protein-
351 derived APAP-cysteine is subject to those of APAP-cysteine, with the step of
352 filtration being the single difference.

353

354 **LLOQ and ULOQ**

355 The results of determination of LLOQ and ULOQ are shown in Table 2.

356

357 **Accuracy, repeatability, and reproducibility**

358 The accuracy, repeatability, and reproducibility data all met the
359 requirement of being less than 15%, except for the APAP-cysteine accuracy
360 of 30.9% (error of measurement) for QC high. The results are shown in Table
361 2.

362

363

364

365 **Stability**

366 Except for APAP-glutathione, the recovery of all QCs was between 90%
367 and 110%, indicating that they were stable for at least 48 hours when stored
368 at 6 °C. APAP-glutathione was only stable for 24 hours. All prepared QCs
369 were stable for at least 48 hours when kept in the autosampler. The effect
370 of drying showed no significant difference between the dried and non-dried
371 standard (Mann Whitney test; $p < 0.05$).

372

373 **Matrix effect and recovery**

374 Matrix effects and absolute recoveries in the development of the LC-
375 MS/MS method are shown in Table 3. The test of Matuszewska showed
376 that APAP, APAP-cysteine, and APAP-sulfate experienced neither matrix
377 effect nor an effect from the sample preparation. Concerning APAP-
378 glucuronide though, 191.2% matrix effect indicates ion enhancement. On
379 the other hand, for APAP-mercapturate, ion suppression was observed;
380 matrix effect was 72.0%. A good recovery was achieved for all analytes,
381 except for APAP-glutathione with mean 18.6%.

382

383 **Application to pediatric pharmacokinetic samples**

384 A total of 162 post-dose samples were collected from a pediatric
385 cohort ($n = 30$), consisting of children with Down's Syndrome ($n = 17$) and
386 without Down's Syndrome ($n = 13$) (data unpublished); median age at

387 surgery was 176 days (range 92-944), median weight at cardiac surgery 6.1
388 kg (4-12.9).

389 For APAP, APAP-cysteine, APAP-glucuronide, and APAP-sulfate, only 1
390 of the 162 (0.6%) samples was measured below LLOQ. For APAP-
391 mercapturate and APAP-glutathione, this was the case in 5 (3.1%) and 161
392 (99.4%) samples, respectively. None of the analytes was measured above
393 the ULOQ in these samples. Due to the small sample volume, it was not
394 possible to differentiate between protein bound and unbound APAP-
395 cysteine.

396

397 **DISCUSSION**

398 We have validated an UPLC-MS method for the quantification of
399 acetaminophen and its metabolites according to US Food and Drug
400 Administration guidelines, with an easy sample preparation, short runtime,
401 and minimal injection volume. Therefore, the assay is very suitable for TDM.
402 The metabolites incorporated in this method are APAP-glucuronide, APAP-
403 sulfate, APAP-glutathione, APAP-cysteine, APAP-mercapturate, and protein-
404 derived APAP-cysteine. Prior reported methods for the quantification of
405 acetaminophen and metabolites in human plasma contained few
406 metabolites, mostly acetaminophen sulfate and/or glucuronide [26-28].
407 Assays with more metabolites were prior validated in animal matrices [29],
408 although Cook et al. recently published a method in human plasma and
409 urine that comes close to the performance of our assay [30]. Our assay is

410 distinguished by a shorter total runtime per injection of 5.3 minutes versus
411 20 minutes, easier sample preparation, and the ability to quantify the toxic
412 metabolite protein-derived APAP-cysteine.

413 Our assay fulfilled the desired criteria for accuracy, repeatability, and
414 reproducibility, except for the 30.9% accuracy of QC high of APAP-cysteine.
415 This QC high concentration of APAP-cysteine was outside the linear range.
416 The overestimation of APAP-cysteine could be caused by transformation
417 from the instable APAP-glutathione. At therapeutic doses, the
418 acetaminophen ULOQ is generally not exceeded, although it may be for
419 toxicology purposes. The ranges for linearity for all other analytes were
420 perfectly suitable for clinical pharmacology and toxicology.

421 Relevant matrix effects were measured for APAP-glucuronide during
422 the experiment, resulting in an increased process efficiency. This is in line
423 with the general problematic behavior of glucuronide-metabolites in LC-
424 MS/MS analyses, due to their susceptibility to interferences from the co-
425 eluting matrix analytes [31]. On the other hand, for APAP-mercapturate,
426 matrix effects lead to observed ion suppression. The coefficients of variation
427 of APAP-mercapturate in the samples spiked before extraction are 5.8% and
428 4.7% for QC-L and QC-H, respectively. Therefore, the effect of the matrix can
429 be considered acceptable. For all other analytes, no matrix effects were
430 measured, which indicates the absence of interferences by matrix
431 compounds, stable-isotope-labeled internal standards, and co-eluting
432 compounds, that may affect ion suppression and ion enhancement. The

433 matrix effect of plasma is relevant, although it does not influence
434 quantification as all analytes and calibrators are prepared in the same
435 plasma-matrix and are subject to influence to the same extent.

436 Except for APAP-glutathione, the stability of all analytes was good,
437 which means they were stable for at least 48 hours when stored at 6 °C.
438 APAP-glutathione was only stable for 24 hours at 6 °C, and therefore the
439 measurement or storage of the plasma sample in a freezer should be aimed
440 for within 24 hours. This instability has also been reported by Cook et al.
441 [30]. Hydrolysis of APAP-glutathione quickly transforms APAP-glutathione to
442 APAP-cysteine, presumably by gamma-glutamyl transferase and
443 dipeptidases. This may lead to an undervaluation of the actual concentration
444 APAP-glutathione at the time of sample collection and may lead to an
445 increased APAP-cysteine concentration. This instability of APAP-glutathione,
446 where APAP-cysteine is formed from APAP-glutathione, may also be
447 responsible for the increased recovery and process efficiency of APAP-
448 cysteine and the lower recovery and process efficiency of APAP-glutathione.
449 For future research, the addition of peptidase inhibitors during sample
450 collection could prevent or reduce this degradation.

451 The assay was successfully validated for clinical practice and research
452 purposes, quantifying acetaminophen and its metabolites in 162 plasma
453 samples from children. APAP-glutathione could only be quantified in one
454 sample, as a result of rapid conversion into APAP-cysteine. This confirms the
455 relevance of the addition of a peptidase inhibitor during sample collection.

456 APAP-mercapturate could not yet be detected in five samples, which were
457 all the first to be collected post dose, as APAP-mercapturate is the last
458 metabolite to be formed. For one sample, which was drawn 4 minutes after
459 the dose, all analytes were below LLOQ. Since acetaminophen was not yet
460 detectable at that time, metabolites could not have been formed either. In
461 conclusion, the assay is performing well for samples in clinical practice.

462 Quantification of APAP-glutathione during therapeutic as well as toxic
463 dosages of acetaminophen may be relevant, as it plays a crucial role in the
464 formation of toxic metabolites, although quantification of in vivo APAP-
465 glutathione levels has only been reported in animals yet [29]. Normally, the
466 reactive metabolite NAPQI is quickly detoxified by conjugation with
467 glutathione and further converted to the cysteine conjugate before it is
468 acetylated to form APAP-mercapturate. However, when the formation of
469 the reactive metabolite exceeds the glutathione-conjugation capacity of the
470 liver, covalent binding of NAPQI to cellular macromolecules may result,
471 which initiates the events ultimately leading to cytotoxicity.

472 Nevertheless, protein-derived APAP-cysteine can be measured with
473 our assay if 130 μ L of plasma sample is available. Generally, this allows the
474 quantification in adults and older infants, but not in neonates. Protein-
475 derived APAP-cysteine is mostly present in hepatocytes and is directly
476 related to toxicity and detectable in serum at therapeutic doses [6]. The
477 interpretation for the treatment or toxicity still remains to be investigated.

478 More research is needed on the toxic effects and characteristics of
479 acetaminophen metabolites in specific populations where different
480 metabolism may be expected. These may include patients with anorexia,
481 patients from different ethnic backgrounds, elderly patients, pregnant
482 women and their fetuses [7], obese adults [5], preterm infants [32-34],
483 patients with possible pharmacokinetic interactions on CYP1A2, and patients
484 subjected to repeated administration of acetaminophen leading to induced
485 CYP enzymes. Repeated administration of acetaminophen at a subtoxic dose
486 may result in an induction of hepatic CYP enzymes CYP2E1, CYP3A, and
487 CYP1A [35].

488 Generally, for toxicology purposes, acetaminophen concentrations
489 have only been considered important to measure for patients who are
490 suspected for intake of a toxic amount, not its metabolites. Extended
491 knowledge is required about metabolite formation following intoxication to
492 optimize treatment by infusion of the antidote, N-acetylcysteine. TDM of
493 metabolites may be indicated in case of toxicity, or as part of standard
494 clinical care in certain populations where metabolites may be used as a
495 marker for suspected liver injury.

497 **CONCLUSION**

498 We have developed a method for the simultaneous quantification of
499 APAP, APAP-glucuronide, APAP-sulfate, APAP-cysteine, APAP-glutathione,
500 APAP-mercapturate, and protein-derived APAP-cysteine in human plasma,

501 which greatly facilitates further research into acetaminophen and
502 metabolites, as well as for TDM purposes, even in the smallest plasma
503 volumes obtained from preterm infants.

504

505

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617 **FIGURE LEGENDS**

618 **Figure 1. Metabolic pathway of acetaminophen**

619 UDP: Uridine 5'-diphospho-glucuronosyltransferase

1 **Tables**

2 **Table 1. Analyte-specific settings**

3 The bold printed product ion mass-to-charge values were chosen.

Analyte	Parent ion (m/z)	Product ion (m/z)	ESI mode	Collision Energy (v)	S-Lens
APAP	152.169	110.16 93.13 65.13	+	15 23 29	77
APAP-D4 (IS)	156.191	114.19 97.16 69.17	+	15 22 30	77
APAP-D3-sulfate	235.017	155.06 113.10 68.10	+	26 37 31	77
APAP-sulfate	232.046	152.06 110.10 65.10	+	13 22 39	77
APAP-glucuronide	328.202	152.14 110.07 93.03	+	15 35 55	80
APAP-cysteine	271.155	182.08 140.07 96.07	+	12 26 37	76
APAP-mercapturate	313.176	208.10 166.10 140.050	+	16 27 31	77
APAP-glutathione	457.245	328.18 181.89 140.01	+	13 22 40	110

4 ESI: Electrospray ionization

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6

7 **Table 2. Validation results**

Analyte	QC	Accuracy [#] (%)	Repeatability RSD (%) <i>(within-run imprecision)</i>	Reproducibility RSD (%) <i>(between-run imprecision)</i>	LLOQ (mg/L)	ULOQ (mg/L)
APAP	QC L	-2.6	2.2	5.0	0.020	25.0
	QC M	4.7	2.4	5.4		
	QC H	1.9	2.0	6.1		
APAP-cysteine	QC L	-4.9	5.9	8.6	0.020	10.0
	QC M	-0.4	3.7	6.9		
	QC H	30.9	4.1	11.7		
APAP-glucuronide	QC L	1.5	5.1	5.7	0.047	47.0
	QC M	-6.4	3.8	5.4		
	QC H	4.6	10.4	12.4		
APAP-glutathione	QC L	-4.0	6.6	13.7	0.022	43.0
	QC M	7.2	3.4	6.4		
	QC H	-0.6	3.6	7.0		
APAP-mercapturate	QC L	2.2	3.6	4.8	0.010	15.0
	QC M	1.8	2.5	4.7		
	QC H	-2.3	3.4	5.2		
APAP-sulfate	QC L	-0.6	3.1	6.4	0.043	43.0
	QC M	0.8	3.4	4.4		
	QC H	2.2	3.1	6.2		

8 QC: Quality control

9 LLOQ: Lower limit of quantification

10 ULOQ: Upper limit of quantification

11 [#] The percentage for accuracy is the error of measurement.

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17 **Table 3. Matrix effect, recovery, and process efficiency**

Analyte	Matrix effect mean (%)	Recovery mean (%)	Process Efficiency mean (%)
APAP	90.3	108.2	97.7
APAP-cysteine	104.5	122.2	127.6
APAP-glucuronide	191.2	105.9	204.0
APAP-glutathione	81.4	18.6	16.2
APAP-mercapturate	72.0	140.3	96.4
APAP-sulfate	95.8	104.5	100.2

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ACCEPTED

