

**STATE OF THE ART OF UNBOUND CEFTRIAXONE AS A PHARMACODYNAMIC TOOL: ARE WE READY
FOR ITS IMPLEMENTATION IN CLINICAL PRACTICE?**

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Running head: Unbound ceftriaxone in clinical practice

SUPPLEMENTARY MATERIALS

- 1) Supplementary Material S1: Total (CEF_{tot}) and unbound (CEF_u) quantitation
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Supplementary Material S1: Total (CEF_{tot}) and unbound (CEF_u) quantitation

Briefly, 50 µL of serum (CEF_{tot}) or filtrate (CEF_u) was mixed with 100 µL of methanol spiked with internal standard [¹³C,²H₃]-Ceftriaxone (5 µg/mL). After centrifugation at 13 200 x g for 5 minutes, 100 µL of supernatant were collected for analysis. The chromatographic system consisted in a Prominence Shimadzu UFLC system (Shimadzu®, Prominence, Kyoto, Japan) in combination with a 4500QTRAP tandem mass spectrometer equipped with an electrospray ionization source operating in positive ion mode (Sciex®, Toronto, Canada). Quantitation was performed using the following precursor ion to product ion transitions: ceftriaxone m/z 554.8→395.8 / 554.8→323.8 (quantitation/qualification) and [¹³C,²H₃]-Ceftriaxone m/z 558.8→399.9. Chromatographic separation was performed on an Alltima HP C18 HL analytical column (3 µm particle size, 150 mm length × 3 mm inner diameter; VWR international®). The auto-sampler temperature was set at 8°C, the column oven at 30°C, the injected volume was 1 µL. The mobile phase consisted in 0.2% formic acid in pure Methanol (solvent A) and 2 mM ammonium formate with 0.2% formic acid in water (solvent B). The flow rate was 600 µL/min and the following elution gradient was performed: 70% B (0–0.1 min), 70–0% B (0.1–1.5 min), 0% B (1.5–3.5 min), 0–70% B (3.5–3.6 min), and 70% B (3.6–5.5 min). Calibration curves ranged from 1 to 20 µg/mL for serum ceftriaxone quantitation. Samples above the upper limit of quantification were 20-fold diluted in ceftriaxone-free serum. Chromatographic data acquisition and processing were performed using the Analyst v1.6.3 software (AB Sciex®). The analytical procedure used in this study met international analytical requirements according to ISO 15189, EMA and FDA guidelines on bioanalytical method validation [1,2]. The lower limit of quantification of the method was 1 µg/mL, the accuracy ranged from 85 to 115% and the intraday and interday precision were lower than 15%.

[1] EMA Guideline on Bioanalytical Method Validation. 2011.

[2] FDA Guidance for Industry Bioanalytical Method Validation. 2001.

Supplementary Table S1 : Comparison of between-model absolute and percentage error using ANOVA followed by Tukey Honest Significant Differences tests

Difference	Lower CI	Upper CI	Adjusted P-value	Error Type	Comparison
13.69	5.24	22.15	2.19e-05	Signed Error	Dreesen-Bos
15.80	7.35	24.26	3.56e-07	Signed Error	Gijzen-Bos
24.95	16.50	33.41	4.10e-10	Signed Error	Gregoire-Bos
18.45	10.00	26.91	1.41e-09	Signed Error	Hartmann-Bos
21.53	13.08	29.98	4.11e-10	Signed Error	Heffernan-Bos
9.56	1.10	18.01	1.37e-02	Signed Error	Leegwater-Bos
-10.46	-18.92	-2.01	4.09e-03	Signed Error	Standing-Bos
2.39	-6.07	10.84	9.94e-01	Signed Error	Ulldemolins-Bos
2.11	-6.34	10.56	9.97e-01	Signed Error	Gijzen-Dreesen
11.26	2.81	19.72	1.28e-03	Signed Error	Gregoire-Dreesen
4.76	-3.69	13.22	7.12e-01	Signed Error	Hartmann-Dreesen
7.84	-0.62	16.29	9.40e-02	Signed Error	Heffernan-Dreesen
-4.13	-12.59	4.32	8.45e-01	Signed Error	Leegwater-Dreesen
-24.16	-32.61	-15.70	4.10e-10	Signed Error	Standing-Dreesen
-11.30	-19.76	-2.85	1.20e-03	Signed Error	Ulldemolins-Dreesen
9.15	0.70	17.60	2.26e-02	Signed Error	Gregoire-Gijzen

Difference	Lower CI	Upper CI	Adjusted P-value	Error Type	Comparison
2.65	-5.80	11.10	9.88e-01	Signed Error	Hartmann-Gijsen
5.73	-2.73	14.18	4.68e-01	Signed Error	Heffernan-Gijsen
-6.24	-14.70	2.21	3.44e-01	Signed Error	Leegwater-Gijsen
-26.27	-34.72	-17.81	4.10e-10	Signed Error	Standing-Gijsen
-13.41	-21.87	-4.96	3.61e-05	Signed Error	Ulldemolins-Gijsen
-6.50	-14.95	1.95	2.89e-01	Signed Error	Hartmann-Gregoire
-3.42	-11.88	5.03	9.42e-01	Signed Error	Heffernan-Gregoire
-15.40	-23.85	-6.94	8.20e-07	Signed Error	Leegwater-Gregoire
-35.42	-43.87	-26.96	4.10e-10	Signed Error	Standing-Gregoire
-22.57	-31.02	-14.11	4.10e-10	Signed Error	Ulldemolins-Gregoire
3.08	-5.38	11.53	9.69e-01	Signed Error	Heffernan-Hartmann
-8.90	-17.35	-0.44	3.05e-02	Signed Error	Leegwater-Hartmann
-28.92	-37.37	-20.46	4.10e-10	Signed Error	Standing-Hartmann
-16.07	-24.52	-7.61	2.06e-07	Signed Error	Ulldemolins-Hartmann
-11.97	-20.43	-3.52	4.22e-04	Signed Error	Leegwater-Heffernan
-31.99	-40.45	-23.54	4.10e-10	Signed Error	Standing-Heffernan
-19.14	-27.60	-10.69	6.03e-10	Signed Error	Ulldemolins-Heffernan
-20.02	-28.48	-11.57	4.32e-10	Signed Error	Standing-Leegwater

Difference	Lower CI	Upper CI	Adjusted P-value	Error Type	Comparison
-7.17	-15.62	1.28	1.72e-01	Signed Error	Ulldemolins-Leegwater
12.85	4.40	21.31	9.71e-05	Signed Error	Ulldemolins-Standing
54.42	2.34	106.50	3.28e-02	Percentage Error	Dreesen-Bos
72.70	20.62	124.78	5.54e-04	Percentage Error	Gijssen-Bos
142.10	90.02	194.18	4.10e-10	Percentage Error	Gregoire-Bos
88.89	36.81	140.97	5.50e-06	Percentage Error	Hartmann-Bos
125.36	73.28	177.45	4.20e-10	Percentage Error	Heffernan-Bos
51.61	-0.47	103.69	5.43e-02	Percentage Error	Leegwater-Bos
-144.27	-196.35	-92.19	4.10e-10	Percentage Error	Standing-Bos
-95.14	-147.22	-43.06	7.43e-07	Percentage Error	Ulldemolins-Bos
18.29	-33.79	70.37	9.75e-01	Percentage Error	Gijssen-Dreesen
87.68	35.60	139.77	7.99e-06	Percentage Error	Gregoire-Dreesen
34.47	-17.61	86.56	5.01e-01	Percentage Error	Hartmann-Dreesen
70.95	18.87	123.03	8.66e-04	Percentage Error	Heffernan-Dreesen
-2.81	-54.89	49.27	1.00e+00	Percentage Error	Leegwater-Dreesen
-198.69	-250.77	-146.61	4.10e-10	Percentage Error	Standing-Dreesen
-149.56	-201.64	-97.48	4.10e-10	Percentage Error	Ulldemolins-Dreesen
69.40	17.32	121.48	1.27e-03	Percentage Error	Gregoire-Gijssen

Difference	Lower CI	Upper CI	Adjusted P-value	Error Type	Comparison
16.19	-35.89	68.27	9.89e-01	Percentage Error	Hartmann-Gijsen
52.66	0.58	104.74	4.51e-02	Percentage Error	Heffernan-Gijsen
-21.10	-73.18	30.98	9.42e-01	Percentage Error	Leegwater-Gijsen
-216.97	-269.06	-164.89	4.10e-10	Percentage Error	Standing-Gijsen
-167.85	-219.93	-115.77	4.10e-10	Percentage Error	Ulldemolins-Gijsen
-53.21	-105.29	-1.13	4.09e-02	Percentage Error	Hartmann-Gregoire
-16.74	-68.82	35.34	9.86e-01	Percentage Error	Heffernan-Gregoire
-90.50	-142.58	-38.41	3.33e-06	Percentage Error	Leegwater-Gregoire
-286.37	-338.45	-234.29	4.10e-10	Percentage Error	Standing-Gregoire
-237.25	-289.33	-185.16	4.10e-10	Percentage Error	Ulldemolins-Gregoire
36.47	-15.61	88.55	4.20e-01	Percentage Error	Heffernan-Hartmann
-37.29	-89.37	14.80	3.88e-01	Percentage Error	Leegwater-Hartmann
-233.16	-285.24	-181.08	4.10e-10	Percentage Error	Standing-Hartmann
-184.04	-236.12	-131.95	4.10e-10	Percentage Error	Ulldemolins-Hartmann
-73.76	-125.84	-21.68	4.21e-04	Percentage Error	Leegwater-Heffernan
-269.64	-321.72	-217.55	4.10e-10	Percentage Error	Standing-Heffernan
-220.51	-272.59	-168.43	4.10e-10	Percentage Error	Ulldemolins-Heffernan
-195.88	-247.96	-143.80	4.10e-10	Percentage Error	Standing-Leegwater

Difference	Lower CI	Upper CI	Adjusted P-value	Error Type	Comparison
-146.75	-198.83	-94.67	4.10e-10	Percentage Error	Ulldemolins-Leegwater
49.13	-2.95	101.21	8.22e-02	Percentage Error	Ulldemolins-Standing

Supplementary Figure S1: Bland-Altman plots for signed differences and relative differences for the 9 models

