1	Supporting Information
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3	Modeling the ionization efficiency of small molecules in positive electrospray
4	ionization using molecular dynamics simulations
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21	Contents: Text S1-S2, Figures S1-S10 and Tables S1-S2
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23 SUPPLEMENTARY TEXT

24	Text S1: Determining the number of H₃O ⁺ ions
25	We extracted the underlying data from the publication ¹ using WebPlotDigitizer and reconstructed
26	the figures from the extracted datapoints (Figure S2). The relationship between diameter and droplet
27	charge was described by fitting a power trendline of the form of:
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29	$y = ax^b$ (eq. S1)
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31	In place of diameter, we used the body diagonal of the 64 nm ³ cube, calculated from:
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33	$D = \sqrt{3}a$ (eq. S2)
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35	which was determined to be 6.9 nm. The number of calculated H_3O^+ molecules in a droplet with a
36	diameter of 6.9 nm was calculated to be 149 in a water droplet and 108 in a methanol droplet. Using
37	the average of these two values we estimated 129 H_3O^+ molecules for our system. In order to verify
38	whether this number is in agreement with other experimental observations and with electrochemical
39	theory, we applied the Nernst equation to calculate the output voltage of a 64 nm ³ droplet with 129
40	H_3O^+ molecules and compared that to the observations of Maze et al. ² For an electrochemical cell with
41	water, where the following reactions take place:
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43	$2H_2O + 2e^- \rightleftharpoons H_2 + 2OH^-$ (eq. S3)
44	$2H^+ + 2e^- \rightleftharpoons H_2$ (eq. S4)

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$$O_2 + 2H_2O + 4e^- \rightleftharpoons 4OH^- \quad (eq. S5)$$

S-2

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$$O_2 + 4H^+ + 4e^- \rightleftharpoons 2H_2O$$
 (eq. S6)

48 the Nernst equation^{3,4} for the half-reaction eq. S4 is as follows:

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 $E = E^{0} - \frac{0.059}{2} \log \frac{p(H_{2})}{\left[H^{+}\right]^{2}}$ (eq. S7)

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where, E is the cell potential (in our case the droplet potential), E⁰ is the standard reduction potential,
p is the pressure of the system.

55 For eq. S7, $E^0 = 0$,³ and for our experiment, p = 1 atm, hence, eq. S7 can be written as 56

$$E = -\frac{0.059}{2}\log\frac{1}{[H^+]^2}$$
 (eq. S8)

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59 Converting the number of 129 H⁺ in 64 nm³ to concentration we get 2 x 10^{24} H⁺/L, which when 60 converted to mol/L using Avogadro's number equals to 3.35 mol/L. Solving E in eq. S8 for $[H^+]$ 3.35 61 mol/L gives E = 0.031 V. Maze et al.² observed an output voltage of 0.5 V for a droplet with 4 µm 62 diameter. Considering that our system is substantially smaller and given that Smith et al.¹ observed a 63 decrease in charge with decreasing diameter, we concluded that our estimated number of 129 H₃O⁺ is 64 within a reasonable range.

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66 Text S2: Determining the number of H+ ions

One of the challenges we faced during the simulations was that when using a small number of H⁺ ions (e.g., 30), the ions moved towards the edges of the water droplet and did not sufficiently interact with the analyte that often remained in the center of the box (Figure S3 and S4). This may be hard to see in a 2D representation (Figure S3 and S4) but it can be clearly seen in the calculations of the Coulomb and Lennard-Jones interactions (Figure S5).

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73 Both System 1 and System 2 were neutralized with the addition of OH⁻ ions equal to the number of 74 H3O⁺ for System 1 and to the number of H⁺ for System 2. It should be noted that although, OH⁻ are not 75 expected to be at such high concentrations in ESI+, neutralizing the system is necessary in order to conduct the simulations. Not neutralizing the system results in an unstable system that crashes before 76 77 the simulation is completed. It should be clarified that although OH⁻ ions were added to the system, their interactions with the analyte were not included in the model for predicting RRF. It should also be 78 79 noted that the addition of OH⁻ ions does not influence the interactions of the analyte with the water, 80 methanol, H3O⁺ and H⁺ molecules. The interactions between the analyte and each group of molecules (e.g., water) are not dependent upon the interactions of the analyte and another group of molecules 81 82 (e.g., OH⁻).

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91 SUPPLEMENTARY FIGURES









96~ HPLC water with 0.1% methanol and 5 mM ammonium acetate. B: methanol with 10% HPLC water and

- 97 5 mM ammonium acetate.
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Figure S2: Reconstructed figures from the publication of Smith et al.¹ showing the relationship between
charge and droplet diameter for electrospray droplets created in positive electrospray ionization for
water and methanol. The trendline is shown in red. The number of calculated H₃O⁺ molecules in a
droplet with a diameter of 6.9 nm was calculated to be 149 in a water droplet and 108 in a methanol
droplet.



112 Figure S3: Caffeine in a solvent box among H^+ ions at t = 0 ps and at t = 1000 ps. The box contained 1

113 molecule of caffeine, 625 water molecules, 699 OH^- ions and 699 H^+ ions. The figure shows only the

114 caffeine molecule and the H^+ ions for simplicity.



Figure S4: Caffeine in a solvent box with H^+ ions at t = 0 ps and at t = 1000 ps. The box contains 1 molecule of caffeine, 30 H^+ ions, 2084 water molecules and 30 OH^- ions. The figure shows only the caffeine molecule and the H^+ ions for simplicity.





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121 Figure S5: Coulomb and Lennard-Jones interactions between caffeine and H⁺ ions during the

simulation. The top figures (A and B) show the Coulomb and Lennard-Jones interactions over time for a system with 699 H⁺ ions and the bottom figures show the interaction energies for a system with 30 H⁺ ions.



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132 ure S6: A: Relationship between the absolute errors (AE) of the model and retention time (RT). B:

133 Relationship between RT and RRF.





Figure S7: R² of the model and number of chemicals as a function of the penalty cutoff point for
including chemicals in the dataset. As the penalty cutoff point decreases so does the number of
chemicals included in the model. The R² of the model appears to be consistent with an increase around
300. At cutoff point 50 the model shows a high R² as a result of overparameterization given the small
number of chemicals remaining in at this cutoff point (n=16).



Figure S8: (A) 10-fold cross-validation showing the individual groups of compounds for each fold. The figure shows the results for each group of compounds when these compounds were not included in the training set. (B) Y-randomization for all the compounds in the dataset. R² is the coefficient of determination and MAE is the mean absolute error between the predictions and the experimental values.

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154 Figure S9: 10-fold cross-validation showing the results for the testing sets including the two outliers 155 that were excluded for being outside the applicability domain of the models that were used to predict 156 their log RRF. The figure shows the results for each group of compounds when these compounds were 157 not included in the training set.









Figure: S10: Experimental and predicted values of log transformed RRF. The diagonal lines show the 1-

163 to-1 agreement line, and the \pm 1 log unit deviation line. The figure shows the results of the model

including all the chemicals in the dataset and pointing the results for Furalaxyl and N,N-

Dibutylthiourea.

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175 SUPPLEMENTARY TABLES

176 Table S1: Statistics for the multi-parameter linear regression model that contained all compounds in

177 the dataset

All compounds

R ²	0.74					
	coefficient	standard error	t	p-value	[0.025	0.975]
const	2.51	0.278	9.049	0.000	1.95	3.078
U _{p1}	0.53	0.328	1.607	0.117	-0.139	1.193
LJ _{p2}	-6.04	1.509	-4.001	0.000	-9.101	-2.974
Ц _{р3}	5.24	1.407	3.725	0.001	2.384	8.096
LJ _{std}	3.47	0.883	3.934	0.000	1.681	5.267
Coul _{p1}	-0.01	0.002	-3.421	0.002	-0.009	-0.002
Coul _{p2}	0.02	0.01	2.195	0.035	0.002	0.043
Coul _{p3}	-0.02	0.01	-1.705	0.097	-0.036	0.003
Coul _{std}	-0.03	0.007	-4.995	0.000	-0.048	-0.02
W Ll _{p2}	0.18	0.054	3.318	0.002	0.07	0.289
W Ll _{p3}	-0.17	0.055	-3.145	0.003	-0.284	-0.061
W LJ _{std}	0.31	0.082	3.761	0.001	0.143	0.477
S _W	0.11	0.071	1.6	0.119	-0.031	0.258

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179 Table S2: Statistics for the multi-parameter linear regression model that contained compounds with a

180 penalty less than 300

Compounds with penalties less than 300

R ²	0.82					
	coefficient	standard error	t	p-value	[0.025	0.975]
const	2.51	0.262	9.607	0.000	1.979	3.049
LJ _{p1}	0.63	0.288	2.204	0.036	0.046	1.222
LJ _{p2}	-5.80	1.317	-4.406	0.000	-8.497	-3.109
LJ _{p3}	4.92	1.222	4.027	0.000	2.421	7.419
LJ _{std}	3.49	0.773	4.52	0.000	1.912	5.074
Coul _{p1}	-0.01	0.001	-4.072	0.000	-0.009	-0.003
Coul _{p2}	0.03	0.009	3.336	0.002	0.012	0.051
Coul _{p3}	-0.03	0.009	-2.779	0.009	-0.045	-0.007

Coul _{std}	-0.04	0.006	-6.235	0.000	-0.05	-0.025
W LJ _{p2}	0.21	0.052	4.025	0.000	0.103	0.317
W LJ _{p3}	-0.20	0.052	-3.843	0.001	-0.307	-0.094
W LJ _{std}	0.35	0.08	4.45	0.000	0.192	0.518
Sw	0.14	0.063	2.163	0.039	0.007	0.265

182 **REFERENCES**

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