## SUPPLEMENTARY INFORMATION

for

## "Using Orbitrap mass spectrometry to assess the isotopic compositions of individual components in mixtures"

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**Figure S1.** Comparison of Total Ion Current (TIC) profiles (gray icons) for mixtures in (a) ESI and (b) EI instruments. Points selected for analysis within the 'para-xylene-only' and 'para-xylene + n-decane' regions are shown in yellow in (b).



**Figure S2.** (a) Chromatogram from a pure para-xylene 'peak capture' experiment on the Q Exactive GC used in this study. Analytical and operating conditions were the same as those used in the para-xylene / n-decane experiments presented in the text. The full TIC trace is presented as gray triangles. Points selected for isotopic analysis (plotted in (b)) are shown in yellow. (b)  ${}^{13}C/{}^{12}C$  ratio of para-xylene over the course of the experiment, corresponding to the yellow points in (a). The red line represents the mean of a 5-minute moving window.



**Figure S3.** Comparison of mass spectra for the two nominally pure alanine solutions from the alanine / glutamate experiment set. (a) Alanine standard G1 with the isotopologue peaks of interest identified. (b) Alanine solution G2, which was prepared as a pure alanine standard, showing a higher relative abundance of contaminating species. (c) Zoomed-in portion of (a) illustrating the relative magnitude of the major peaks in the mass spectrum. (d) Zoomed-in portion of (b) showing the presence of the most abundant contaminant species.



**Figure S4. (a)** Chromatogram from the 11152017 para-xylene + n-decane experiment. Data from the xylene + decane portion of the chromatogram were binned into 2-minute intervals, shown here as regions 1–5. (b) Average  ${}^{13}C/{}^{12}C$  ratio of the analyte as a function of the xylene:decane mixing ratio (represented here as the absolute intensity of the  ${}^{12}C$  ion beam divided by the total ion current). The red line represents a linear regression to the five regions, weighted by errors in the  ${}^{13}C/{}^{12}C$  ratio of each point. The black circle is the calculated average of all five regions. Error bars represent two standard error on the  ${}^{13}C/{}^{12}C$  ratio and one standard deviation on the  ${}^{12}C$  Absolute Intensity / TIC ratio. Retention time as shown in (a) increases to the left in (b). For comparison, the  ${}^{13}C/{}^{12}C$  ratio for the C<sub>7</sub> fragment in the para-xylene-only portion of the same chromatogram is 0.07504 (Table 2) at a  ${}^{12}C/TIC$  value of 0.39 (Table 1).



Figure S5. Comparison of spectral noise on the <sup>12</sup>C- and <sup>13</sup>C-bearing isotopologue peaks from two para-xylene / n-decane experiments performed using the same solution and the same analytical conditions except resolution. Portions of the experiment during which only paraxylene was present in the mass analyzer (corresponding to the selected analysis time windows described in Section 2.3.1) are denoted by the yellow dashed lines and labeled 'xylene only'; the corresponding window for the para-xylene / n-decane mixture is denoted by the blue dashed lines and labeled 'xylene + decane'. (a) Noise (in ion intensities) on the  ${}^{13}$ C-bearing C<sub>7</sub> molecular ion from the 60,000 resolution experiment, showing the increase in noise when the n-decane 'contaminant' enters the mass analyzer. (b) Noise (in ion intensities) on the <sup>13</sup>Cbearing C<sub>7</sub> molecular ion from the 120,000 resolution experiment, showing the increase in noise when the n-decane 'contaminant' enters the mass analyzer. Note that the absolute magnitude of the peak noise on both the 'xylene only' and the 'xylene + decane' portions of the experiment is less than the corresponding noise magnitude in the 60,000 resolution experiment. (c) Noise ratio between the  ${}^{13}C$  peak and the  ${}^{12}C$  peak in the 60,000 resolution experiment showing greater noise on the minor (i.e., <sup>13</sup>C-bearing) species, yet an essentially constant noise ratio throughout the experiment regardless of whether para-xylene was alone in the mass analyzer or not. (d) Same as (c) but for the 120,000 resolution experiment.



**Table S1.** Plateau times and parameter values for empirical fits to the para-xylene / n-decane elution curves.

Identification	Start Time (min)	End Time (min)	С	ct	т	A	w
11142017_xylene	5.96	9.40	3.98E6	5.79	3.67E5	2.25E6	0.138
11142017_xylene+decane	9.77	20.02	7.38E6	9.57	1.92E5	4.50E5	0.138
11152017_xylene	6.14	9.63	2.22E7	6.01	5.04E6	1.79E7	0.115
11152017_xylene+decane	9.96	20.02	4.50E7	9.80	-1.05E6	6.74E6	0.133
01112018_xylene	5.78	9.48	6.55E7	5.61	1.06E6	5.58E7	0.128
01112018_xylene+decane	9.80	19.88	1.31E8	9.64	-2.30E6	1.95E7	0.131
01112018_60K_xylene	5.81	9.54	9.75E7	5.66	3.41E5	8.78E7	0.127
01112018 60K xylene+decane	9.91	20.01	1.93E8	9.73	-8.36E6	3.11E7	0.150