

Electronic Supplementary Information

Rapid Determination of Tumour Stroma Ratio in Squamous Cell Carcinomas with Desorption Electrospray Ionization Mass Spectrometry (DESI-MS): A Proof-Of-Concept Demonstration

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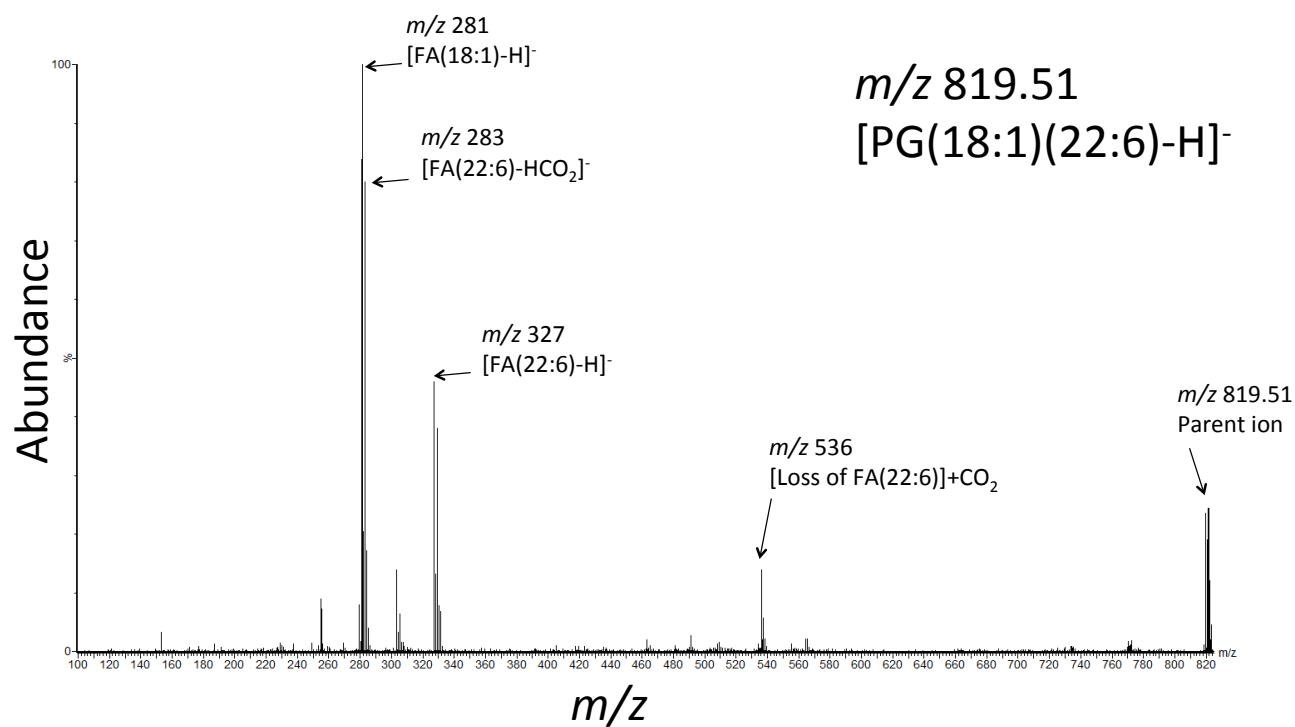
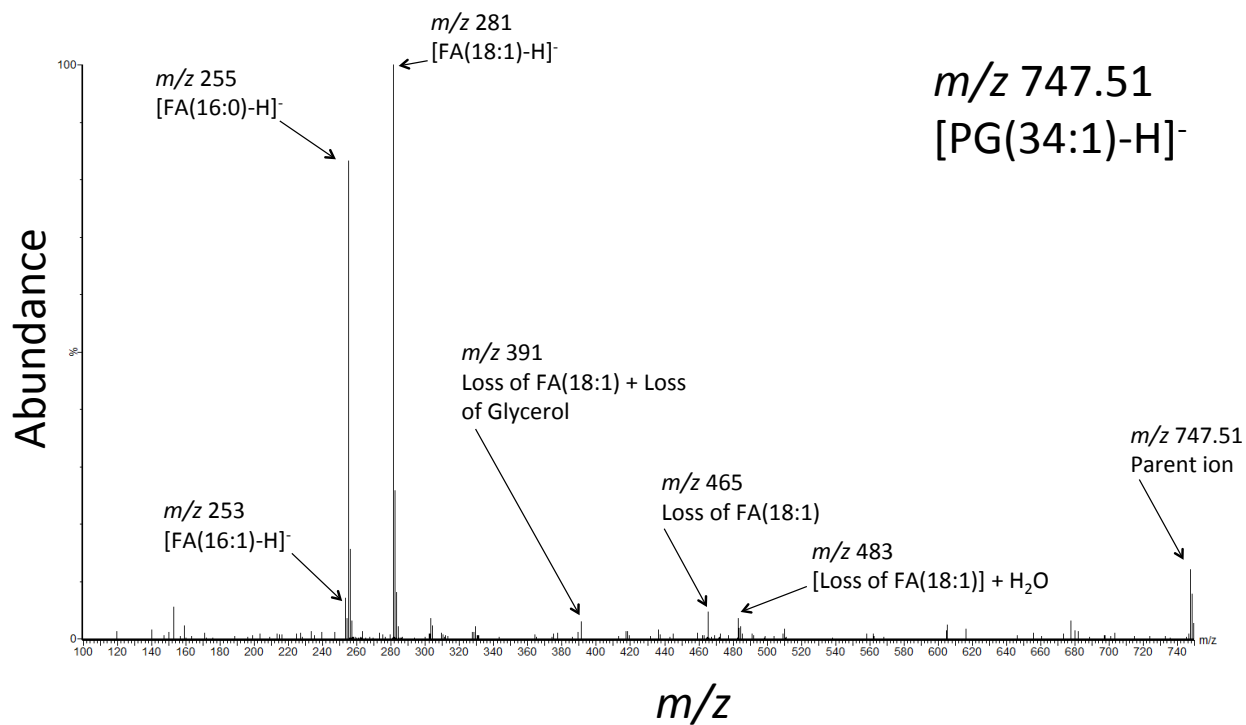
Table of Contents

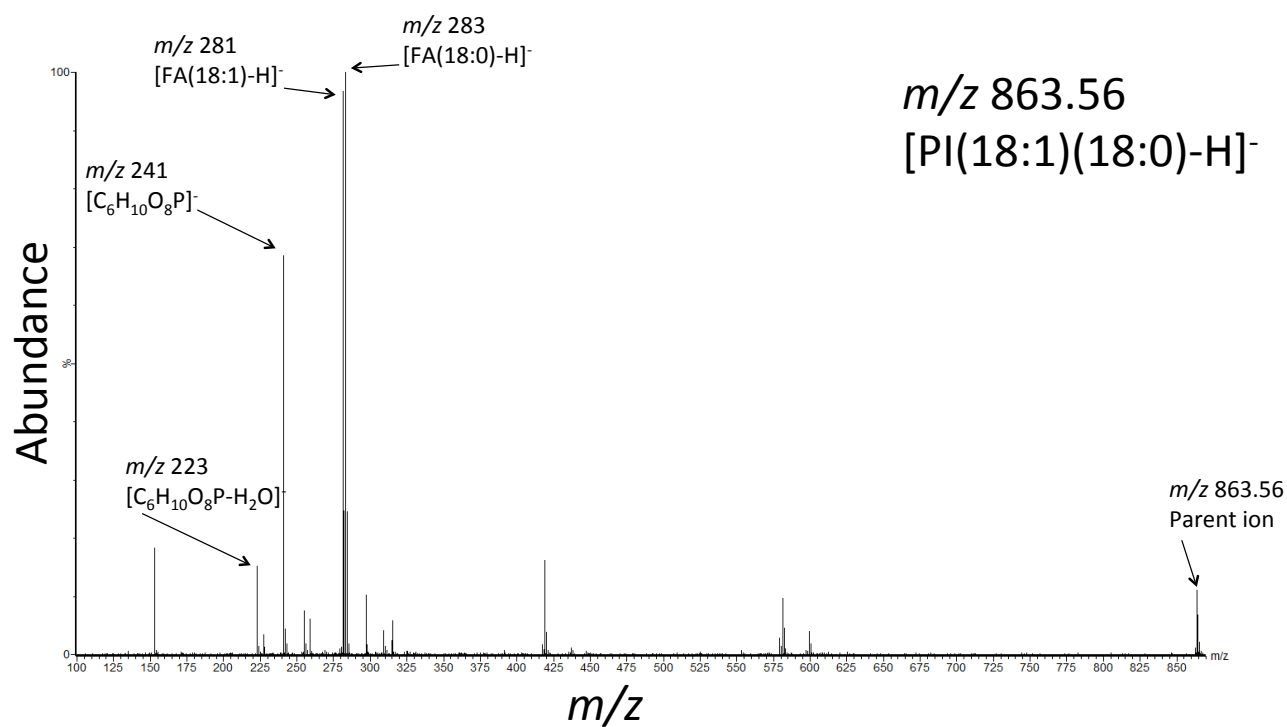
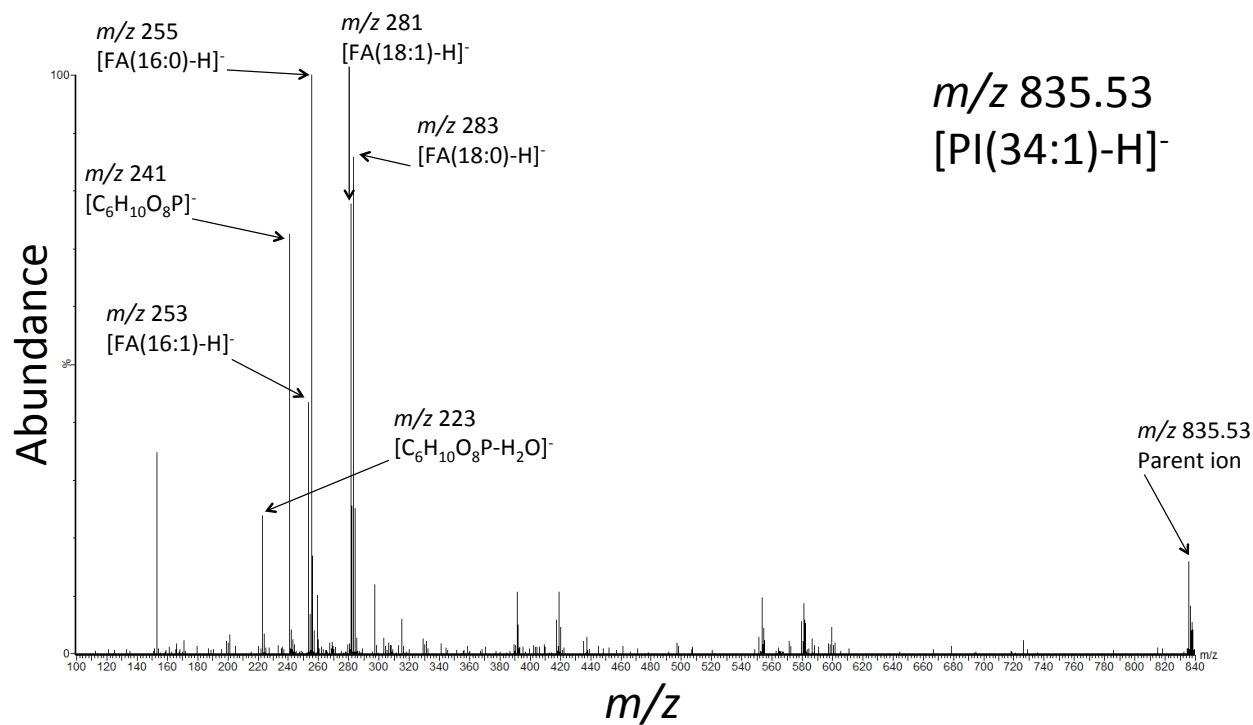
Table S1-----Page 3
Figure S1-----Pages 4-7
Figure S2-----Page 8
Figure S3-----Page 9

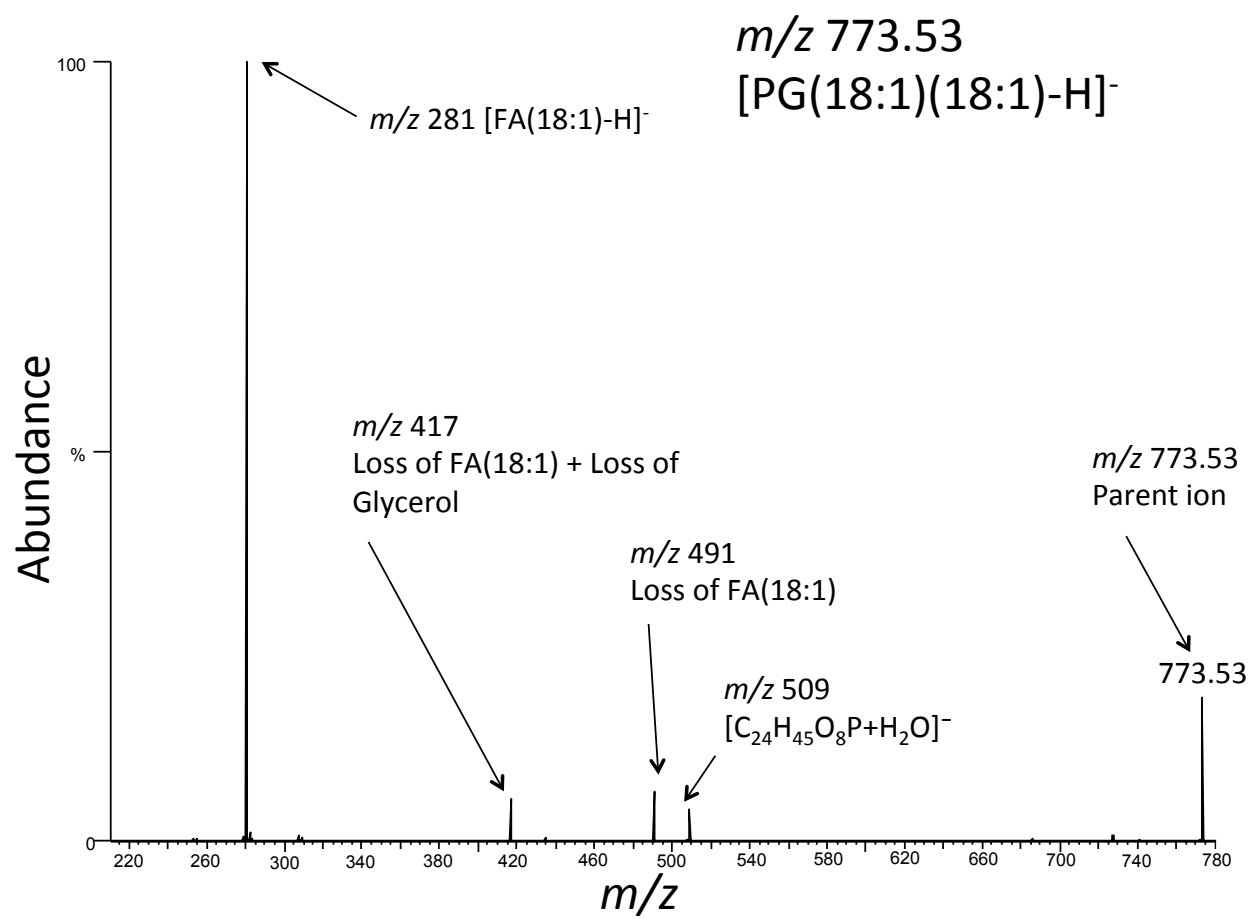
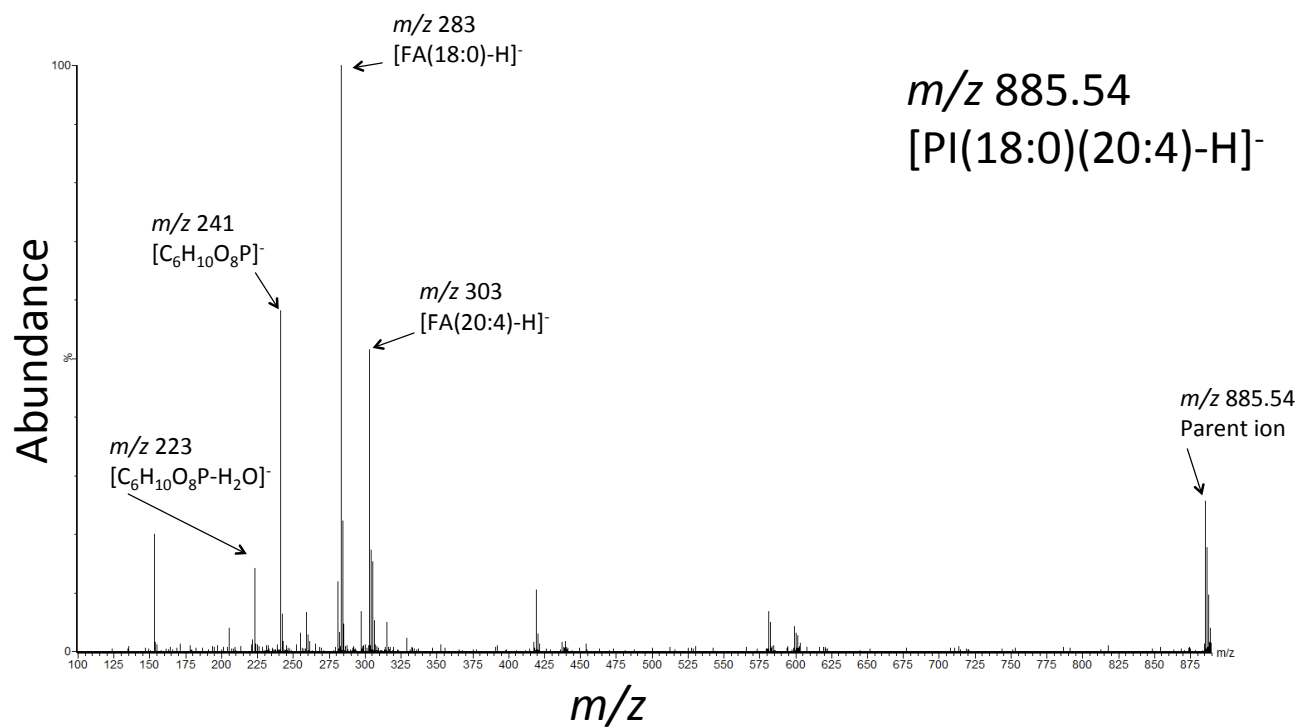
<i>m/z</i>	Intensity FaDu Extract	Intensity 1:1 FaDu CAF Mixture	
773.53	7.32 X 10 ⁵	2.04 X 10 ⁵	
835.53	2.69 X 10 ⁵	9.44 X 10 ⁴	
863.56	2.89 X 10 ⁵	8.84 X 10 ⁴	
TIC	2.25 X 10 ⁷	9.49 X 10 ⁶	
<i>m/z</i> Ratios	FaDu Extract	1:1 FaDu CAF Mixture	% Deviation in Ratios
863.56 / 835.53	1.07	0.94	13
773.53 / 863.56	2.54	2.31	10

Table S1. The abundance of FaDu biomarker ions in pure cancer cell lipid extract, and in 1:1 cancer stroma mixture extract. This table summarizes the intensity values obtained from representative spectrum of lipid extract show in Fig. 1. As detailed in the manuscript, minor ion suppression on the order of ~10% was seen affecting the relative abundance of cancer biomarker ions in mixed cancer, stroma sample. The total ion count (TIC) values are also given. The percent deviation in peak intensity ratios for FaDu markers between mixture and pure cancer cell extract, likely arising due to signal suppression, is also provided.

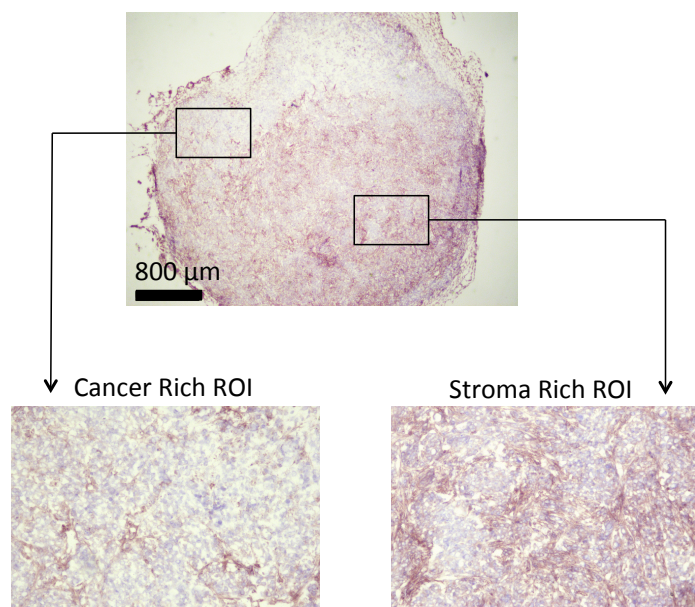
Figure S1. The MS/MS analysis of the biomarker ions described in this study. Below, we have included the ms/ms spectra that summarize the ion identity assignments presented in Table 1. The majority of ions were assigned using DESI-MS/MS analysis. The MS/MS analysis of the ion of m/z 773.53 [PG(18:1)(18:1)-H]⁻ was performed on LTQ-OrbiTrap system. The other MS/MS analyses used the TOF platform utilizing a DESI-MS ion source as described in the manuscript text.







The spatial distribution of cancer and stroma cells



The spatial distribution of cancer and stroma cells on the length scale of DESI-MS pixel

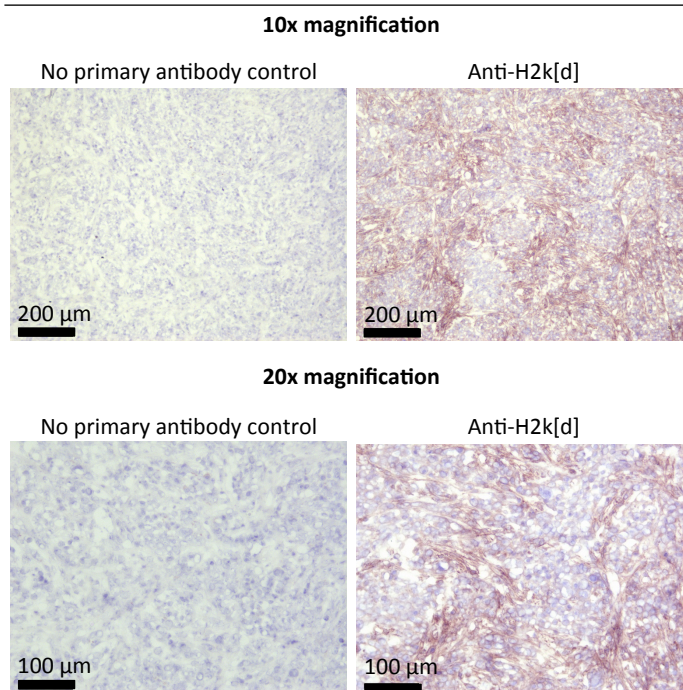


Figure S2. The Immunohistochemistry image of FaDu xenograft tumour slice. A 10 μm slice of FaDu xenograft tumour was subjected to immunostaining. This image illustrates the degree of mixing and infiltration between stroma and cancer cells in FaDu head & neck cancer model. The mixing between cancer and stroma cells is on the order of a single DESI-MS pixel. Both cancer rich and stroma rich ROIs contain cancer and stroma cells.

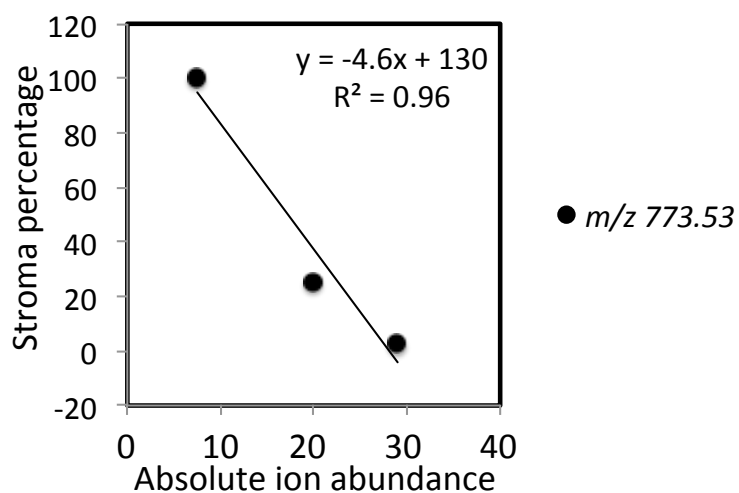
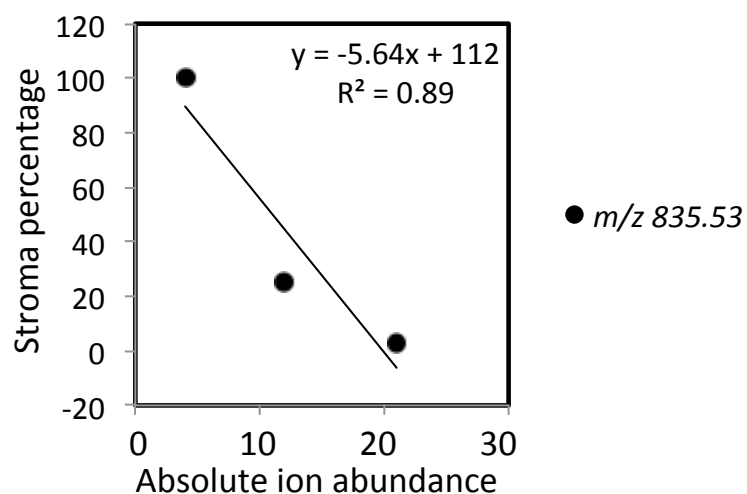
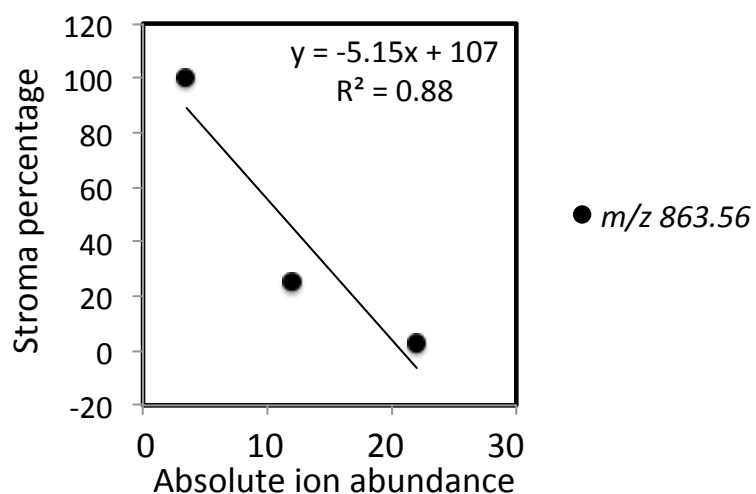


Figure S3. The quantitative concordance between stroma percentage and absolute values of FaDu biomarker ion abundance from lipid extract experiment, averaged from the repetitions shown in Fig. 5. A simple linear relationship between absolute ion abundance values and stroma percentage is assumed. The coefficient of determination R^2 values is provided for each of the FaDu markers listed in Table 1. With the caveat of limited data points, we used a liner fit model to predict stroma percentages of FaDu tumour slices from ROIs shown in Fig. 4.