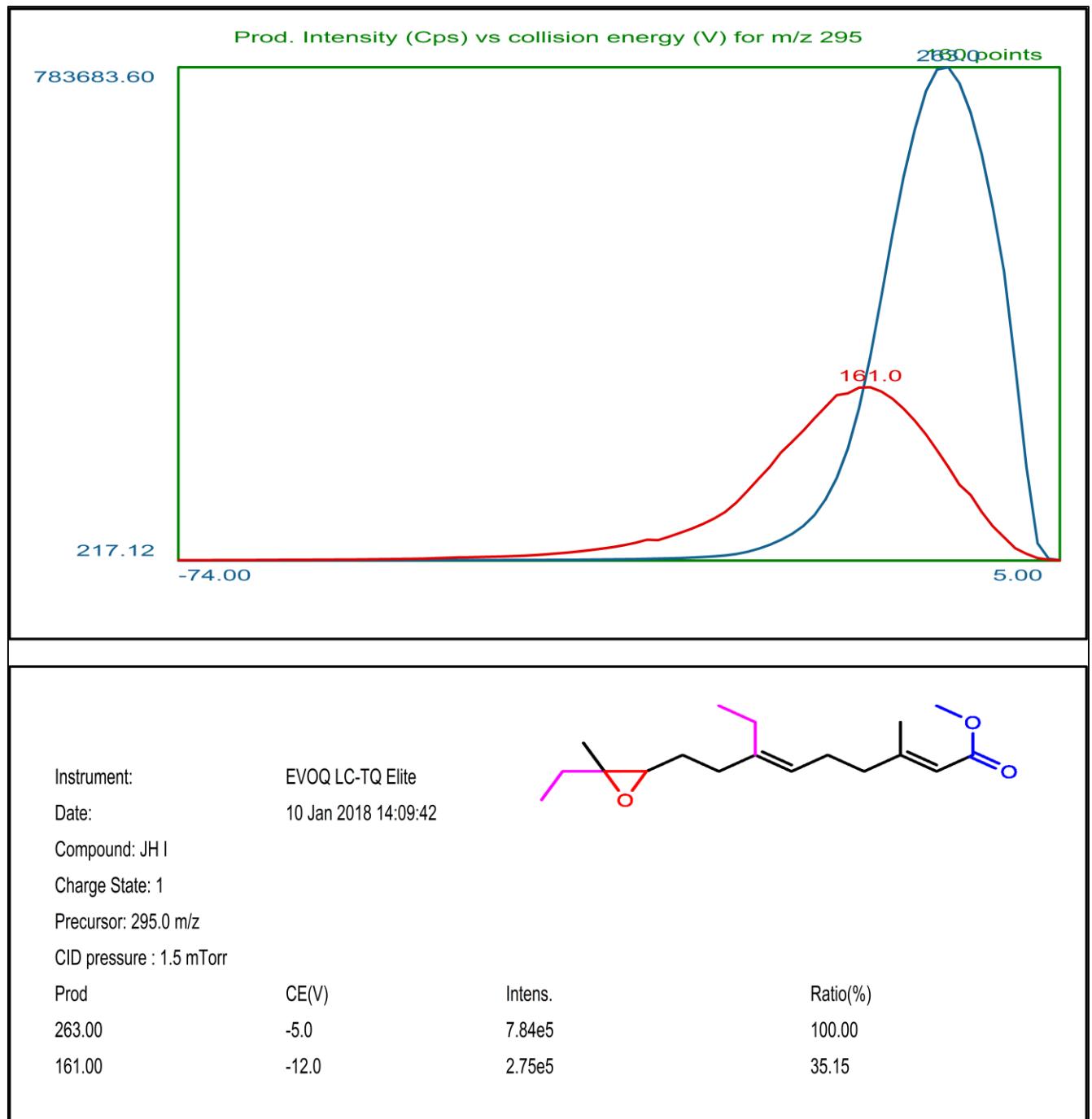


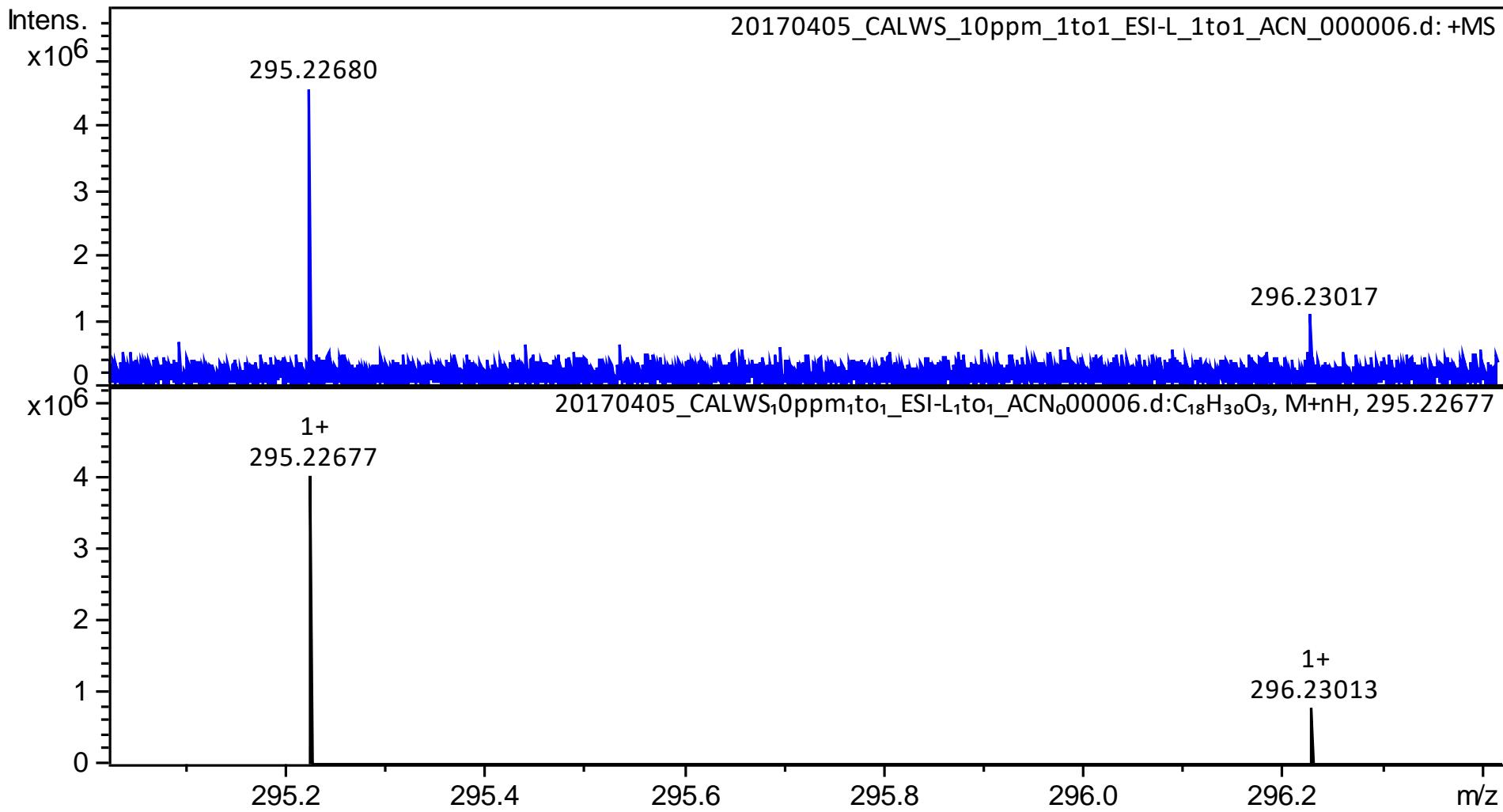
Supplemental Figure 1. Optimization of multiple reaction monitoring (MRM) detection on the Bruker EvoQ LC-TQ Elite Triple Quadrupole Mass Spectrometer and confirmation experiments using the Bruker Solarix 7T Fourier-transform ion cyclotron resonance (FT-ICR) ultra high-resolution mass spectrometer (UHRMS).

- A) Optimization of MRM collision-induced dissociation (CID). The most abundant MRM transitions were selected for LC-MS/MS quantitative analysis, while the second highest were selected for confirmation. In order to add an additional differentiation feature between the isomers JHB₃ and JHSB₃, an exception for this rule was in which the third most abundant MRM transition for JHB₃ (m/z 283→251) was used instead of the second most abundant (m/z 283→119) which was common between them.
- B) Comparison of UHRMS signals for protonated pseudomolecular ions ($[M+H]^+$) with theoretical (according to molecular formulas). Internal calibration was used. Very low mass errors (<0.2 ppm) were observed for all compounds.
- C) Fragmentation spectra (CID, MS/MS) obtained by direct infusion of individual standards using the FT-ICR UHRMS.
- D) Accurate masses of protonated pseudomolecular ion ($[M+H]^+$) and MS/MS fragments, with mass errors according to proposed structures.
- E) Proposed fragmentation pathways for JHs and MF based on collision-induced dissociation (CID) MS/MS experiments using UHRMS.

A: JH I



B: JH I



Ultra high resolution mass spectra of the standard obtained by direct infusion using the Solarix 7T FT-ICR MS .

C: JH I

Fourier-transformation cyclotron resonance (FT-ICR) ultra high-resolution MS/MS.

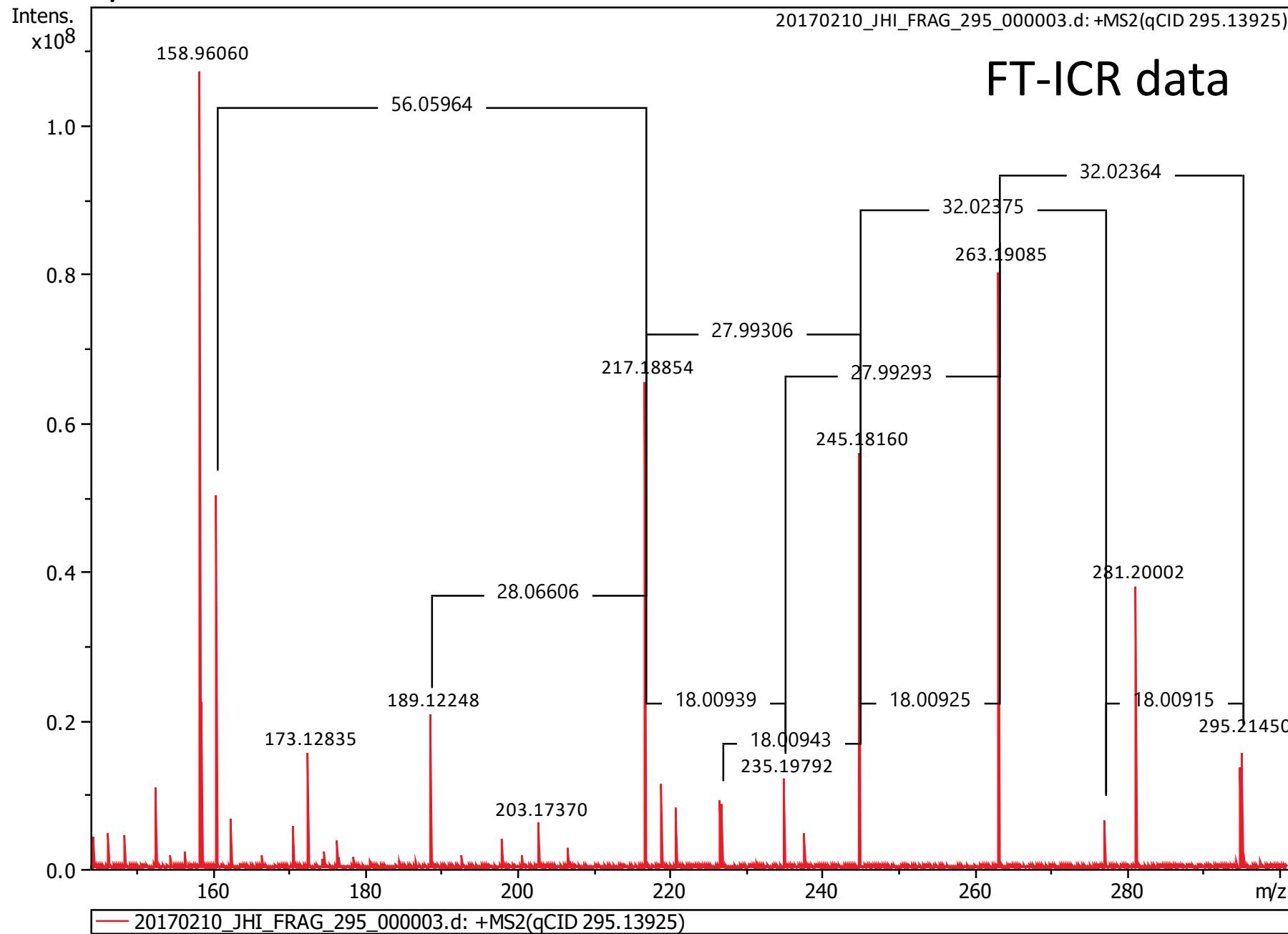


Table showing the MS/MS fragments of the hormone using the Solarix 7T FT-ICR MS.

UHRMS data	m/z	Formula	$\Delta m/z$ (ppm)
FT-ICR MS	295.22680	[C18H30O3+H] ⁺	0.098
FT-ICR MS ² (CID on 295.2268)	277.21593	[C18H29O2+H] ⁺	-0.999
	263.20045	C17H27O2 ⁺	-0.407
	245.18993	C17H25O ⁺	-0.253
	235.20559	C16H27O ⁺	-0.221
	217.19508	C16H25 ⁺	0.014
	161.13249	C12H17 ⁺	0.081

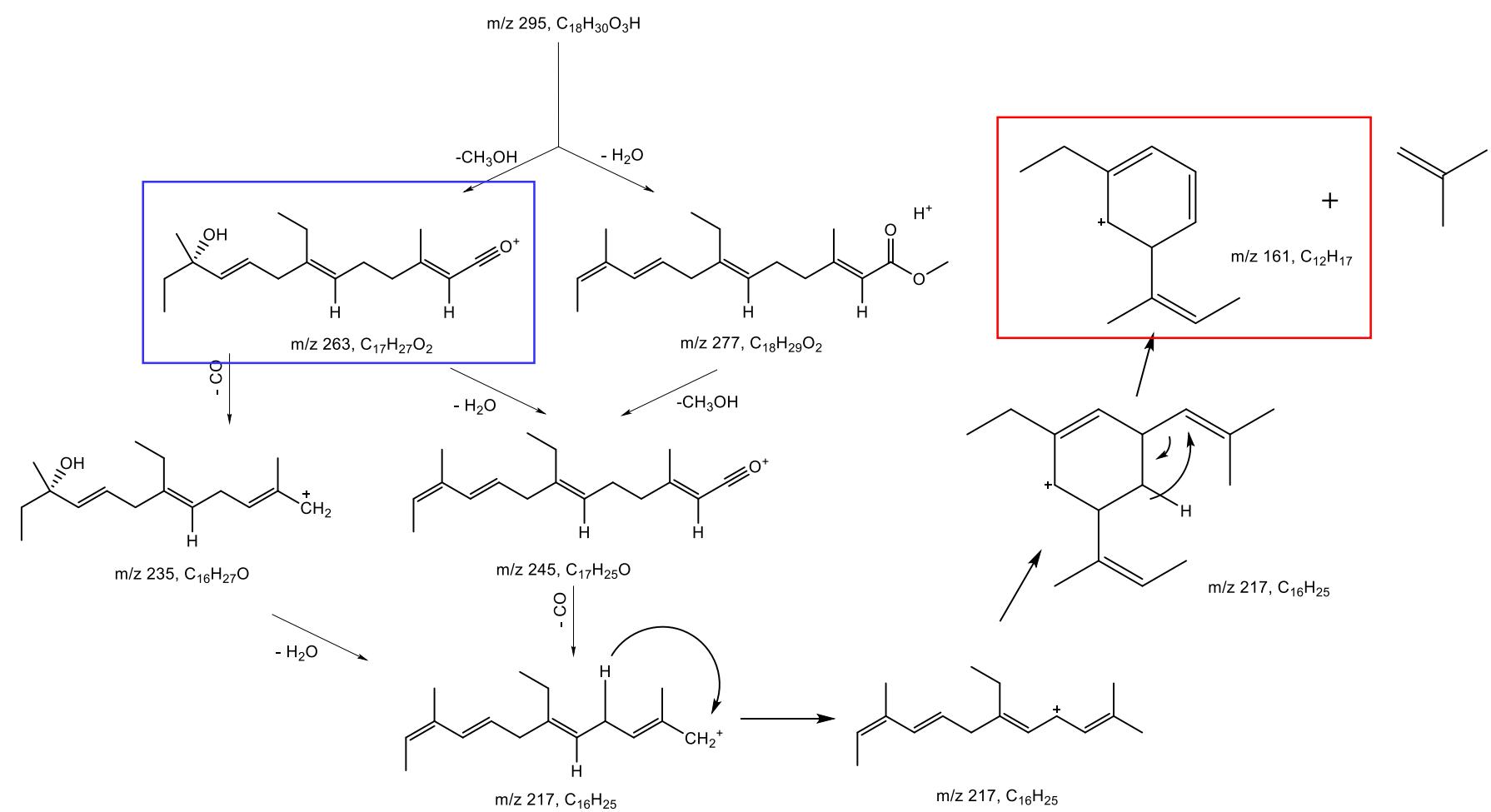
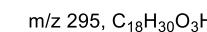
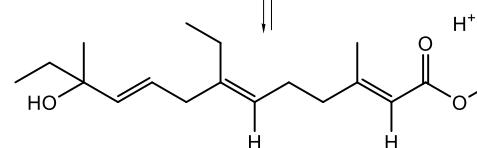
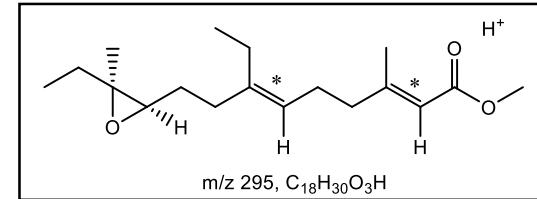
E: JH I

Proposed fragmentation pathways according to the FT-ICR results.

Black box: parent ion

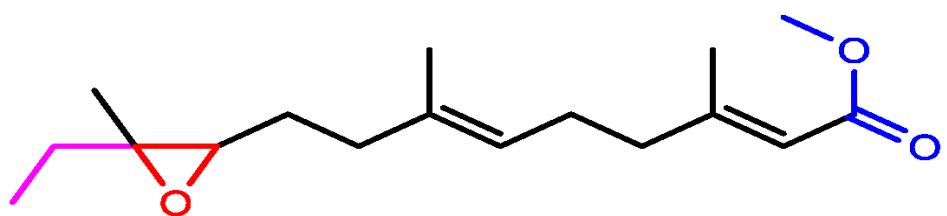
Blue box: primary transition

Red box: secondary transition

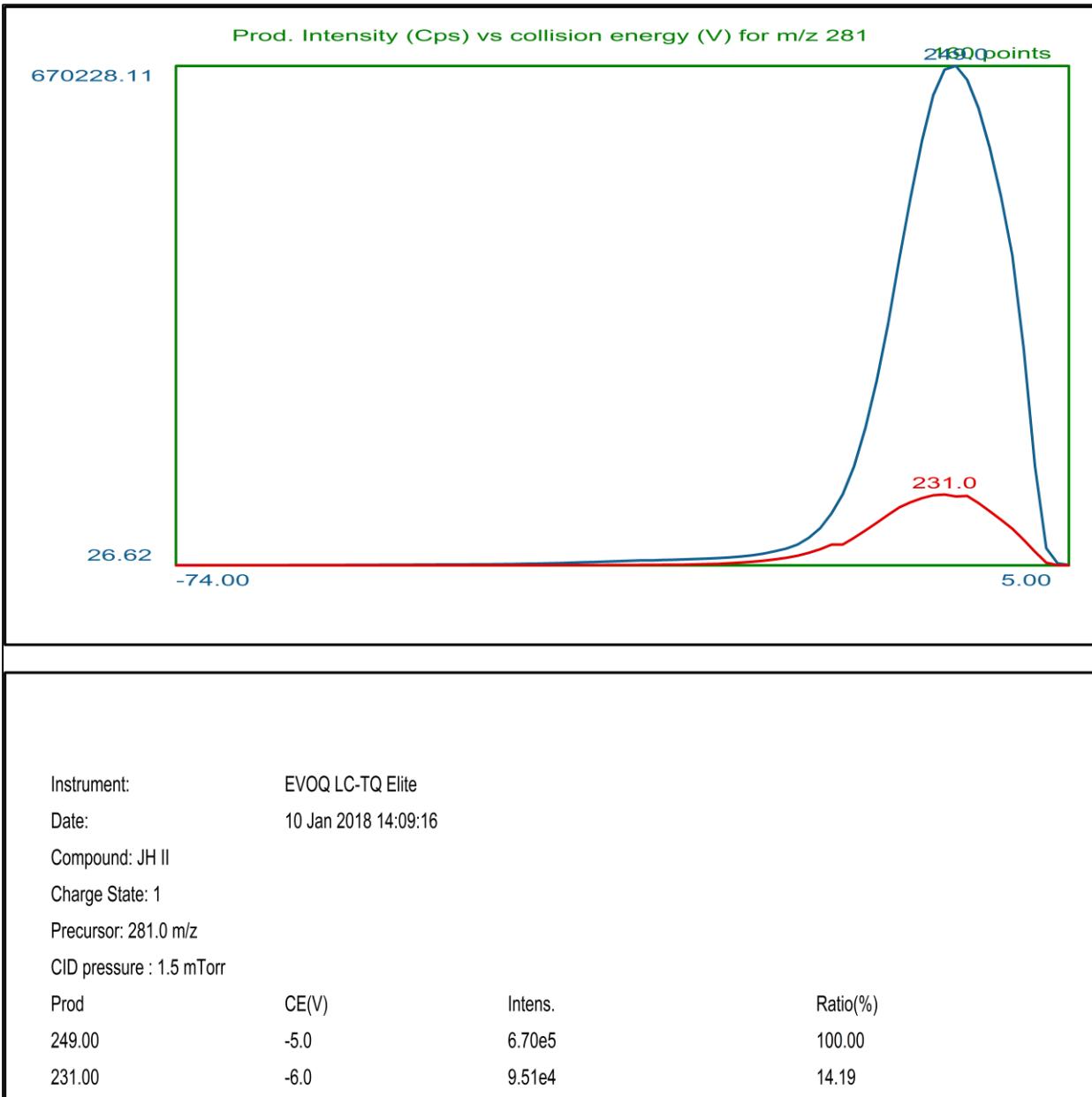


A: JH II

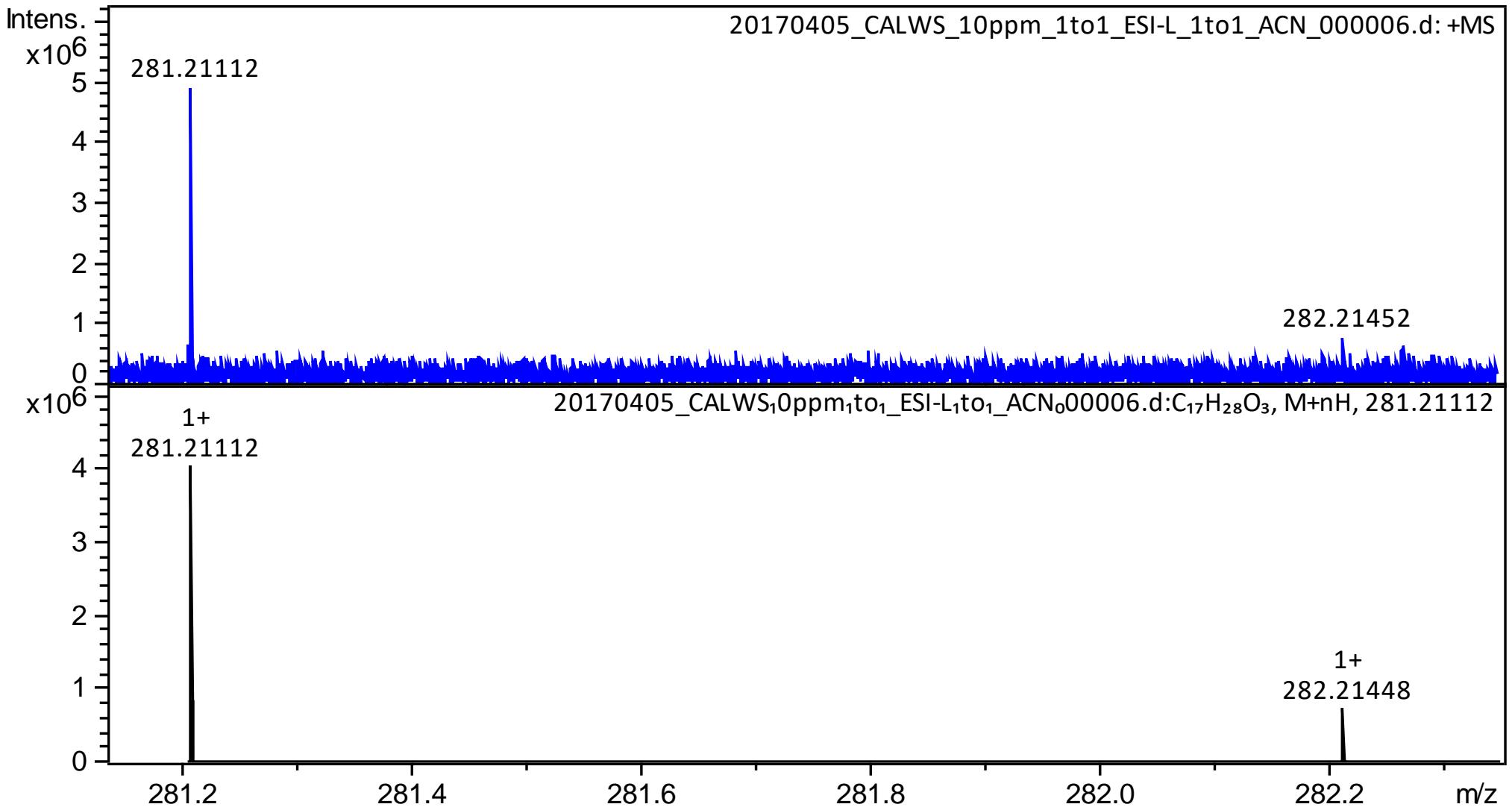
Bruker LC-TQ MRM optimization output



Multiple reaction monitoring (MRM) optimization output using the Bruker EVOQ LC-TQ Elite Triple Quadrupole Mass Spectrometer.



B: JH II



Ultra high resolution mass spectra of the standard obtained by direct infusion using the Solarix 7T FT-ICR MS .

C: JH II

Fourier-transformation cyclotron resonance (FT-ICR) ultra high-resolution MS/MS.

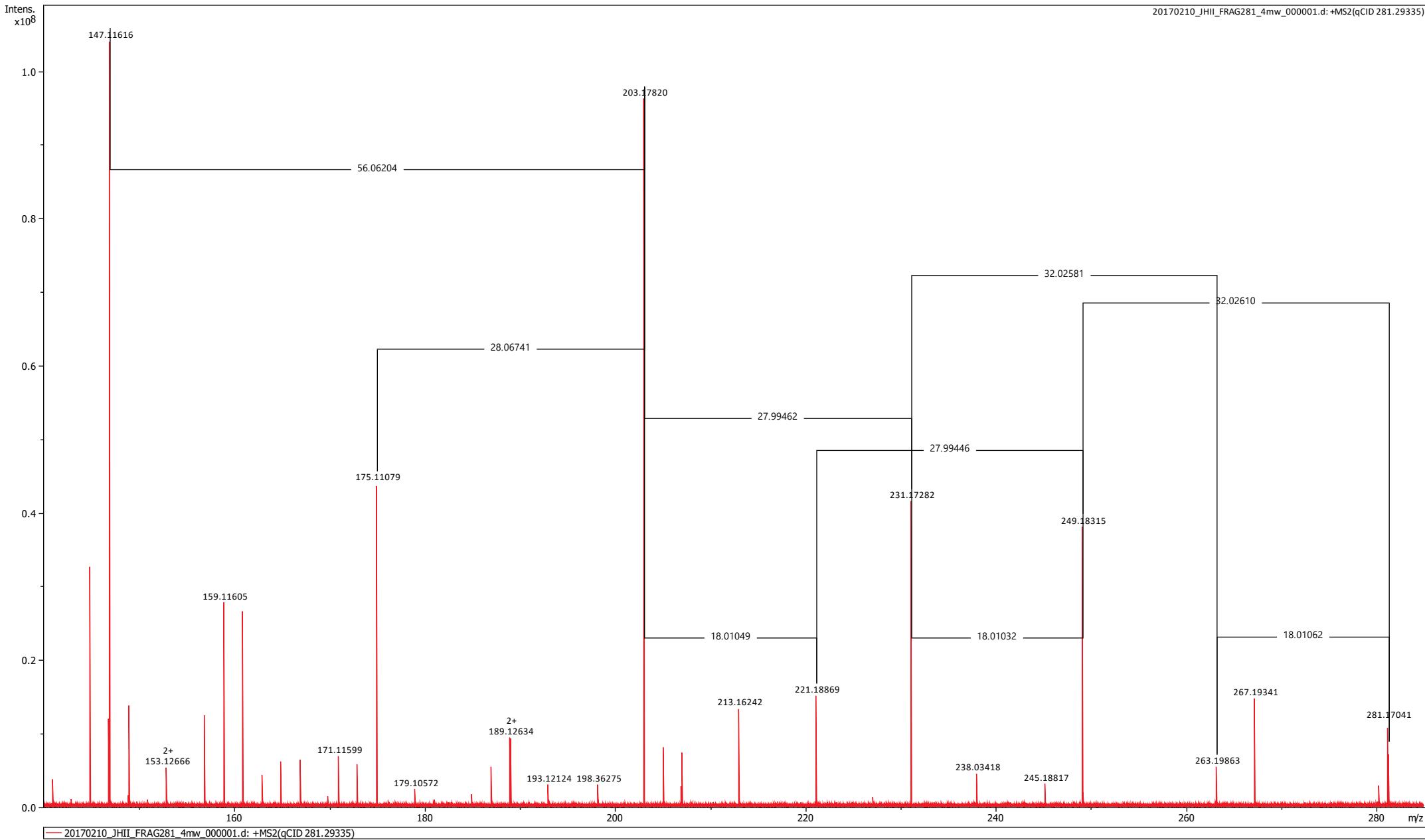
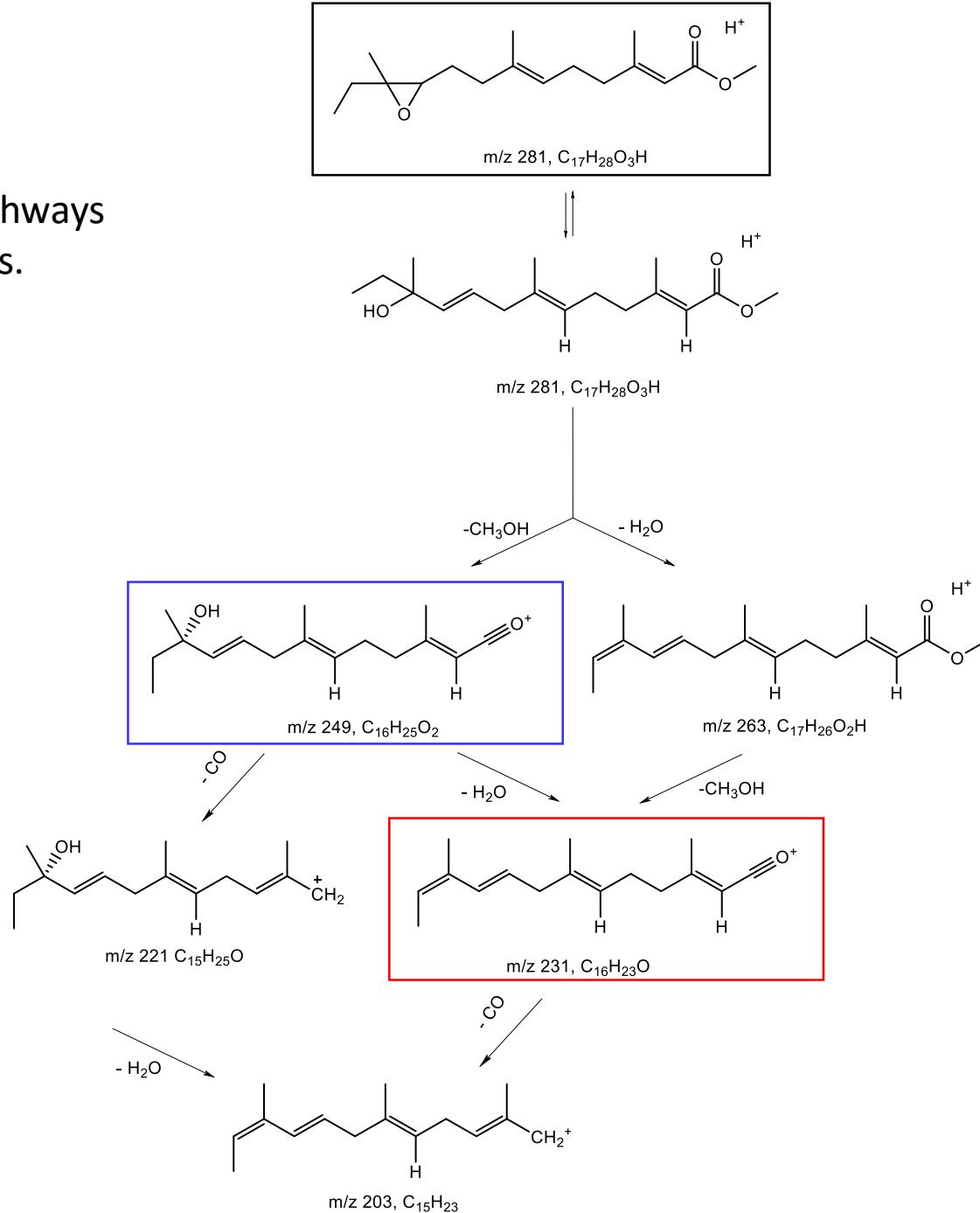


Table showing the MS/MS fragments of the hormone using the Solarix 7T FT-ICR MS.

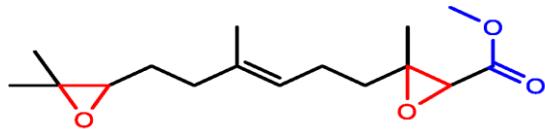
UHRMS data	m/z	Formula	$\Delta m/z$ (ppm)
FT-ICR MS	281.21112	$[C_{17}H_{28}O_3+H]^+$	<0.001
FT-ICR MS ² (CID on 281.21112)	263.200557	$[C_{17}H_{26}O_2+H]^+$	0.581
	249.184906	$C_{16}H_{25}O_2^+$	0.337
	231.174342	$C_{16}H_{23}O^+$	0.294
	221.189992	$C_{15}H_{25}O^+$	0.669
	203.179427	$C_{15}H_{23}^+$	0.015
	263.200557	$C_{11}H_{15}^+$	0.581

E: JH II

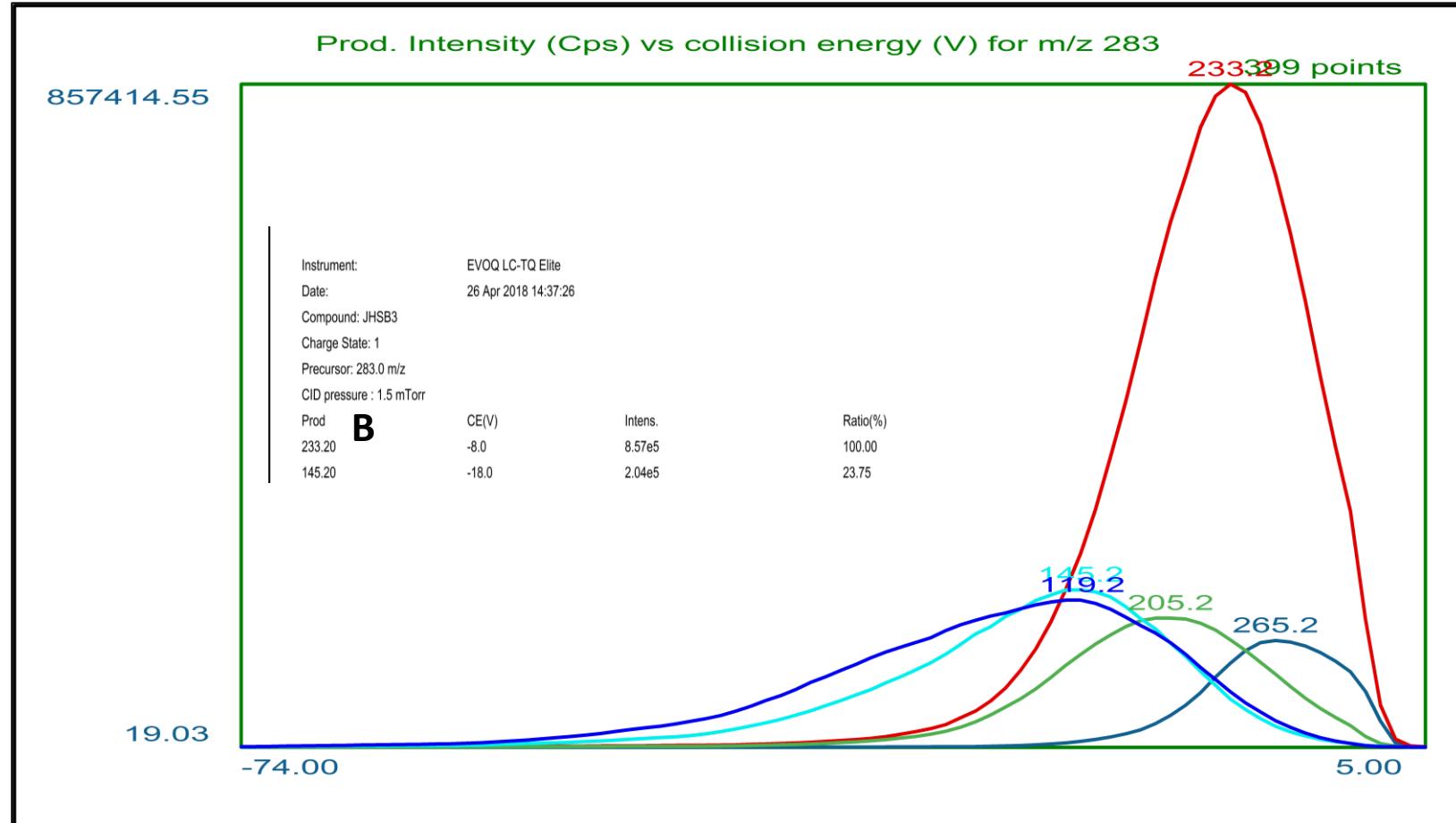
Proposed fragmentation pathways
according to the FT-ICR results.



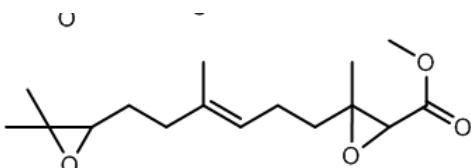
A: JHSB3



Bruker LC-TQ MRM optimization output



Multiple reaction monitoring (MRM) optimization output using the Bruker EvoQ LC-TQ Elite Triple Quadrupole Mass Spectrometer.

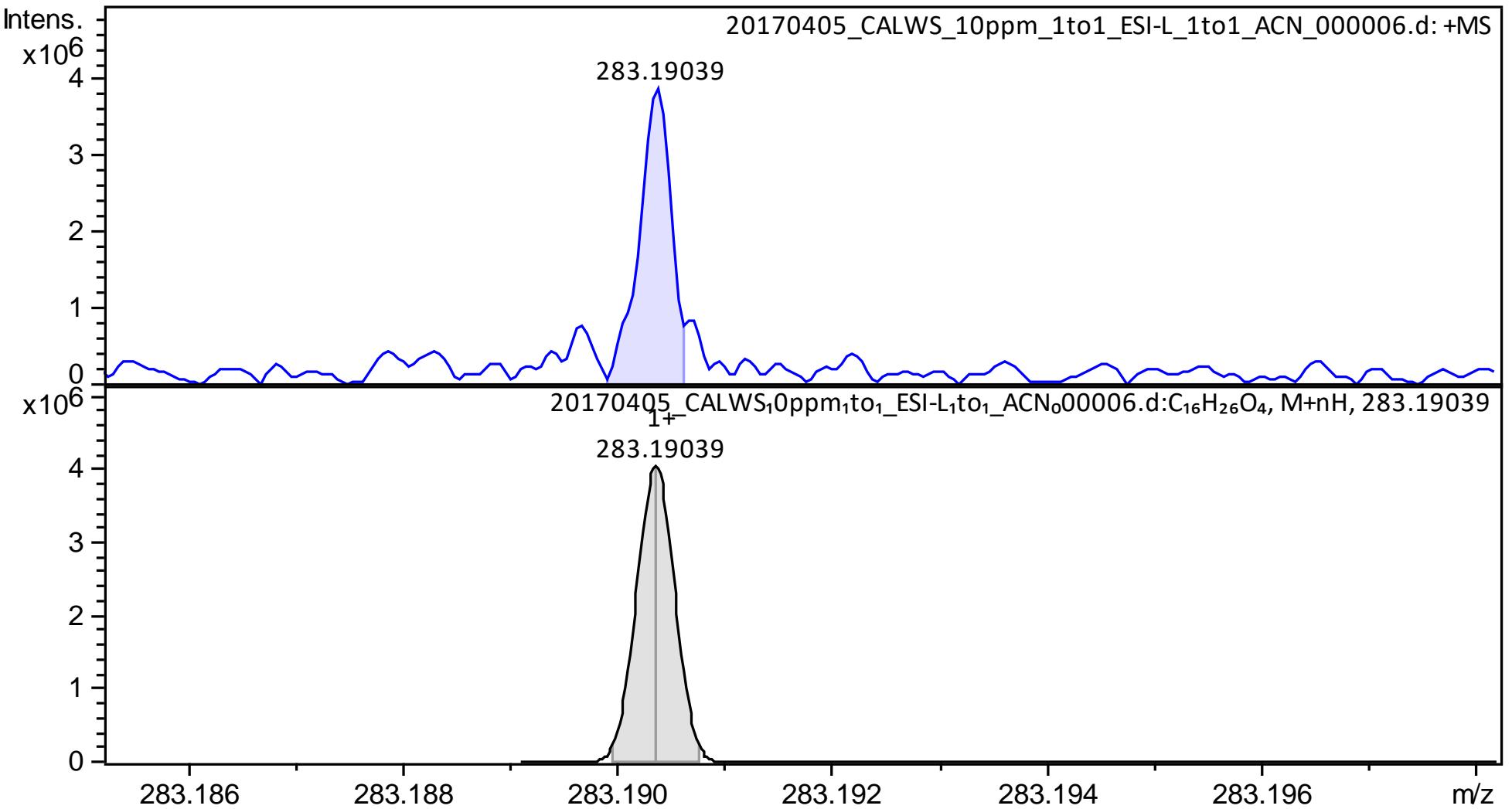


JHSB₃

283.3 → 233.2 ; 8.0
283.3 → 145.2 ; 18.0

8.8

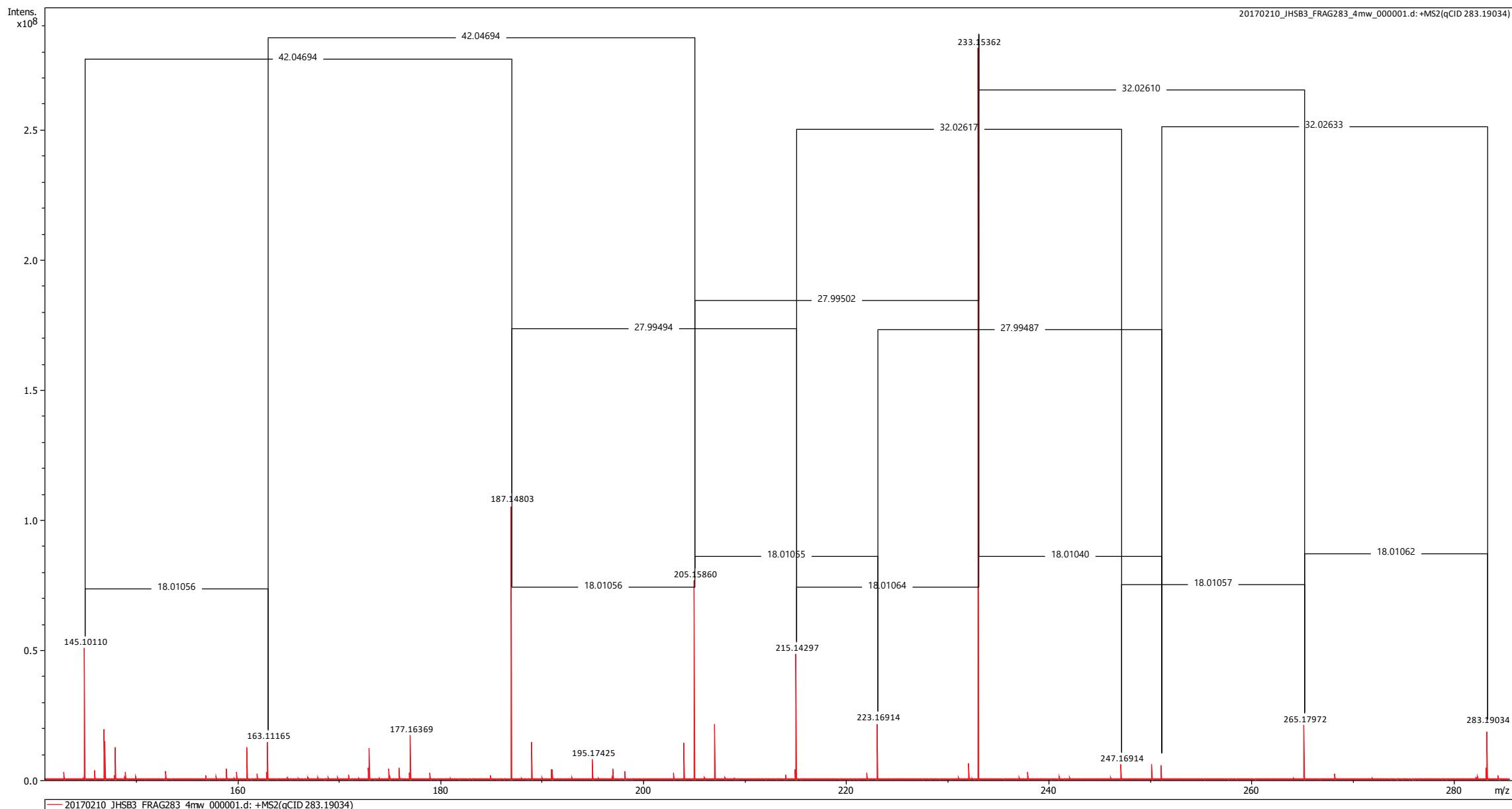
B: JHSB3



Ultra high resolution mass spectra of the standard obtained by direct infusion using the Solarix 7T FT-ICR MS .

C: JHSB3

Fourier-transformation cyclotron resonance (FT-ICR) ultra high-resolution MS/MS.



D: JHSB3

Table showing the MS/MS fragments of the hormone using the Solarix 7T FT-ICR MS.

UHRMS data	m/z	Ion	Formula	$\Delta m/z$ (ppm)
FT-ICR MS	283.19039	[M+H] ⁺	[C ₁₆ H ₂₆ O ₄ +H] ⁺	0.014
FT-ICR MS ² (CID on 283.19039)	265.17972	[M+H] ⁺ - H ₂ O	[C ₁₆ H ₂₄ O ₃ +H] ⁺	-0.381
	251.16407	[M+H] ⁺ - CH ₃ OH	C ₁₅ H ₂₃ O ₃ ⁺	-0.402
	247.16914	[M+H] ⁺ - H ₂ O - H ₂ O	[C ₁₆ H ₂₂ O ₂ +H] ⁺	-0.469
	233.15362	[M+H] ⁺ - H ₂ O - CH ₃ OH [M+H] ⁺ - CH ₃ OH - H ₂ O	C ₁₅ H ₂₁ O ₂ ⁺	0.060
	223.16914		C ₁₄ H ₂₃ O ₂ ⁺	-0.520
	215.14297	[M+H] ⁺ - H ₂ O - CH ₃ OH- CO	C ₁₅ H ₁₉ O ⁺	-0.335
	205.15860	[M+H] ⁺ - H ₂ O - CH ₃ OH- CO	C ₁₄ H ₂₁ O ⁺	-0.445
	187.14803		C ₁₄ H ₁₉ ⁺	-0.518
	145.10110		C ₁₁ H ₁₃ ⁺	-0.530

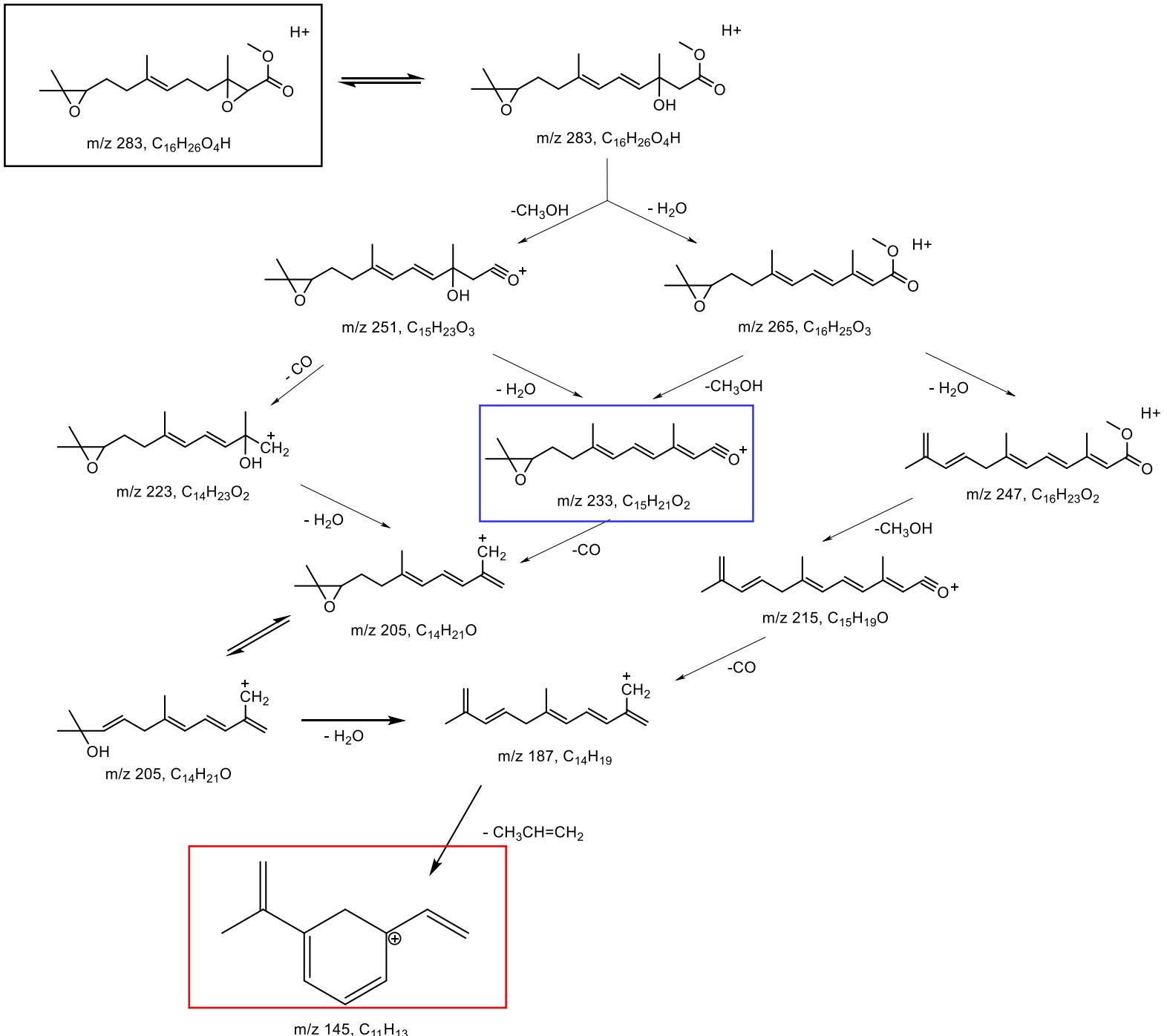
E: JHSB3

Proposed fragmentation pathways
according to the FT-ICR results.

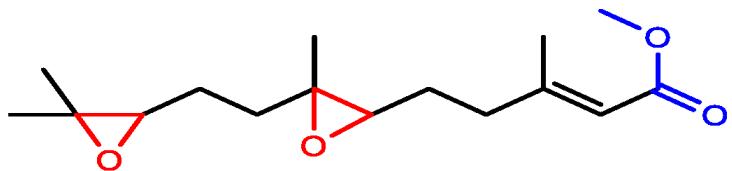
Black box: parent ion

Blue box: primary transition

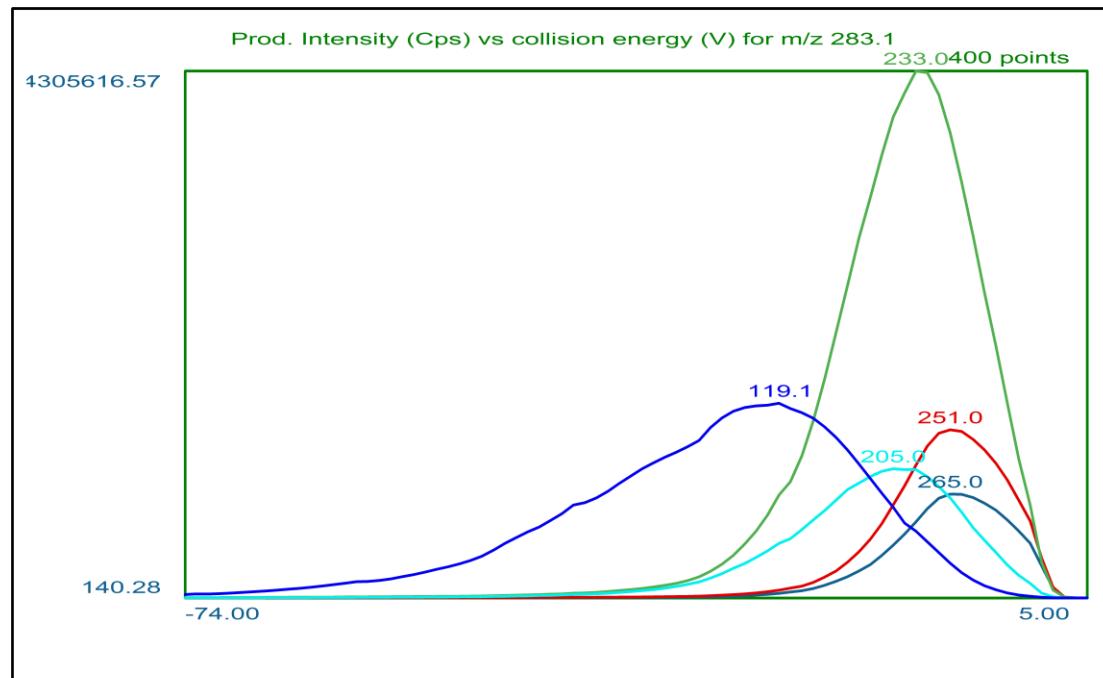
Red box: secondary transition



A: JHB3



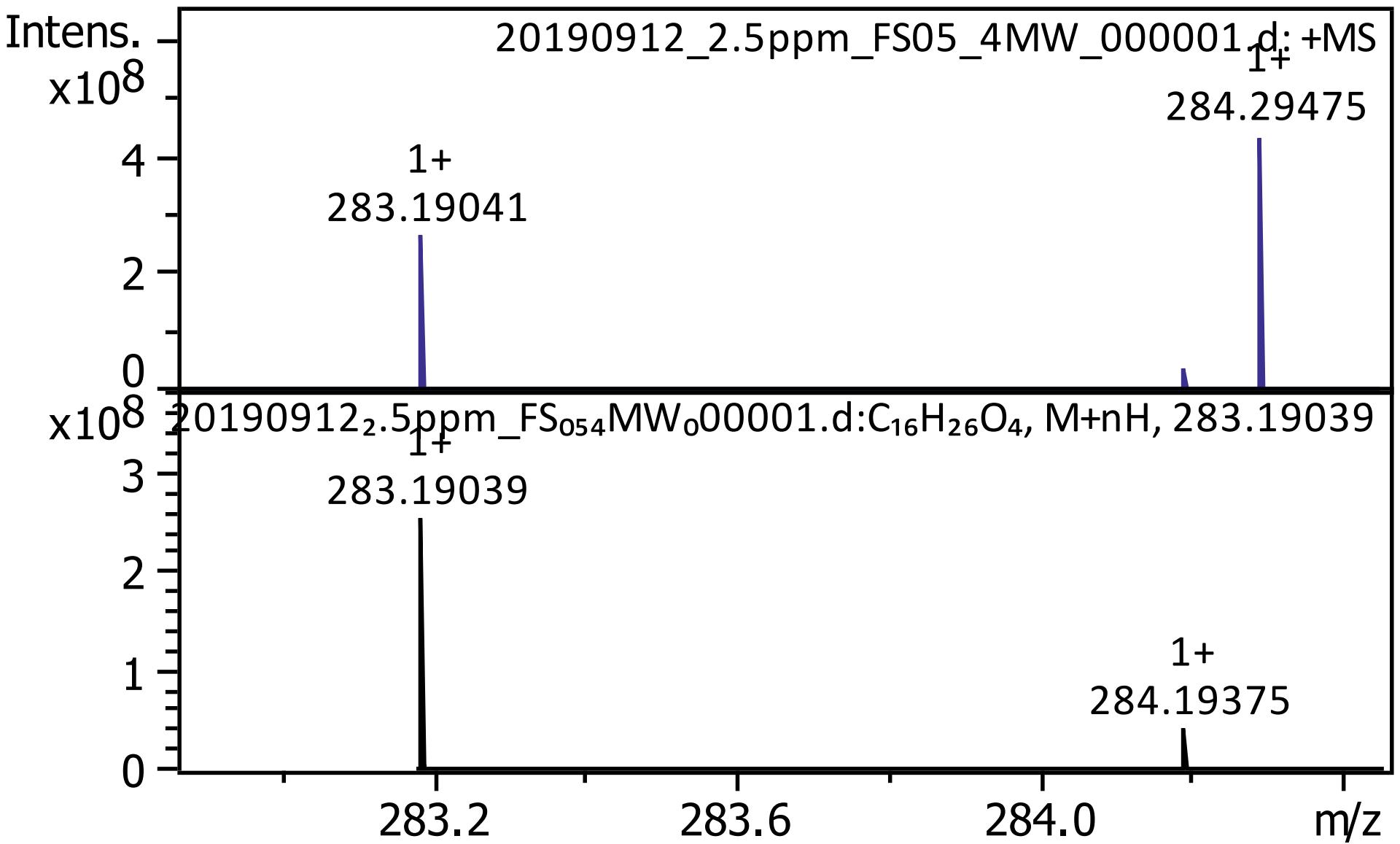
Multiple reaction monitoring (MRM) optimization output using the Bruker EvoQ LC-TQ Elite Triple Quadrupole Mass Spectrometer.



m/z 251 was selected as secondary (instead of m/z 119) because it is not abundant for JHSB3

Instrument:	EVOQ LC-TQ Elite		
Date:	27 Aug 2019 9:02:16		
Compound:	JHB3		
Charge State:	1		
Precursor:	283.1 m/z		
CID pressure :	1.5 mTorr		
Prod	CE(V)	Intens.	Ratio(%)
233.00	-10.0	4.31e6	100.00
119.10	-22.0	1.59e6	36.96
251.00	-7.0	1.37e6	31.93
205.00	-12.0	1.06e6	24.51
265.00	-7.0	8.49e5	19.72

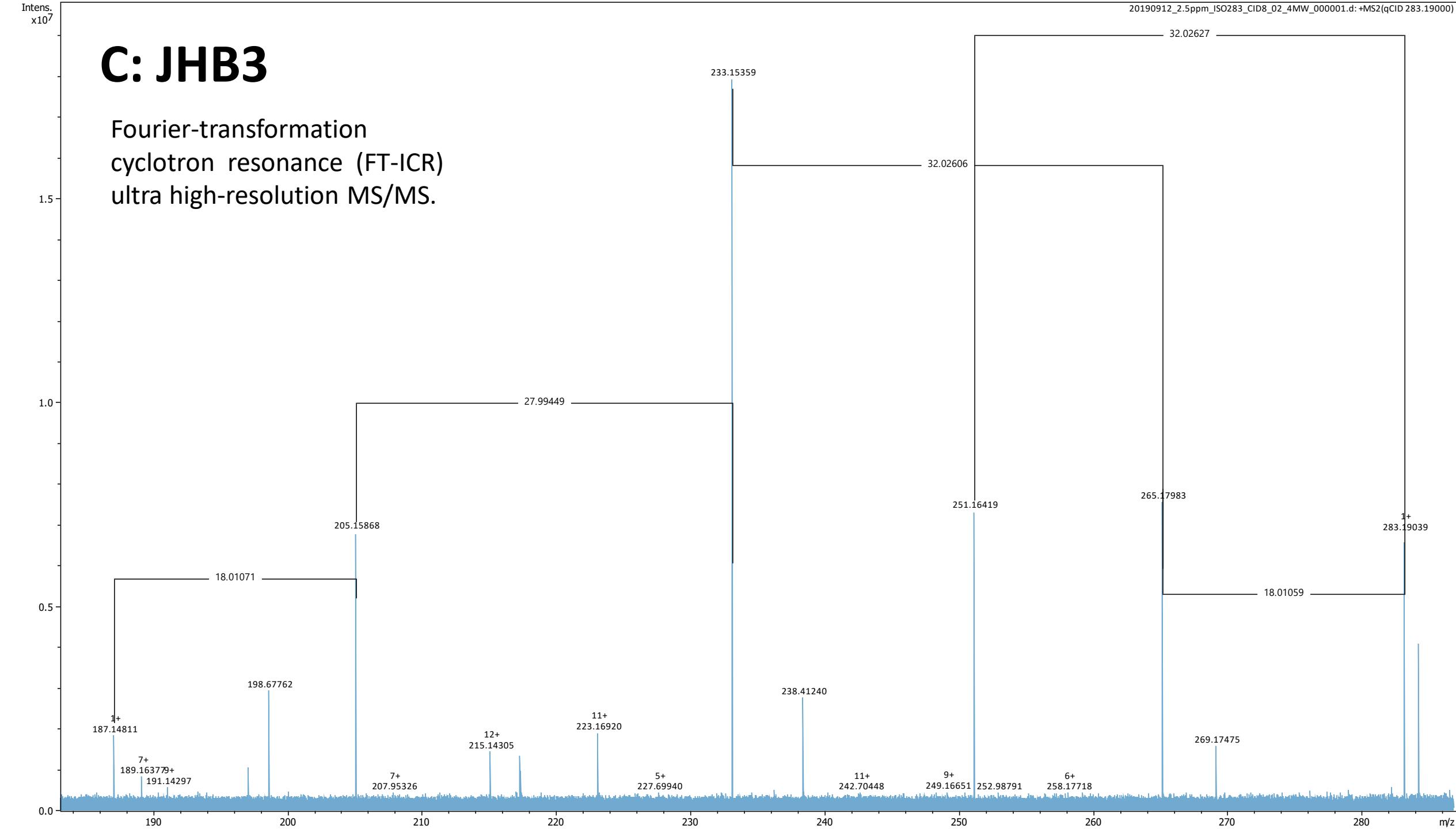
B: JHB3



Ultra high resolution mass spectra of the standard obtained by direct infusion using the Solarix 7T FT-ICR MS .

C: JHB3

Fourier-transformation
cyclotron resonance (FT-ICR)
ultra high-resolution MS/MS.



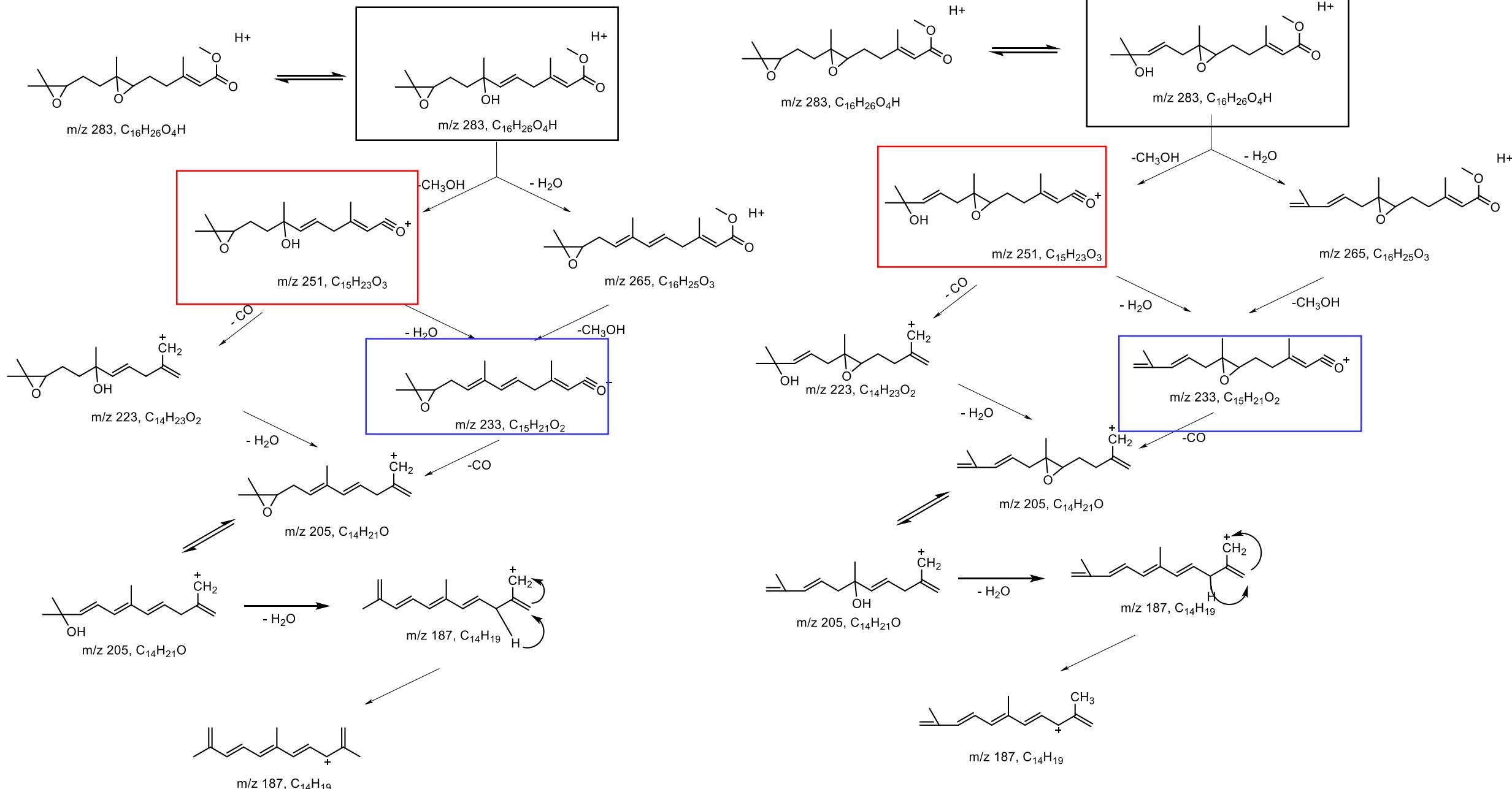
C: JHB3

Table showing the MS/MS fragments of the hormone using the Solarix 7T FT-ICR MS.

UHRMS data	m/z	Ion	Formula	Δm/z (ppm)
FT-ICR MS	283.19041	[M+H] ⁺	[C ₁₆ H ₂₆ O ₄ +H] ⁺	0.085
FT-ICR MS ² (CID on 283.19039)	265.17983	[M+H] ⁺ - H ₂ O	[C ₁₆ H ₂₄ O ₃ +H] ⁺	0.034
	251.16419	[M+H] ⁺ - CH ₃ OH	C ₁₅ H ₂₃ O ₃ ⁺	0.076
	233.15359	[M+H] ⁺ - H ₂ O - CH ₃ OH [M+H] ⁺ - CH ₃ OH - H ₂ O	C ₁₅ H ₂₁ O ₂ ⁺	-0.069
	223.1692		C ₁₄ H ₂₃ O ₂ ⁺	-0.251
	215.14305	[M+H] ⁺ - H ₂ O - CH ₃ OH - CO	C ₁₅ H ₁₉ O ⁺	0.037
	205.15868	[M+H] ⁺ - H ₂ O - CH ₃ OH - CO	C ₁₄ H ₂₁ O ⁺	-0.058
	187.14811		C ₁₄ H ₁₉ ⁺	-0.091

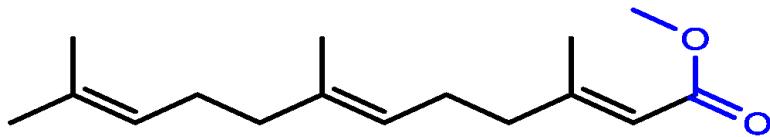
E: JHB3

Proposed fragmentation pathways according to the FT-ICR results.

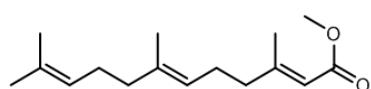
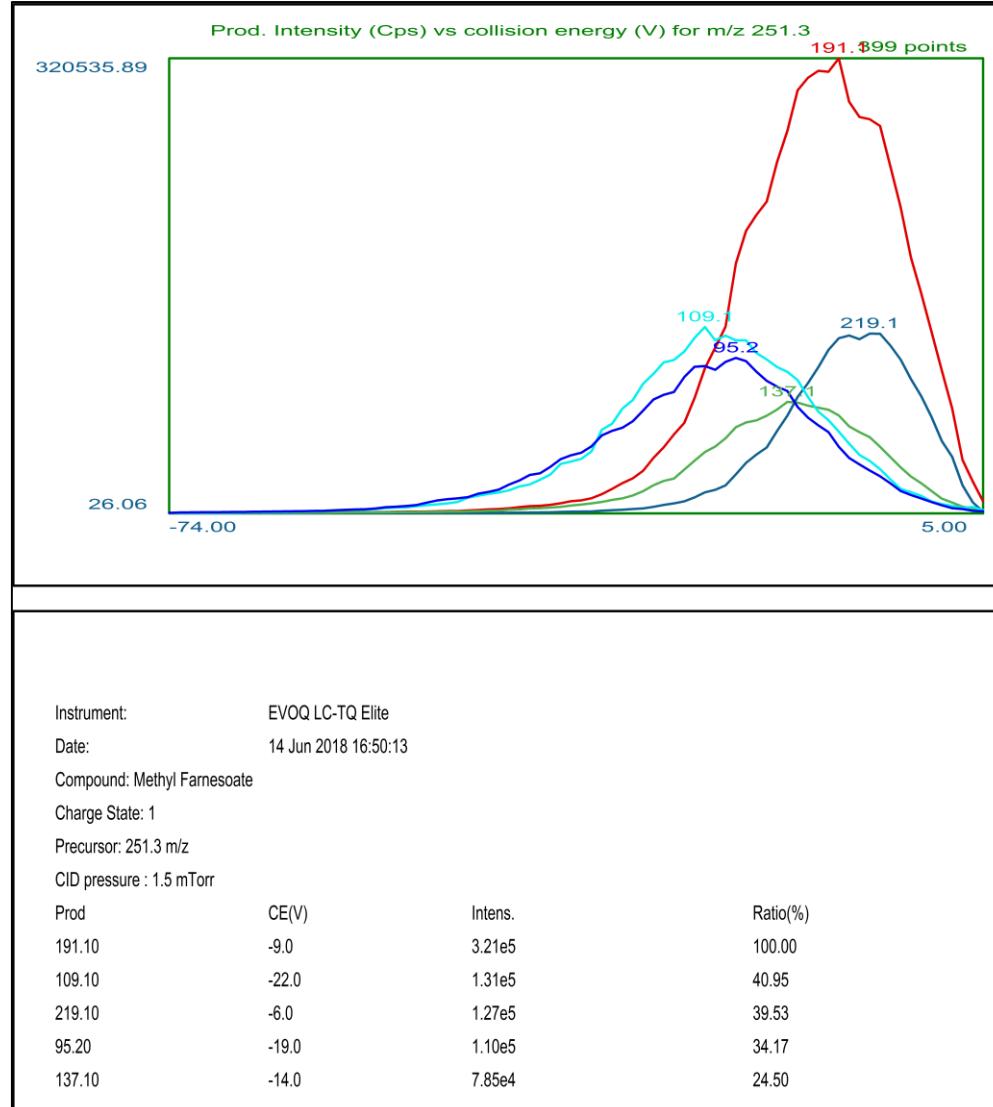


Bruker LC-TQ MRM OPT

A: MF



Multiple reaction monitoring (MRM) optimization output using the Bruker EVOQ LC-TQ Elite Triple Quadrupole Mass Spectrometer.

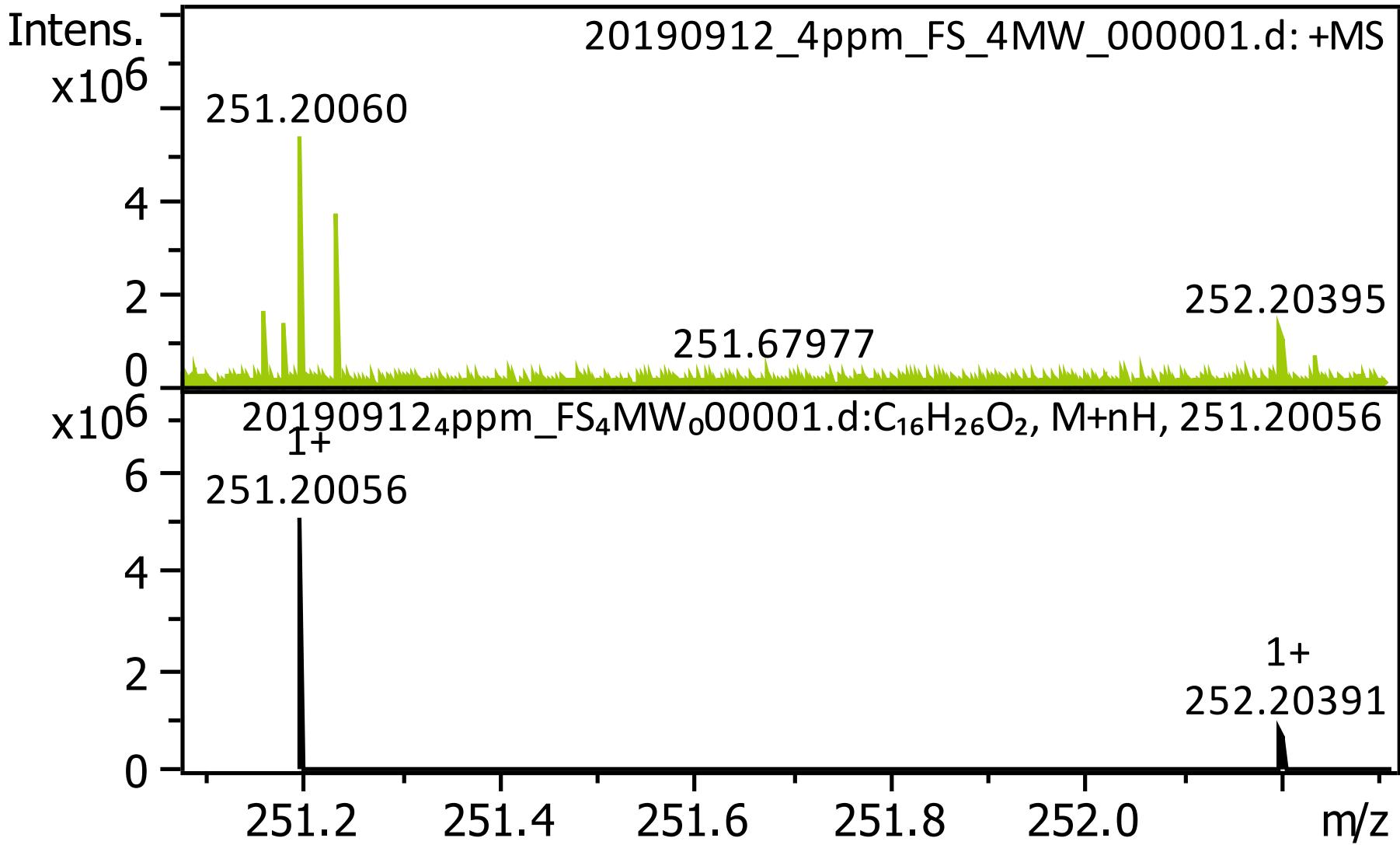


MF

251.3 → 191.2 ; 9.0
295.2 → 219.2 ; 6.0

10.0

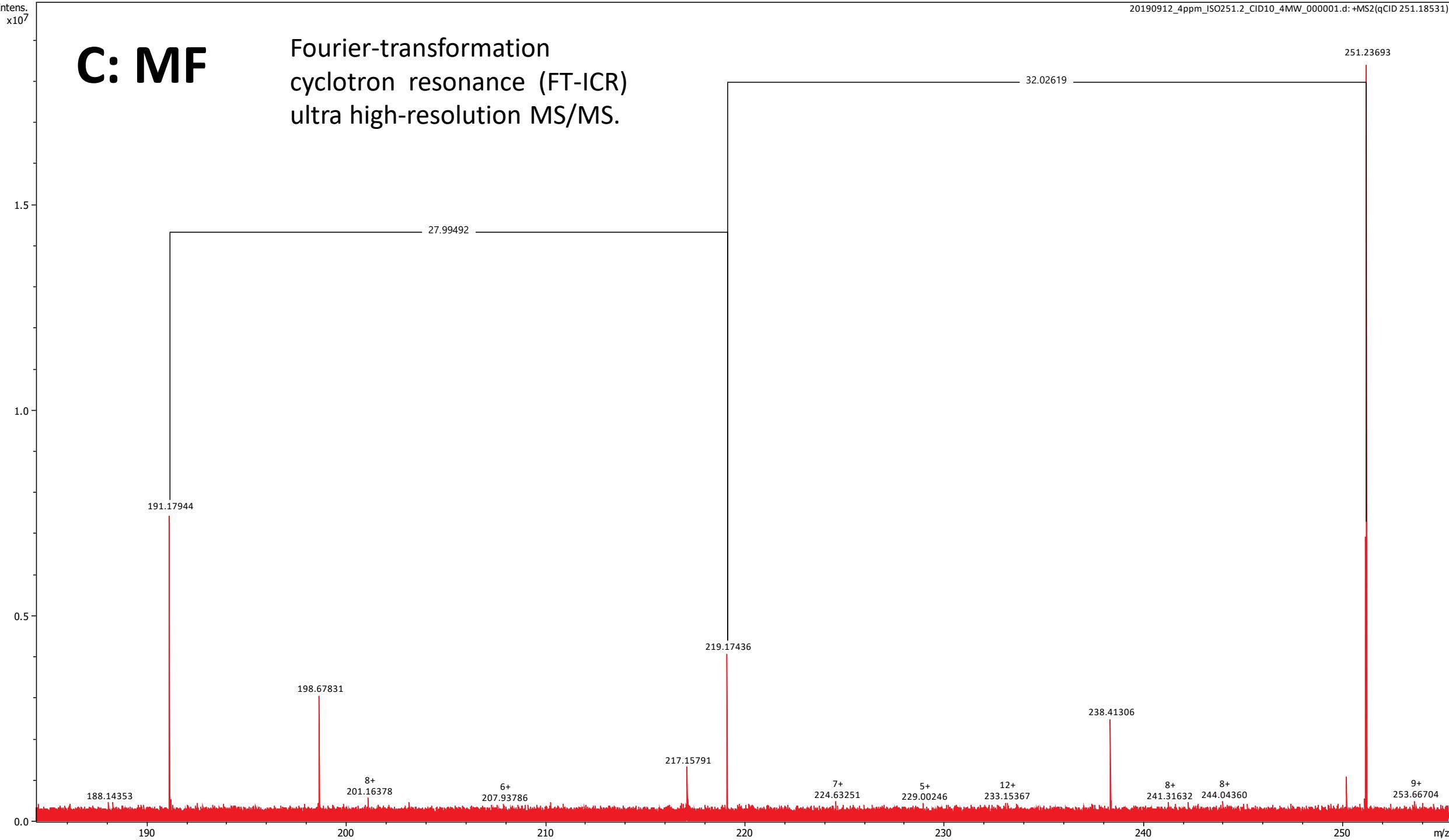
B: MF



Ultra high resolution mass spectra of the standard obtained by direct infusion using the Solarix 7T FT-ICR MS .

C: MF

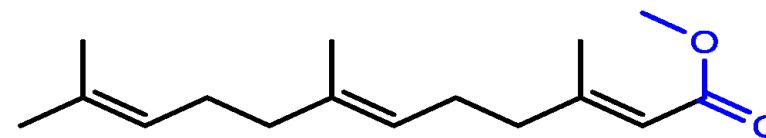
Fourier-transformation
cyclotron resonance (FT-ICR)
ultra high-resolution MS/MS.



C: MF

Table showing the MS/MS fragments of the hormone using the Solarix 7T FT-ICR MS.

UHRMS data	m/z	Ion	Formula	Δm/z (ppm)
FT-ICR MS	251.2006	[M+H] ⁺	[C ₁₆ H ₂₆ O ₄ +H] ⁺	0.171
FT-ICR MS ² (CID on 251.2006)	219.17436	[M+H] ⁺ - H ₂ O	[C ₁₆ H ₂₂ O ₂ +H] ⁺	0.082
	191.17944	[M+H] ⁺ - CH ₃ OH	C ₁₅ H ₂₃ O ₃ ⁺	0.068



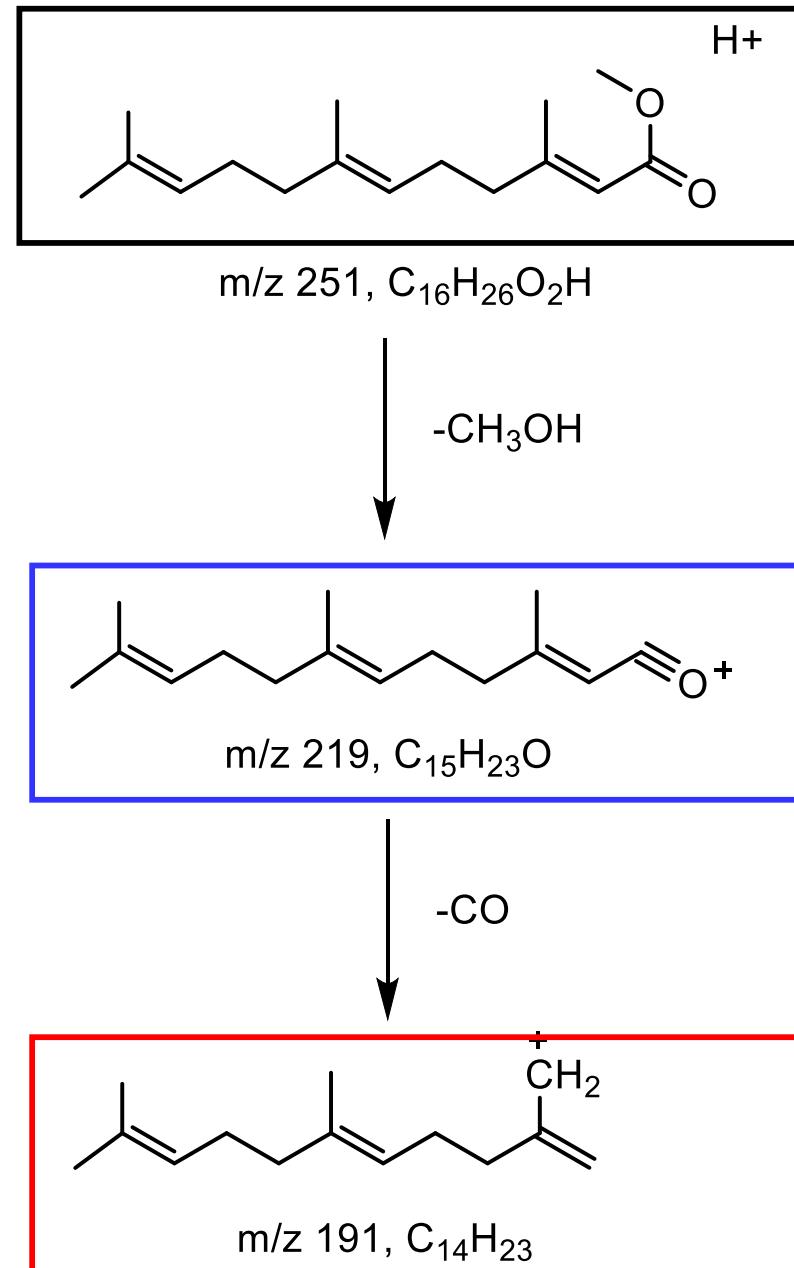
E: MF

Proposed fragmentation pathways according to the FT-ICR results.

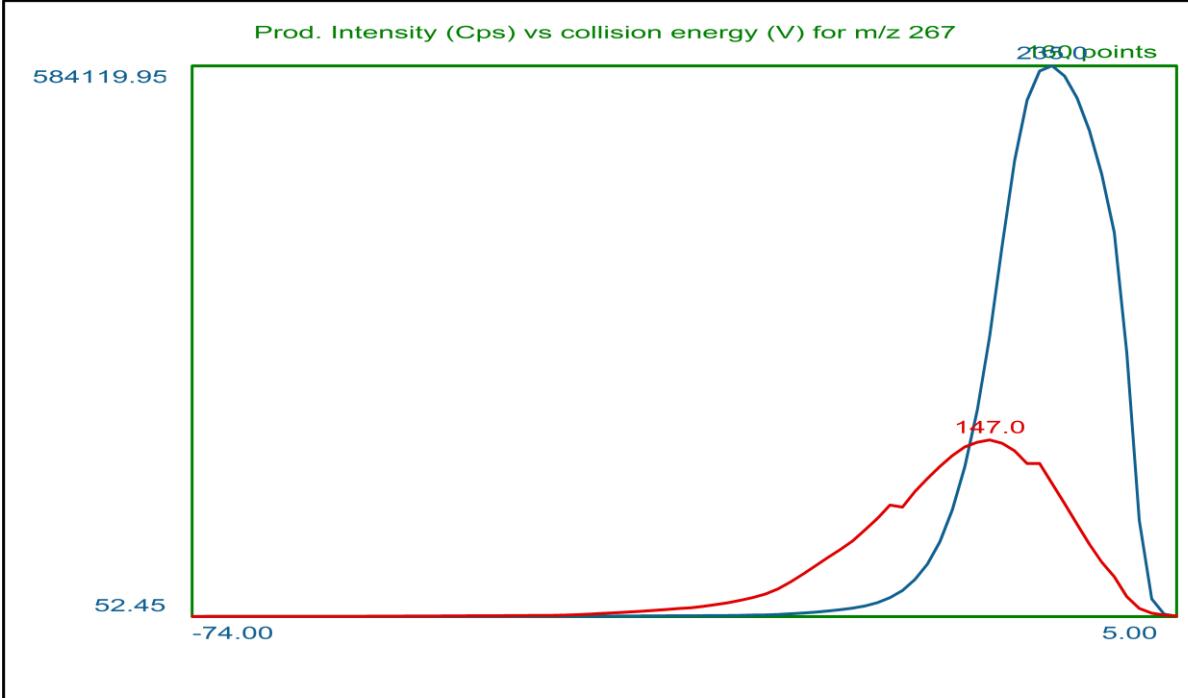
Black box: parent ion

Blue box: primary transition

Red box: secondary transition



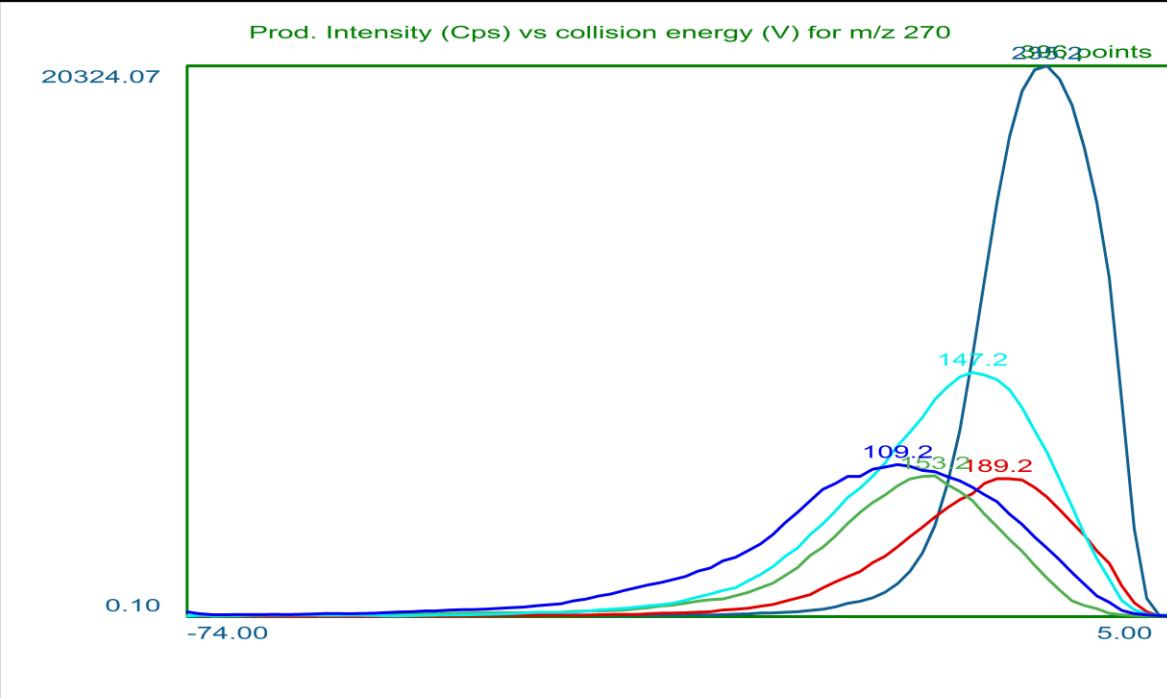
A: JH III



Instrument: EVOQ LC-TQ Elite
Date: 10 Jan 2018 14:08:51
Compound: JH III
Charge State: 1
Precursor: 267.0 m/z
CID pressure : 1.6 mTorr

Prod	CE(V)	Intens.	Ratio(%)
235.00	-5.0	5.84e5	100.00
147.00	-10.0	1.87e5	32.08

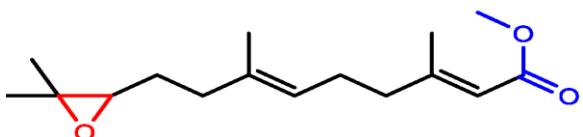
A: JH III-D3



Instrument:	EVOQ LC-TQ Elite		
Date:	23 Jan 2018 15:34:21		
Compound:	JH III-D3		
Charge State:	1		
Precursor:	270.0 m/z		
CID pressure :	1.5 mTorr		
Prod	CE(V)	Intens.	Ratio(%)
235.20	-5.0	2.03e4	100.00
147.20	-11.0	9.01e3	44.34
109.20	-17.0	5.61e3	27.61
153.20	-14.0	5.20e3	25.58
189.20	-9.0	5.09e3	25.05

Fragmentation of JH III (from Ramirez et al, 2016)

	Parent <i>m/z</i>	Fragment <i>m/z</i>
In-source MS/MS	267	249, 235, 217, 189, 147
	249	217, 189, 147
	235	217, 189, 147
	217	189, 147
	189	147
MS/MSⁿ	267 -> 235	217, 207, 189
	267 -> 249	217, 189, 147
	267 -> 217	189, 147
	267 -> 189	147
FT-ICR MS/MS	267.195461 C ₁₆ H ₂₇ O ₃	249.18492 C ₁₆ H ₂₅ O ₂ 235.169298 C ₁₅ H ₂₃ O ₂ 217.158786 C ₁₅ H ₂₁ O 207.174417 C ₁₄ H ₂₃ O 189.163858 C ₁₄ H ₂₁ 147.116827 C ₁₁ H ₁₅



Fragmentation of JH III (from Ramirez et al, 2016)

