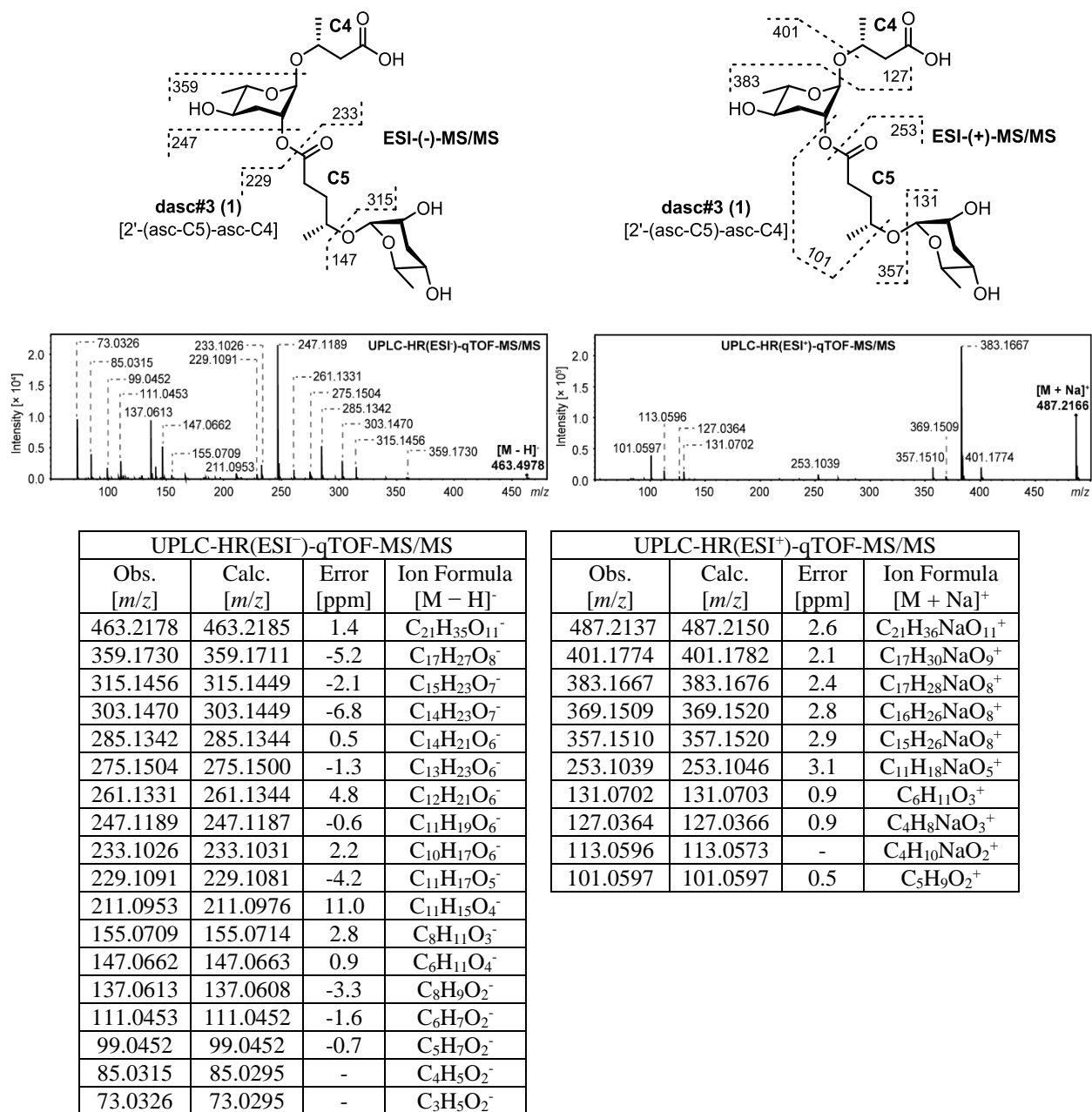
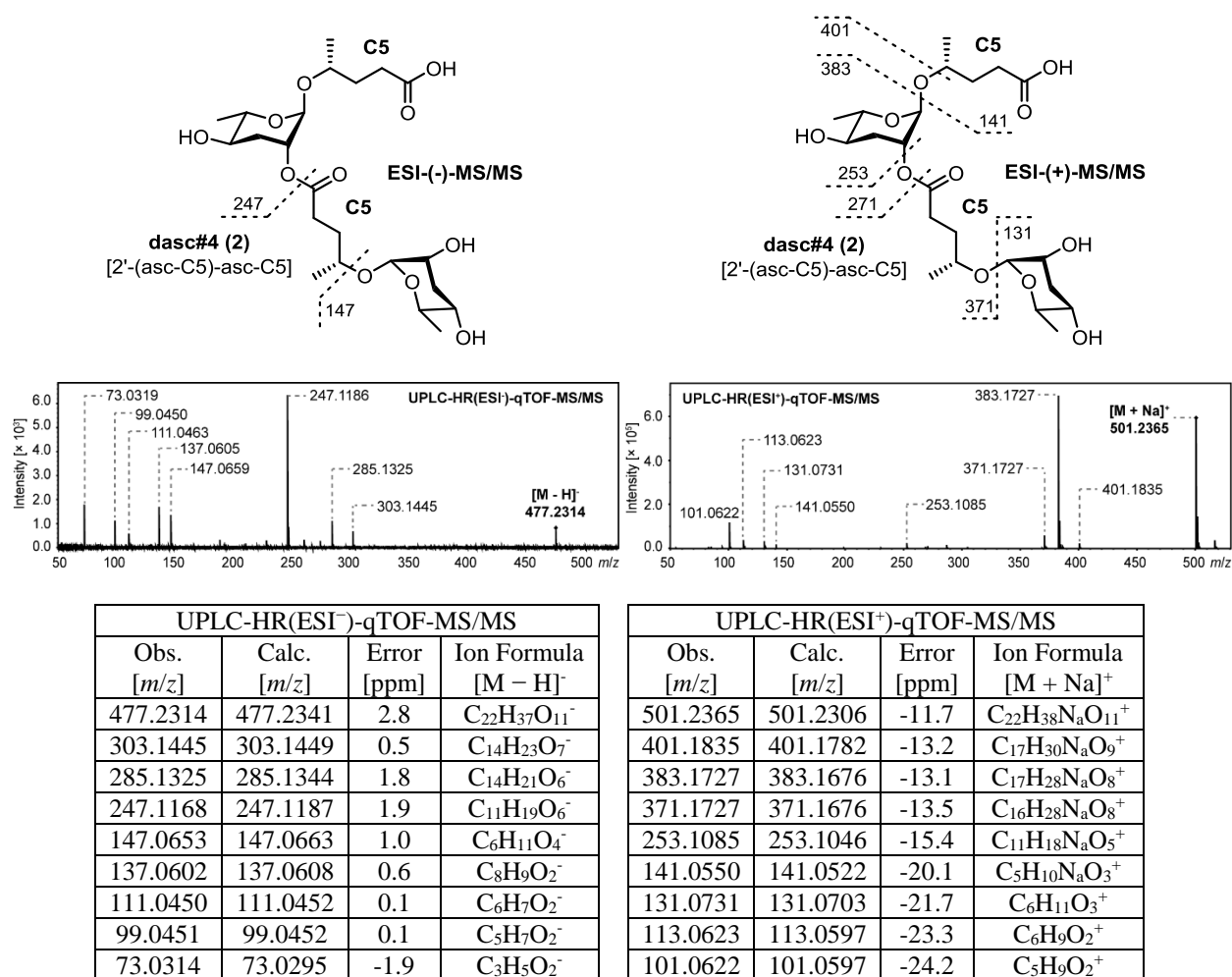


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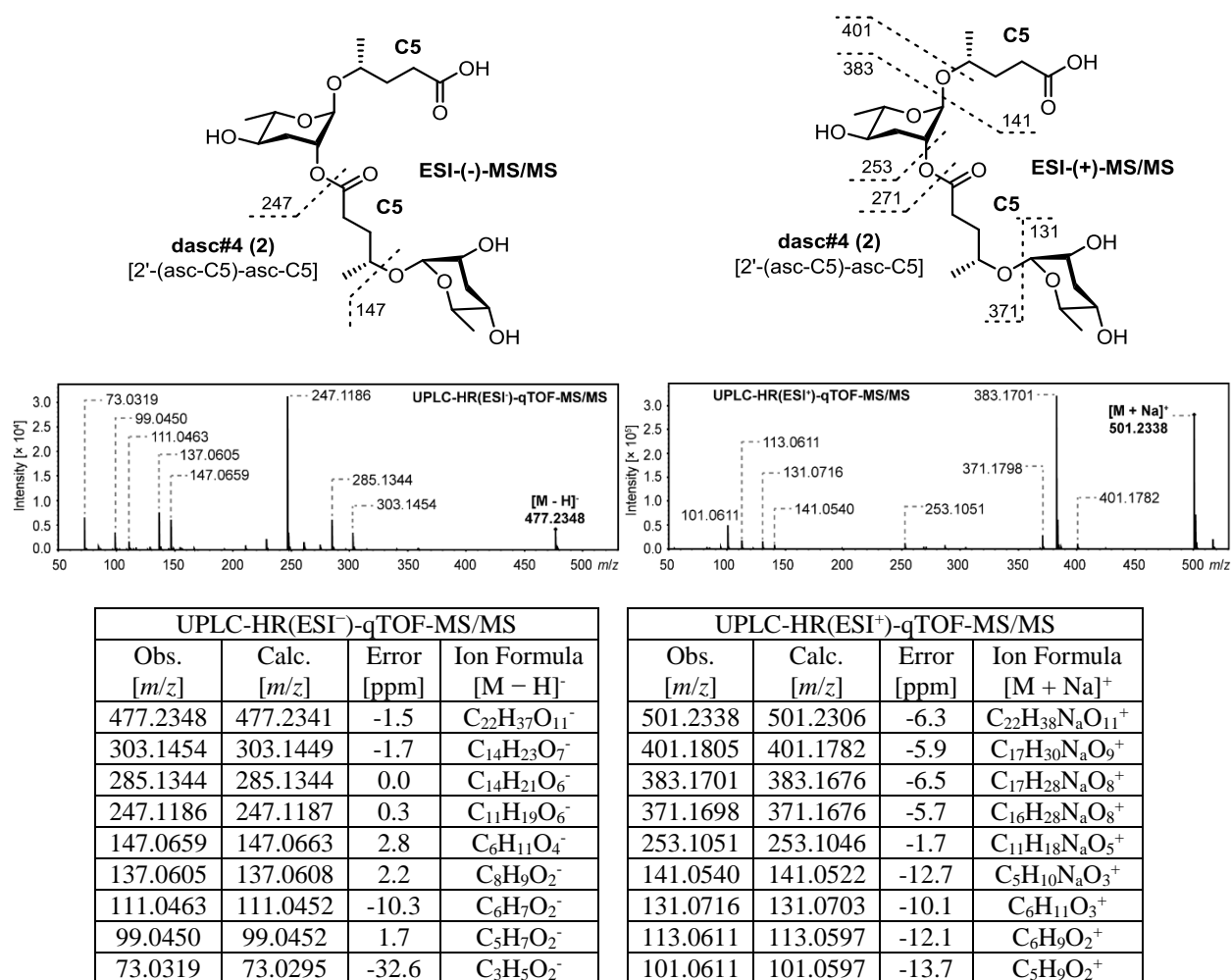
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<b>Figure 5.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of dasc#5 ( <b>3</b> ) from <i>P. mayeri</i> .	S7
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<b>Figure 7.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of dasc#6 [2'-(asc-C6)-asc-C5, <b>4</b> ].	S9
<b>Figure 8.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of dasc#9 [2'-(asc-C7)-asc-C5].	S10
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<b>Figure 11.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of dasc#1 [4'-(asc-C7)-asc-C7].	S13
<b>Figure 12.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of dasc#2 [2'/4'-(asc-C4)-asc-C4].	S14
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<b>Figure 17.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of dasc#14 [4'-(asc- $\Delta$ C7)-asc-C5, <b>6</b> ].	S19
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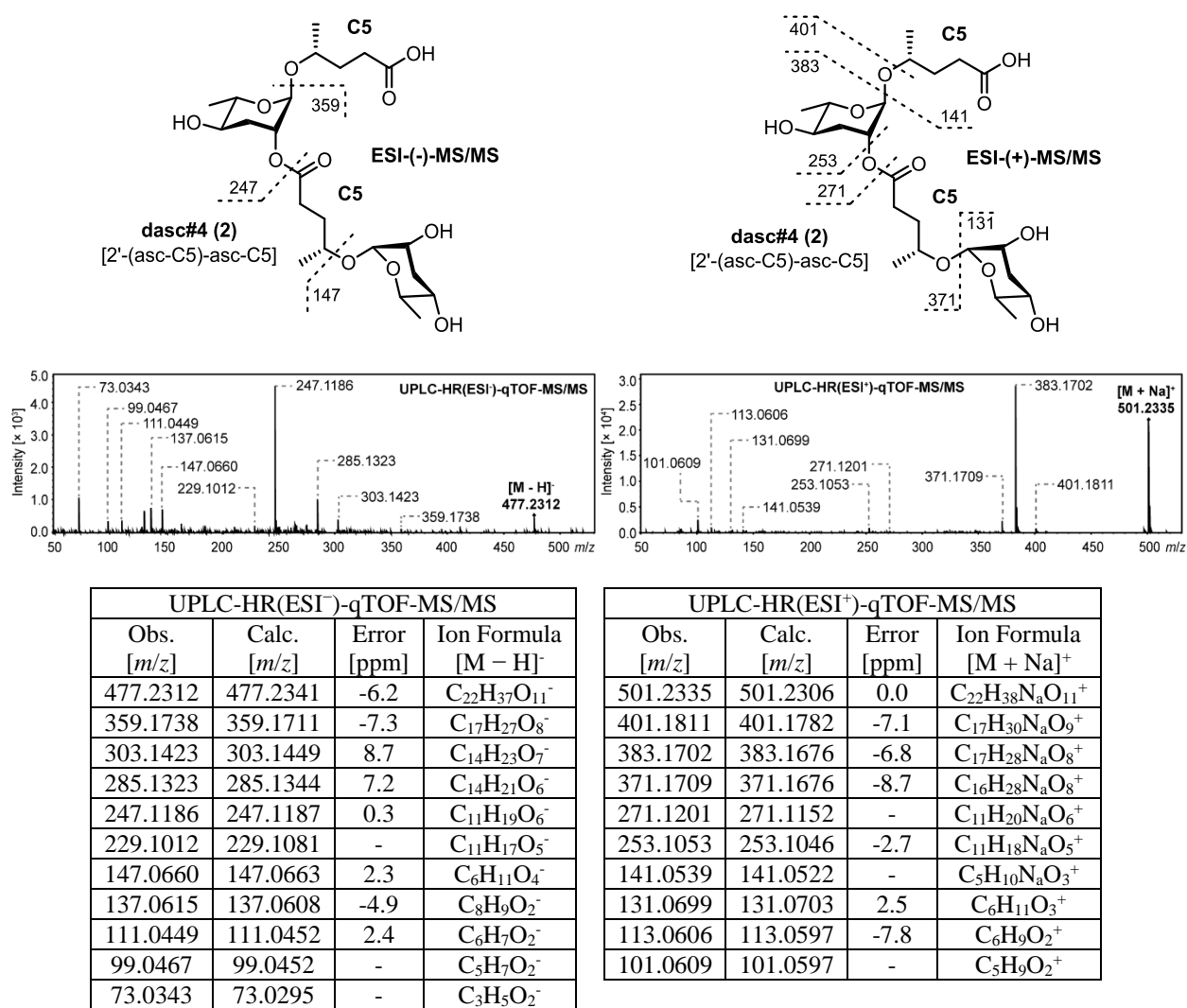
**Figure 1.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#3 [2'-(asc-C5)-asc-C4, **1**] in both negative and positive ion modes. MS/MS fragmentation in positive ion mode produced an intensive ion signal of C<sub>17</sub>H<sub>28</sub>NaO<sub>8</sub><sup>+</sup> ( $m/z = 383.1667$ ) suggesting that the first ascaroside (LC-MS/MS fragmentation pattern of dimeric ascarosides were described in **Figure 4 – figure supplement 1C**) in dasc#3 [2'-(asc-C5)-asc-C4, **1**] should be ascr#9 [asc-C5] or oasc#9 [asc-ωC5] (*von Reuss et al., 2012*). *dqf*-COSY spectrum of an HPLC enriched sample containing dasc#3 [2'-(asc-C5)-asc-C4, **1**] confirmed that a unit of ascr#9 [asc-C5] was linked the 2'-position of ascr#11 [asc-C4] (*supplementary file 1b – Figure 1*).



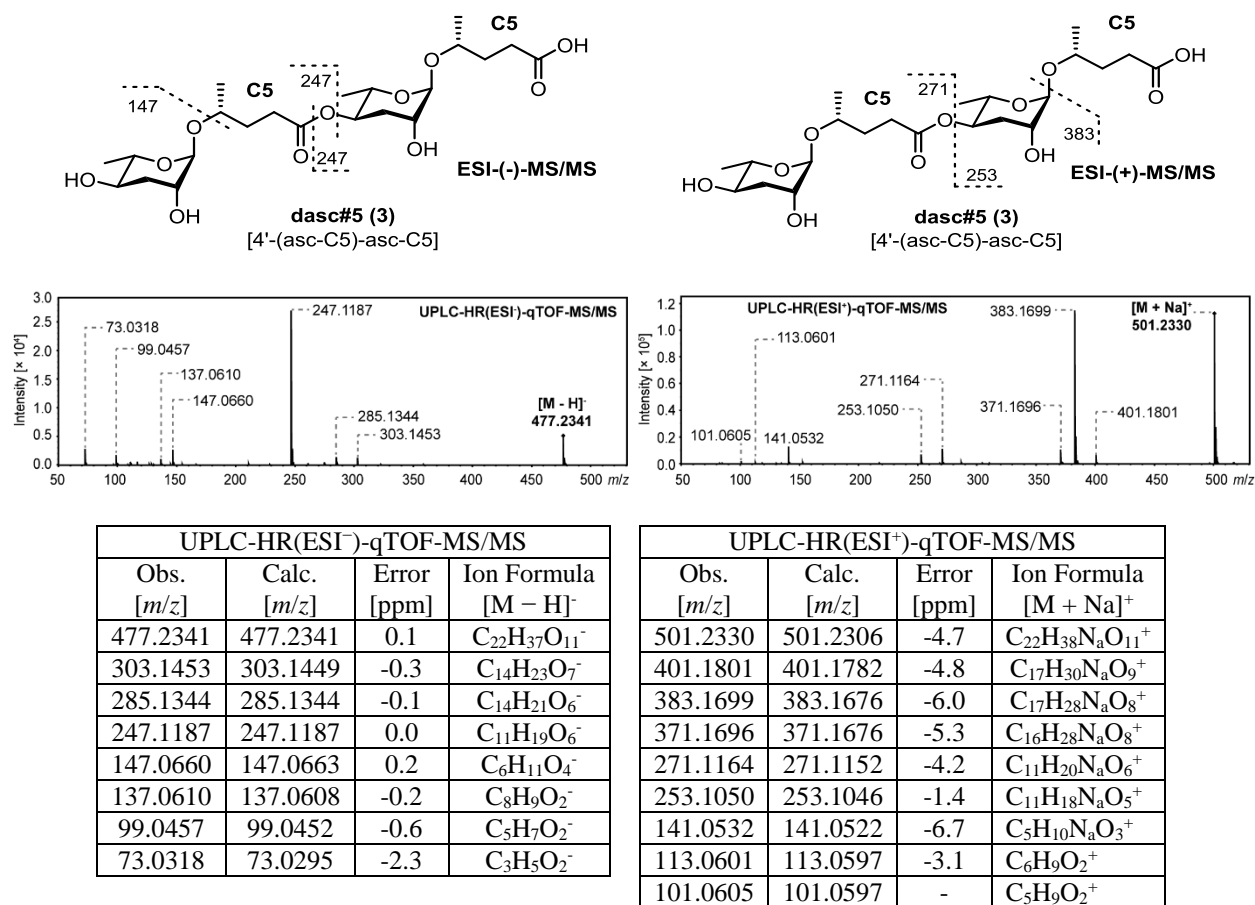
**Figure 2.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#4 [2'-(asc-C5)-asc-C5, 2] from *P. taiwanensis* in both negative and positive ion modes.



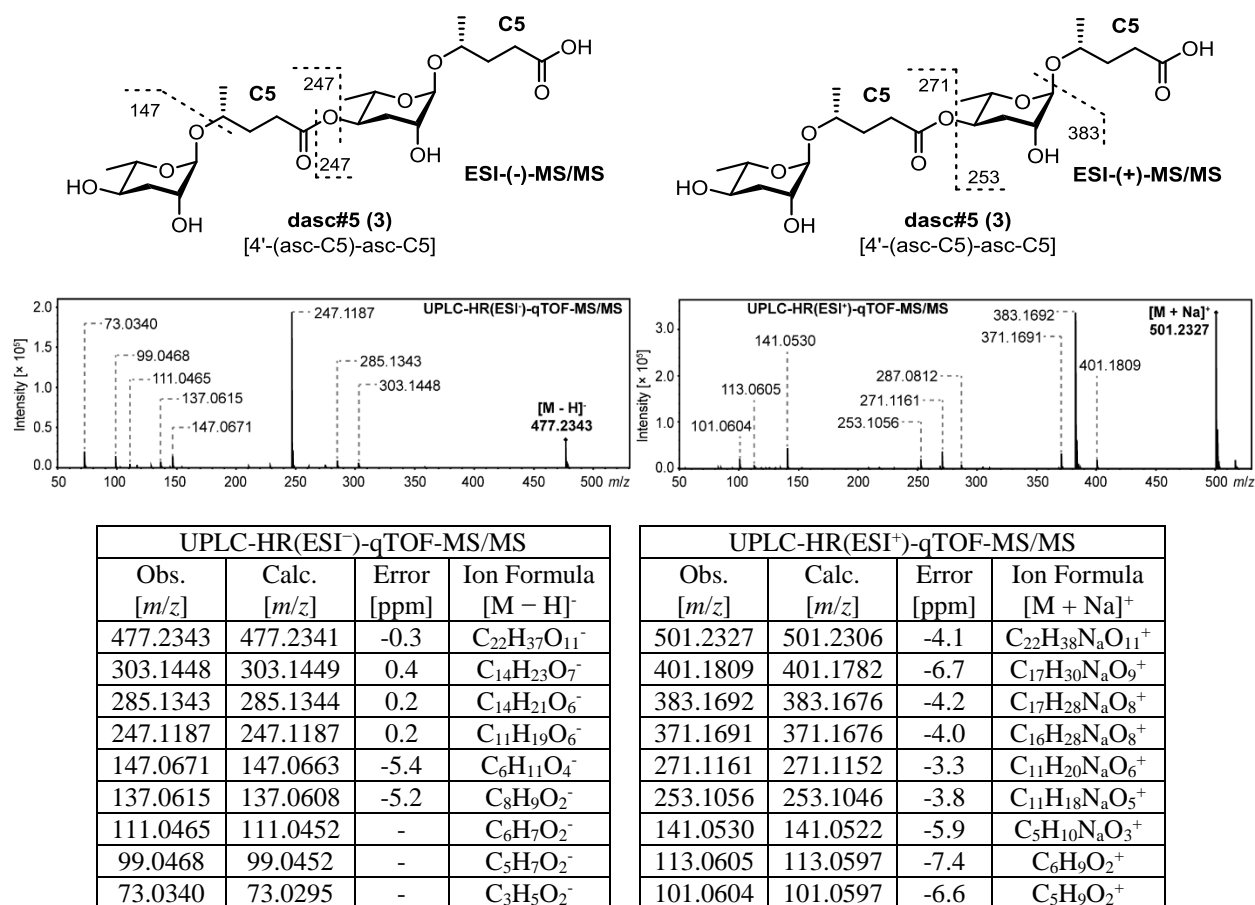
**Figure 3.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#4 [2'-(asc-C5)-asc-C5, 2] from *P. entomophagus* in both negative and positive ion modes.



**Figure 4.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#4 [2'-(asc-C5)-asc-C5, 2] from *P. pacificus* RS2333 in both negative and positive ion modes.

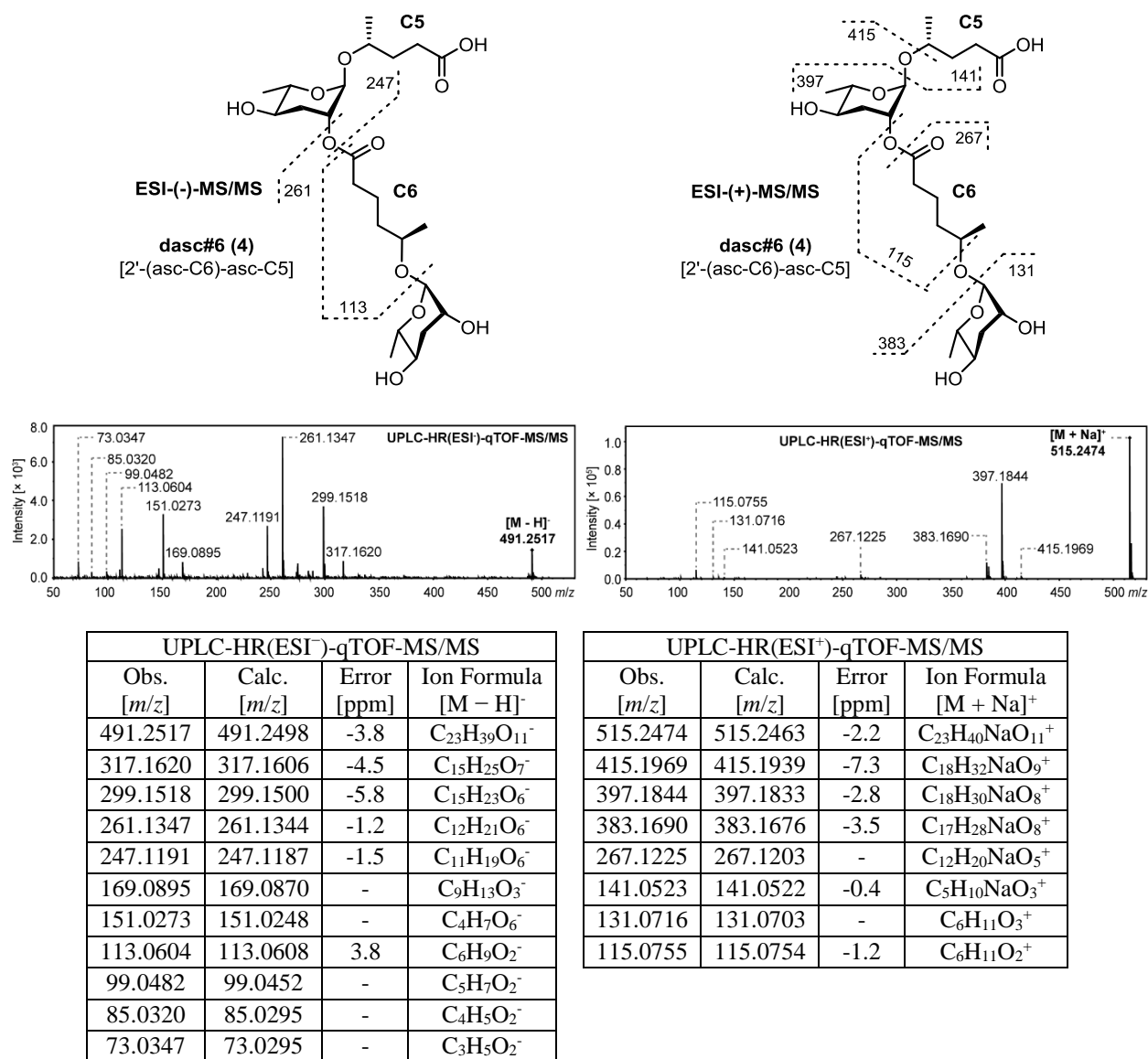


**Figure 5.** UPLC-HR(ESI<sup>-/+</sup>)-qTOF-MS/MS mass spectral data of dasc#5 [4'-(asc-C5)-asc-C5, 3] from *P. mayeri* in both negative and positive ion modes.

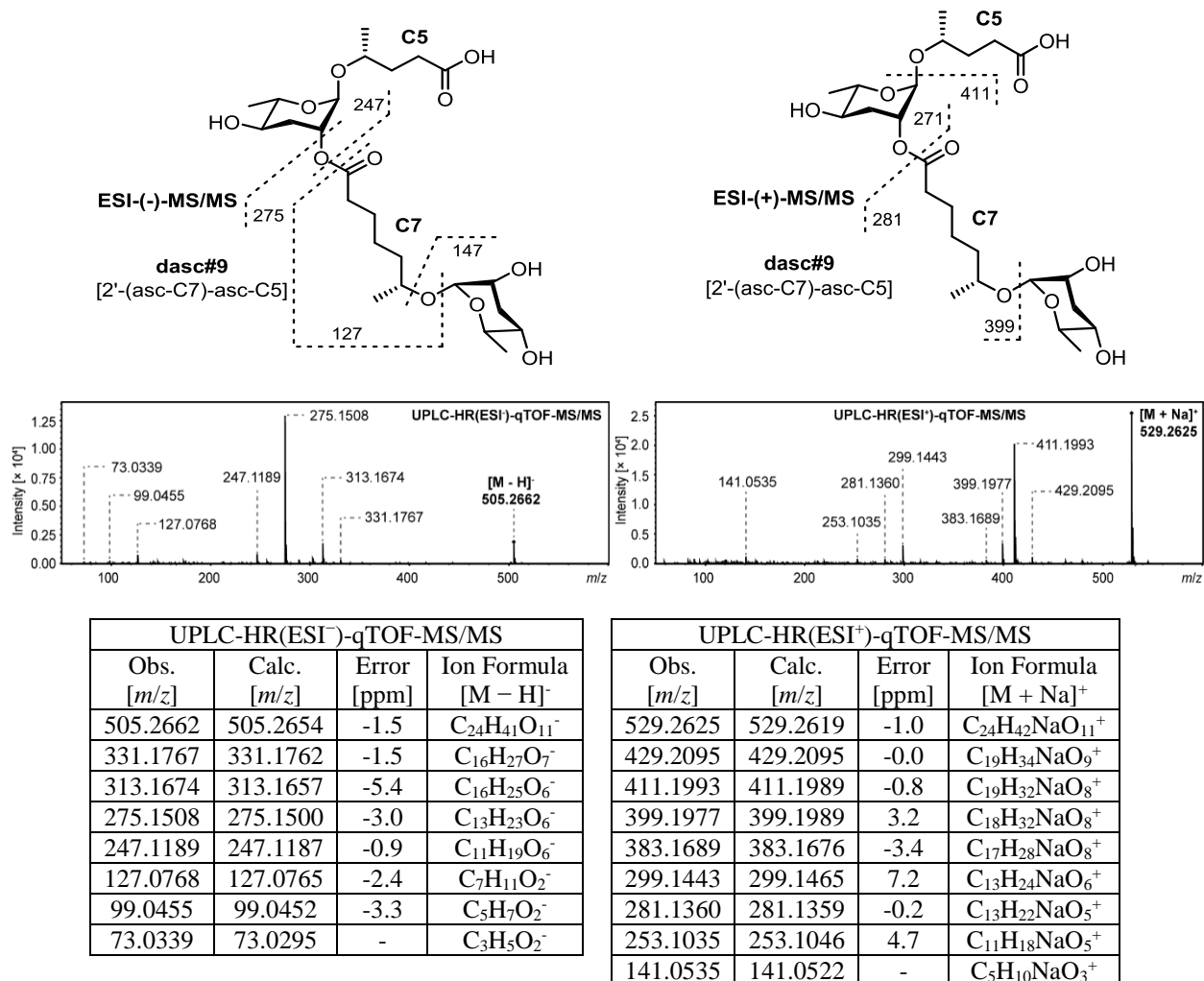


**Figure 6.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#5 [4'-(asc-C5)-asc-C5, 3] from *D. magnus* in both negative and positive ion modes.

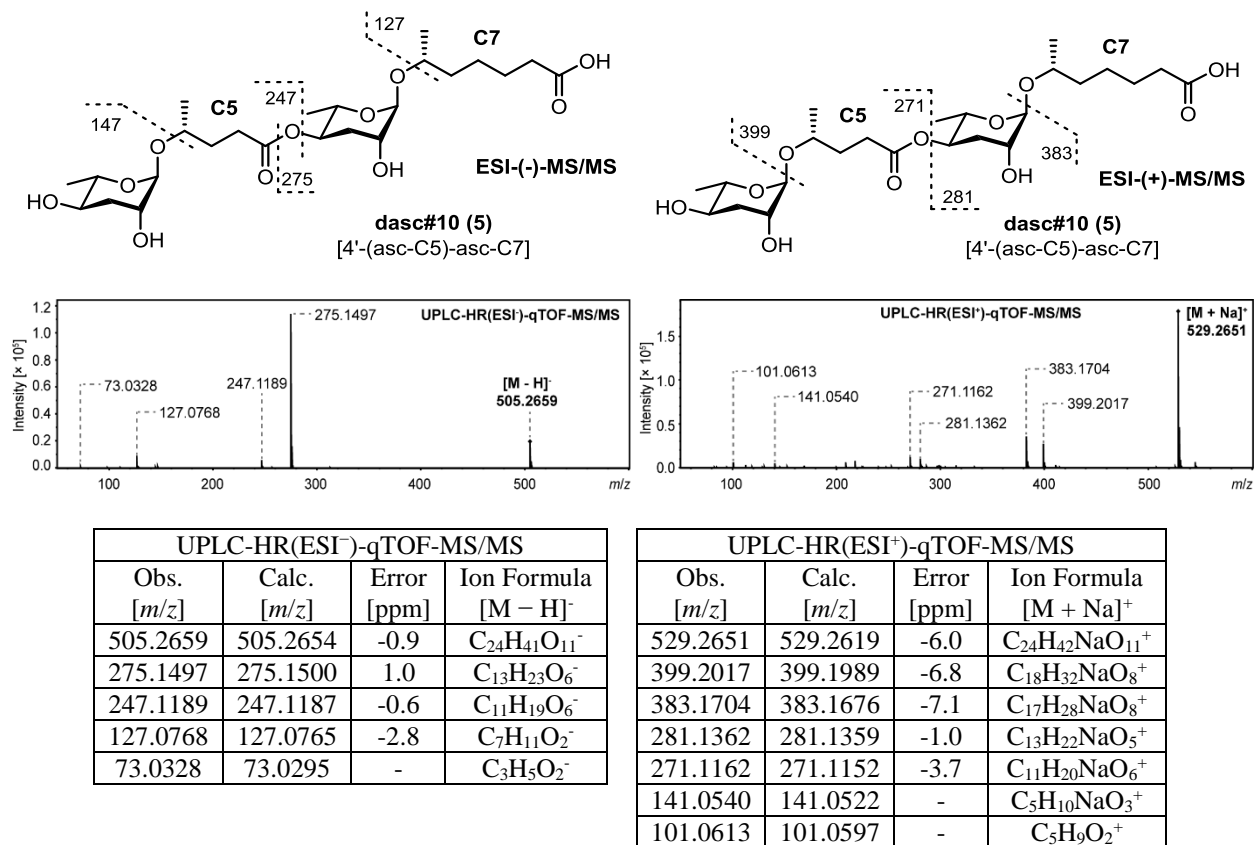




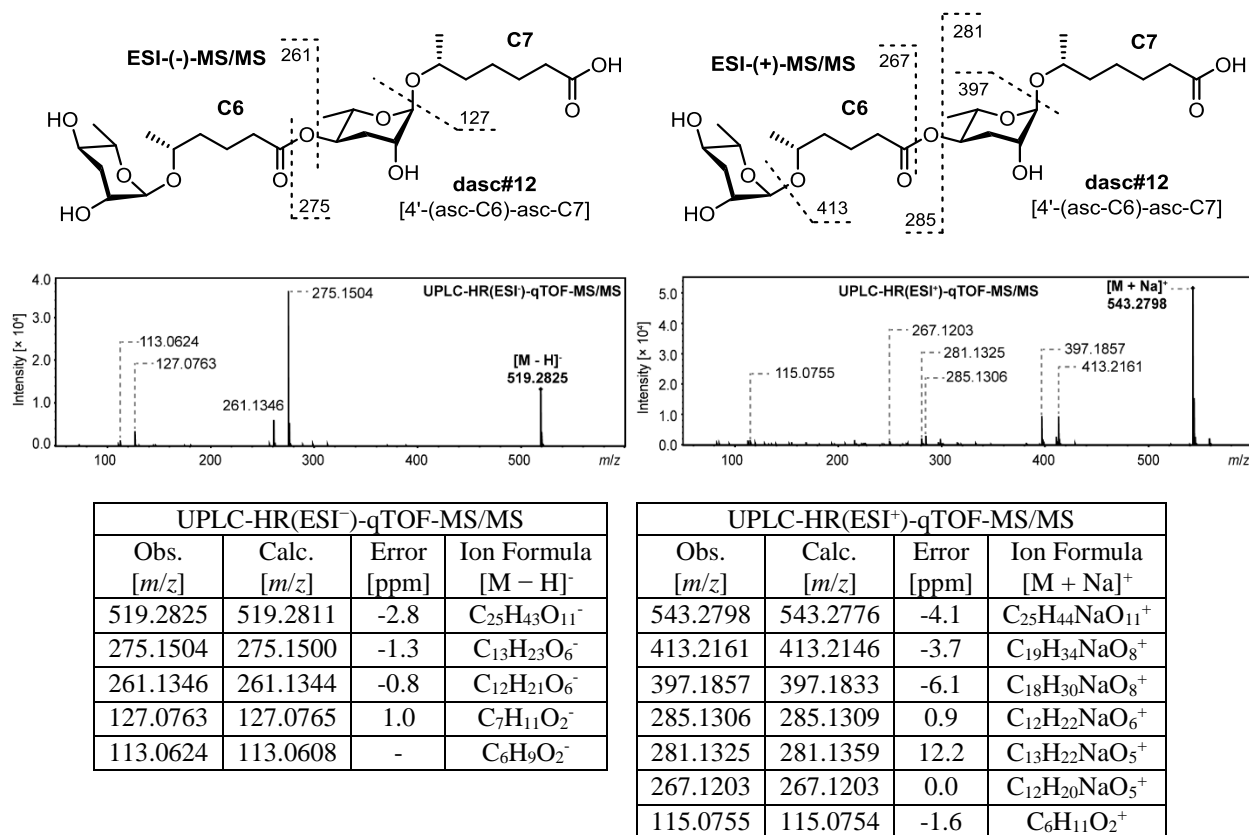
**Figure 7.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#6 [2'-(asc-C6)-asc-C5, 4] in both negative and positive ion modes.



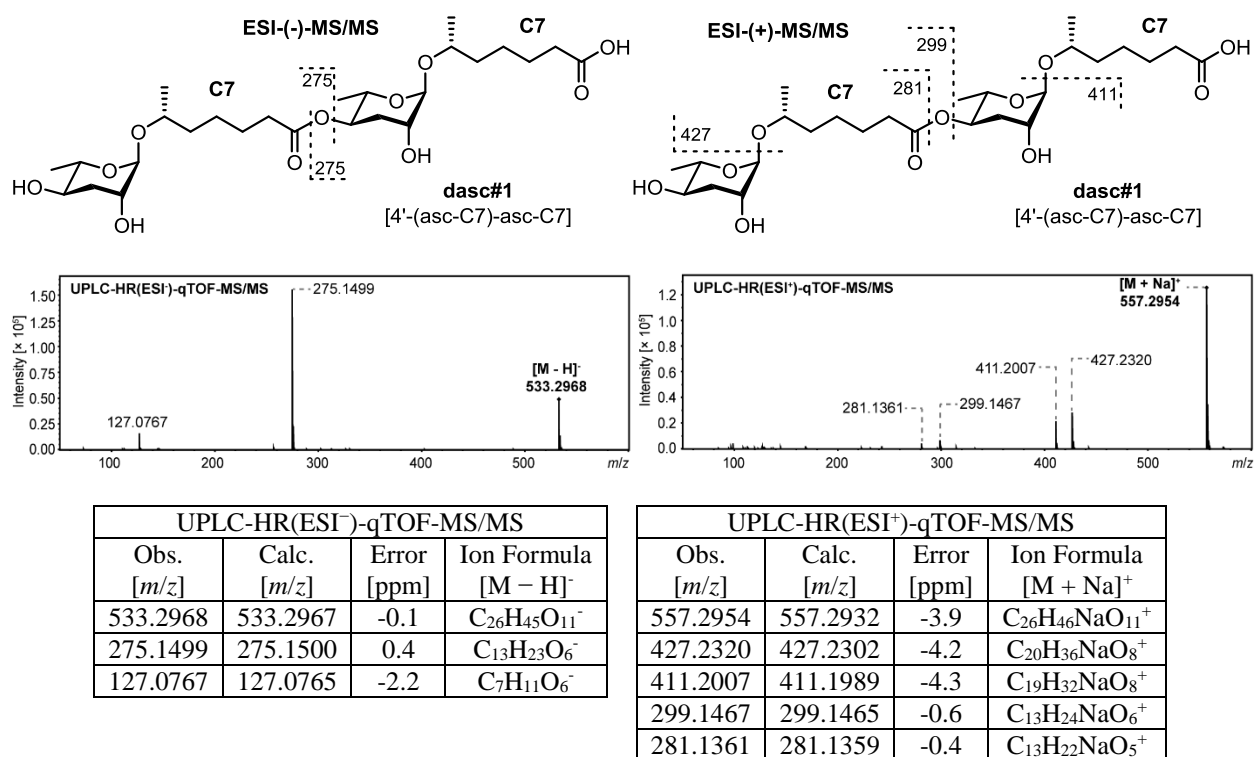
**Figure 8.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#9 [2'-(asc-C7)-asc-C5] in both negative and positive ion modes. Note that dasc#9 [2'-(asc-C7)-asc-C5] is tentatively assigned to be linked at the 2'-position, but its final structure has not yet been elucidated.



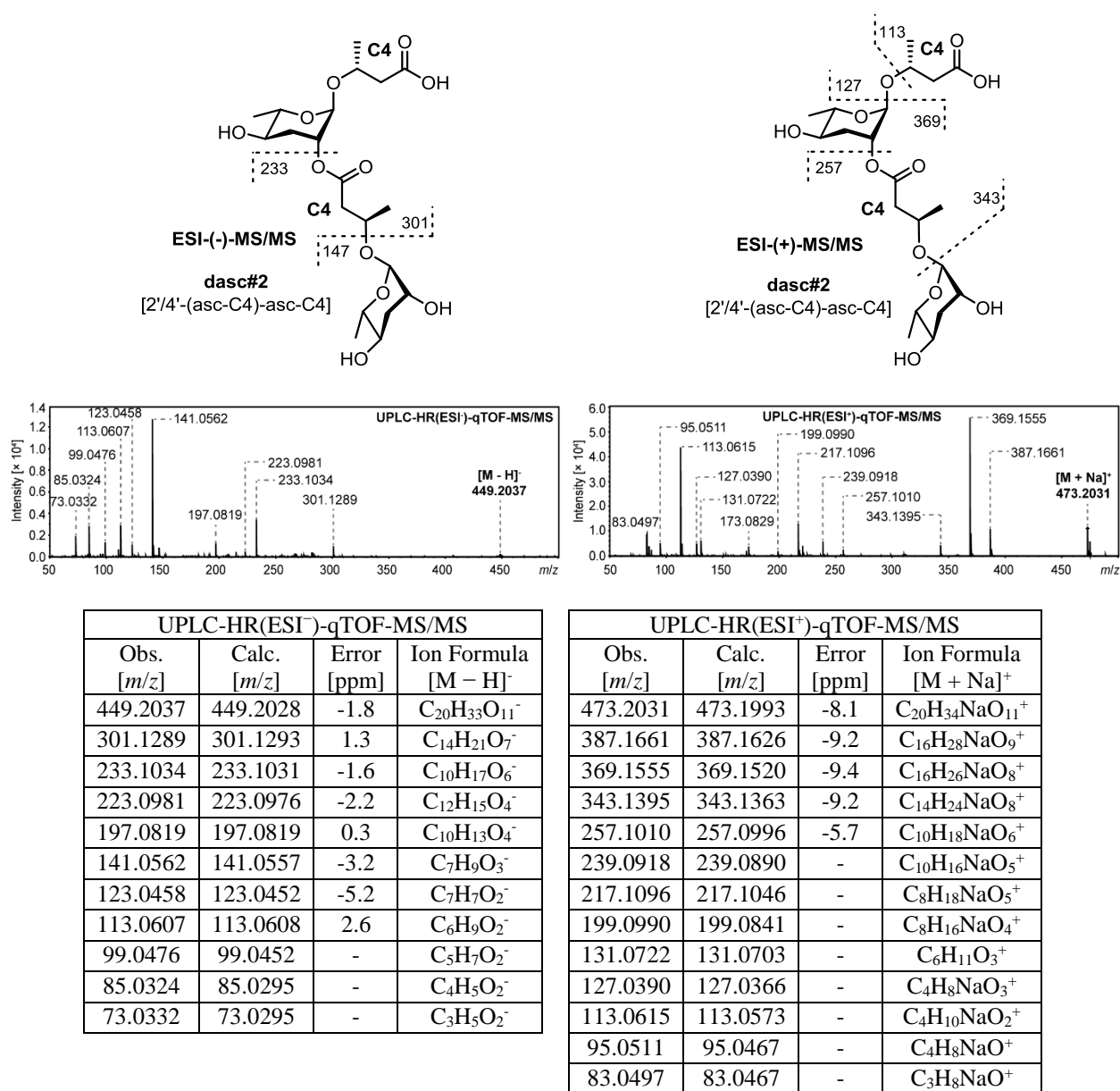
**Figure 9.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#10 [4'-(asc-C5)-asc-C7, **5**] in both negative and positive ion modes.



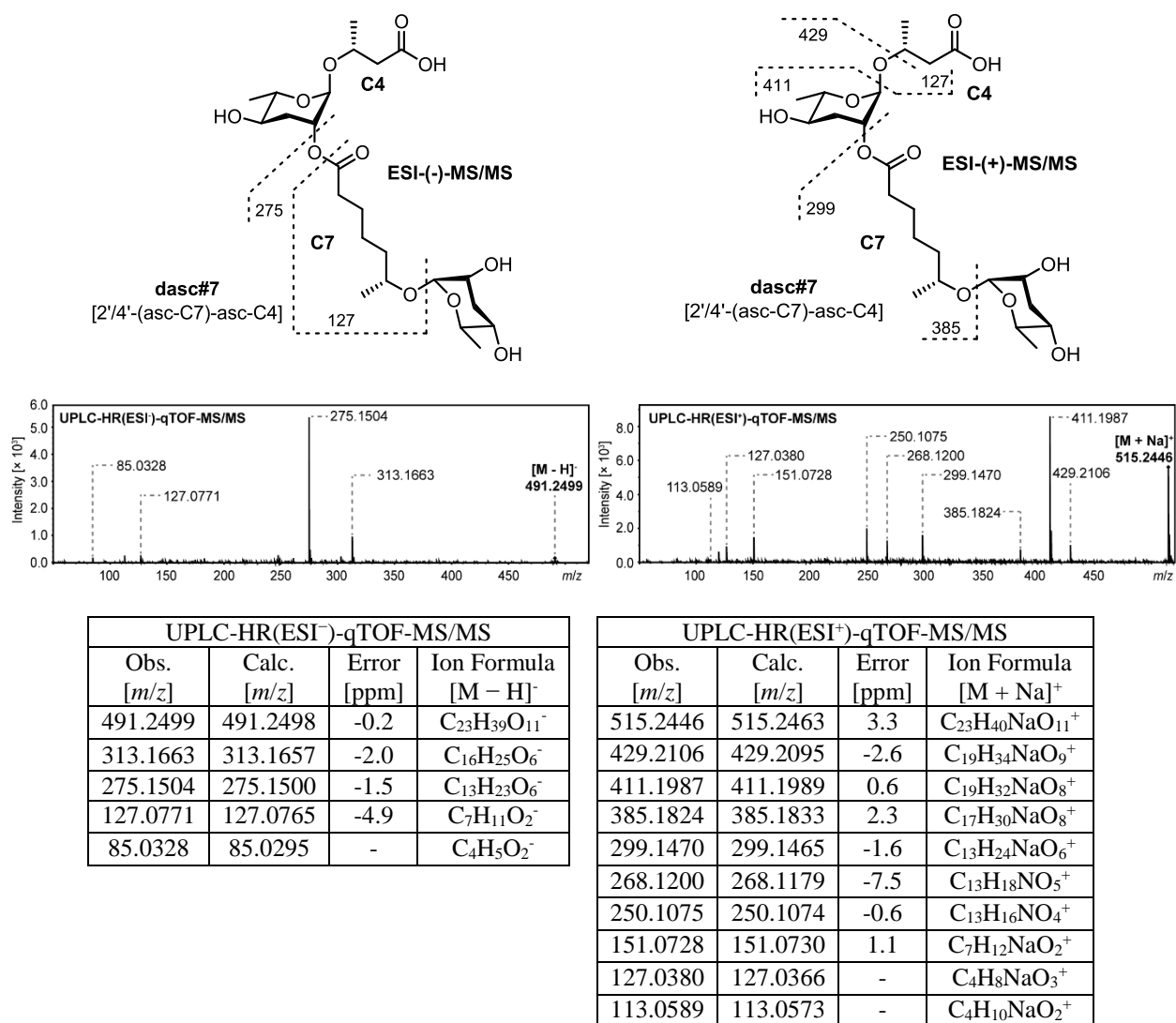
**Figure 10.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#12 [4'-(asc-C6)-asc-C7] in both negative and positive ion modes. Note that dasc#12 [4'-(asc-C6)-asc-C7] is tentatively assigned to be linked at the 4'-position, but its absolute structure has not yet been elucidated.



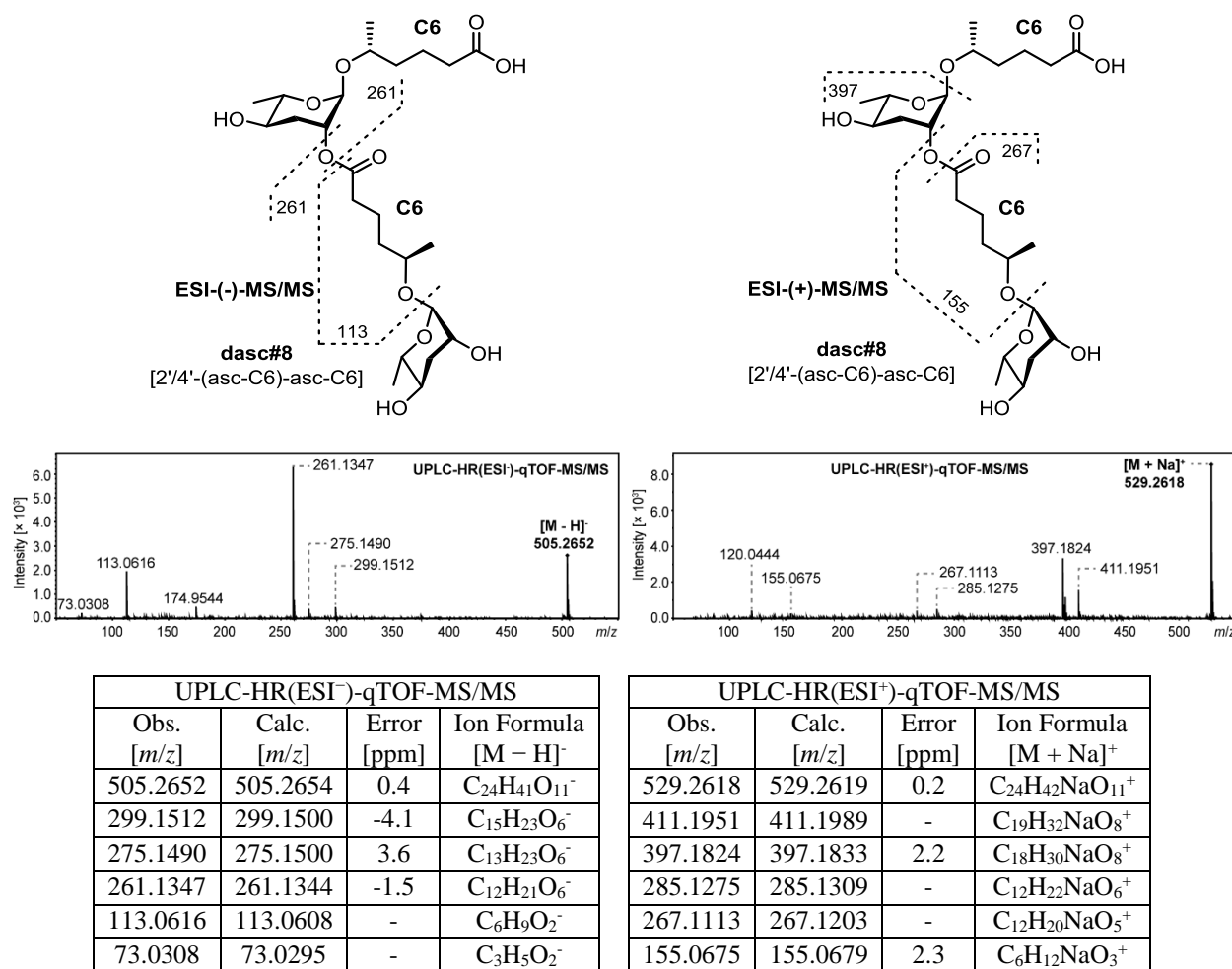
**Figure 11.** UPLC-HR(ESI<sup>-/+</sup>)-qTOF-MS/MS mass spectral data of known dasc#1 [4'-(asc-C7)-asc-C7] (Bose *et al.*, 2012) in both negative and positive ion modes.



**Figure 12.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#2 [2'/4'-(asc-C4)-asc-C4] in both negative and positive ion modes. Note that dasc#2 [2'/4'-(asc-C4)-asc-C4] is proposed and illustrated to be modified at the 2'-position, but its absolute structure has not yet been elucidated.

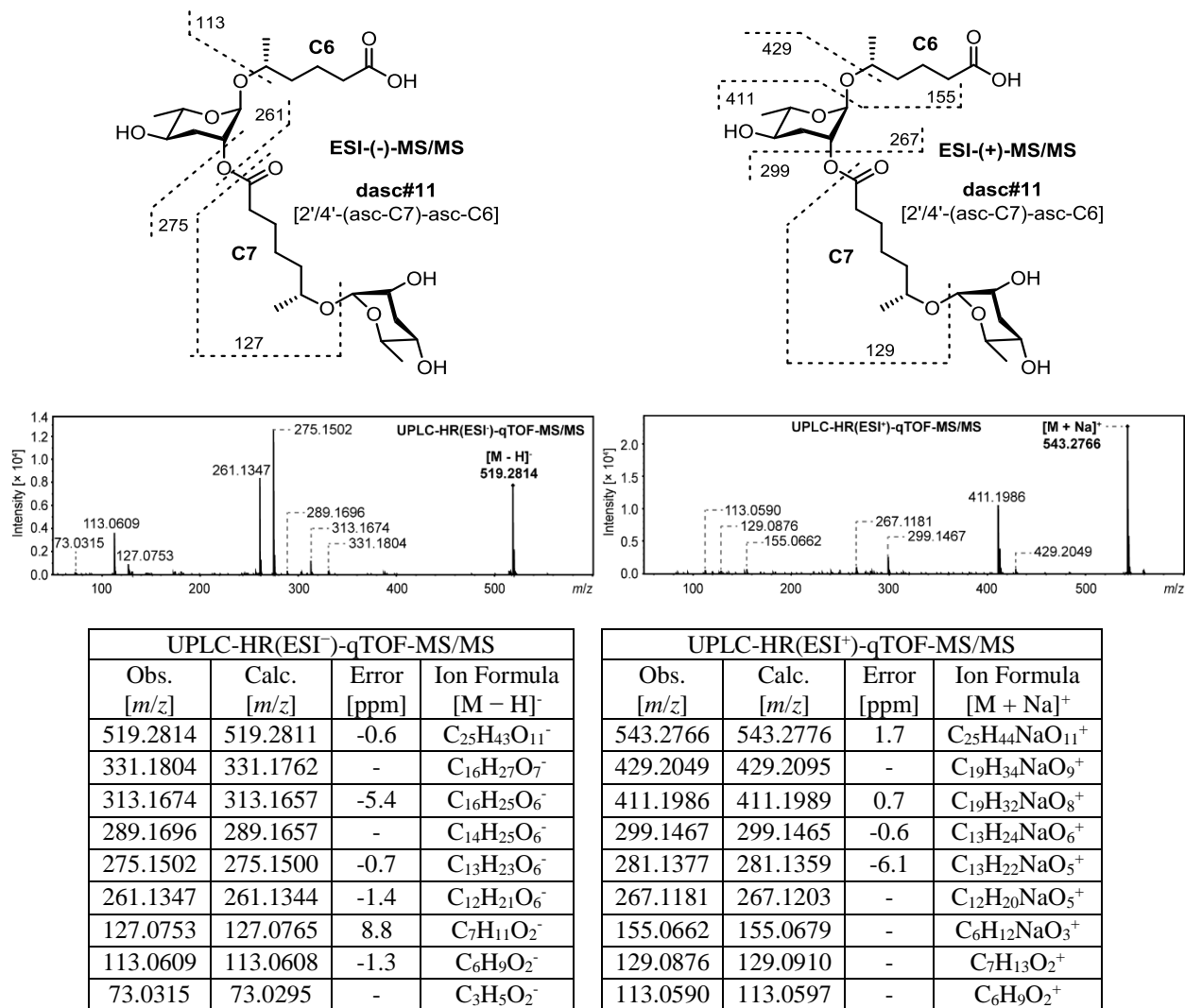


**Figure 13.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#7 [2'/4'-(asc-C7)-asc-C4] in both negative and positive ion modes. Note that dasc#7 [2'/4'-(asc-C7)-asc-C4] is tentatively assigned to be linked at the 2'-position, but its absolute structure has not yet been elucidated.

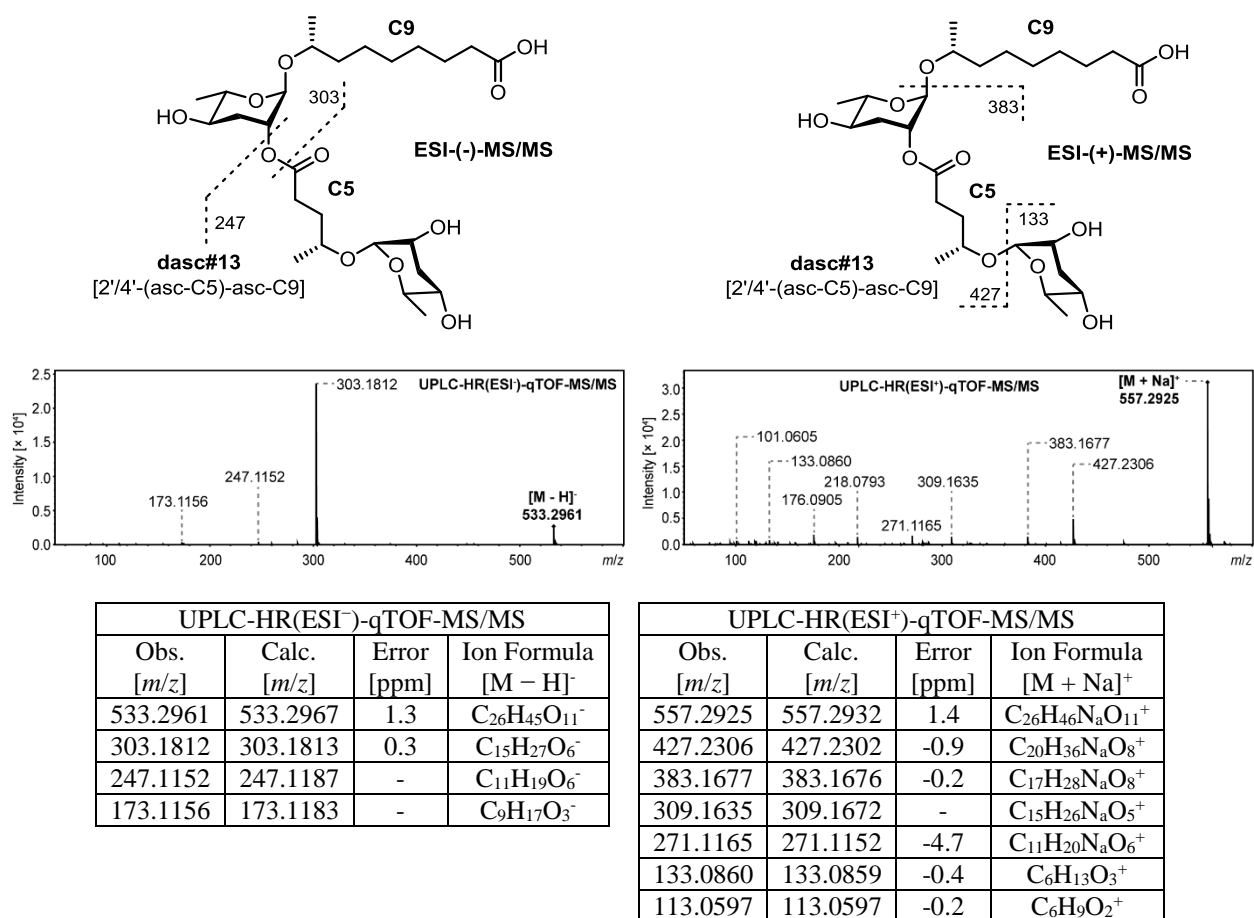


**Figure 14.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of dasc#8 [2'/4'-(asc-C6)-asc-C6] in both negative and positive ion modes. Note that dasc#8 [2'/4'-(asc-C6)-asc-C6] is tentatively assigned to be linked at the 2'-position, but its absolute structure has not yet been elucidated.

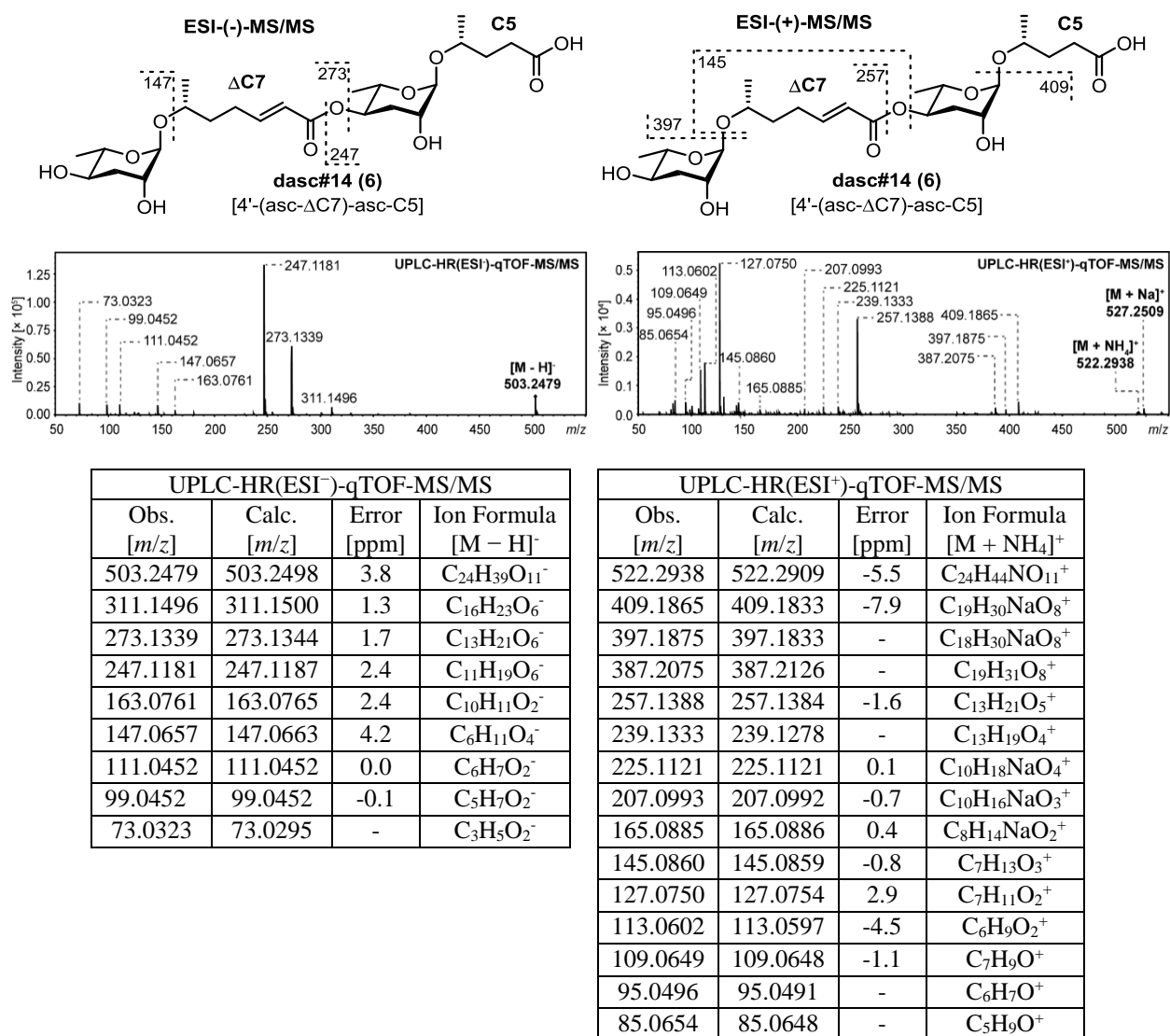




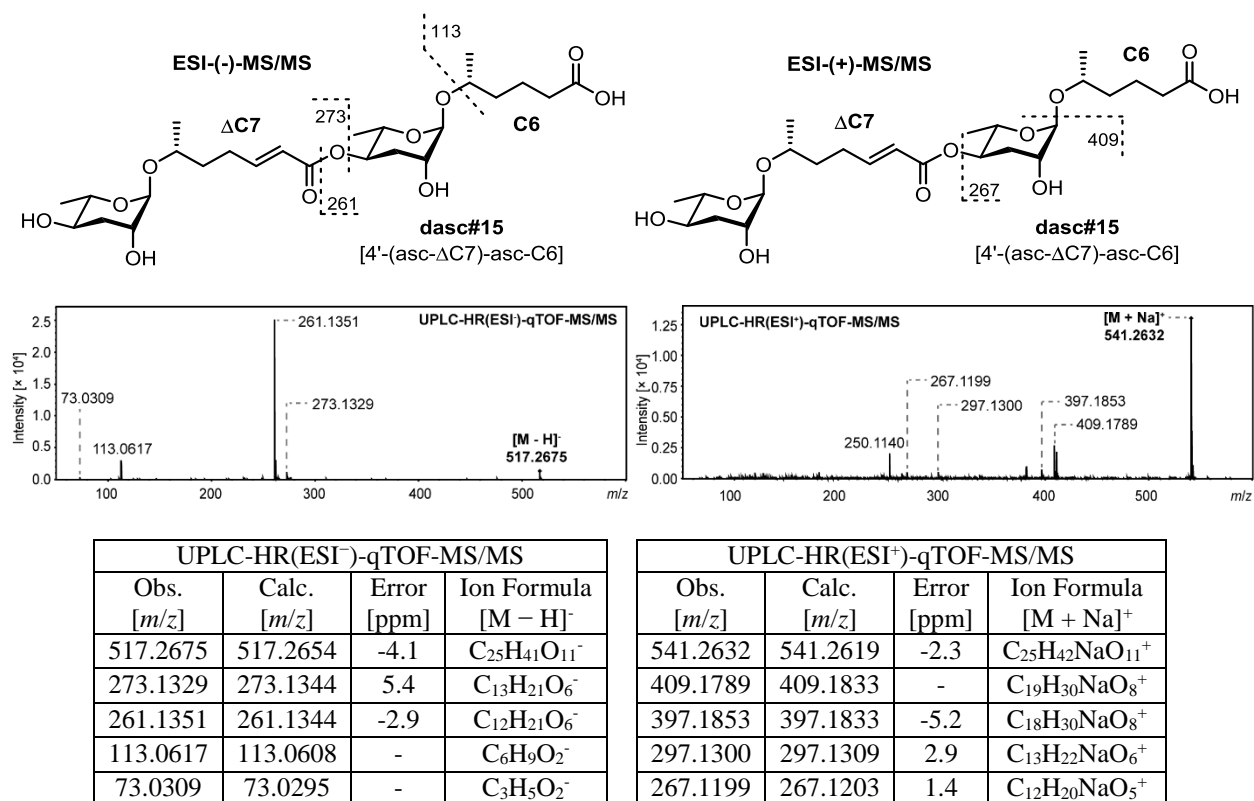
**Figure 15.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of dasc#11 [2'/4'-(asc-C7)-asc-C6] in both negative and positive ion modes. Note that dasc#11 [2'/4'-(asc-C7)-asc-C6] is tentatively assigned to be linked at the 2'-position, but its absolute structure has not yet been elucidated.



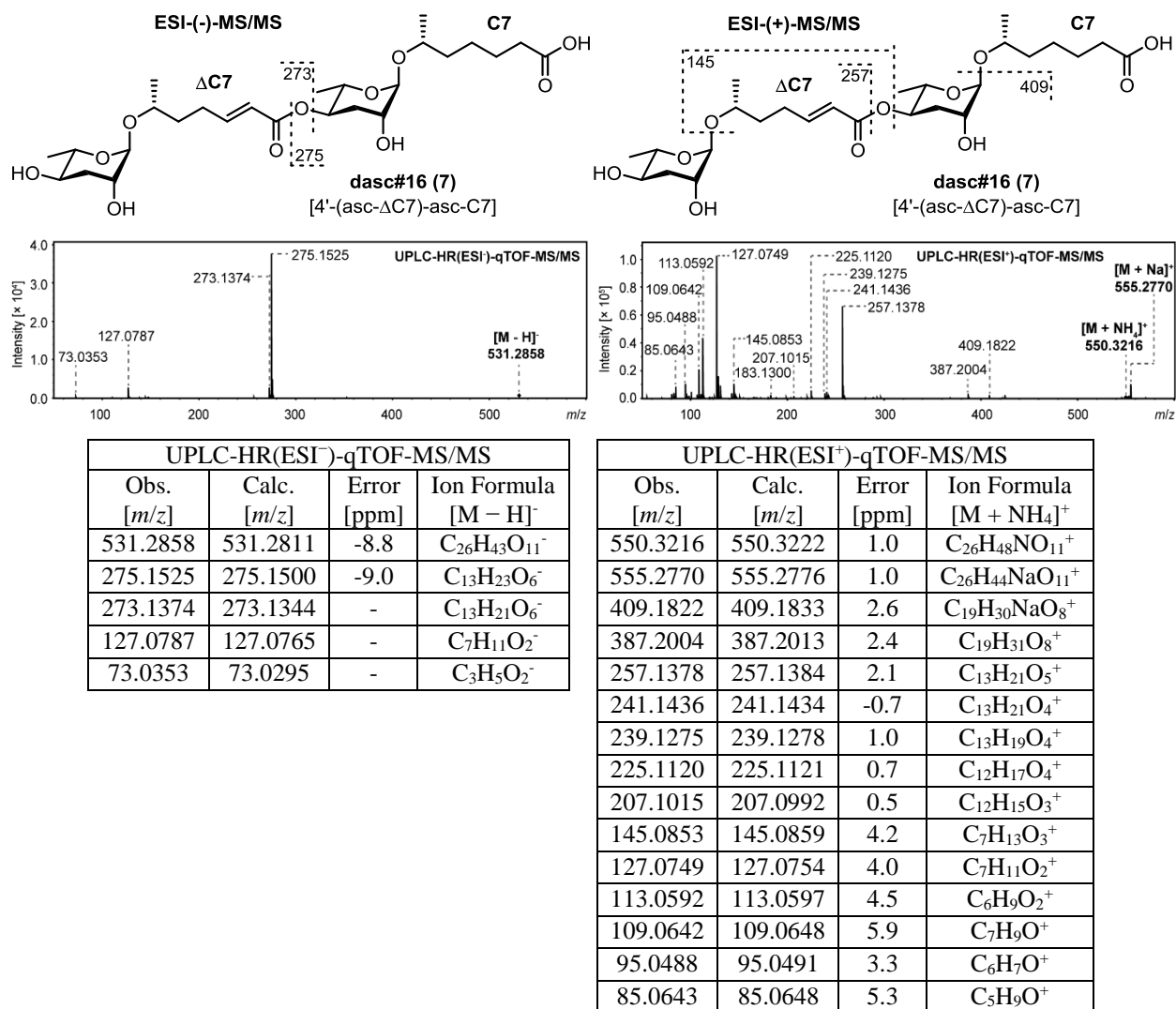
**Figure 16.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of dasc#13 [2'/4'-(asc-C5)-asc-C9] in both negative and positive ion modes. Note that dasc#13 [2'/4'-(asc-C5)-asc-C9] is tentatively assigned to be linked at the 2'-position, but its absolute structure has not yet been elucidated.



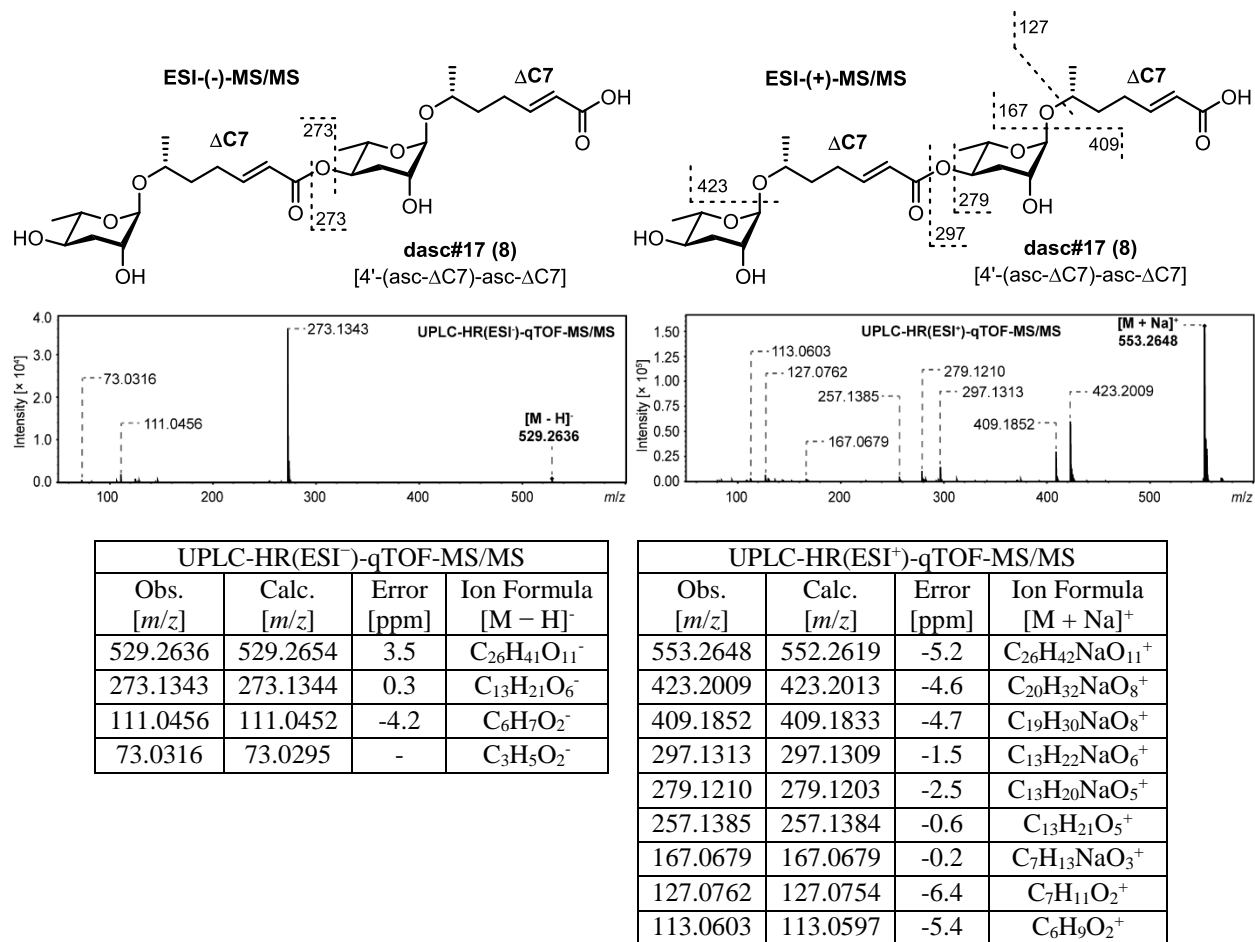
**Figure 17.** UPLC-HR(ESI<sup>+/−</sup>)-qTOF-MS/MS mass spectral data of dasc#14 [4'-(asc-ΔC7)-asc-C5, **6**] in both negative and positive ion modes.



**Figure 18.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of dasc#15 [4'-(asc-ΔC7)-asc-C6] in both negative and positive ion modes. Note that dasc#15 [4'-(asc-ΔC7)-asc-C6] is proposed to be linked at the 4'-position, but its absolute structure has not yet been elucidated.



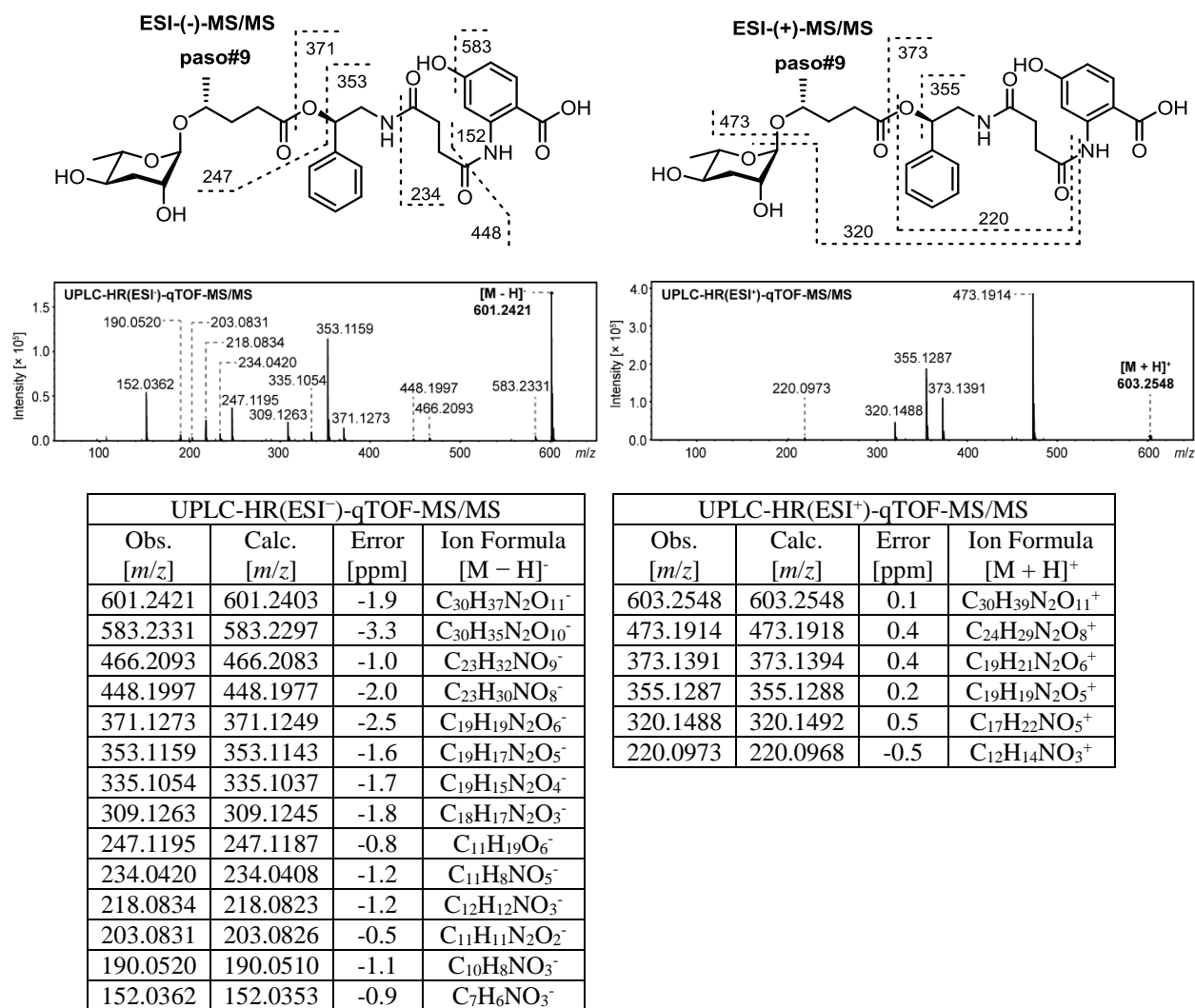
**Figure 19.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of dasc#16 [4'-(asc-ΔC7)-asc-C7, 7] in both negative and positive ion modes.



**Figure 20.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of dasc#17 [4'-(asc- $\Delta$ C7)-asc- $\Delta$ C7, **8**] in both negative and positive ion modes.

**supplementary file 2b: MS/MS spectral data of paso#9**  
**Figure 1.** LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of paso#9.

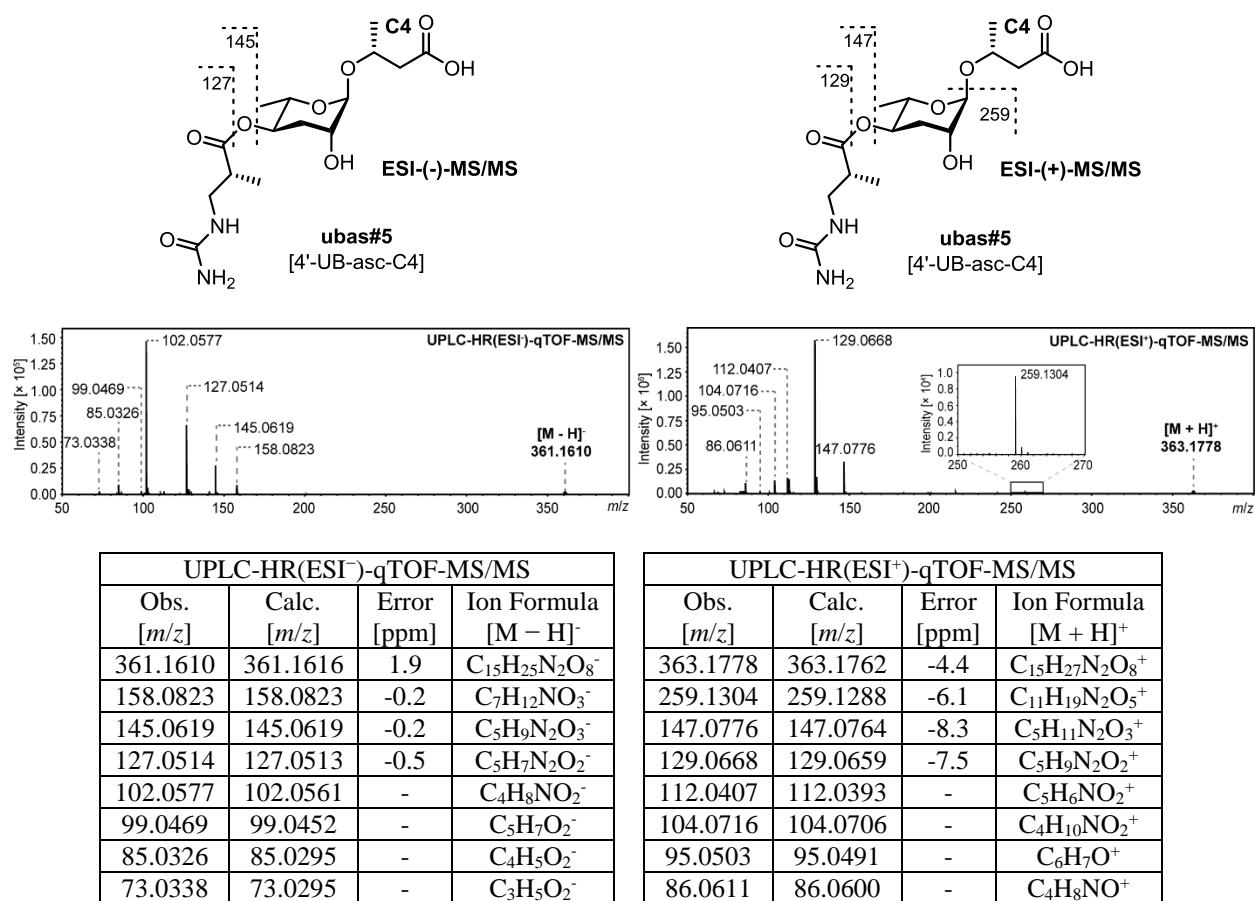
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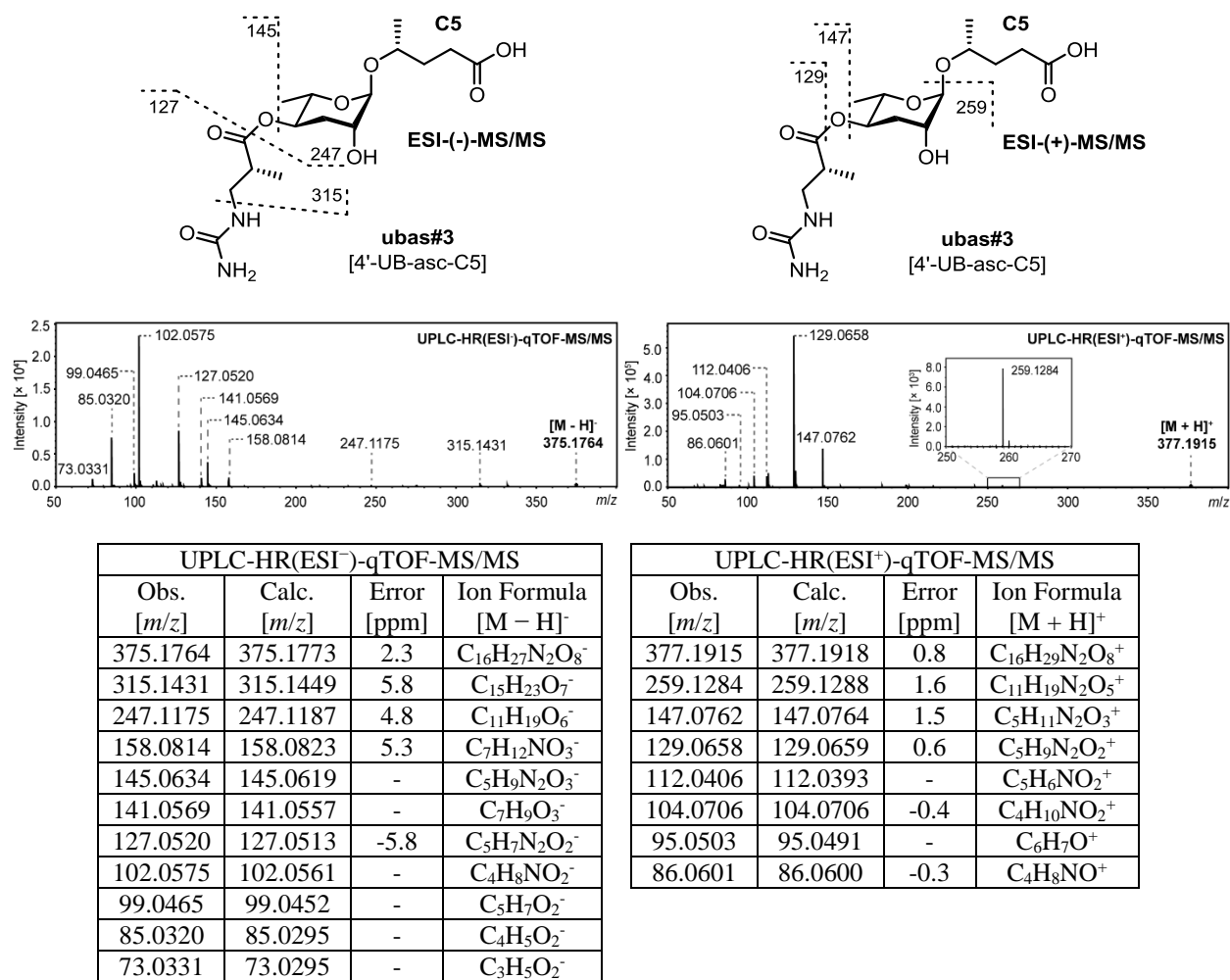
**Figure 1.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of paso#9 in both negative and positive ion modes. Analysis of NMR data of paso#9 suggests a 4-hydroxyanthranilic acid moiety in paso#9 (*supplementary file 1c – Figures 5-13 and supplementary file 3 – Table 14*).



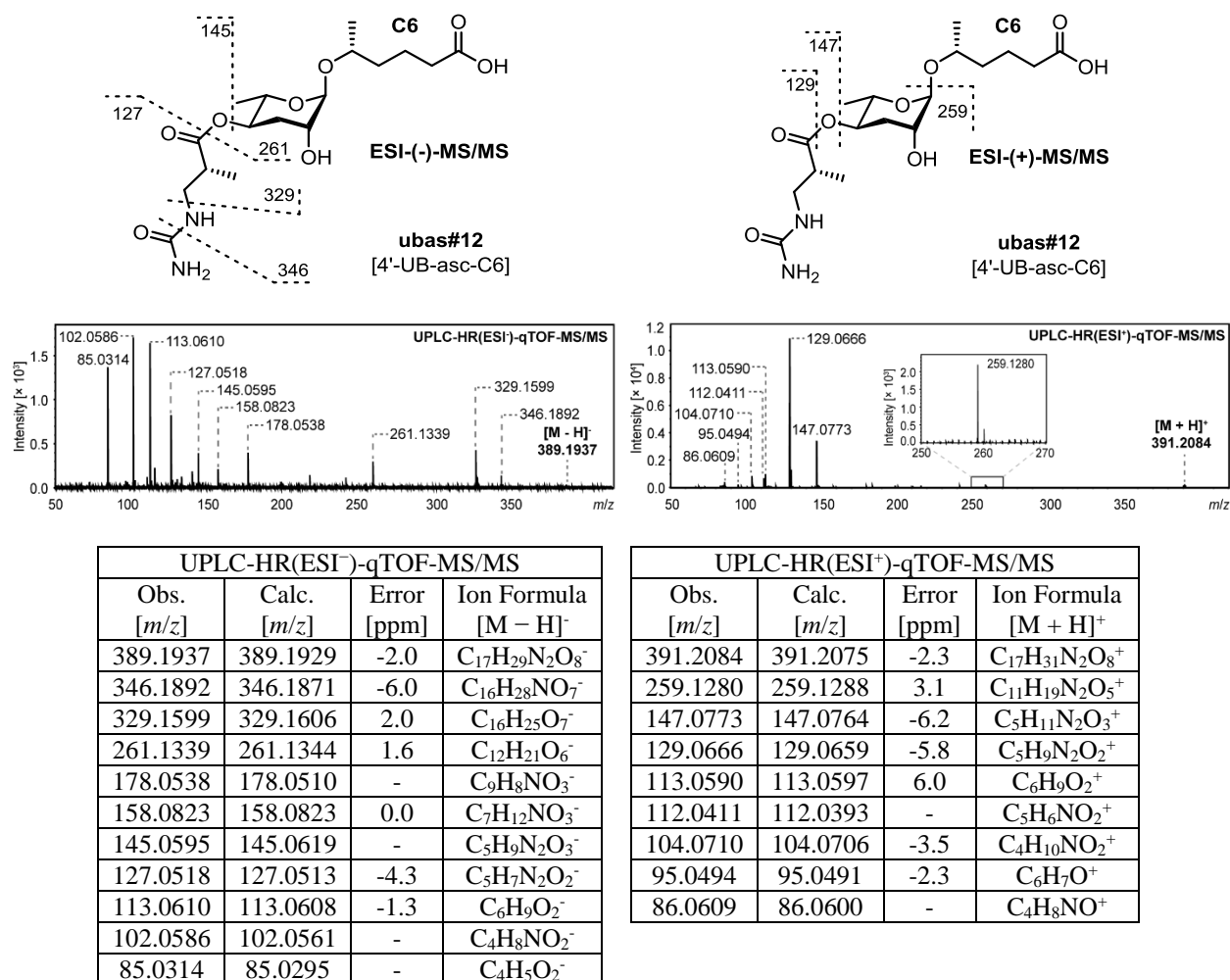
<b>supplementary file 2c: MS/MS spectral data of UBAS chemicals</b>	<b>Pages</b>
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<b>Figure 2.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#3 [4'-UB-asc-C5] from <i>P. pacificus</i> .	S27
<b>Figure 3.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#12 [4'-UB-asc-C6] from <i>P. pacificus</i> .	S28
<b>Figure 4.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#4 [4'-UB-asc-C7] from <i>P. pacificus</i> .	S29
<b>Figure 5.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#1 from <i>P. pacificus</i> .	S30
<b>Figure 6.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#2 from <i>P. pacificus</i> .	S31
<b>Figure 7.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#26 [4'-UB-asc- $\omega$ C3].	S32
<b>Figure 8.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#33 [4'-UB-2'-(asc- $\omega$ C3)-asc-C4].	S33
<b>Figure 9.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#34 [4'-UB-2'-(asc-C4)-asc-C4, <b>11</b> ].	S34
<b>Figure 10.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#35 [4'-UB-2'-(asc-C4)-asc-C5, <b>12</b> ].	S35
<b>Figure 11.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#36 [4'-UB-2'-(asc-C4)-asc-C6].	S36
<b>Figure 12.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#37 [4'-UB-2'-(asc-C4)-asc-C7].	S37
<b>Figure 13.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#27 [4'-UB-2'-(asc-C5)-asc- $\omega$ C3].	S38
<b>Figure 14.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#28 [4'-UB-2'-(asc-C5)-asc-C4, <b>13</b> ].	S39
<b>Figure 15.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#1 from <i>P. taiwanensis</i> .	S40
<b>Figure 16.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#1 from <i>P. laevicollis</i> .	S41
<b>Figure 17.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#30 ( <b>9</b> ) from <i>P. maxplancki</i> .	S42
<b>Figure 18.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#30 ( <b>9</b> ) from <i>P. fukushimae</i> .	S43
<b>Figure 19.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#29 [4'-UB-2'-(asc-C6)-asc-C4].	S44
<b>Figure 20.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#31 [4'-UB-2'-(asc-C6)-asc-C6].	S45
<b>Figure 21.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of ubas#32 [4'-UB-2'-(asc-C5)-asc-C7, <b>10</b> ].	S46



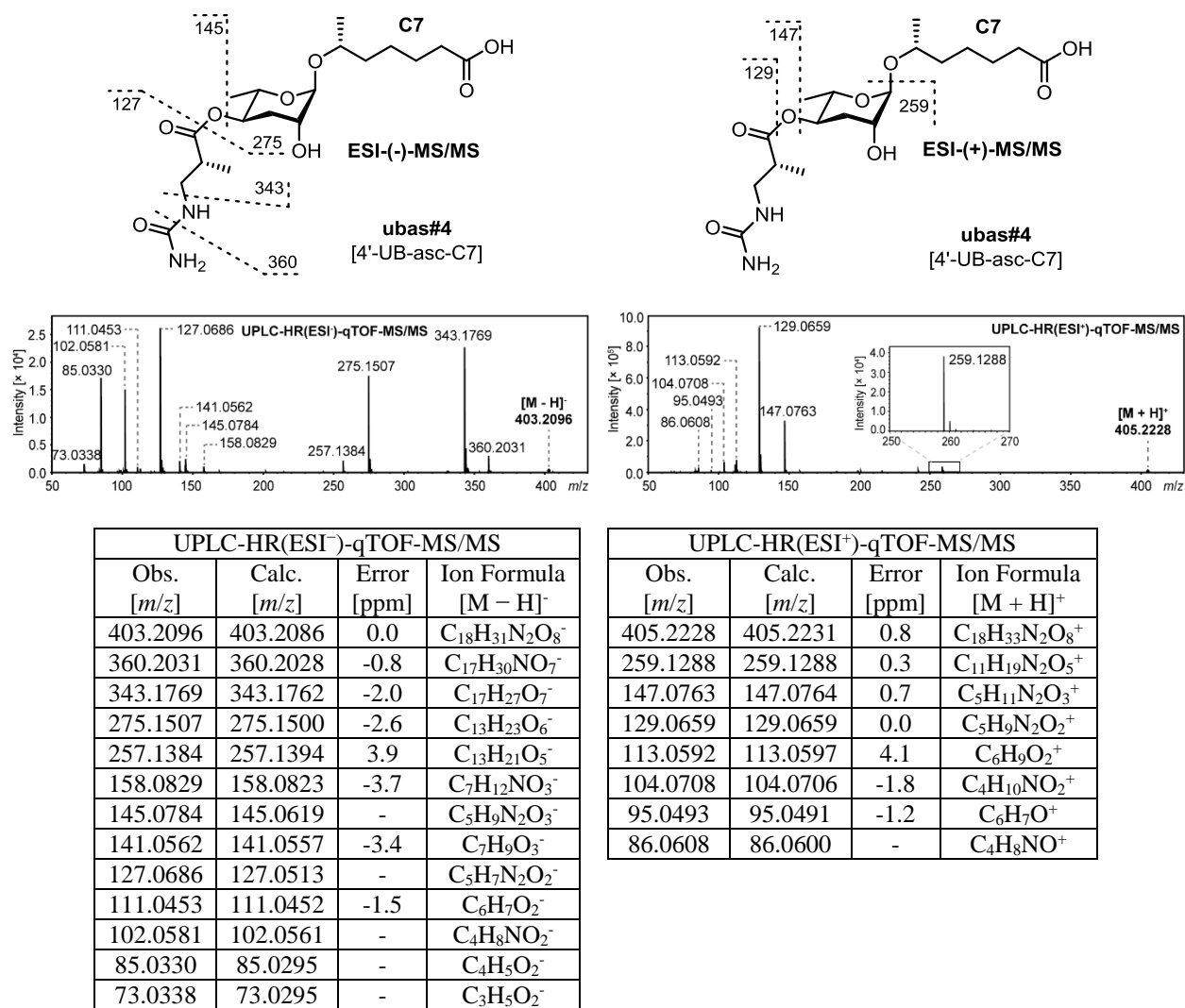
**Figure 1.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of known ubas#5 [4'-UB-asc-C4] (*Falcke et al., 2018*) from *P. pacificus* in both negative and positive ion modes. Characteristic ion signal for *m/z* 259.1304 containing an ureidoisobutyric acid moiety and an ascarylose unit was used for targeted MS/MS screening. *P. triformis* also produced large amounts of ubas#5 [4'-UB-asc-C4] (*supplementary file 1d – Figures 30-35*).



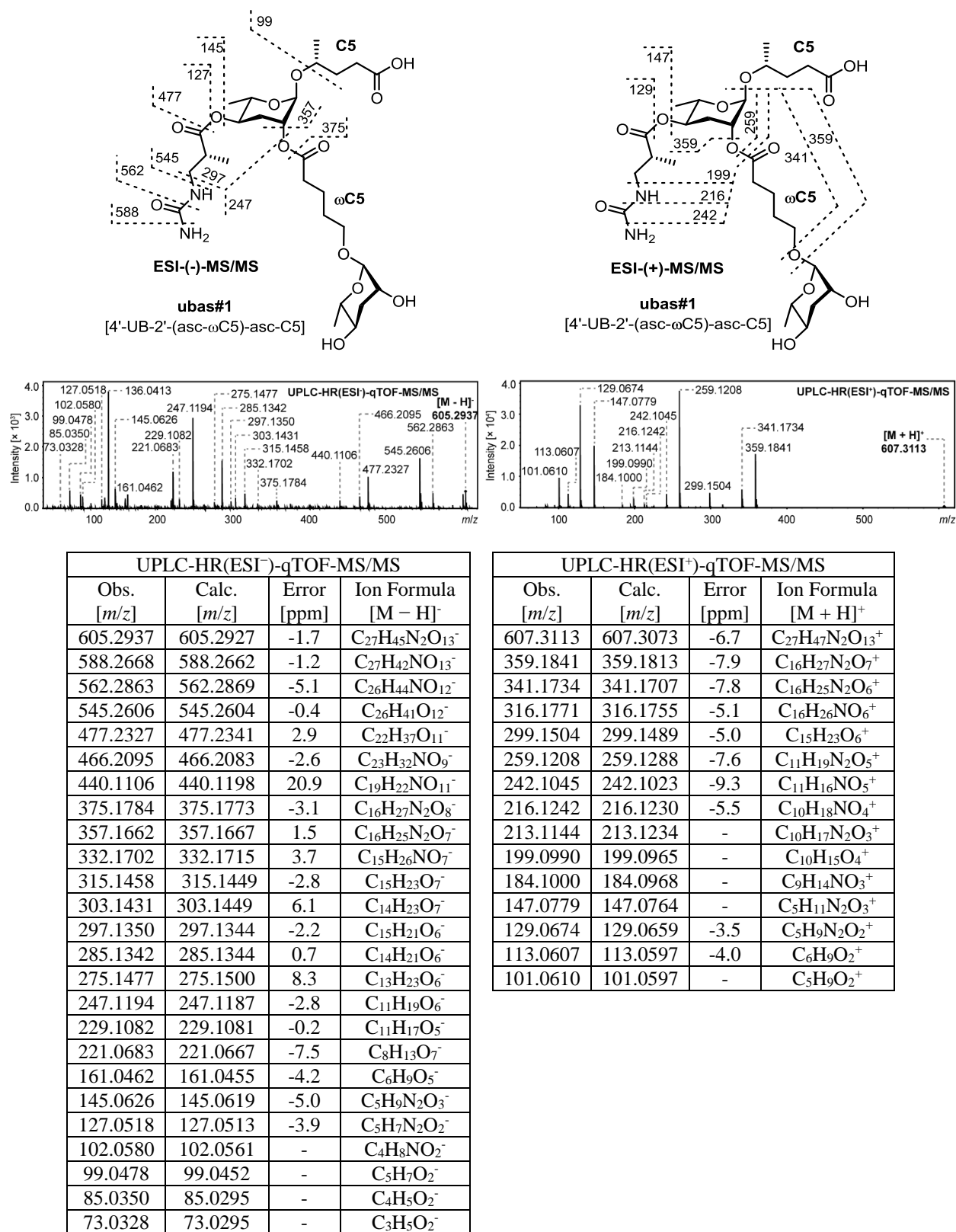
**Figure 2.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of known ubas#3 [4'-UB-asc-C5] (Falcke *et al.*, 2018) from *P. pacificus* in both negative and positive ion modes. Characteristic ion signal for *m/z* 259.1304 containing an ureidoisobutyric acid moiety and an ascarbylose unit was used for targeted MS/MS screening.



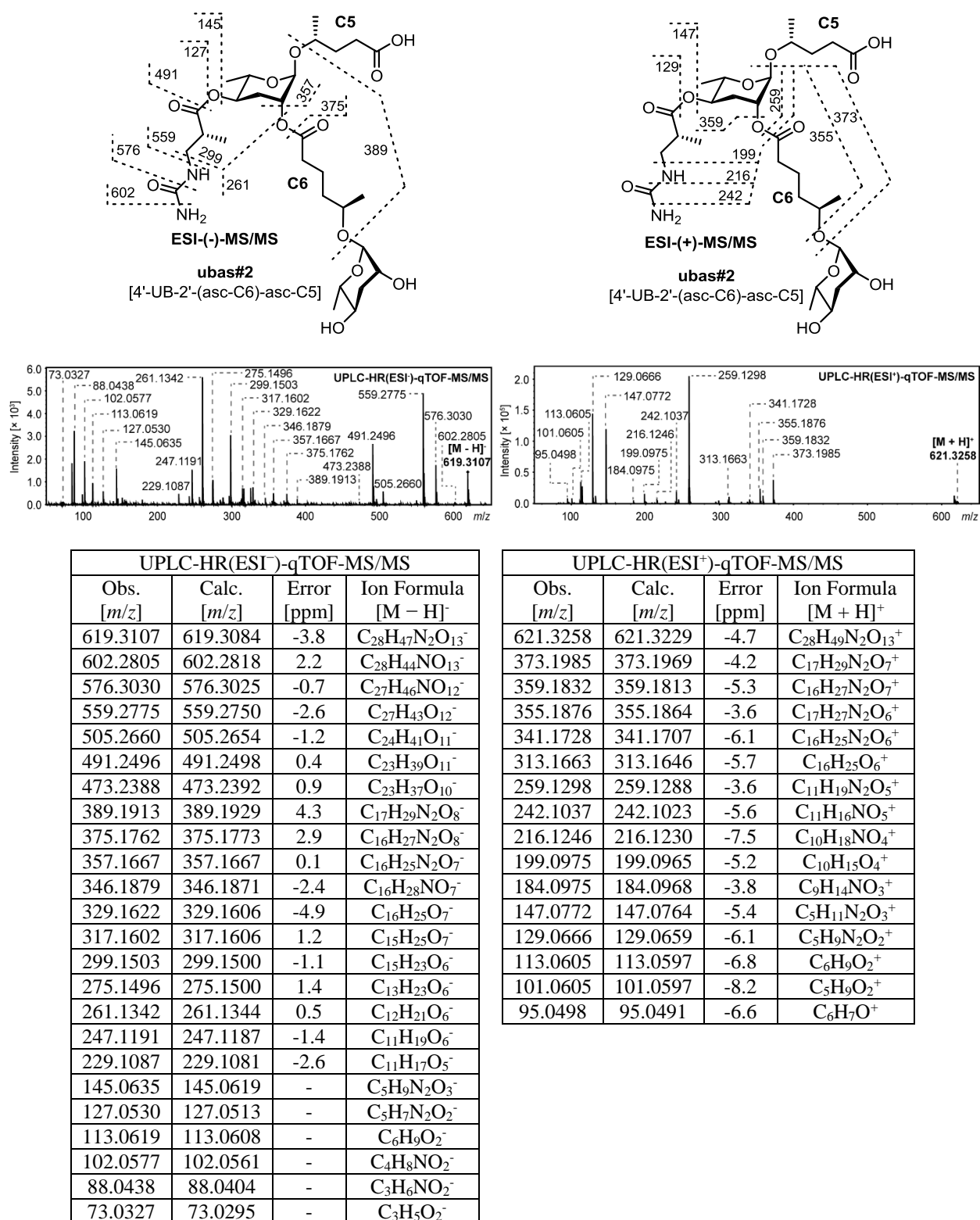
**Figure 3.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of known ubas#12 [4'-UB-asc-C6] (*Falcke et al., 2018*) from *P. pacificus* in both negative and positive ion modes. Characteristic ion signal for *m/z* 259.1304 containing an ureidoisobutyric acid moiety and an ascarylose unit was used for targeted MS/MS screening.



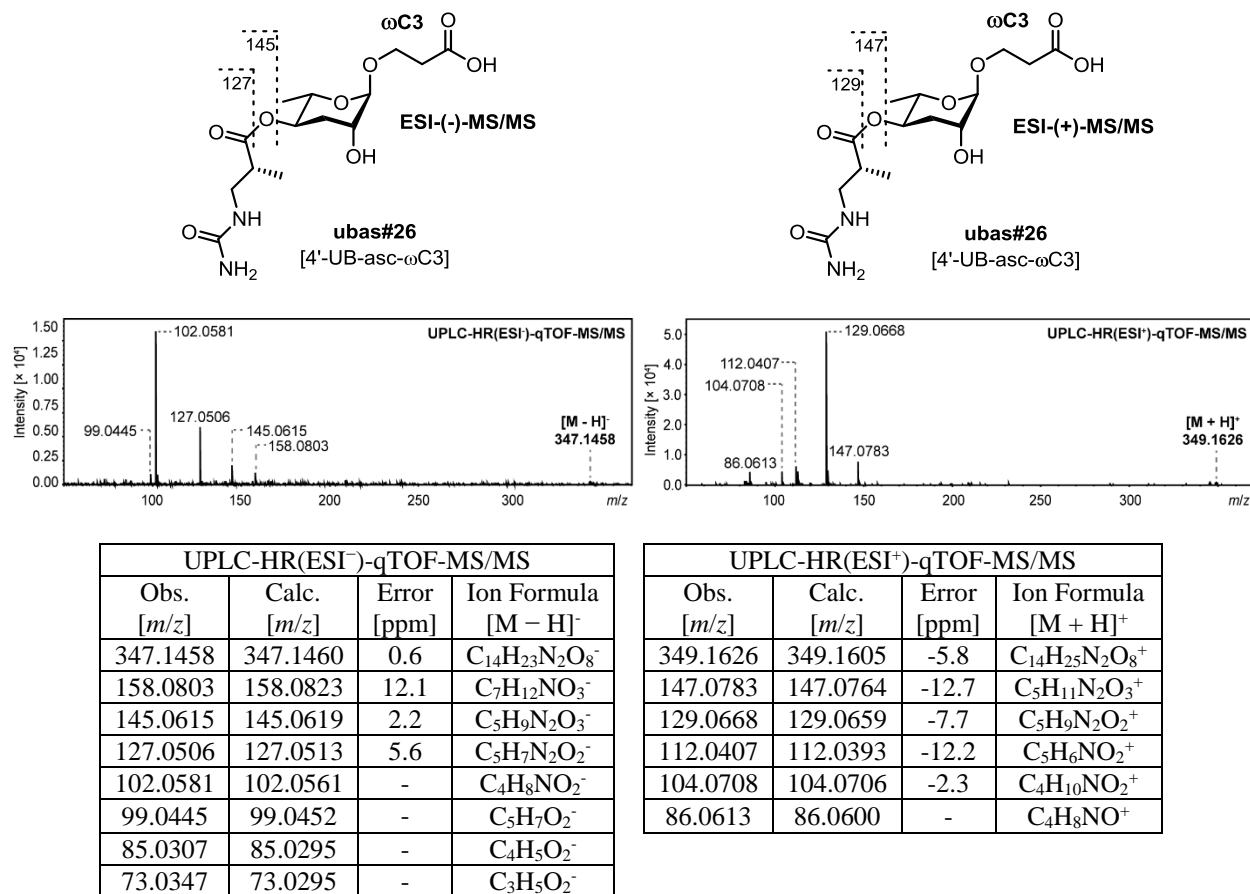
**Figure 4.** UPLC-HR(ESI<sup>-/+</sup>)-qTOF-MS/MS mass spectral data of known ubas#4 [4'-UB-asc-C7] (*Falcke et al., 2018*) from *P. pacificus* in both negative and positive ion modes. Characteristic ion signal for *m/z* 259.1304 containing an ureidoisobutyric acid moiety and an ascaryle unit was used for targeted MS/MS screening.



**Figure 5.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of known ubas#1 [4'-UB-2'-(asc-ωC5)-asc-C5] (supplementary file 1d – Figures 9-10) (Bose et al., 2012) from *P. pacificus*.

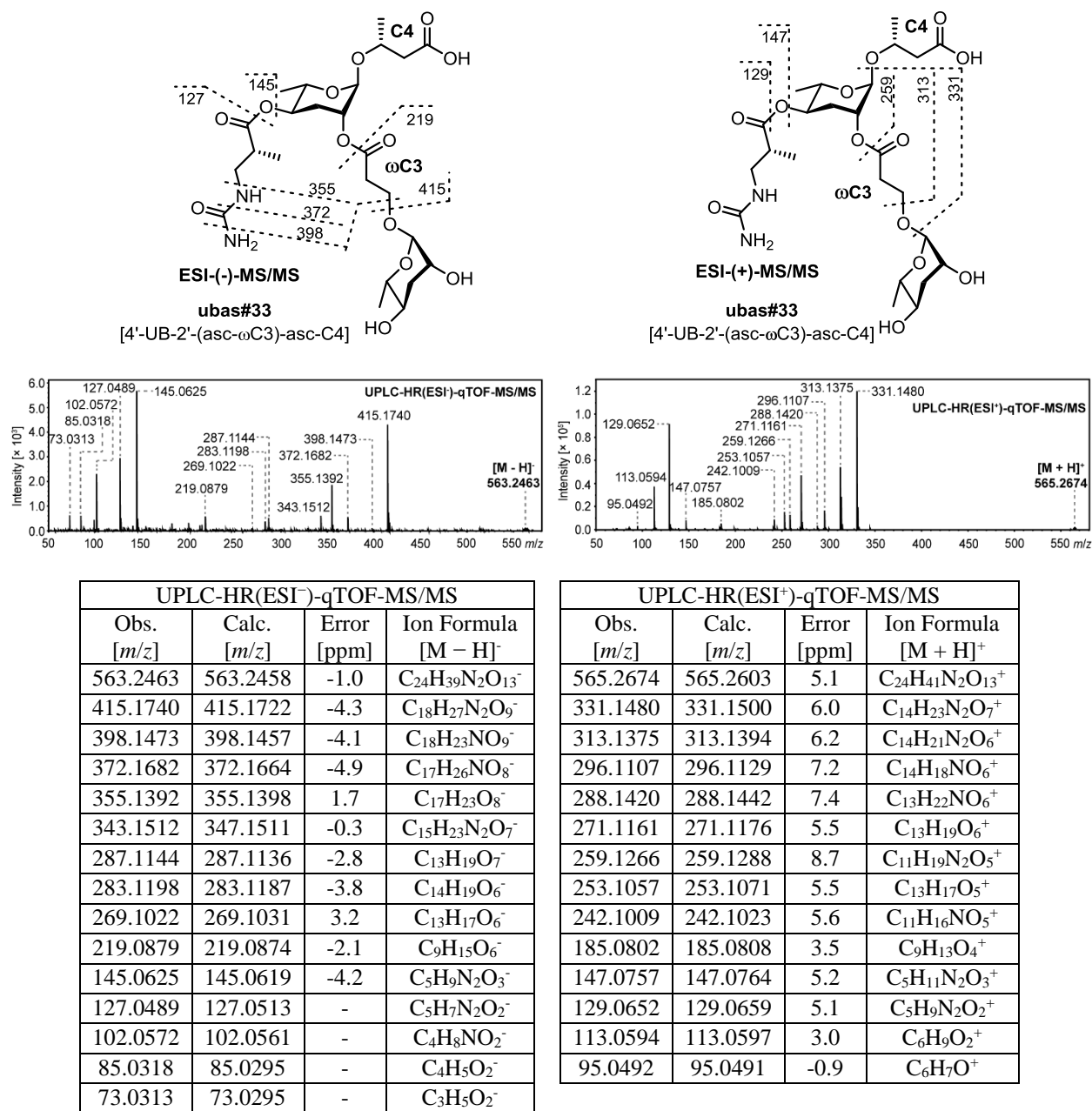


**Figure 6.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of known ubas#2 [4'-UB-2'-(asc-C6)-asc-C5) (*Bose et al., 2012*) from *P. pacificus* in both negative and positive ion modes.

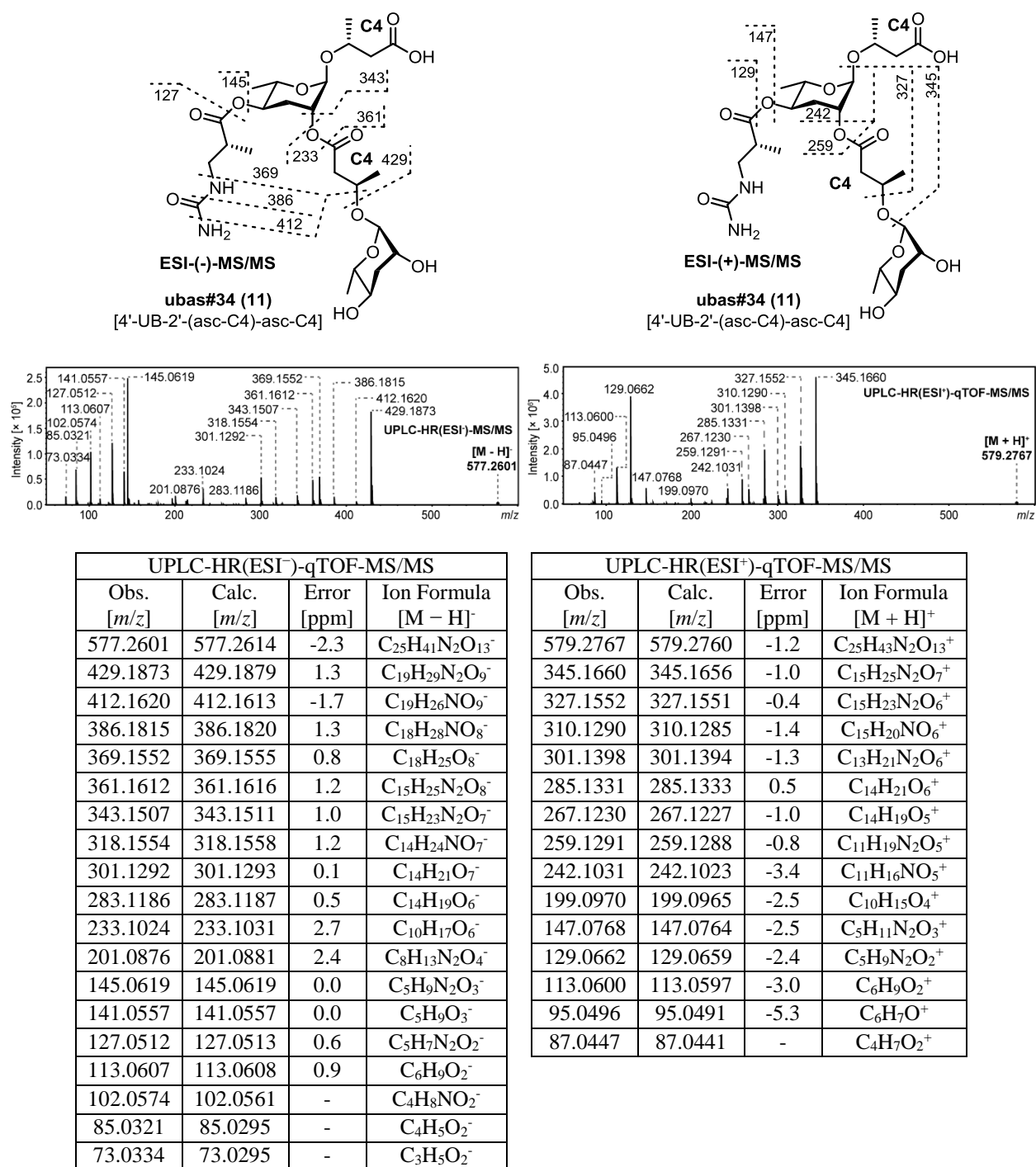


**Figure 7.** UPLC-HR(ESI<sup>-/+</sup>)-qTOF-MS/MS mass spectral data of newly identified ubas#26 [4'-UB-asc- $\omega$ C3] in both negative and positive ion modes.

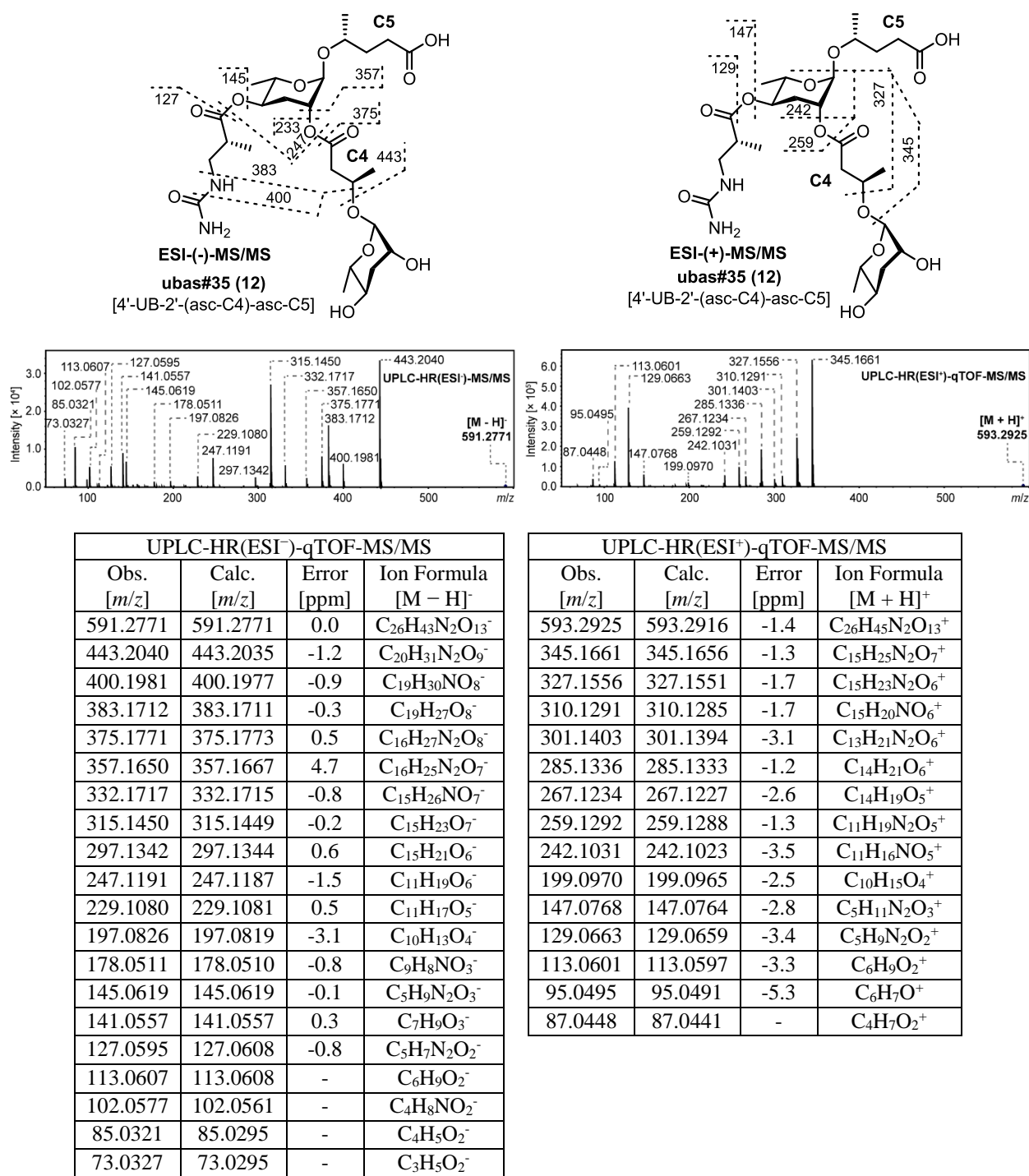




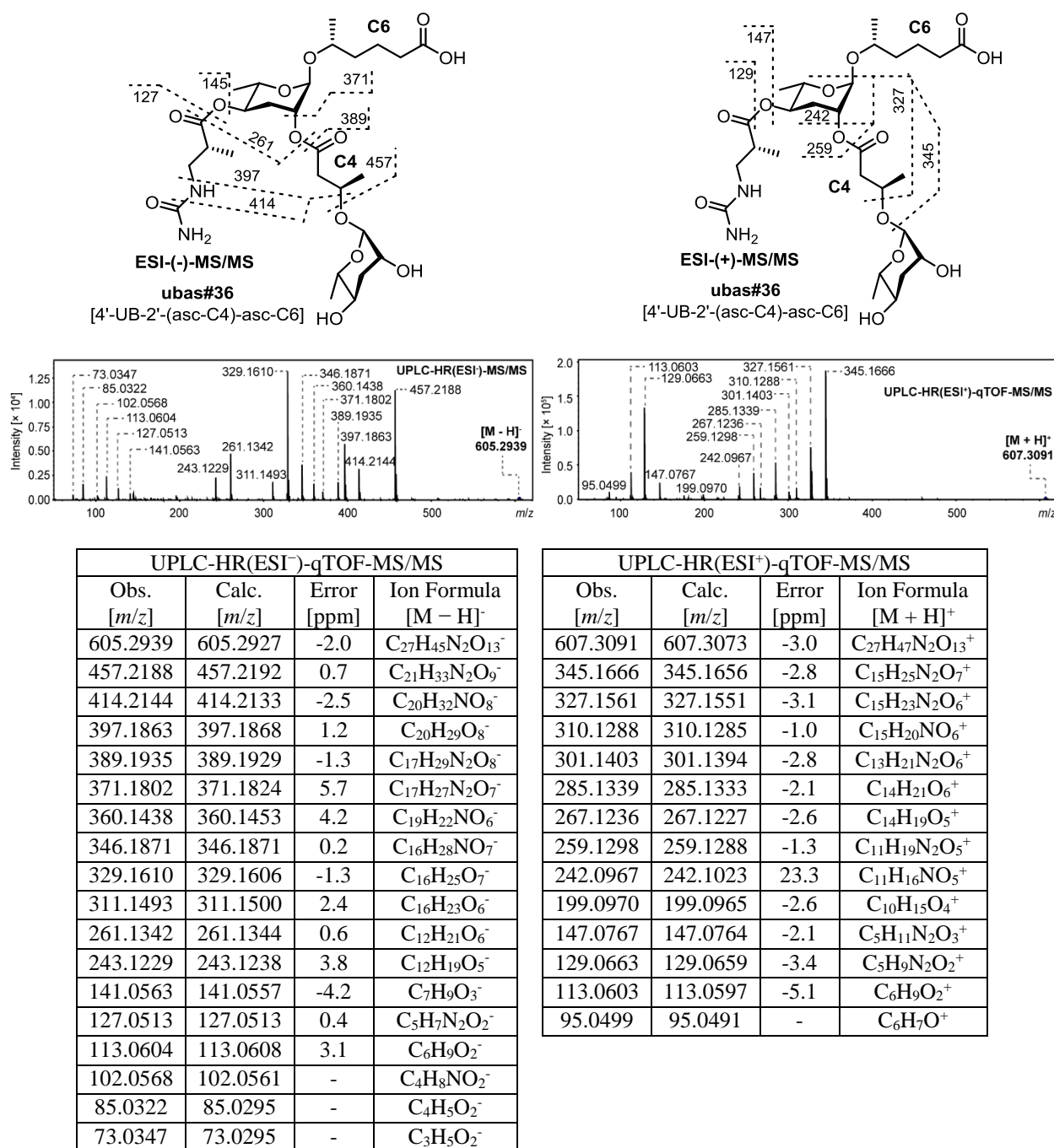
**Figure 8.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#33 [4'-UB-2'-(asc- $\omega$ C3)-asc-C4] in both negative and positive ion modes.



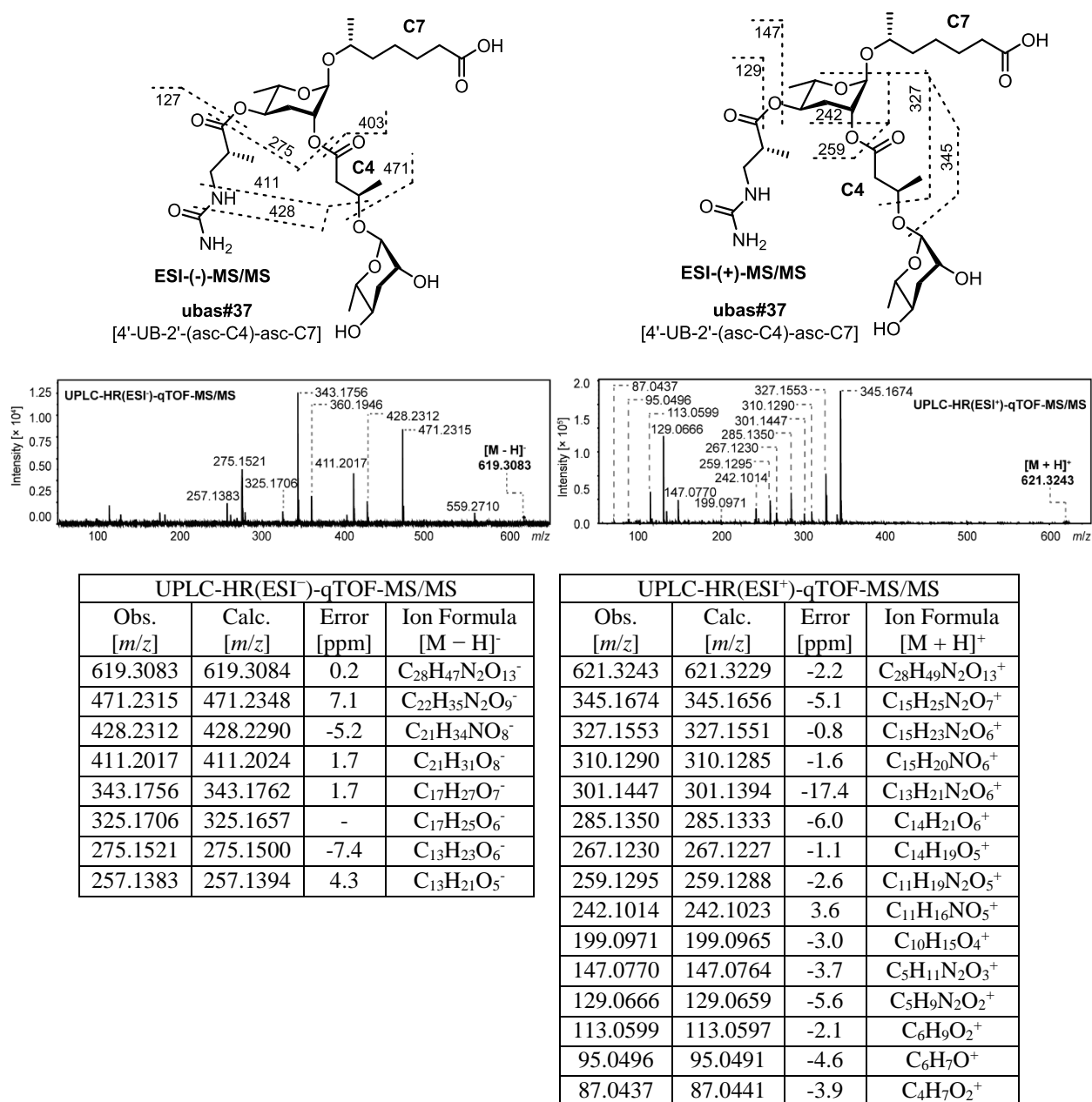
**Figure 9.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#34 [4'-UB-2'-(asc-C4)-asc-C4, **11**] in both negative and positive ion modes. UPLC-HR(ESI<sup>+</sup>)-qTOF-MS/MS fragmentation produced a fragment ion signal for C<sub>15</sub>H<sub>25</sub>N<sub>2</sub>O<sub>7</sub><sup>+</sup> ( $m/z$  345.1660 [M + H]<sup>+</sup>,  $\Delta$  -1.0 ppm) (**Figure 5 – figure supplement 3**), indicating that ascr#11 [asc-C4] represents the first ascaroside in ubas#34 [4'-UB-2'-(asc-C4)-asc-C4, **11**].



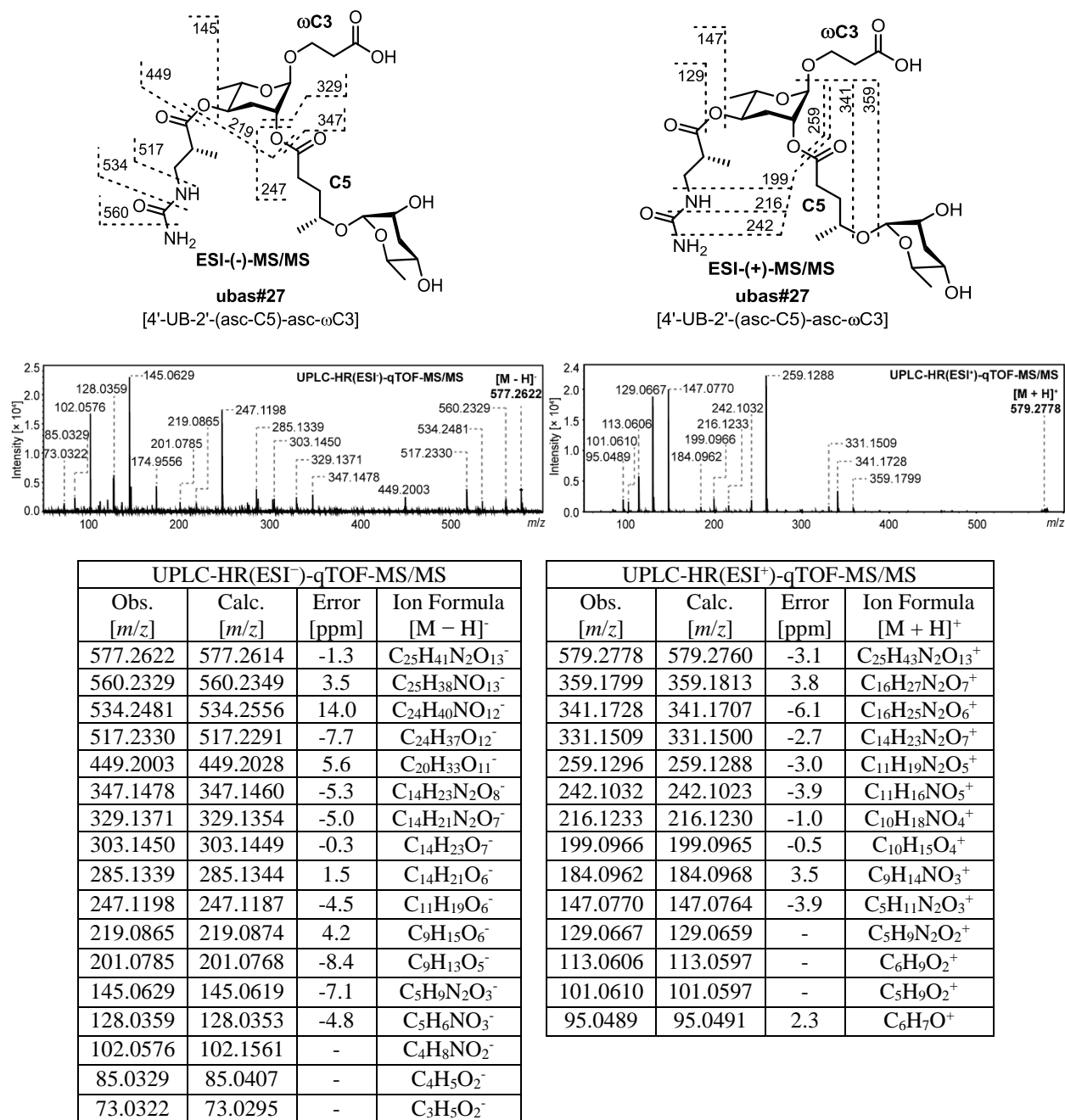
**Figure 10.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#35 [4'-UB-2'-(asc-C4)-asc-C5, 12] in both negative and positive ion modes. UPLC-HR(ESI<sup>+</sup>)-qTOF-MS/MS fragmentation produced a fragment ion signal for C<sub>15</sub>H<sub>25</sub>N<sub>2</sub>O<sub>7</sub><sup>+</sup> ( $m/z$  345.1660 [M + H]<sup>+</sup>,  $\Delta$  -1.0 ppm) (*Figure 5 – figure supplement 3*), indicating that ascr#11 [asc-C4] represents the first ascaroside in ubas#35 [4'-UB-2'-(asc-C4)-asc-C5, 12].



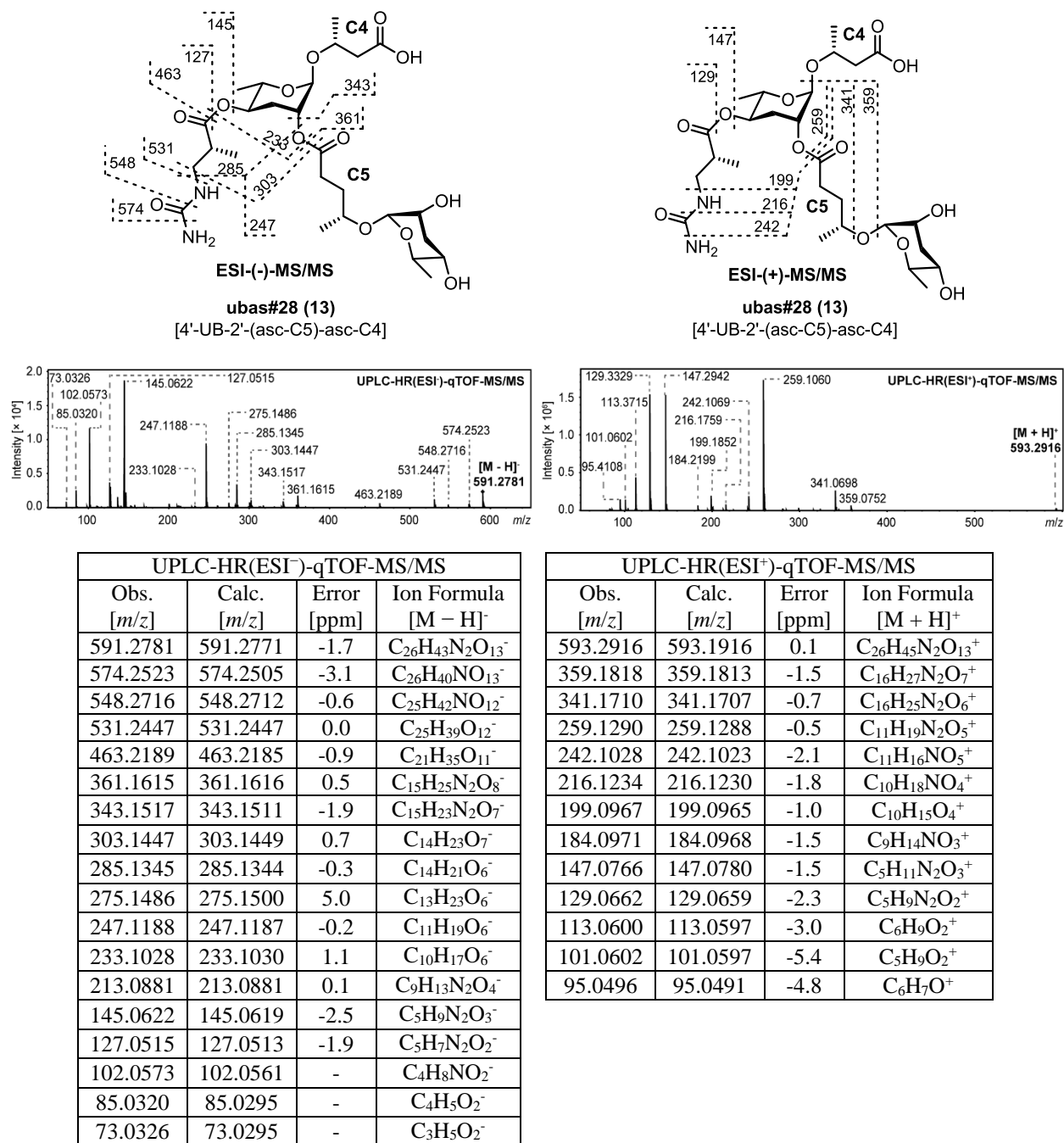
**Figure 11.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of ubas#36 [4'-UB-2'-(asc-C4)-asc-C6] in both negative and positive ion modes. UPLC-HR(ESI<sup>+</sup>)-qTOF-MS/MS fragmentation produced a fragment ion signal for C<sub>15</sub>H<sub>25</sub>N<sub>2</sub>O<sub>7</sub><sup>+</sup> (*m/z* 345.1660 [M + H]<sup>+</sup>, Δ -1.0 ppm) (**Figure 5 – figure supplement 3**), indicating that ascr#11 [asc-C4] represents the first ascaroside in ubas#36 [4'-UB-2'-(asc-C4)-asc-C6].



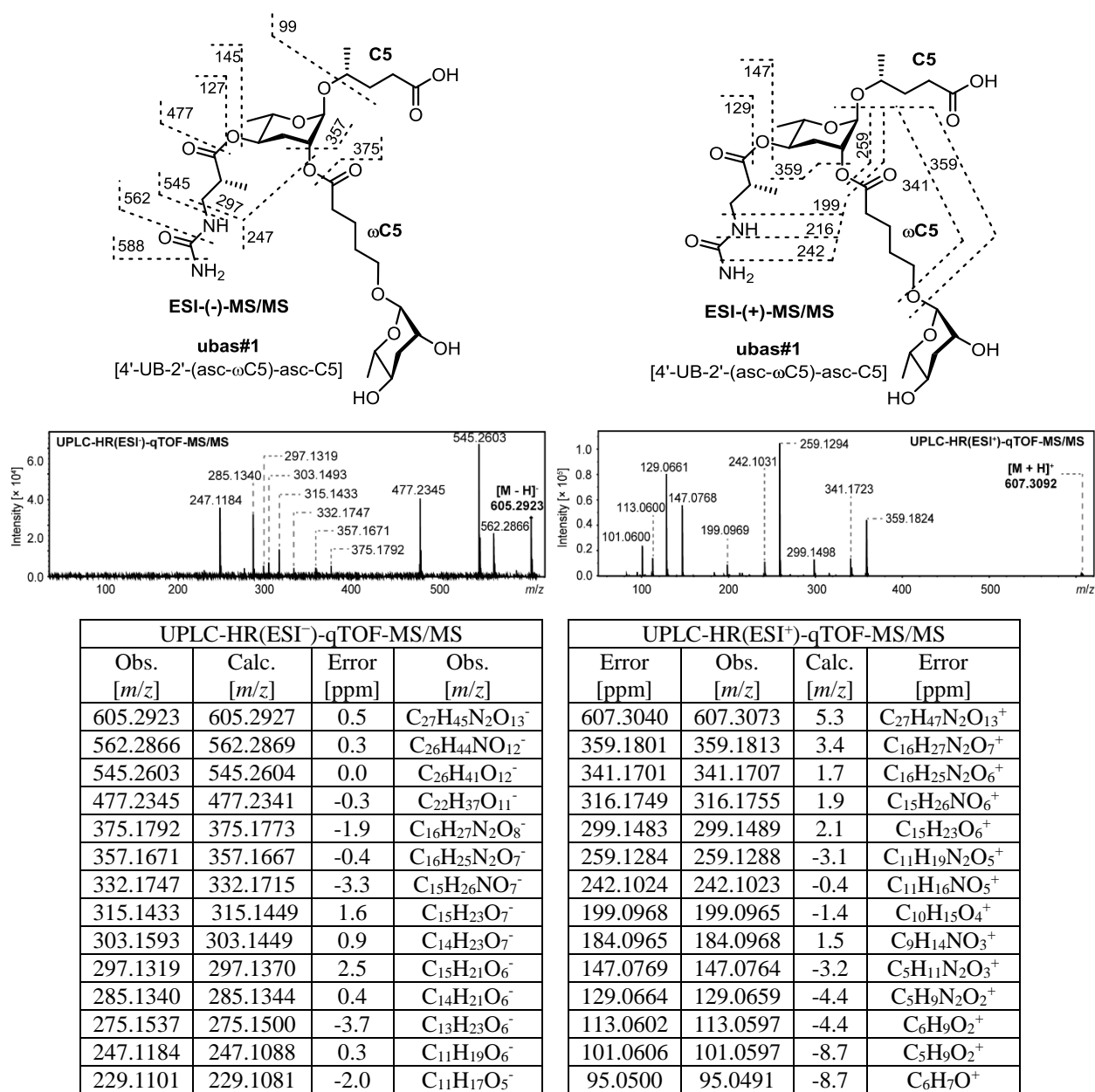
**Figure 12.** UPLC-HR(ESI<sup>+/−</sup>)-qTOF-MS/MS mass spectral data of ubas#37 [4'-UB-2'-(asc-C4)-asc-C7] in both negative and positive ion modes. UPLC-HR(ESI<sup>+</sup>)-qTOF-MS/MS fragmentation produced a fragment ion signal for C<sub>15</sub>H<sub>25</sub>N<sub>2</sub>O<sub>7</sub><sup>+</sup> (*m/z* 345.1660 [M + H]<sup>+</sup>, Δ -1.0 ppm) (**Figure 5 – figure supplement 3**), indicating that ascr#11 [asc-C4] represents the first ascaroside in ubas#37 [4'-UB-2'-(asc-C4)-asc-C7].



**Figure 13.** UPLC-HR(ESI<sup>-/+</sup>)-qTOF-MS/MS mass spectral data of ubas#27 [4'-UB-2'-(asc-C5)-asc-ωC3] in both negative and positive ion modes.

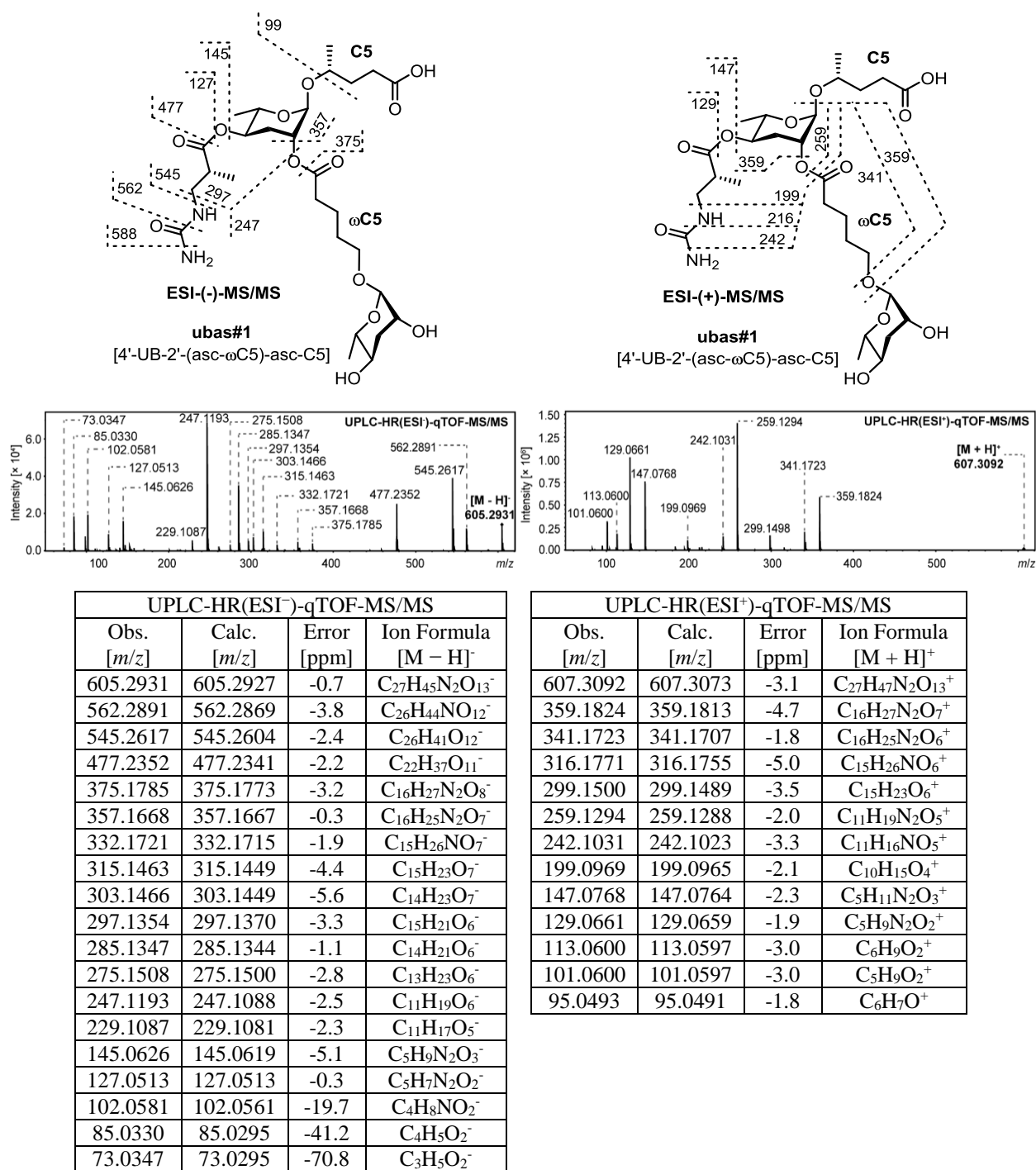


**Figure 14.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#28 [4'-UB-2'-(asc-C5)-asc-C4, 13] in both negative and positive ion modes.

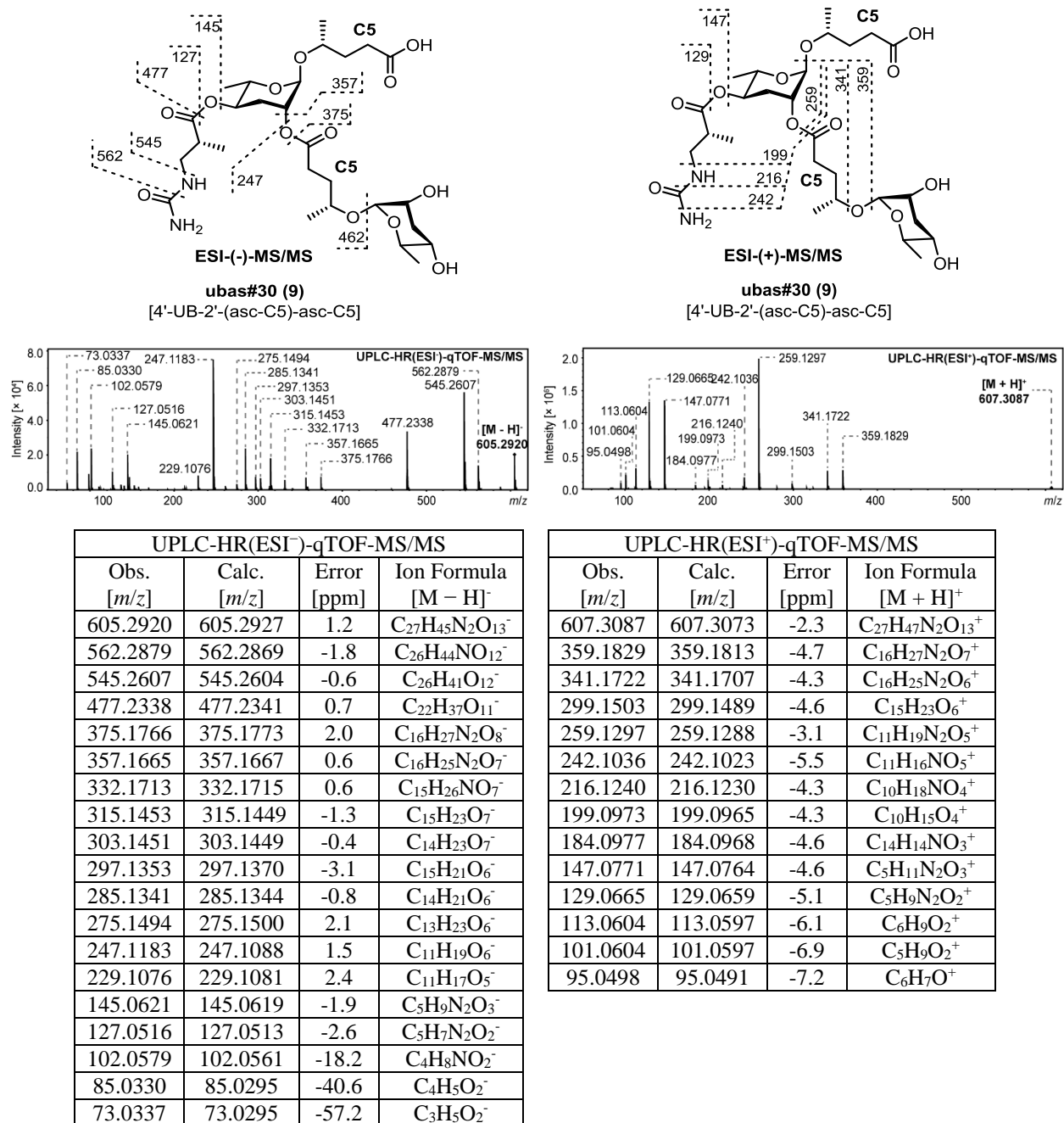


**Figure 15.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#1 [4'-UB-2'-(asc- $\omega$ C5)-asc-C5] (from *P. taiwanensis*) in both negative and positive ion modes. Analysis of NMR data of ubas#1 isolated from the *exo*-metabolome of *P. taiwanensis* has confirmed its chemical identity (*supplementary file 1d – Figures 12-15*).

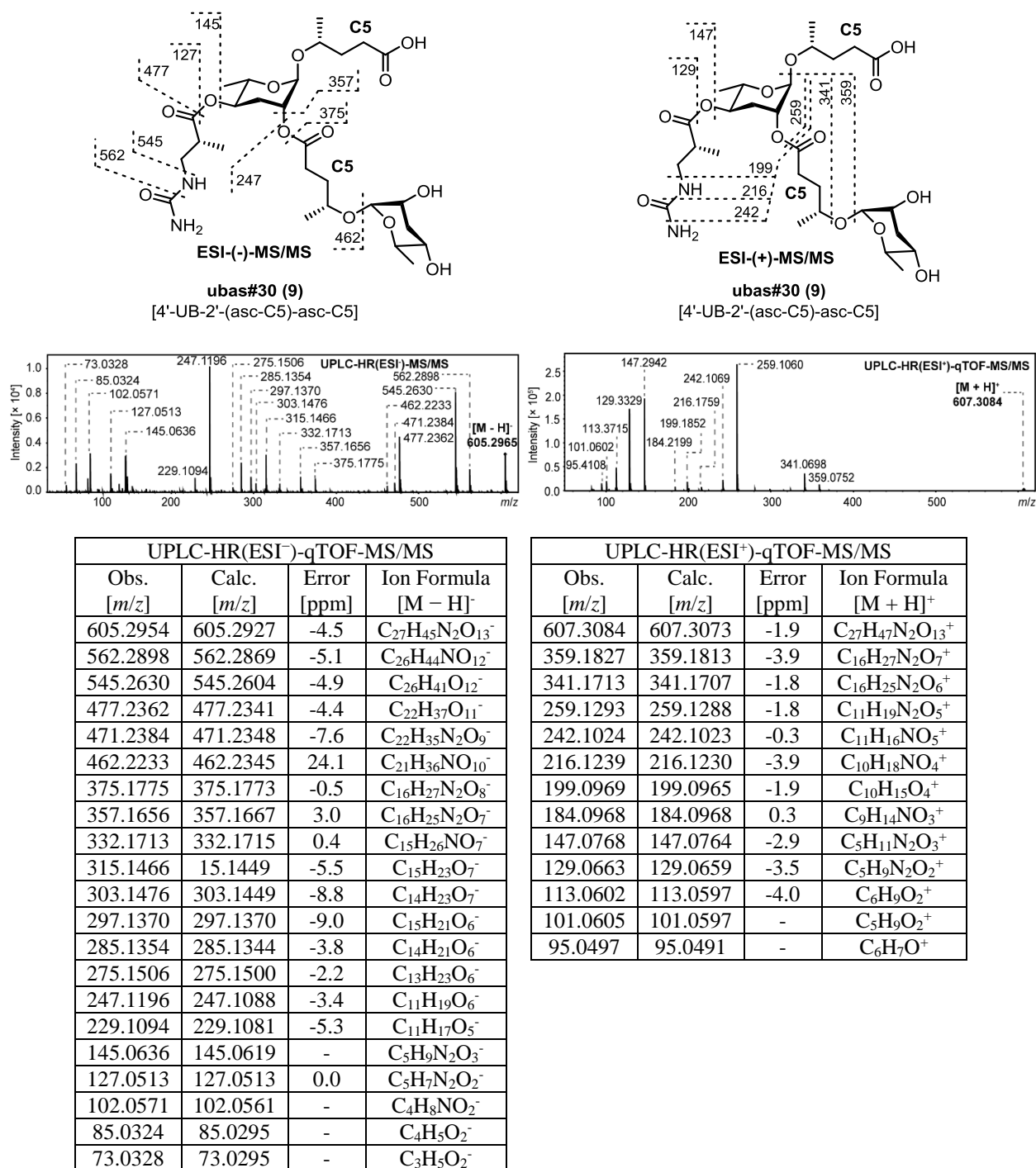




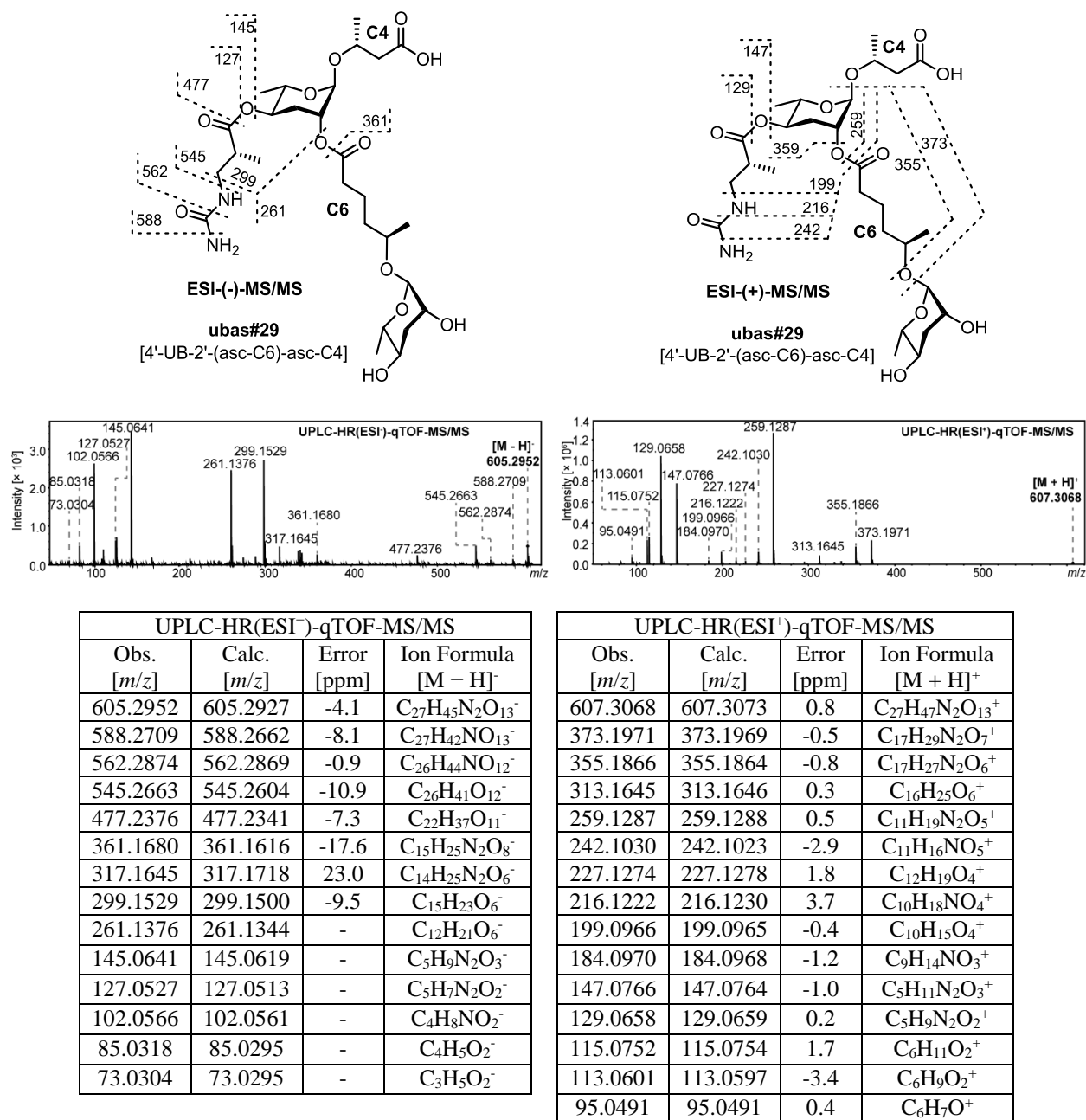
**Figure 16.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of ubas#1 [4'-UB-2'-(asc-ωC5)-asc-C5] (from *P. laeviscolis*) in both negative and positive ion modes. Analysis of NMR data of ubas#1 isolated from the *exo*-metabolome of *P. laeviscolis* confirmed its chemical identity (*supplementary file 1d – Figures 16-19*).



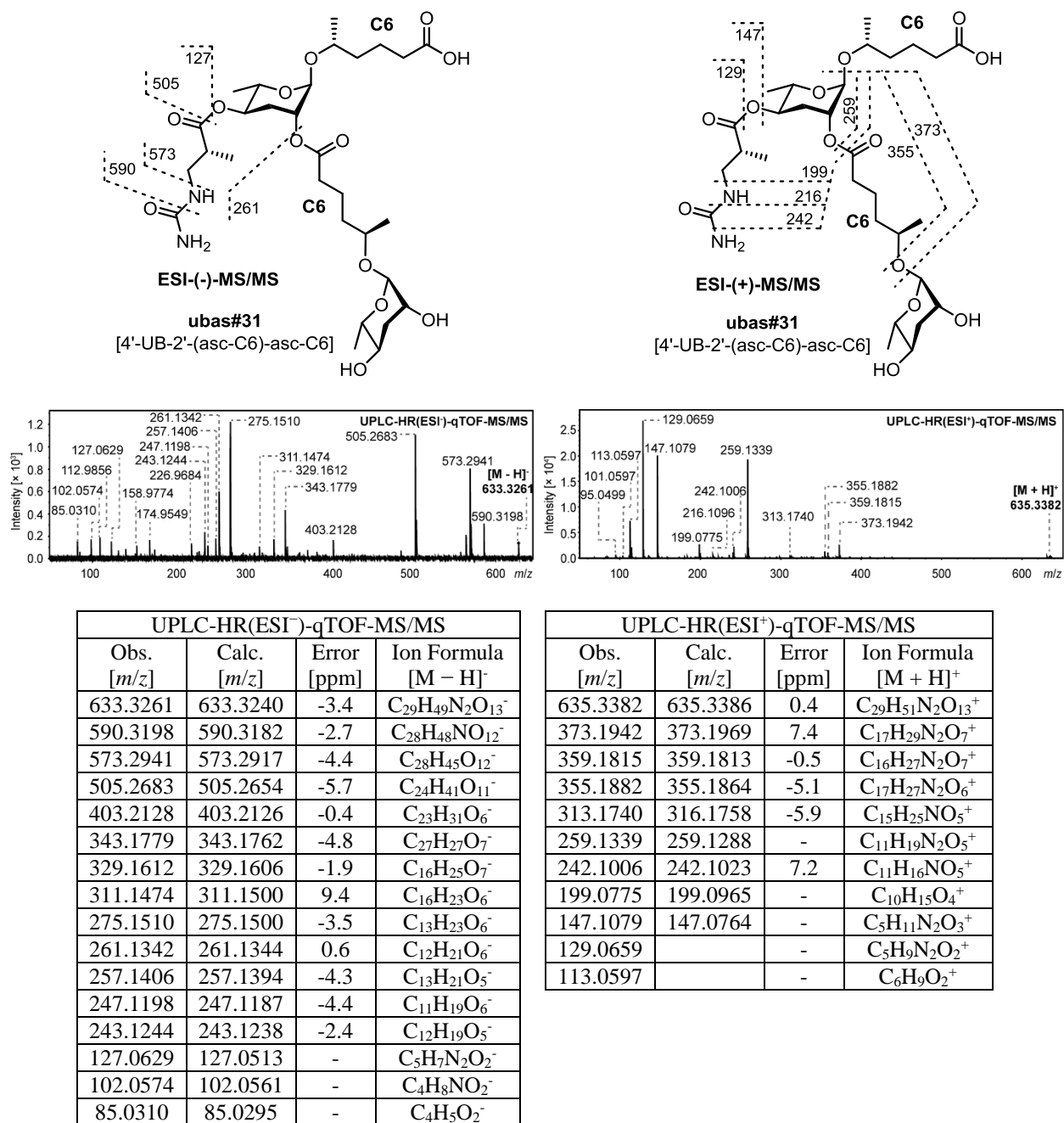
**Figure 17.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#30 [4'-UB-2'-(asc-C5)-asc-C5, 9] (from *P. maxplancki*) in both negative and positive ion modes.



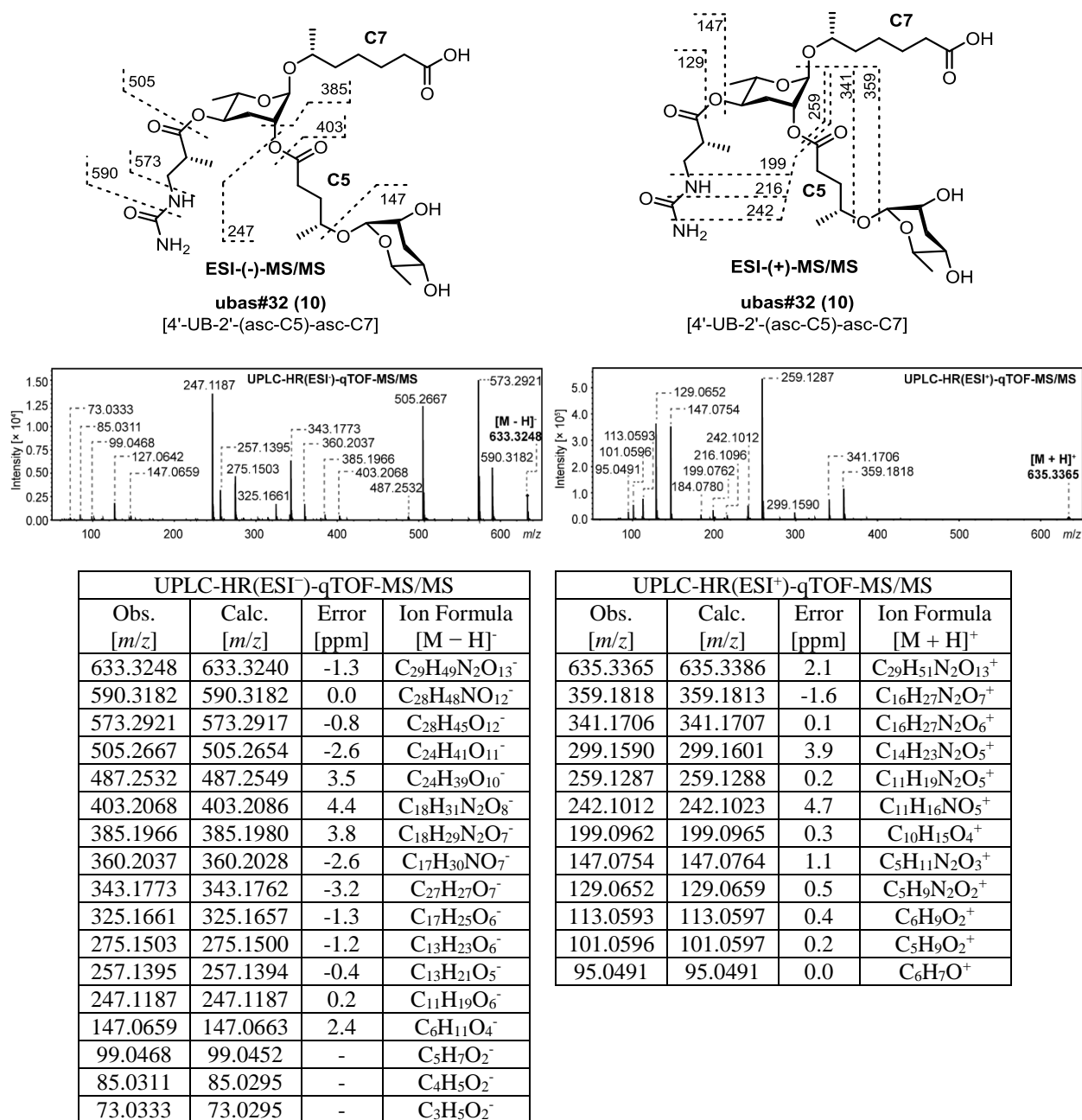
**Figure 18.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#30 [4'-UB-2'-(asc-C5)-asc-C5, **9**] (from *P. fukushima*) in both negative and positive ion modes. Peak at *m/z* 605.2954 was tentatively assigned to be ubas#30 [4'-UB-2'-(asc-C5)-asc-C5, **9**].



**Figure 19.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of ubas#29 [4'-UB-2'-(asc-C6)-asc-C4] in both negative and positive ion modes.

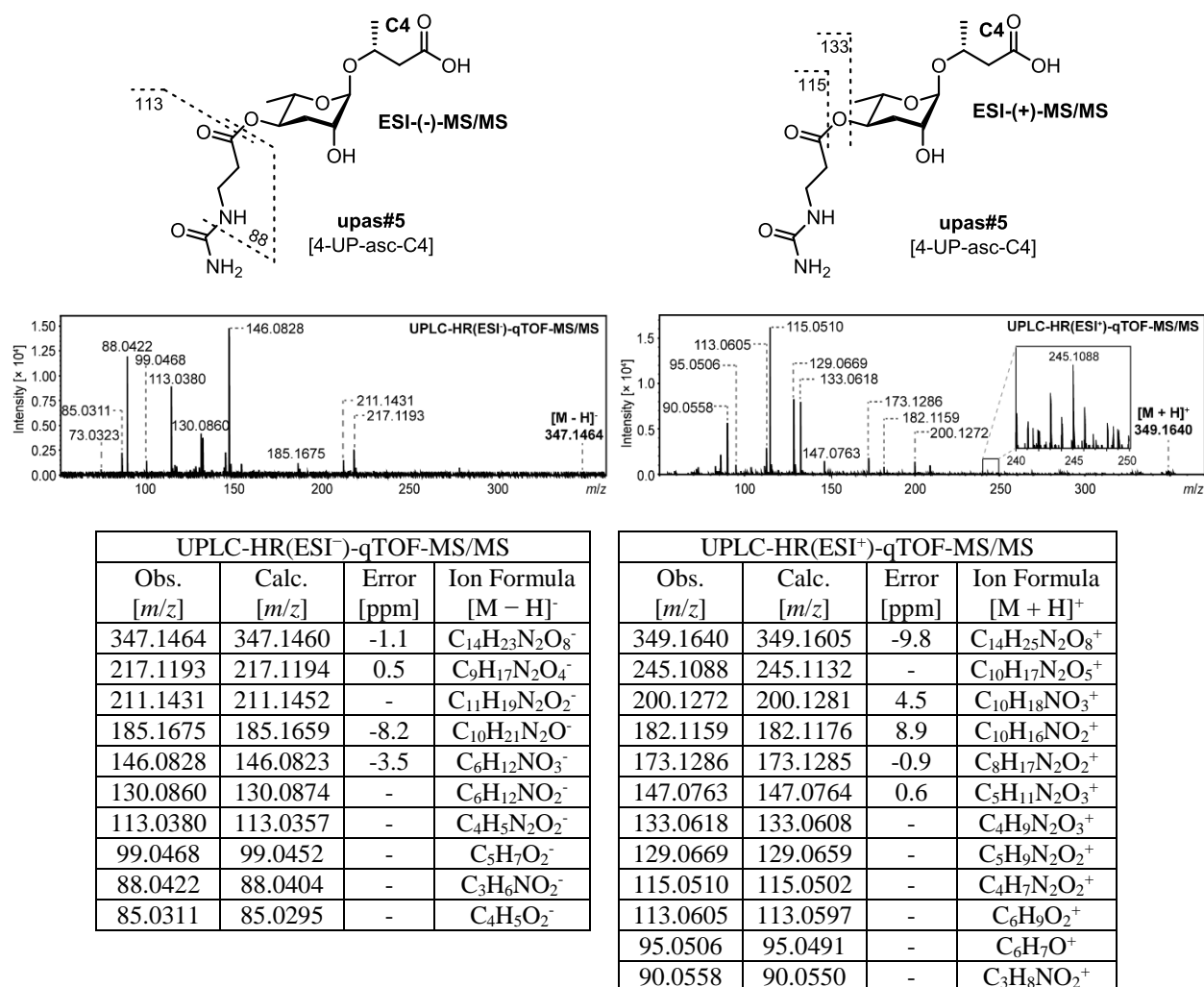


**Figure 20.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of ubas#31 [4'-UB-2'-(asc-C6)-asc-C6] in both negative and positive ion modes.



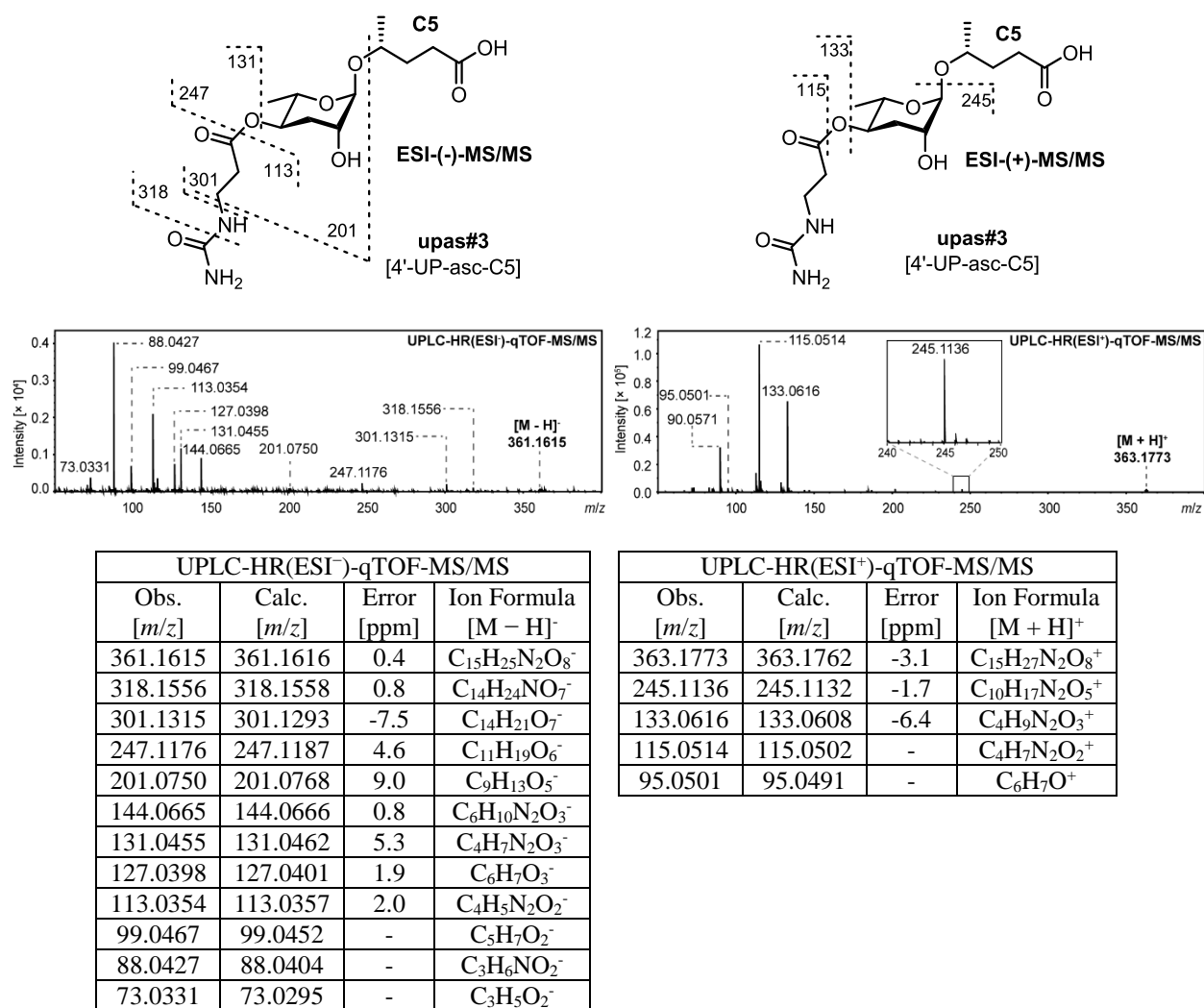
**Figure 21.** UPLC-HR(ESI<sup>+/−</sup>)-qTOF-MS/MS mass spectral data of ubas#32 [4'-UB-2'-(asc-C5)-asc-C7, **10**] in both negative and positive ion modes. Analysis of the *dqf*-COSY spectrum of ubas#32 [4'-UB-2'-(asc-C5)-asc-C7, **10**] (800 MHz, CD<sub>3</sub>OD) enriched from the *exo*-metabolome of *P. maxplancki* confirmed the presence of one 2'- and 4'-substituted ascarylose sugar, an unsubstituted ascarylose moiety, one (ω-1)-hydroxylated C5 side chain and one (ω-1)-hydroxylated C7 side chain in ubas#32 (*supplementary file 1d* – *Figure 20*).

<b>supplementary file 2d: MS/MS spectral data of UPAS chemicals</b>	Pages
<b>Figure 1.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of upas#5 [4'-UP-asc-C4].	S48
<b>Figure 2.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of upas#3 [4'-UP-asc-C5].	S49
<b>Figure 3.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of upas#4 [4'-UP-asc-C7].	S50
<b>Figure 4.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of upas#34 [4'-UP-2'-(asc-C4)-asc-C4, <b>14</b> ].	S51
<b>Figure 5.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of upas#28 [4'-UP-2'-(asc-C5)-asc-C4, <b>15</b> ].	S52
<b>Figure 6.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of upas#30 [4'-UP-2'-(asc-C5)-asc-C5].	S53
<b>Figure 7.</b> LC-HR( $\text{ESI}^{-/+}$ )-MS/MS spectral data of upas#32 [4'-UP-2'-(asc-C5)-asc-C7].	S54

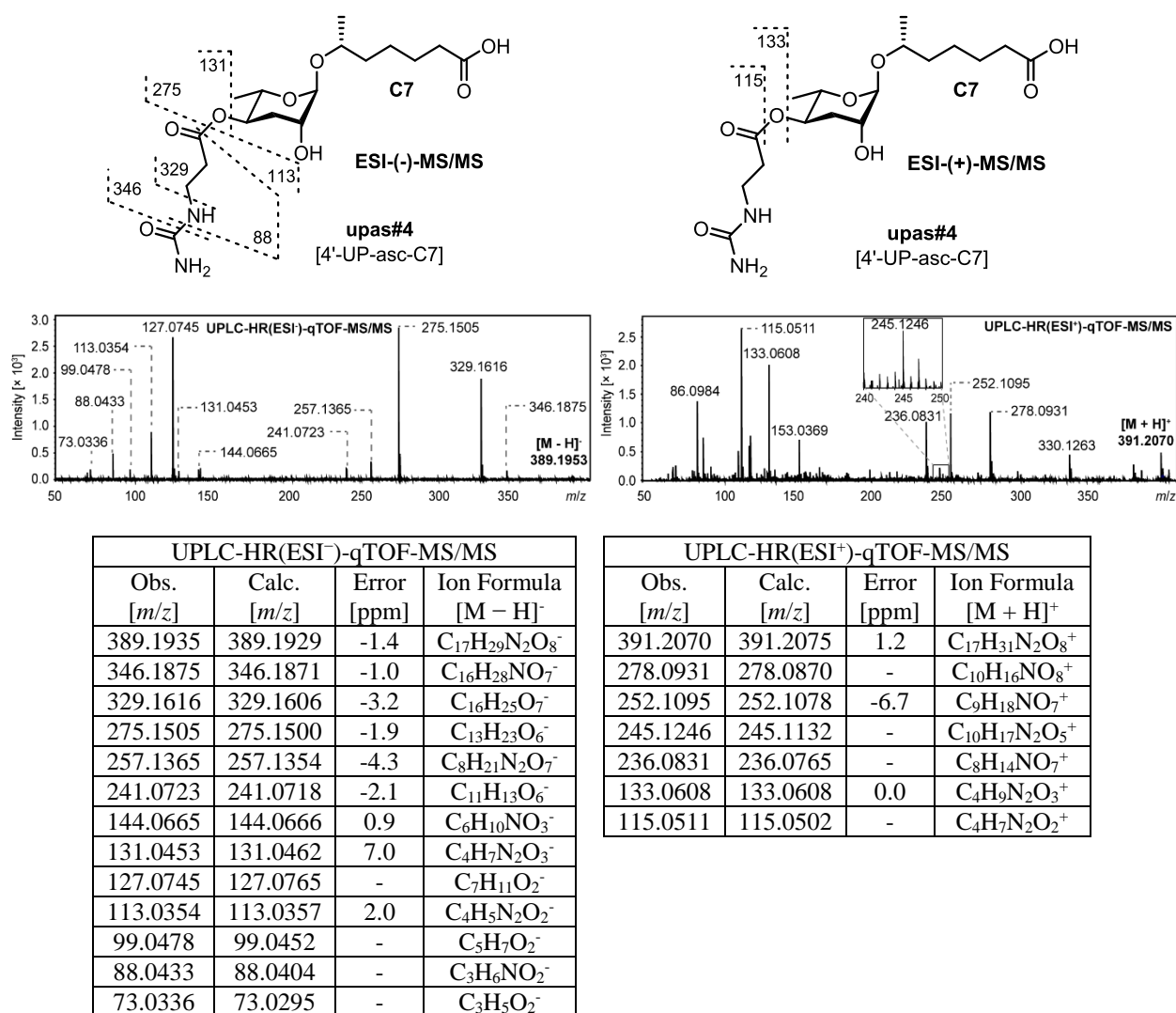


**Figure 1.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of upas#5 [4'-UP-asc-C4] in both negative and positive ion modes.

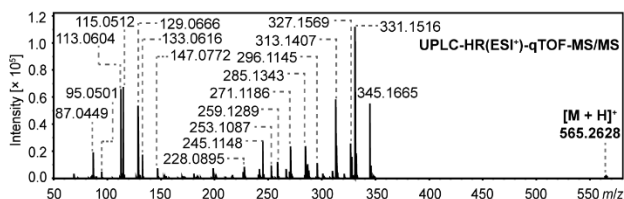
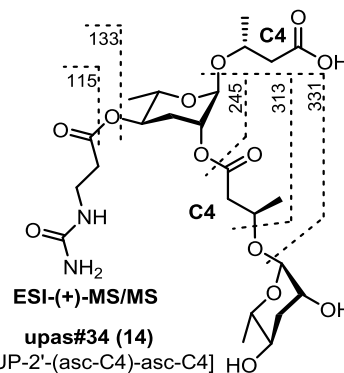




**Figure 2.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of upas#3 [4'-UP-asc-C5] in both negative and positive ion modes.

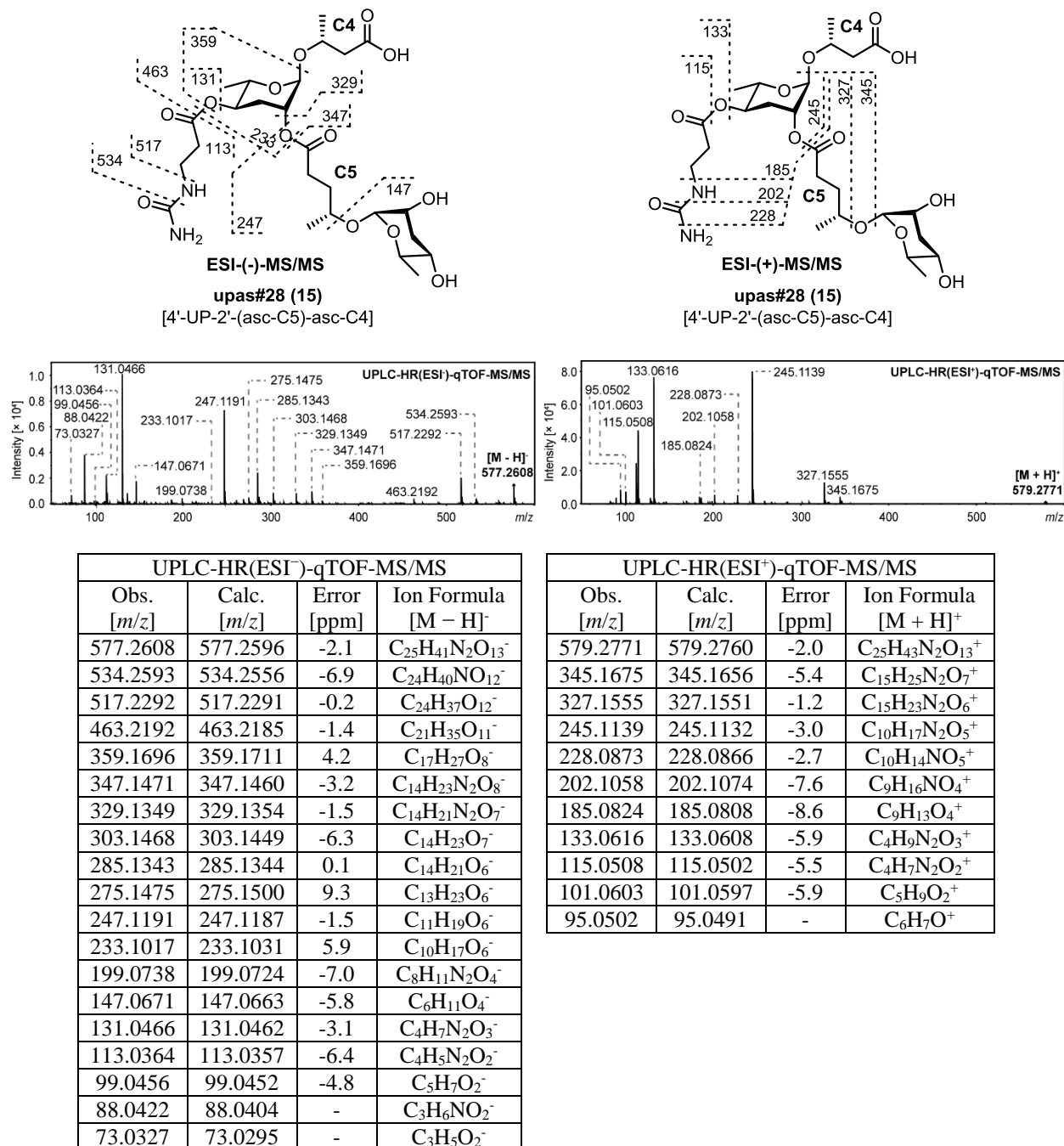


**Figure 3.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of upas#4 [4'-UP-asc-C7] in both negative and positive ion modes.

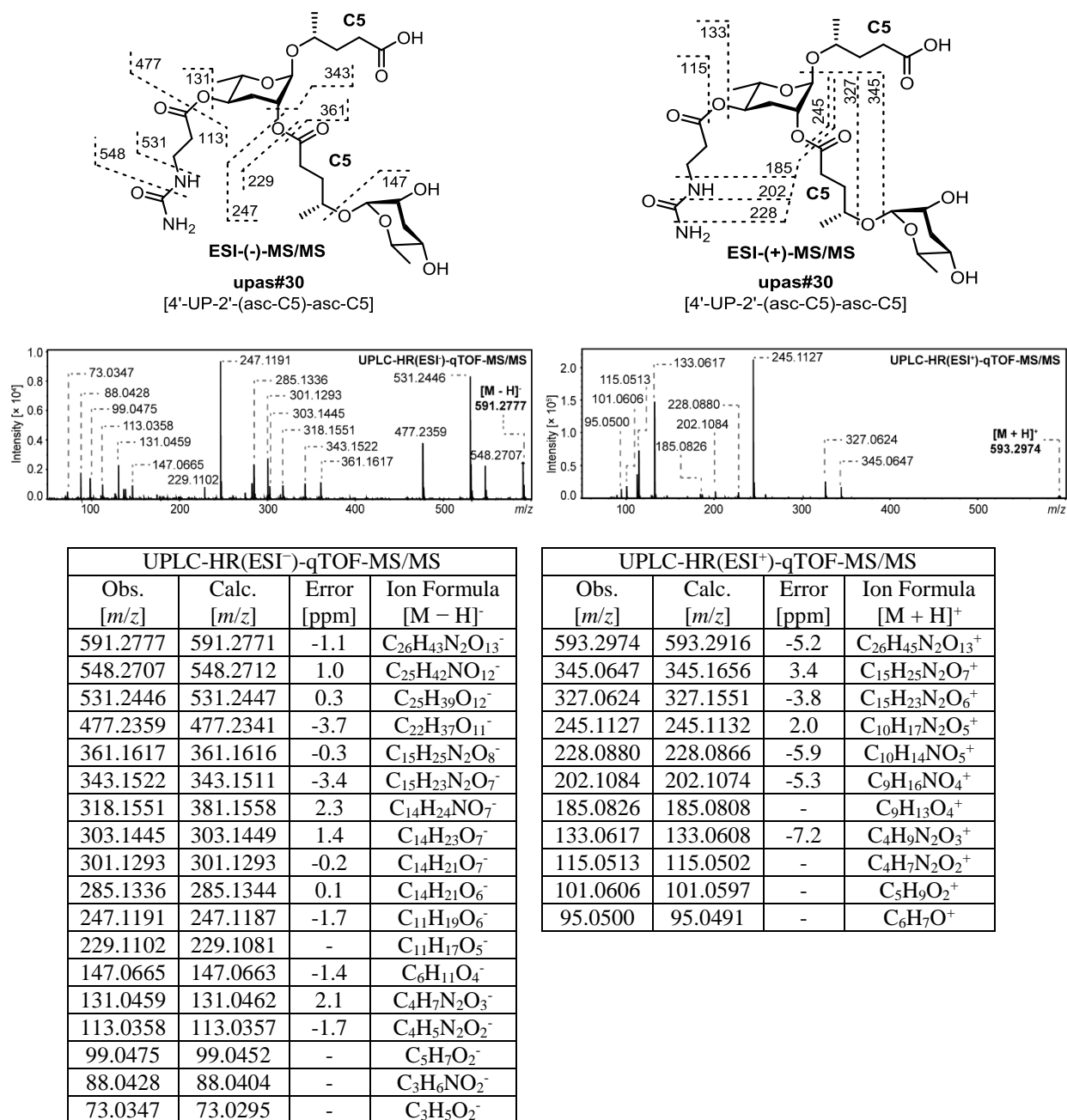


UPLC-HR(ESI <sup>+</sup> )-qTOF-MS/MS			
Obs. [m/z]	Calc. [m/z]	Error [ppm]	Ion Formula [M + H] <sup>+</sup>
565.2628	565.2603	-4.3	C <sub>24</sub> H <sub>41</sub> N <sub>2</sub> O <sub>13</sub> <sup>+</sup>
345.1665	345.1656	-2.4	C <sub>15</sub> H <sub>25</sub> N <sub>2</sub> O <sub>7</sub> <sup>+</sup>
331.1516	331.1500	-4.8	C <sub>14</sub> H <sub>23</sub> N <sub>2</sub> O <sub>7</sub> <sup>+</sup>
327.1569	327.1551	-5.5	C <sub>15</sub> H <sub>23</sub> N <sub>2</sub> O <sub>6</sub> <sup>+</sup>
313.1407	313.1394	-4.1	C <sub>14</sub> H <sub>21</sub> N <sub>2</sub> O <sub>6</sub> <sup>+</sup>
296.1145	296.1129	-5.6	C <sub>14</sub> H <sub>18</sub> NO <sub>6</sub> <sup>+</sup>
285.1343	285.1333	-3.5	C <sub>14</sub> H <sub>21</sub> O <sub>6</sub> <sup>+</sup>
271.1186	271.1176	-3.8	C <sub>13</sub> H <sub>19</sub> O <sub>6</sub> <sup>+</sup>
259.1289	259.1288	-0.3	C <sub>11</sub> H <sub>19</sub> N <sub>2</sub> O <sub>5</sub> <sup>+</sup>
253.1087	253.1071	-6.6	C <sub>13</sub> H <sub>17</sub> O <sub>5</sub> <sup>+</sup>
245.1148	245.1132	-6.4	C <sub>10</sub> H <sub>17</sub> N <sub>2</sub> O <sub>5</sub> <sup>+</sup>
228.0895	228.0866	-	C <sub>10</sub> H <sub>14</sub> NO <sub>5</sub> <sup>+</sup>
217.1083	217.1071	-5.9	C <sub>10</sub> H <sub>17</sub> O <sub>5</sub> <sup>+</sup>
199.0985	199.0965	-	C <sub>10</sub> H <sub>15</sub> O <sub>4</sub> <sup>+</sup>
181.0870	181.0859	-6.1	C <sub>10</sub> H <sub>13</sub> O <sub>3</sub> <sup>+</sup>
147.0772	147.0764	-5.1	C <sub>5</sub> H <sub>11</sub> N <sub>2</sub> O <sub>3</sub> <sup>+</sup>
133.0616	133.0608	-6.0	C <sub>4</sub> H <sub>9</sub> N <sub>2</sub> O <sub>3</sub> <sup>+</sup>
115.0512	115.0502	-	C <sub>4</sub> H <sub>7</sub> N <sub>2</sub> O <sub>2</sub> <sup>+</sup>
113.0604	113.0597	-	C <sub>6</sub> H <sub>9</sub> O <sub>2</sub> <sup>+</sup>
95.0501	95.0491	-	C <sub>6</sub> H <sub>7</sub> O <sup>+</sup>
87.0449	87.0441	-	C <sub>4</sub> H <sub>7</sub> O <sub>2</sub> <sup>+</sup>

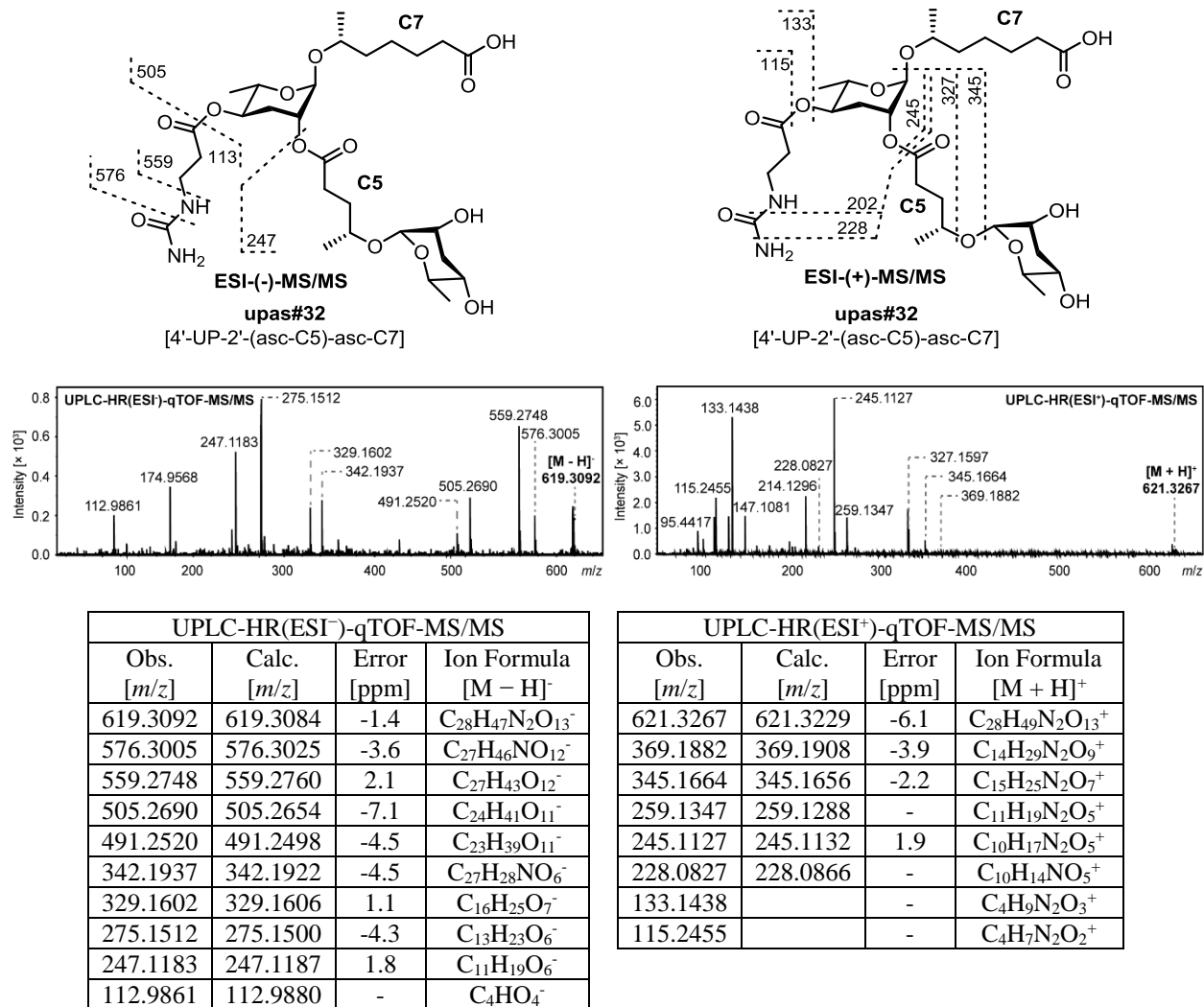
S51



**Figure 5.** UPLC-HR(ESI<sup>-/+</sup>)-qTOF-MS/MS mass spectral data of upas#28 [4'-UP-2'-(asc-C5)-asc-C4, **15**] in both negative and positive ion modes. Observation of the fragment ion signal for C<sub>4</sub>H<sub>7</sub>N<sub>2</sub>O<sub>3</sub><sup>-</sup> (*m/z* 131.0466 [M - H]<sup>-</sup>) indicated the presence of a ureidopropionic acid group in upas#28 [4'-UP-2'-(asc-C5)-asc-C4, **15**]. Observation of two other ion signals of C<sub>15</sub>H<sub>23</sub>N<sub>2</sub>O<sub>6</sub><sup>-</sup> (*m/z* 327.1555 [M + H]<sup>+</sup>) and C<sub>15</sub>H<sub>25</sub>N<sub>2</sub>O<sub>7</sub><sup>-</sup> (*m/z* 345.1675 [M + H]<sup>+</sup>) indicated that the first ascaroside in upas#28 [4'-UP-2'-(asc-C5)-asc-C4, **15**] should carry a C5 side chain.



**Figure 6.** UPLC-HR(ESI<sup>+/+</sup>)-qTOF-MS/MS mass spectral data of upas#30 [4'-UP-2'-(asc-C5)-asc-C5] in both negative and positive ion modes.



**Figure 7.** UPLC-HR(ESI<sup>±</sup>)-qTOF-MS/MS mass spectral data of upas#32 [4'-UP-2'-(asc-C5)-asc-C7] in both negative and positive ion modes.

## References

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