

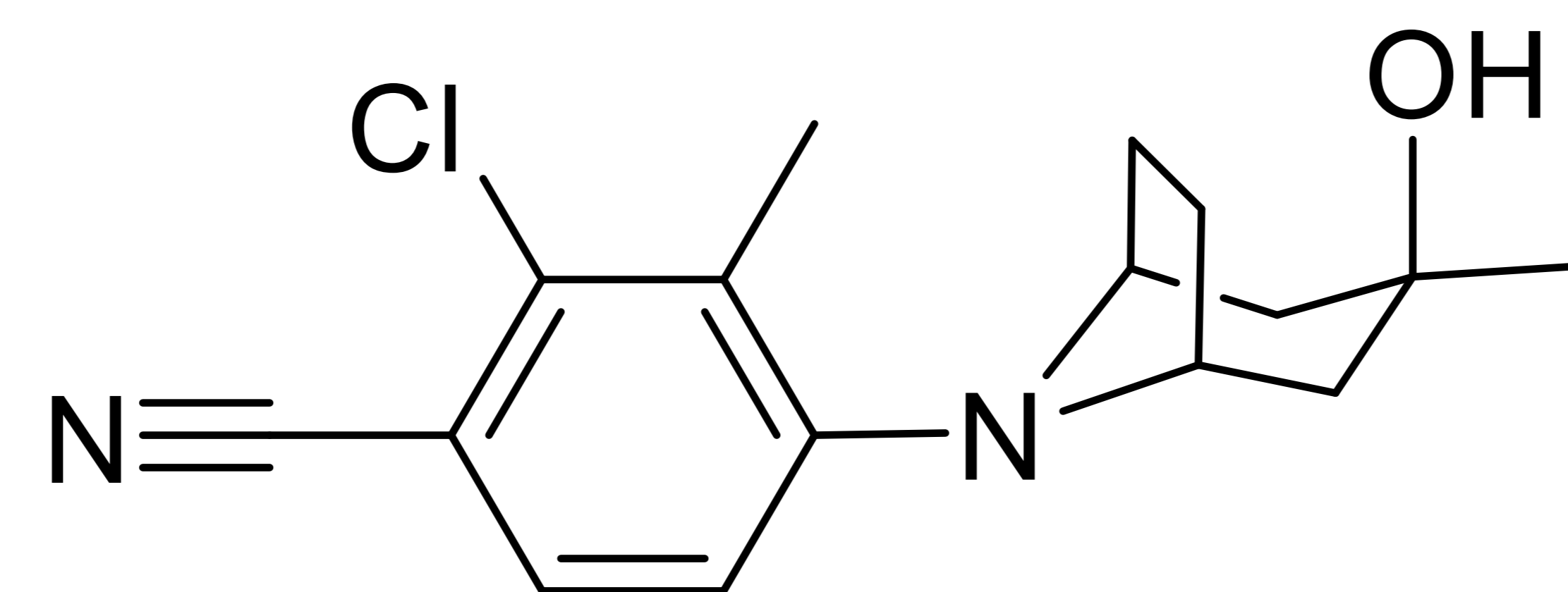
S1: Supplementary material

Structural elucidation and fragmentation of ACP-105 and tentatively identified metabolites

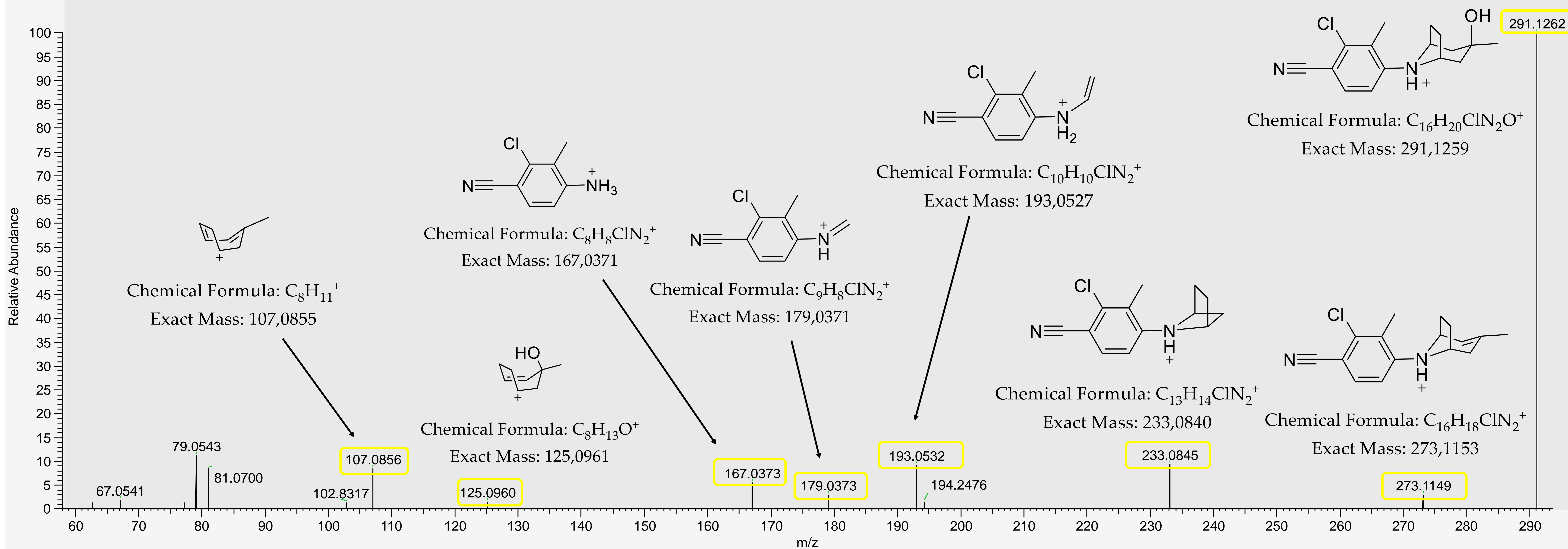
Metabolite	Metabolic transformation	Elemental composition [M+H] ⁺	Theoretical m/z
ACP-105	Administered substance	C ₁₆ H ₂₀ ClN ₂ O ⁺	291.1259
M1a-c	Monohydroxylation	C ₁₆ H ₂₀ ClN ₂ O ₂ ⁺	307.1208
M2a-b	Dihydroxylation	C ₁₆ H ₂₀ ClN ₂ O ₃ ⁺	323.1157
M3a-b	Loss of 2H	C ₁₆ H ₁₈ ClN ₂ O ⁺	289.1102
M4a-b	Monohydroxylation + Loss of 2H	C ₁₆ H ₁₈ ClN ₂ O ₂ ⁺	305.1051
M5a-b	Dihydroxylation + Loss of 2H	C ₁₆ H ₁₈ ClN ₂ O ₃ ⁺	321.1000
M6a-c	Trihydroxylation + Loss of 2H	C ₁₆ H ₁₈ ClN ₂ O ₄ ⁺	337.0950
M7	Glucuronidation	C ₂₂ H ₂₈ ClN ₂ O ₇ ⁺	467.1580
M8a-d	Monohydroxylation + Glucuronidation	C ₂₂ H ₂₈ ClN ₂ O ₈ ⁺	483.1529
M9a-b	Dihydroxylation + Glucuronidation	C ₂₂ H ₂₈ ClN ₂ O ₉ ⁺	499.1478

ACP-105

Administrated substance
Retention time: 11.59 min
m/z: 291.1262



MNB_20200429_03 #4841 RT: 11.62 AV: 1 NL: 2.65E+006
T: FTMS + p ESI d Full ms2 291.1261 @hcd35.00 [50.0000-315.0000]

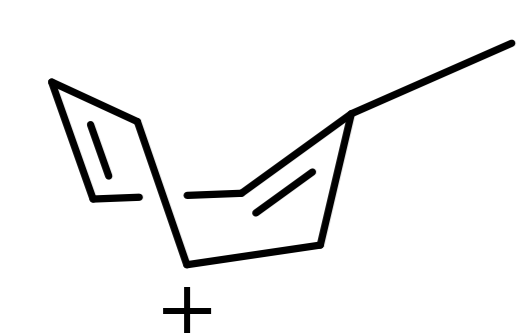
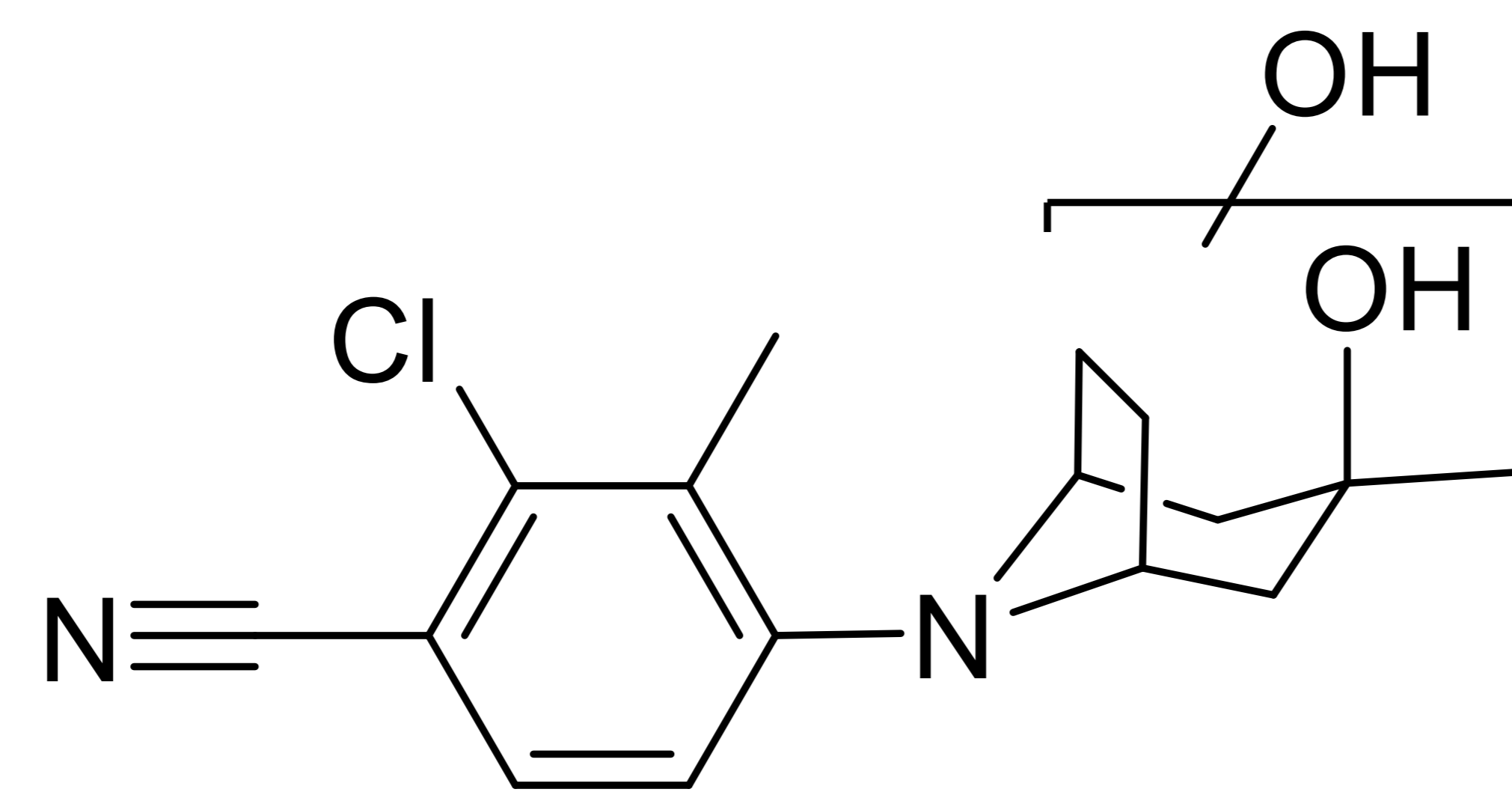


M1a

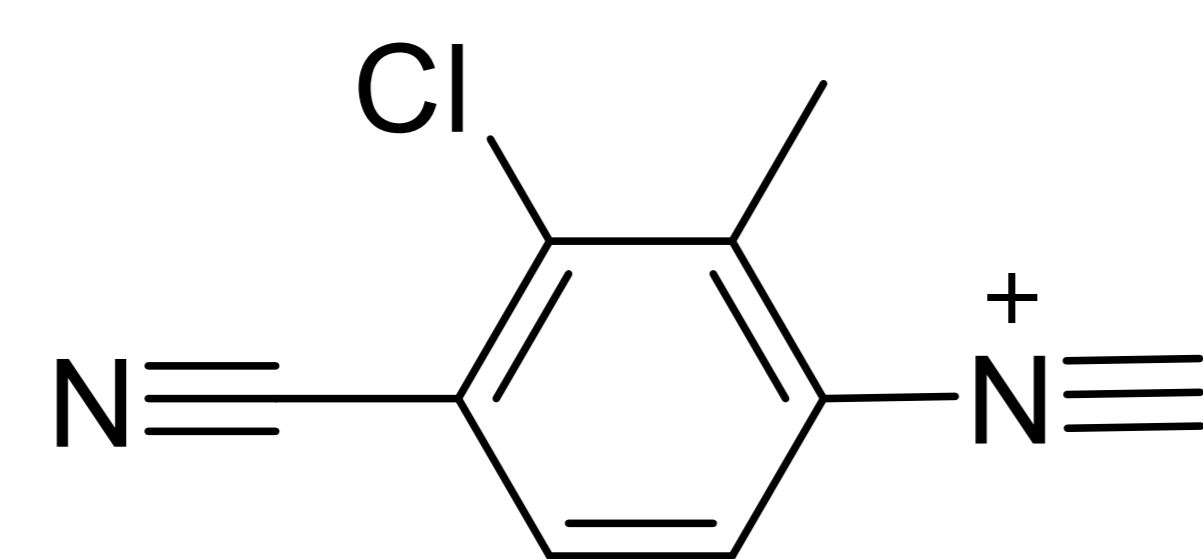
Monohydroxylation

Retention time: 8.65 min

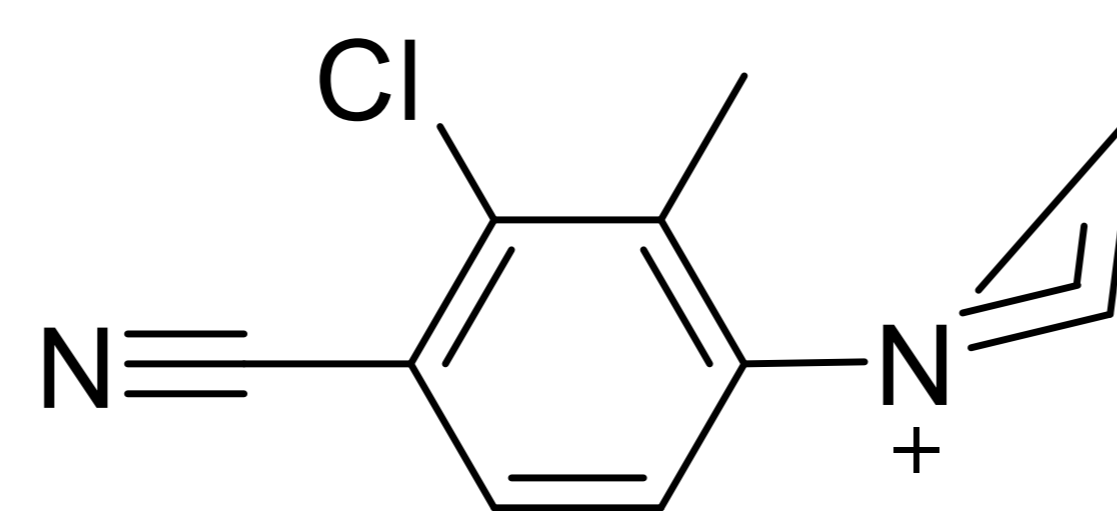
m/z: 307.1209



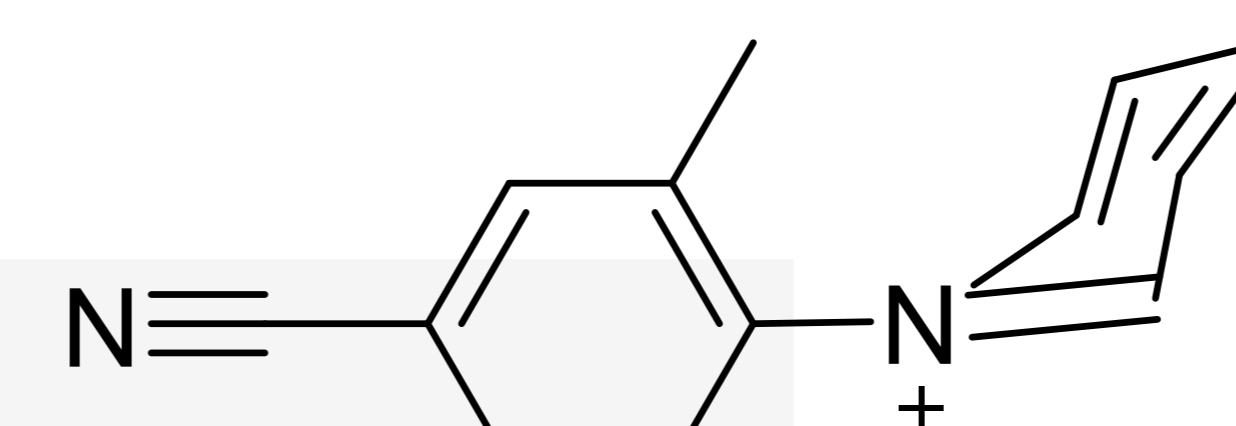
Chemical Formula: $C_8H_{11}^+$
Exact Mass: 107,0855



Chemical Formula: $C_9H_6ClN_2^+$
Exact Mass: 177,0214

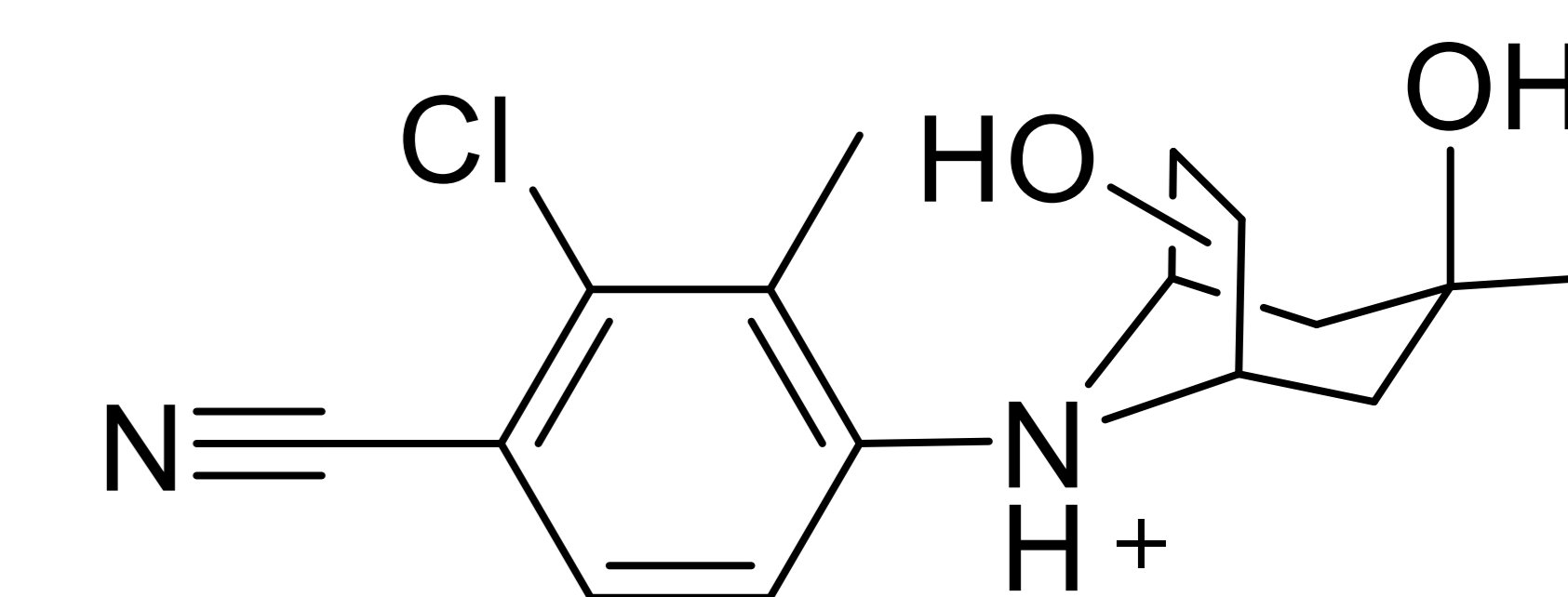
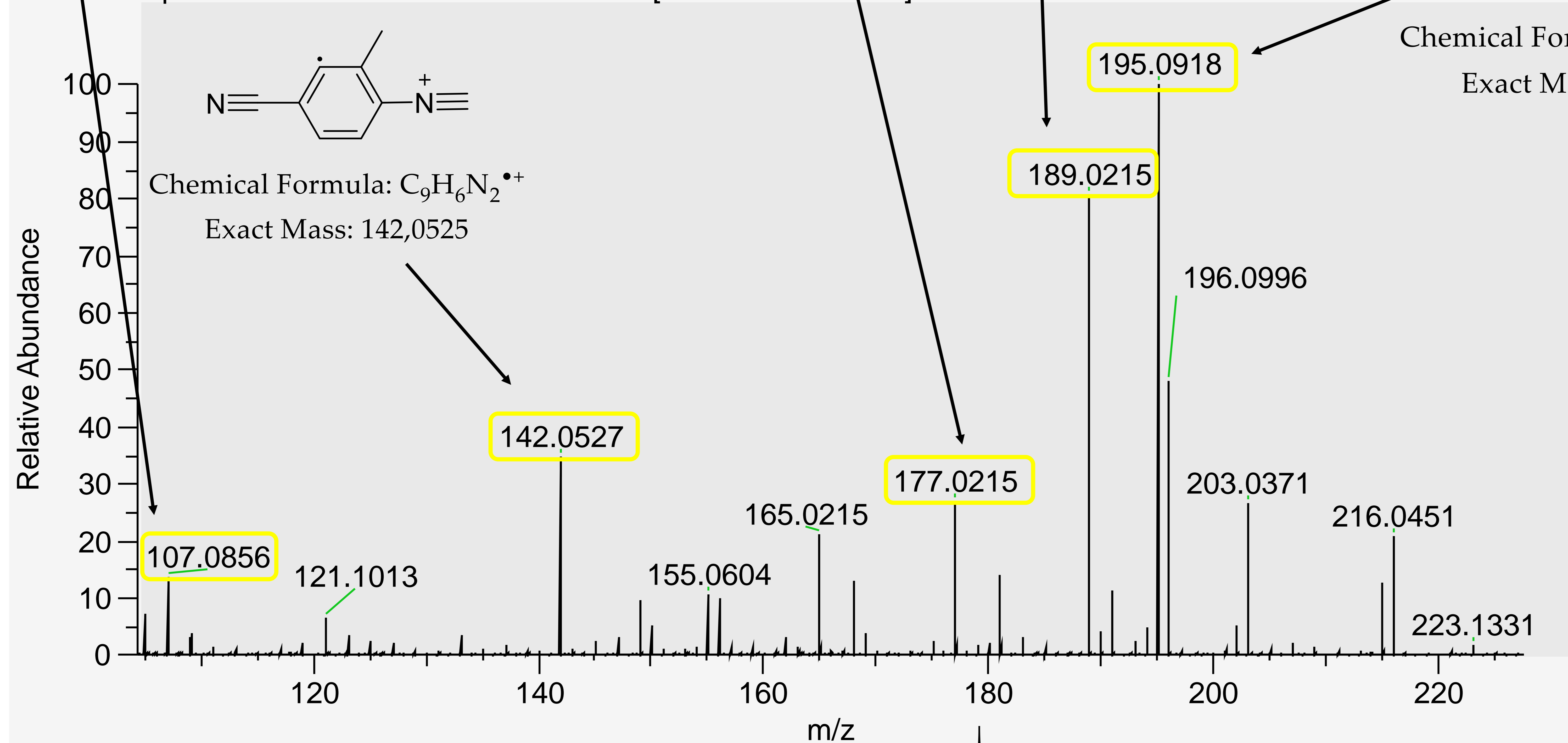


Chemical Formula: $C_{10}H_6ClN_2^+$
Exact Mass: 189,0214

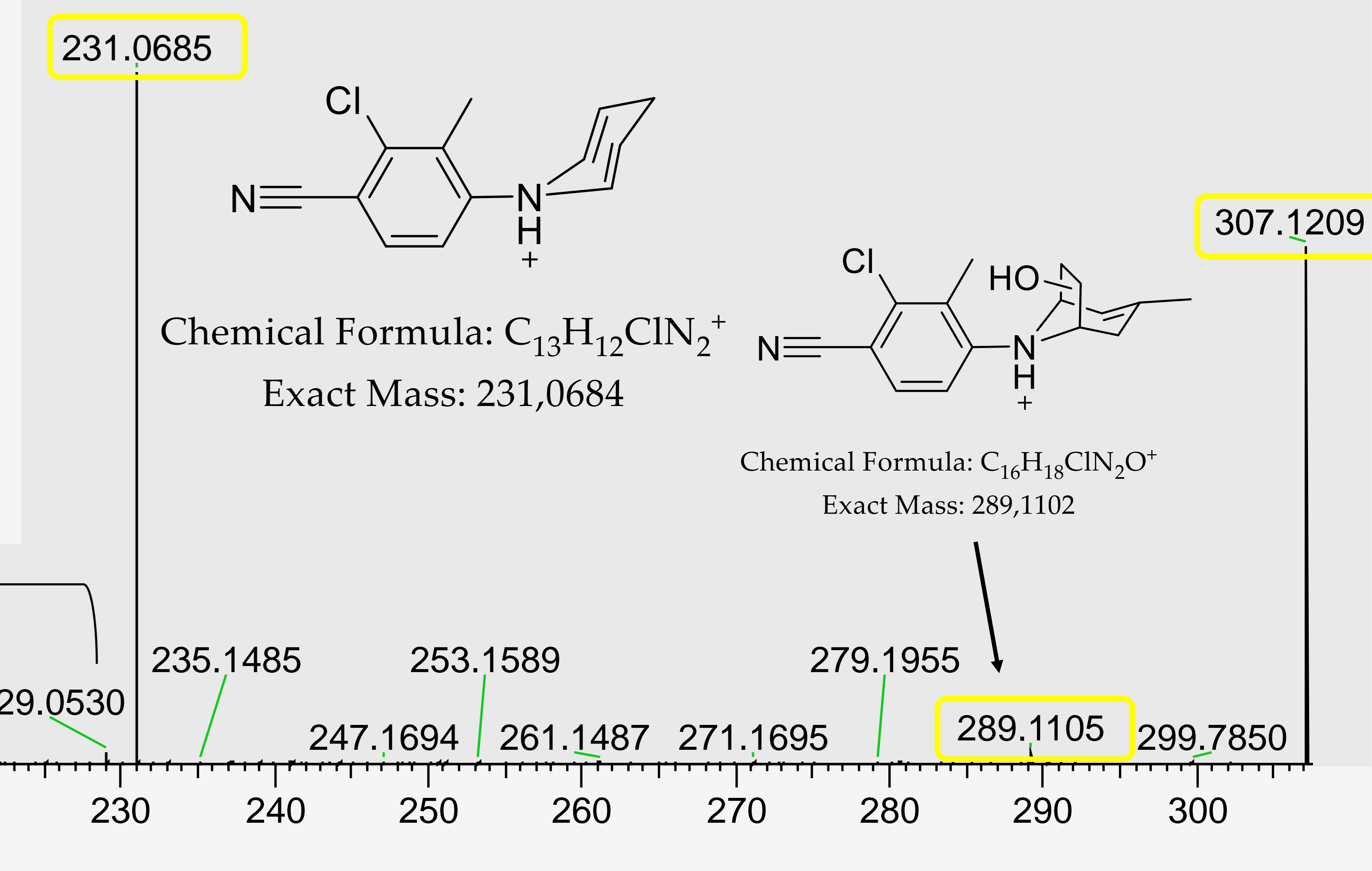


Chemical Formula: $C_{13}H_{11}N_2^+$
Exact Mass: 195,0917

MNE_20200730_014 #1810-1826 RT: 8.62-8.69 AV: 17 SB: 15 8.59-8.62, 8.69-8.72 NL: 5.20E+005
T: FTMS + p ESI Full ms2 307.1208@hcd38.33 [50.0000-330.0000]



Chemical Formula: $C_{16}H_{20}ClN_2O_2^+$
Exact Mass: 307,1208

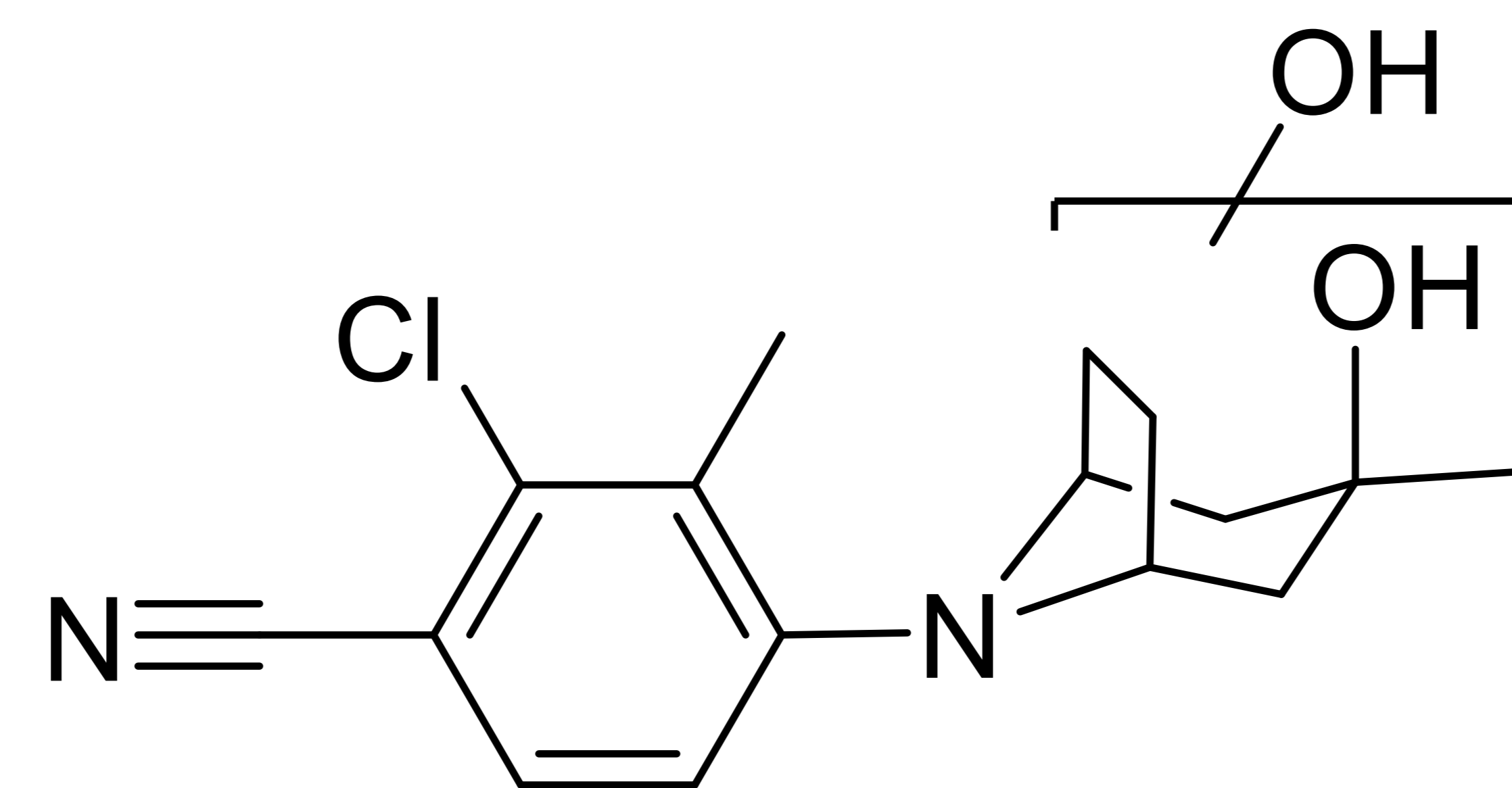


M1b

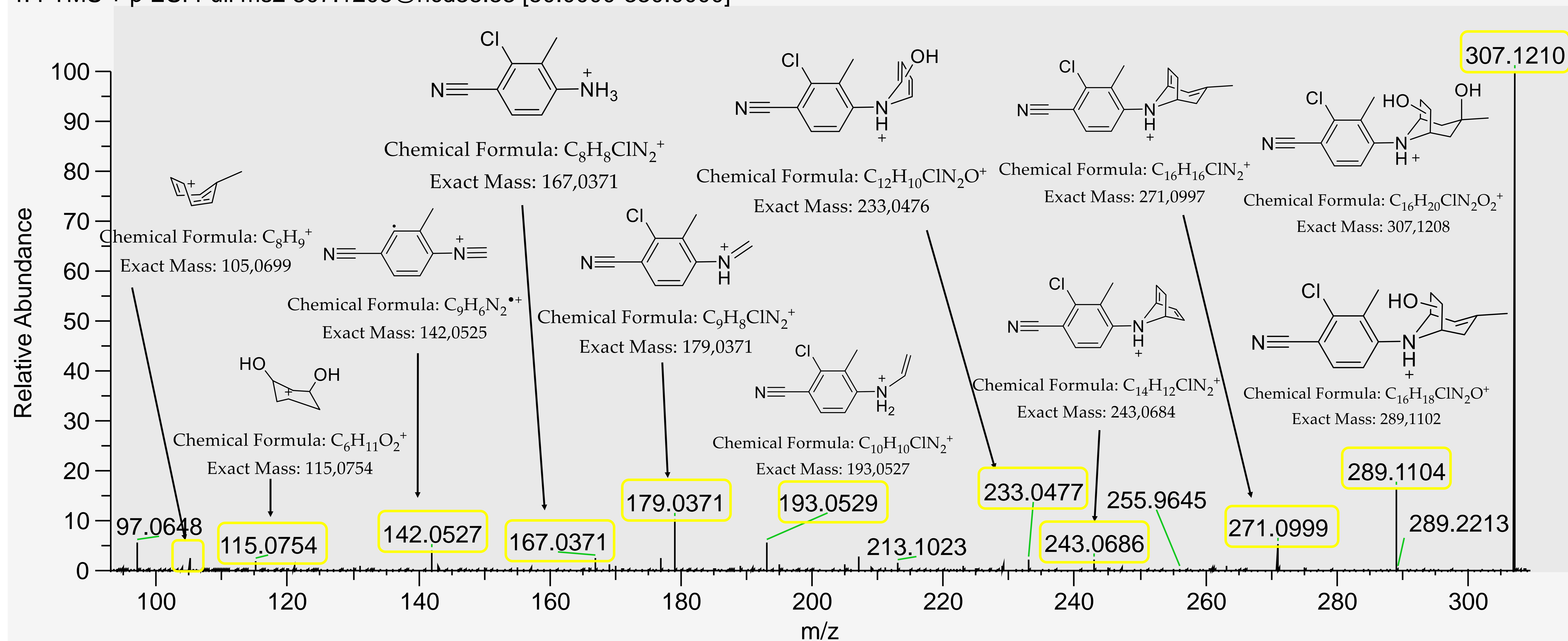
Monohydroxylation

Retention time: 9.00 min

m/z: 307.1210



MNB_20200730_014 #1894-1908 RT: 9-9.06 AV: 15 SB: 26 8.93-9.00 , 9.06-9.10 NL: 3.51E+005
T: FTMS + p ESI Full ms2 307.1208@hcd38.33 [50.0000-330.0000]

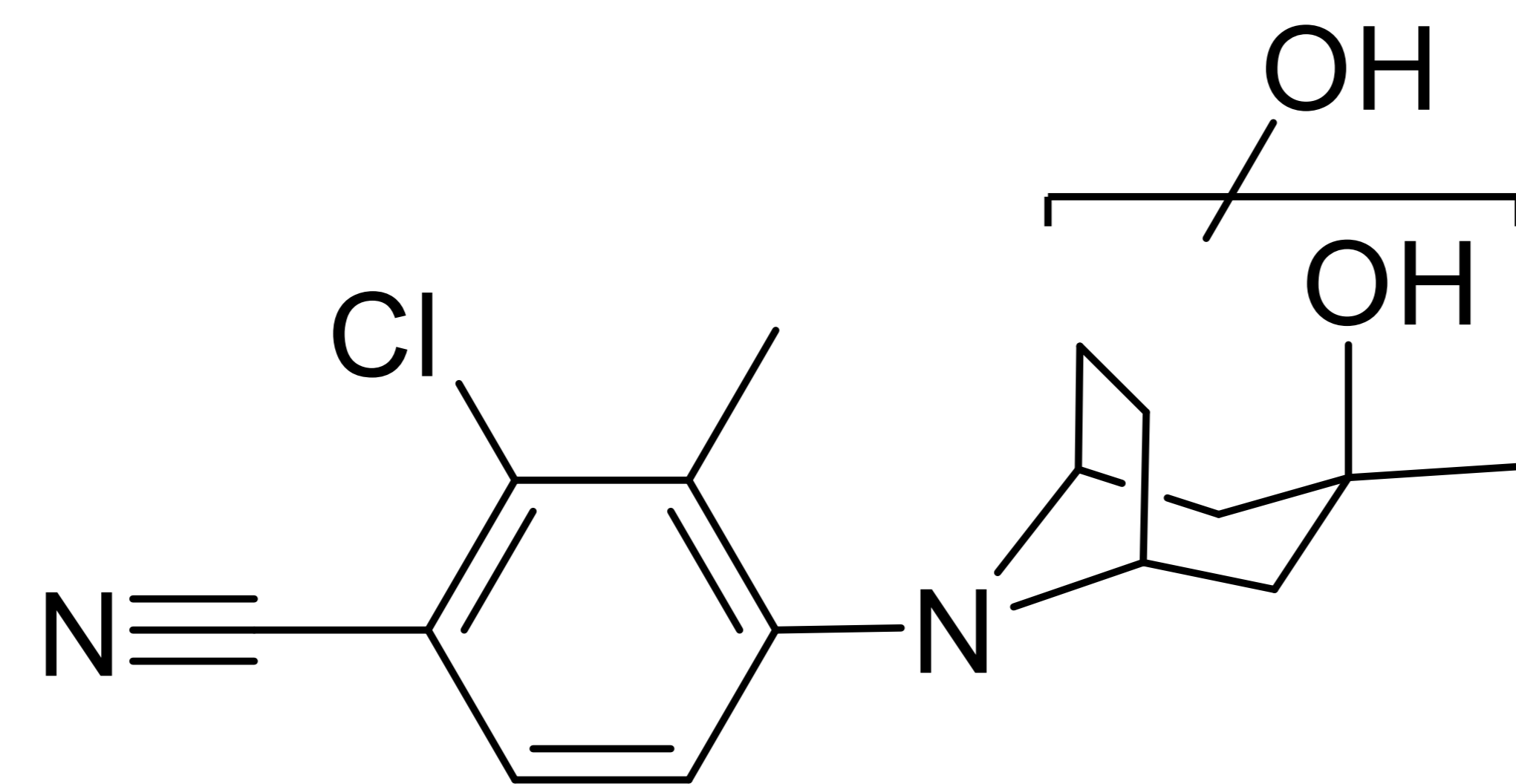


M1c

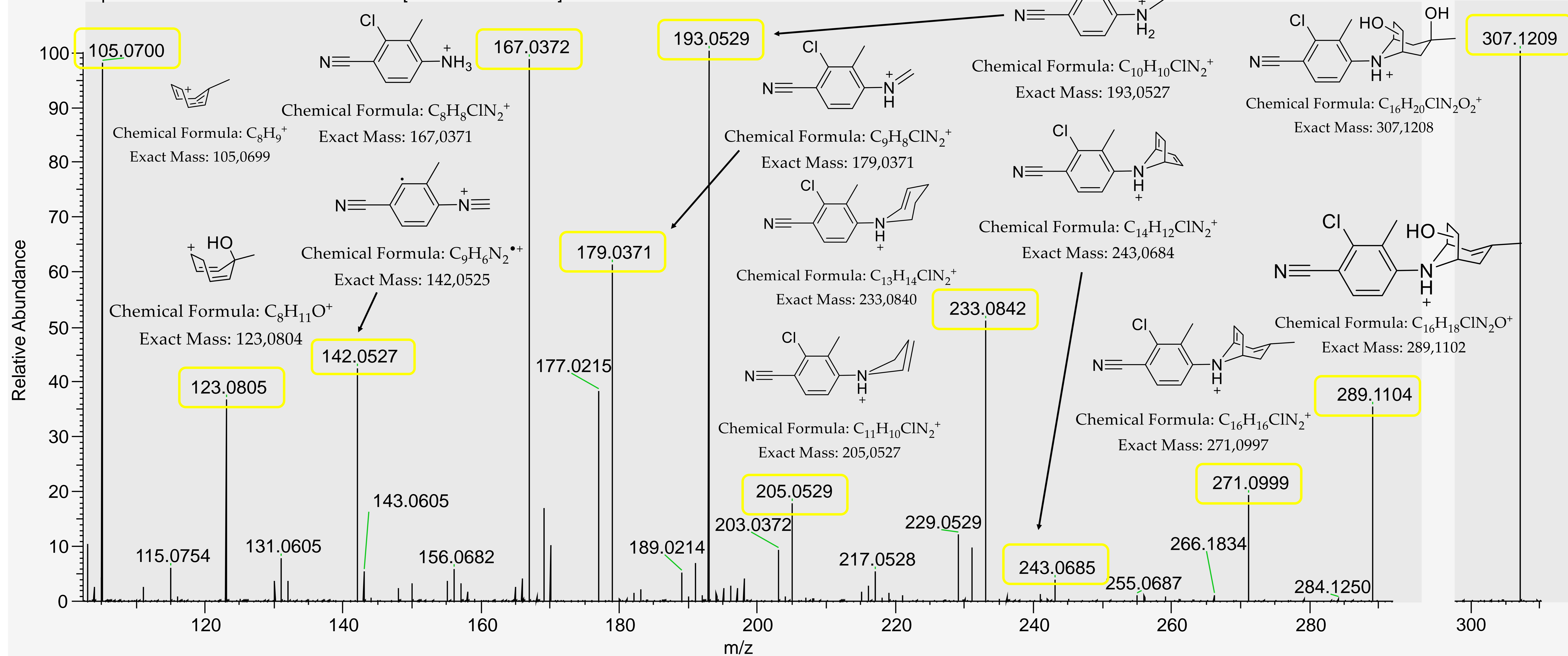
Monohydroxylation

Retention time: 9.14 min

m/z: 307.1209



MNB_20200730_014 #1921-1937 RT: 9.12-9.19 AV: 17 SB: 24 9.07-9.12 , 9.19-9.24 NL: 8.09E+005
T: FTMS + p ESI Full ms2 307.1208@hcd38.33 [50.0000-330.0000]

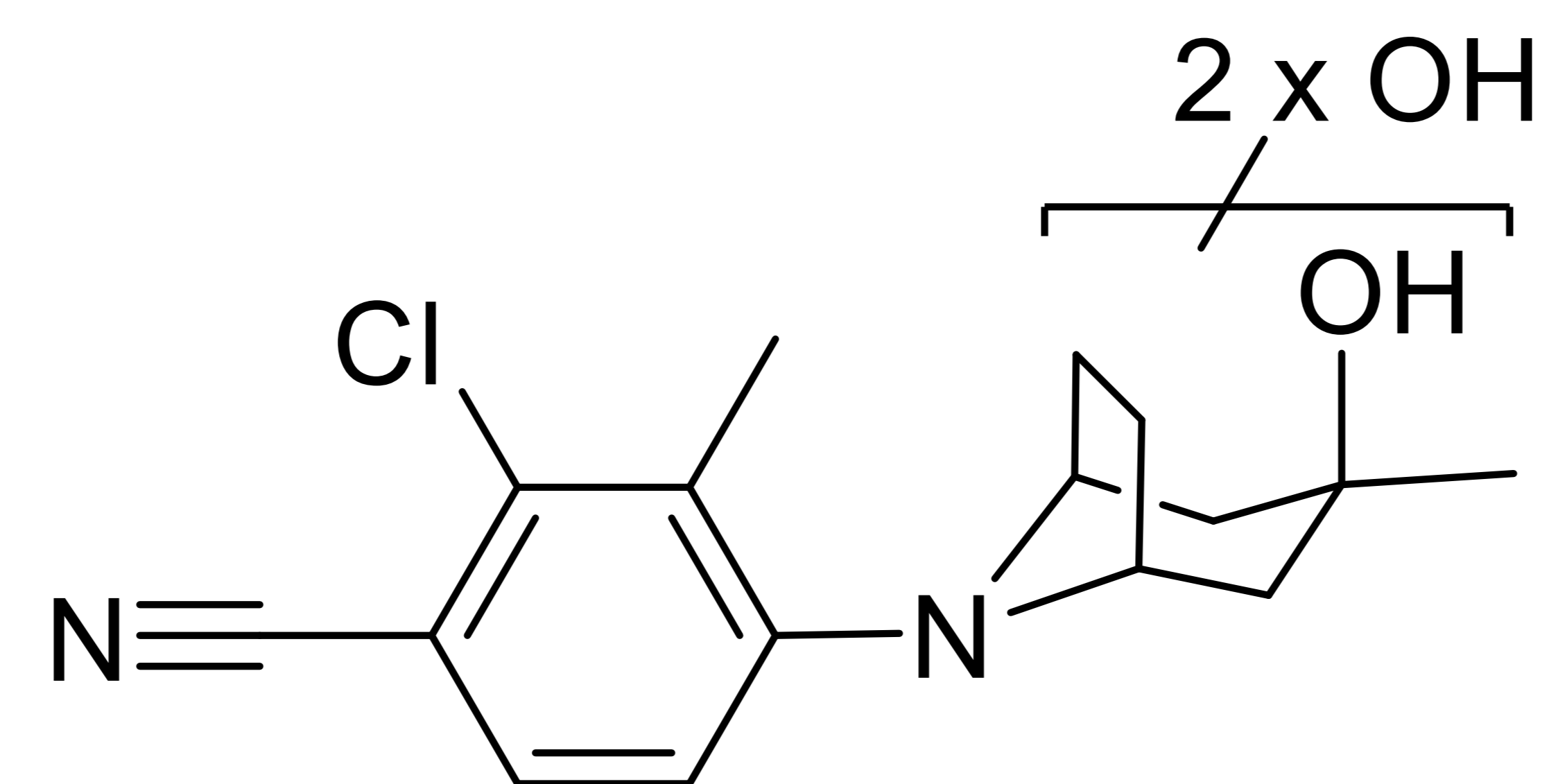


M2a

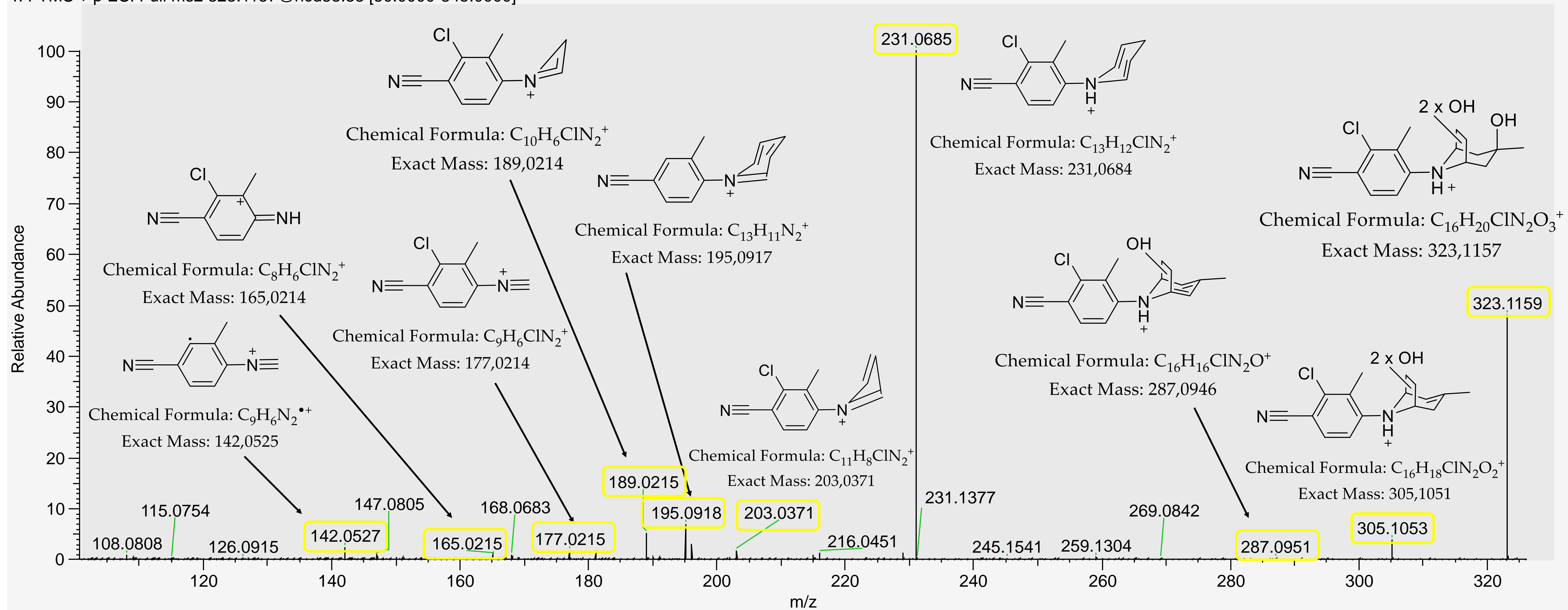
Dihydroxylation

Retention time: 7.03 min

m/z: 323.1159



MNB_20200730_013 #1449-1465 RT: 7.01-7.08 AV: 17 SB: 24 6.96-7.01 , 7.08-7.13 NL: 1.18E+006
T: FTMS + p ESI Full ms2 323.1157@hcd38.33 [50.0000-345.0000]

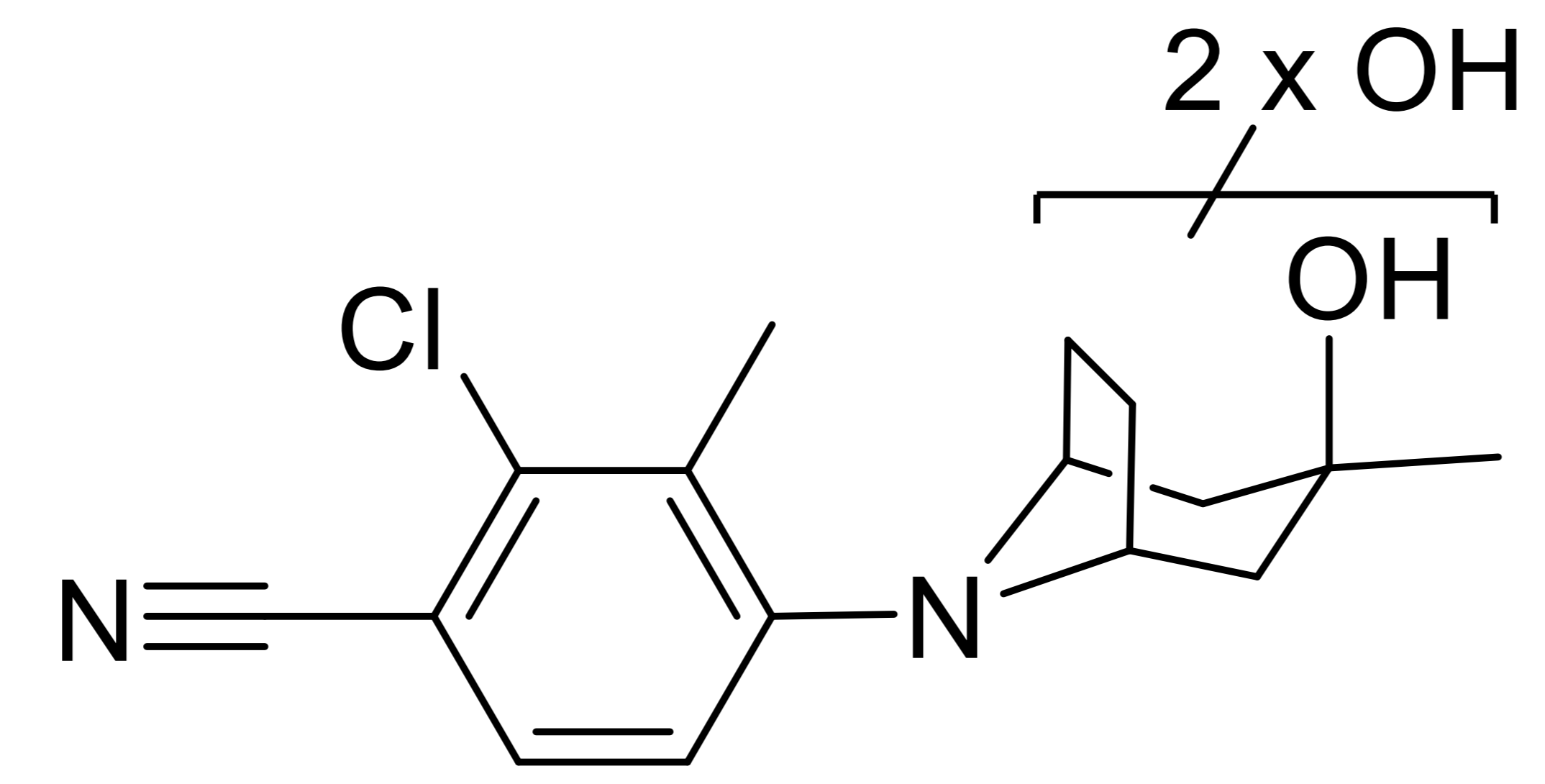


M2b

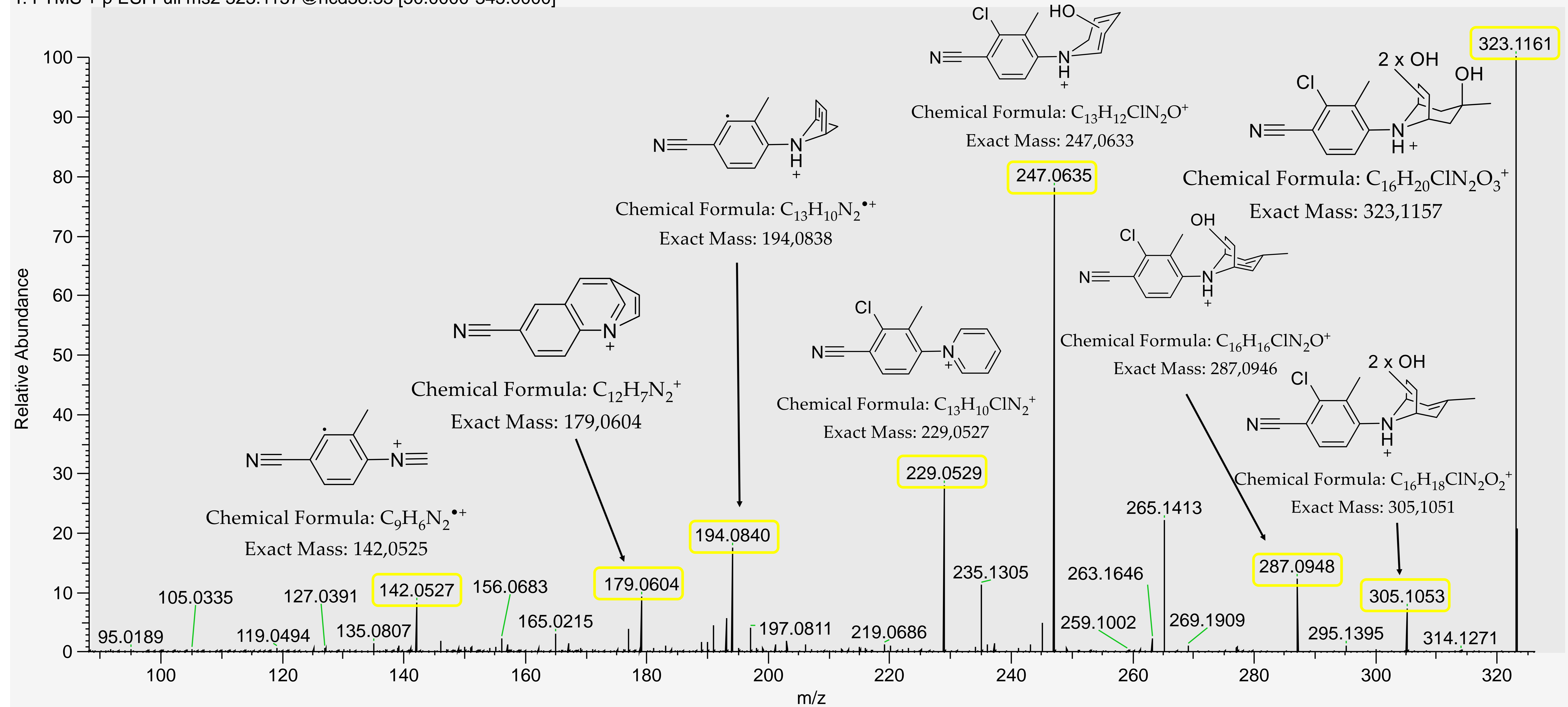
Dihydroxylation

Retention time: 7.63 min

m/z: 323.1161



MNB_20200730_013 #1583-1595 RT: 7.61-7.66 AV: 13 SB: 14 7.58-7.61 , 7.66-7.69 NL: 2.28E+005
T: FTMS + p ESI Full ms2 323.1157@hcd38.33 [50.0000-345.0000]

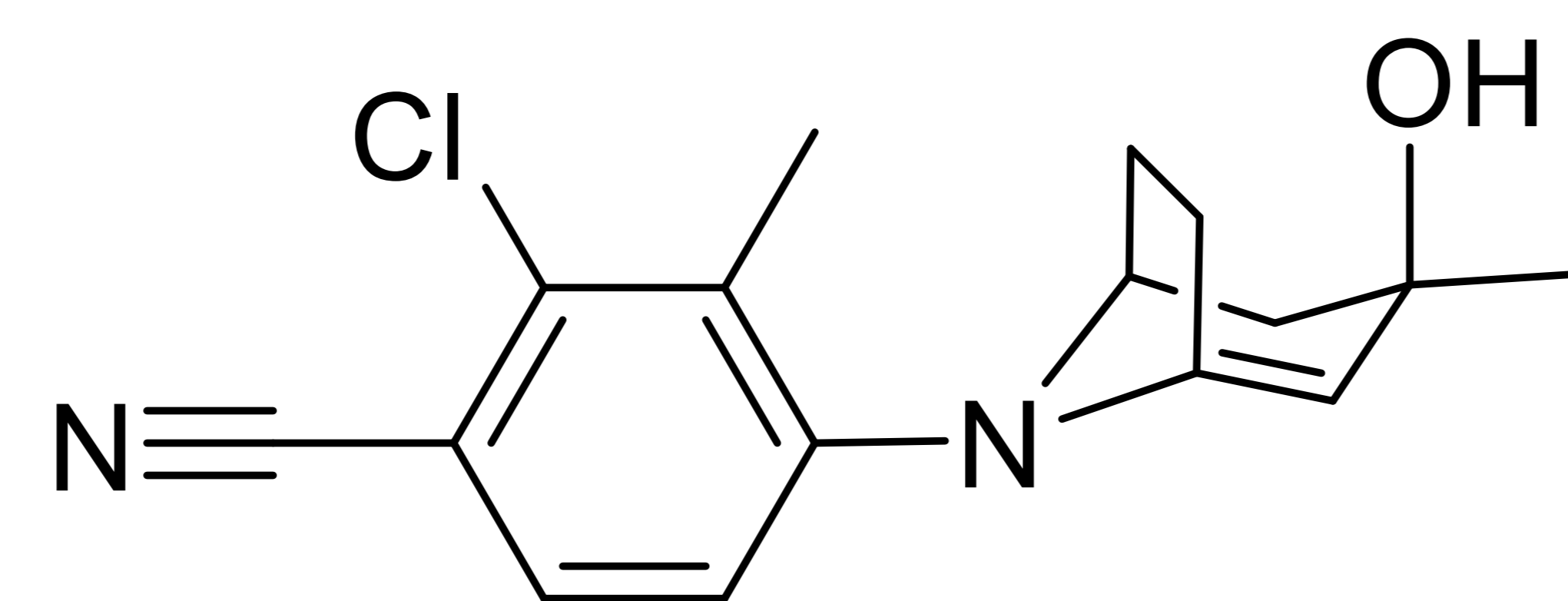


M3a

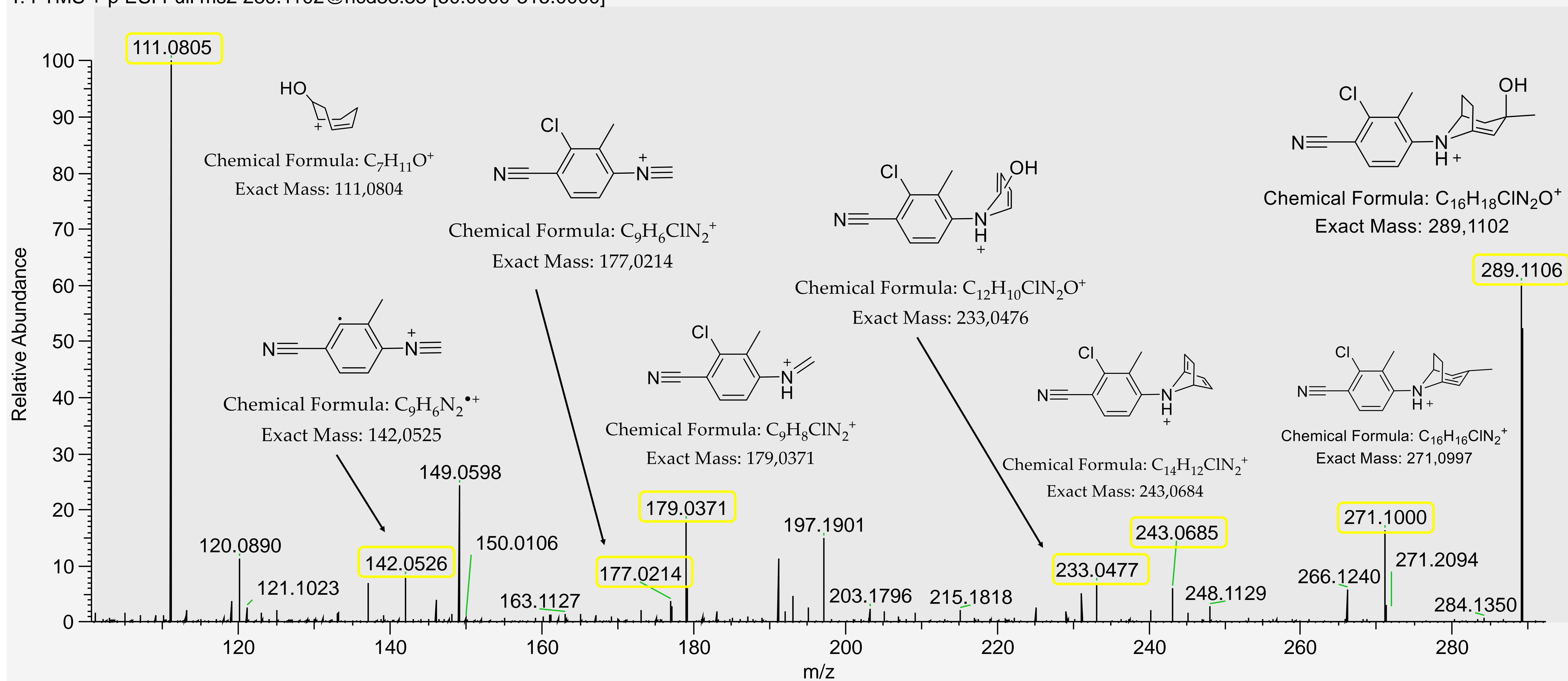
Loss of 2H

Retention time: 10.11 min

m/z: 289.1106



MNB_20200730_012 #2097-2108 RT: 10.11-10.16 AV: 12 SB: 74 9.90-10.11 , 10.16-10.30 NL: 3.57E+004
T: FTMS + p ESI Full ms2 289.1102@hcd38.33 [50.0000-315.0000]

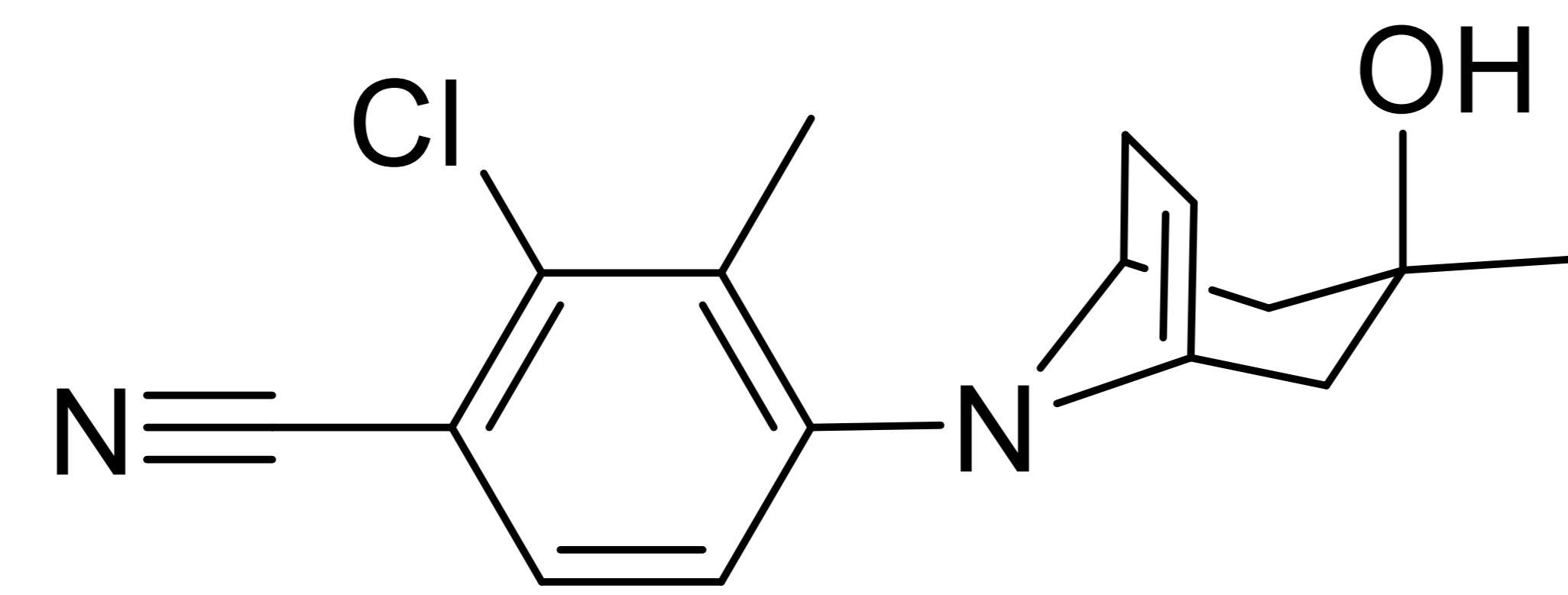


M3b

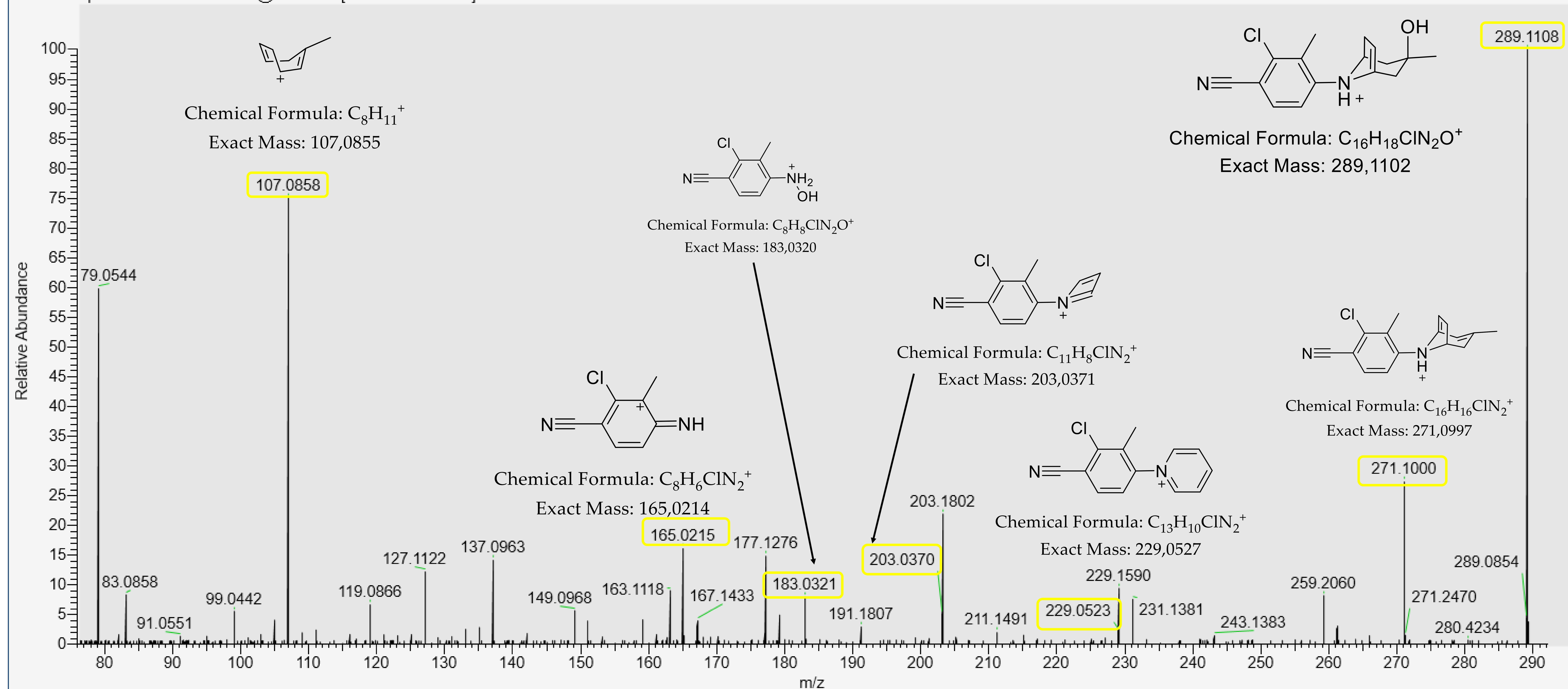
Loss of 2H

Retention time: 11.56 min

m/z: 289.1108



MNB_20200730_012 #2391-2405 RT: 11.53-11.6 AV: 15 SB: 40 11.45-11.53, 11.60-11.70 NL: 9.36E+003
T: FTMS + p ESI Full ms2 289.1102@hcd38.33 [50.0000-315.0000]

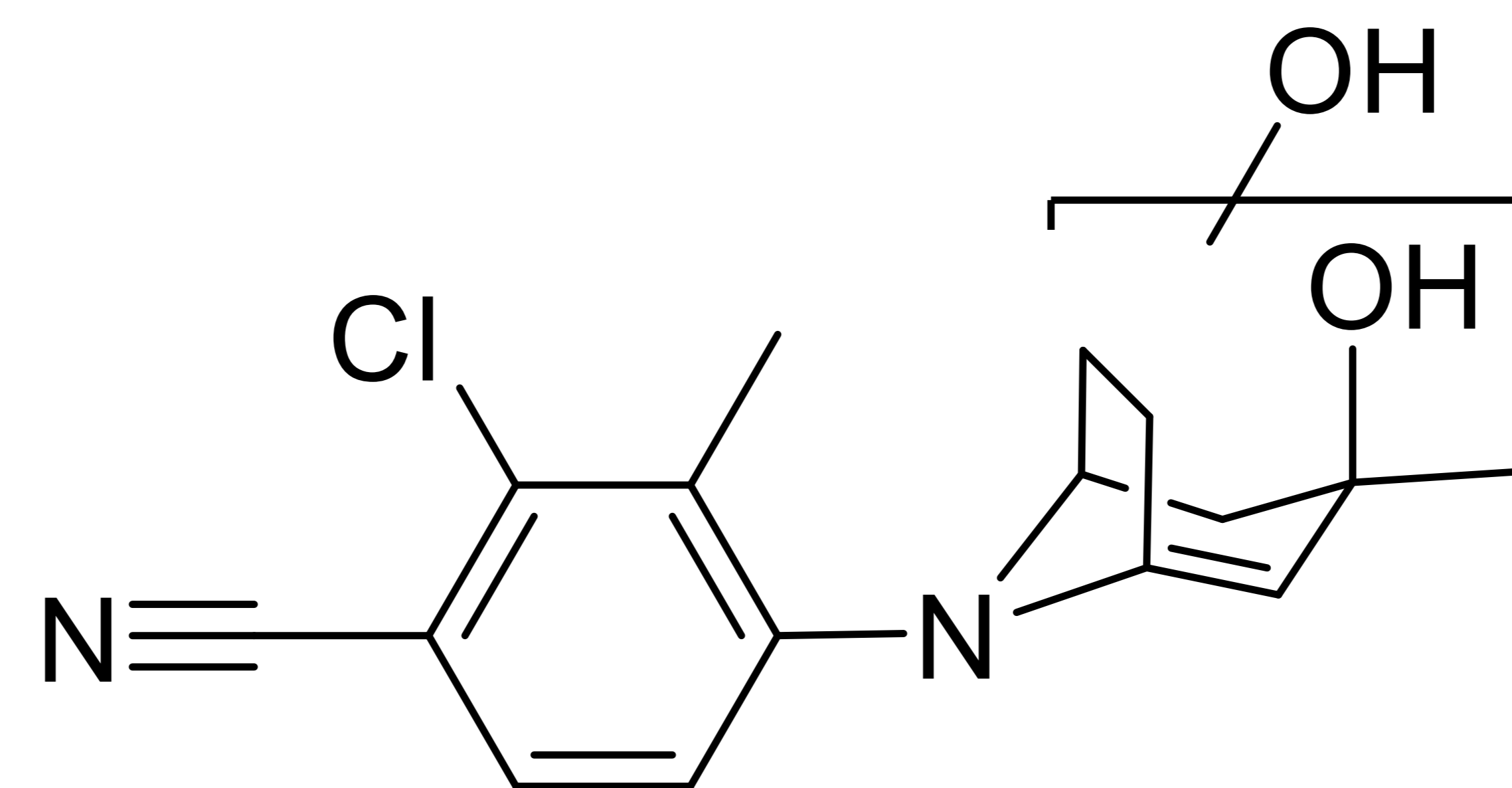


M4a

Monohydroxylation + loss of 2H

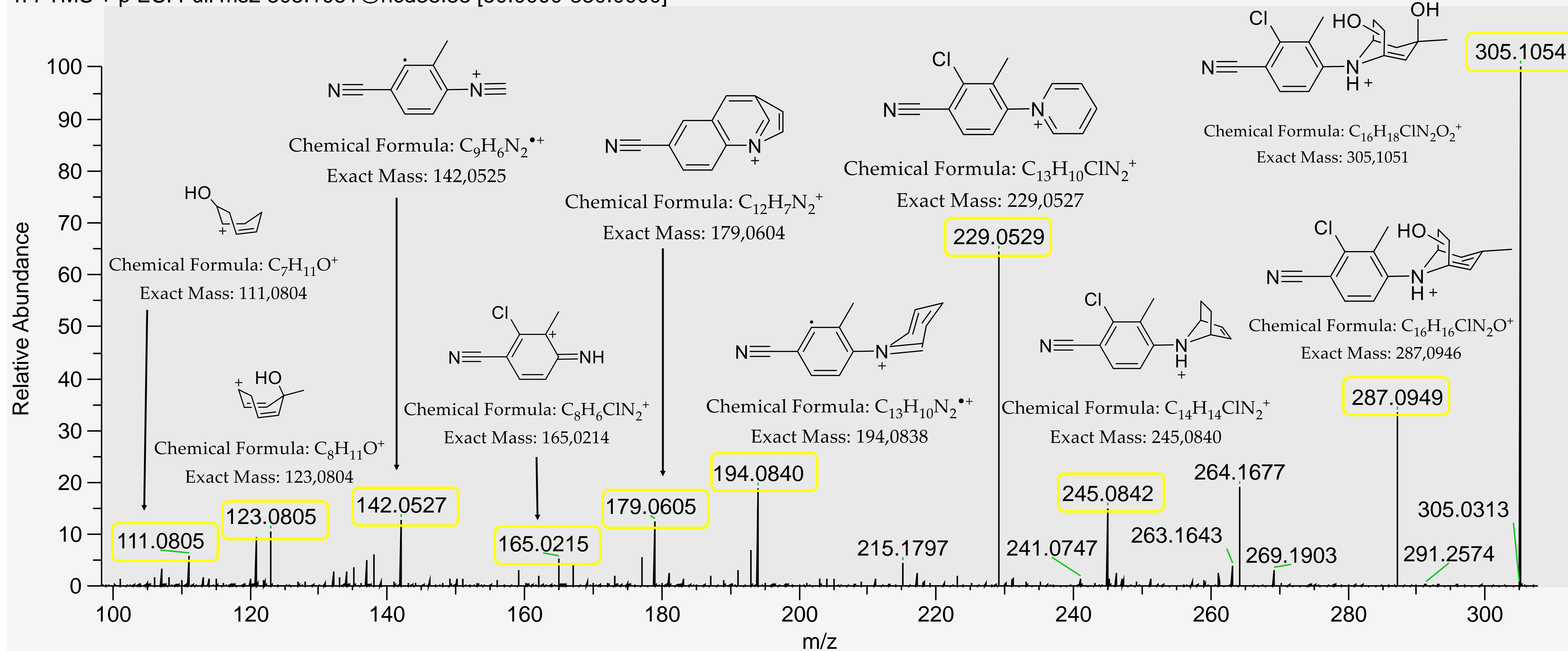
Retention time: 8.88 min

m/z: 305.1054



MNB_20200730_009 #1855-1871 RT: 8.86-8.93 AV: 17 SB: 28 8.80-8.86 , 8.93-8.99 NL: 9.49E+004

T: FTMS + p ESI Full ms2 305.1051 @hcd38.33 [50.0000-330.0000]

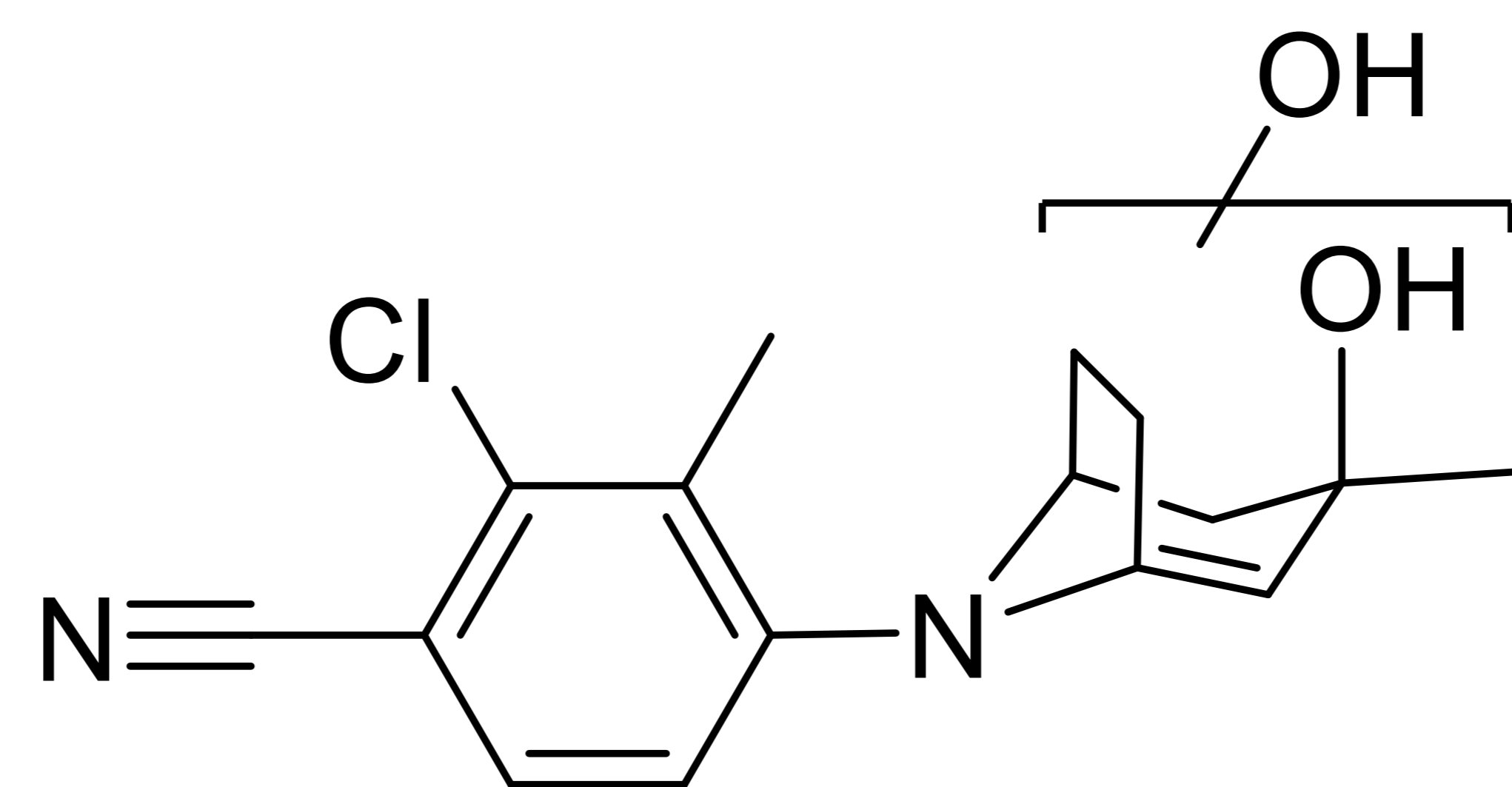


M4b

Monohydroxylation + loss of 2H

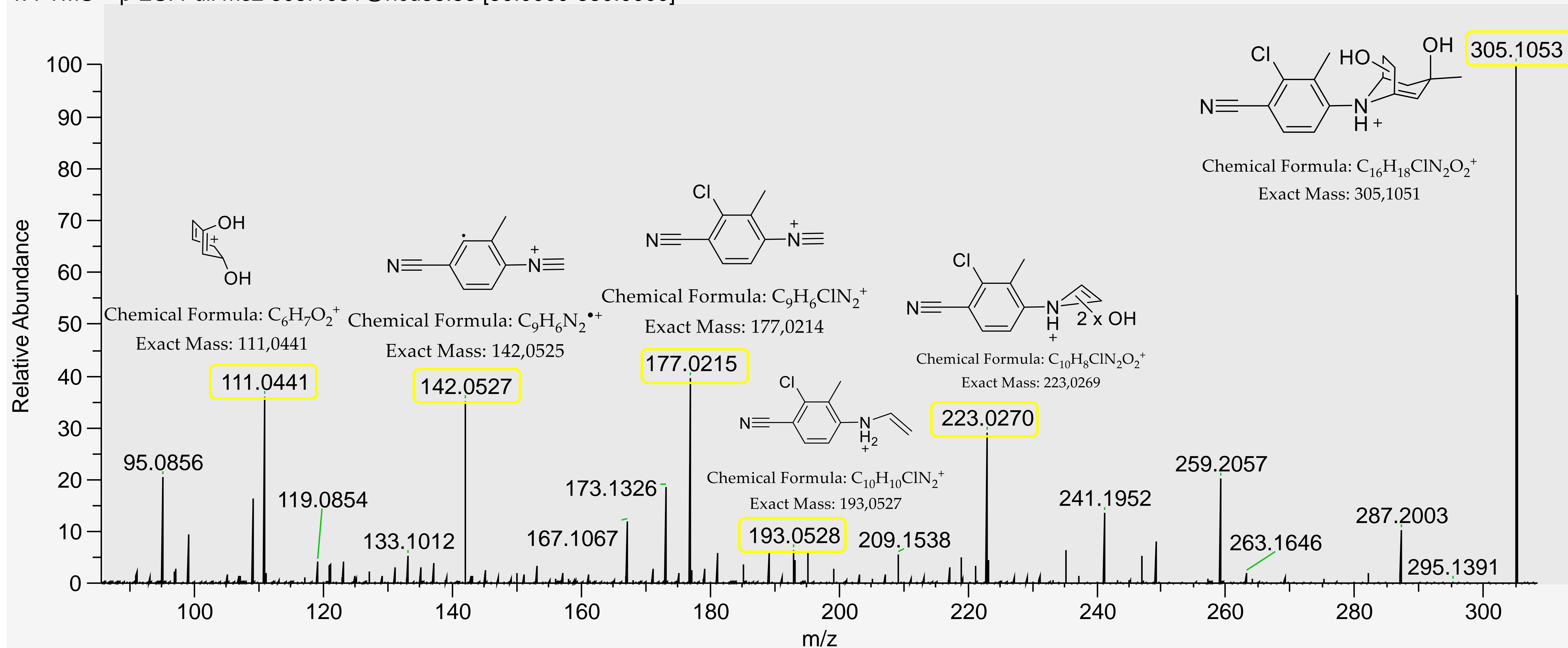
Retention time: 9.83 min

m/z: 305.1053



MNB_20200730_009 #2067-2080 RT: 9.82-9.88 AV: 14 SB: 28 9.75-9.82 , 9.88-9.93 NL: 1.65E+005

T: FTMS + p ESI Full ms2 305.1051 @hcd38.33 [50.0000-330.0000]

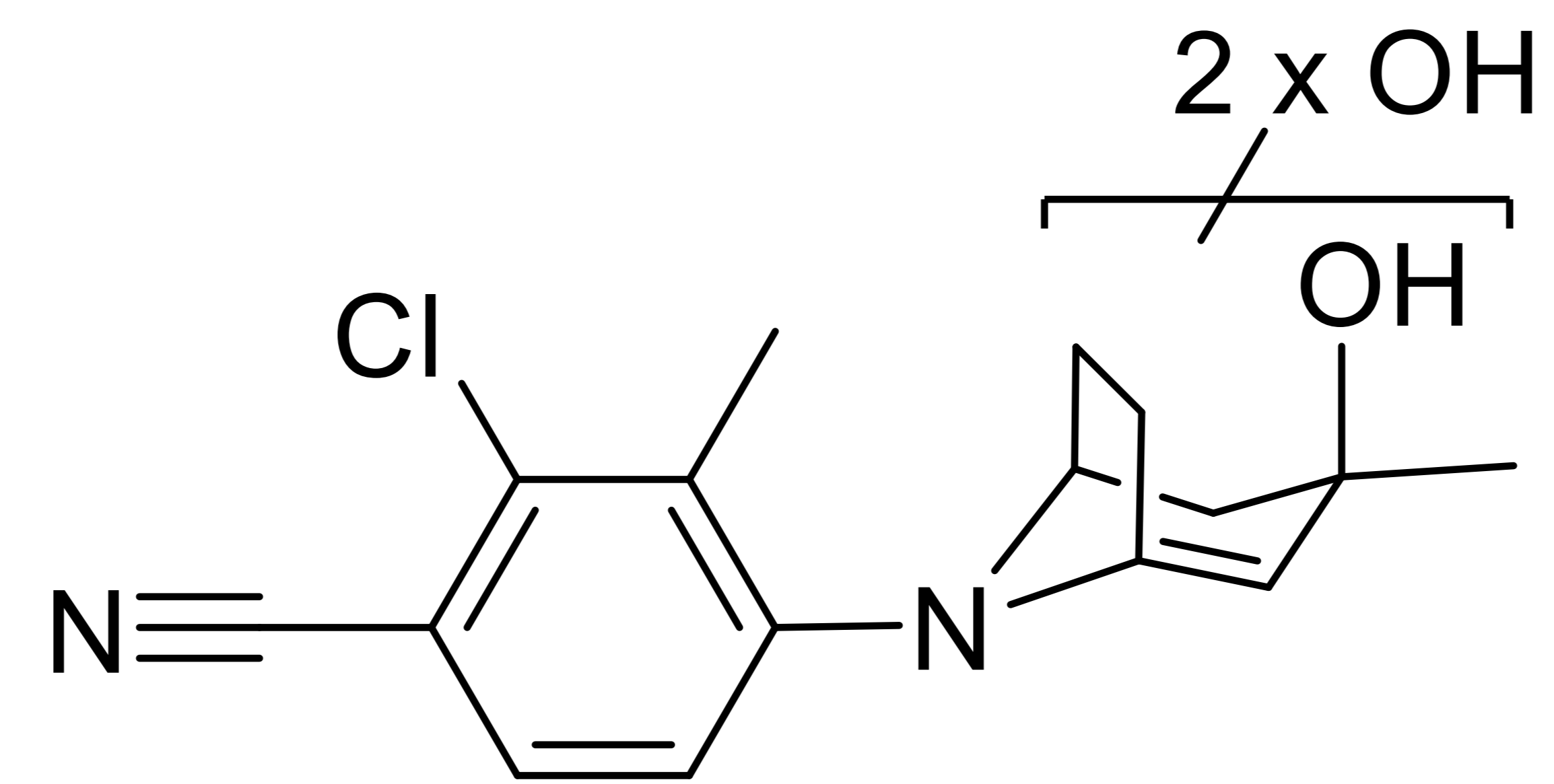


M5a

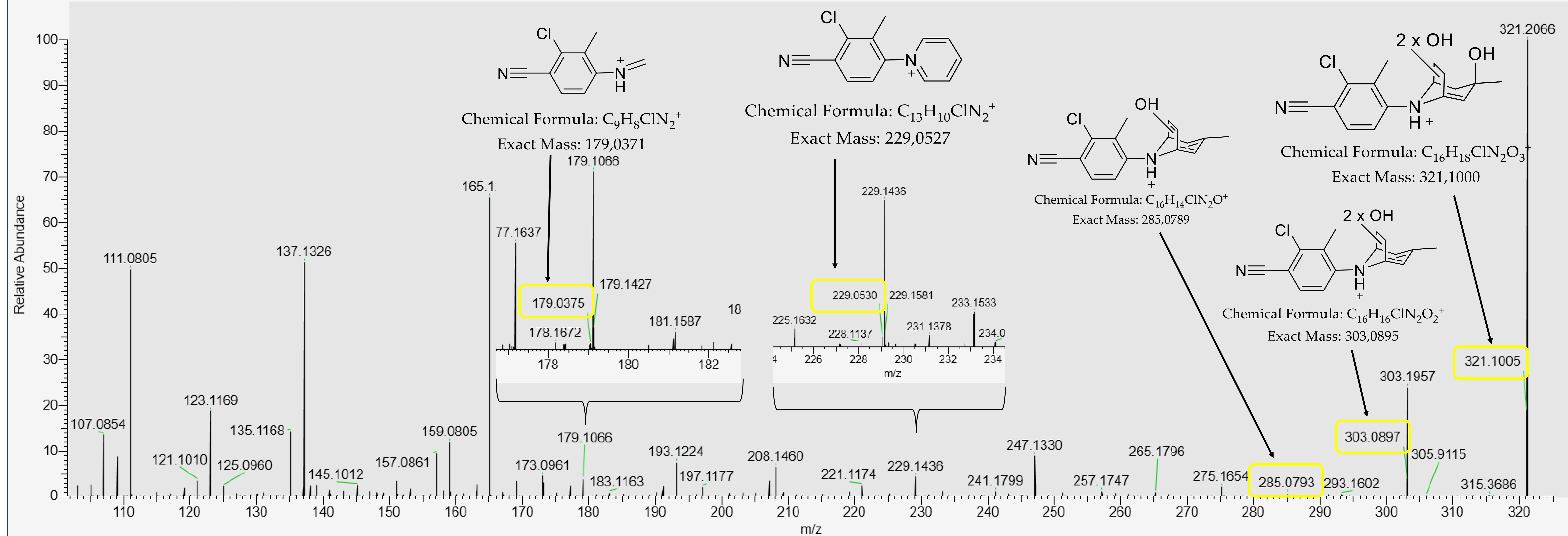
Dihydroxylation + loss of 2H

Retention time: 9.17 min

m/z: 321.1005



MNB_20200730_011 #1922-1933 RT: 9.15-9.2 AV: 12 SB: 78 9.00-9.15 , 9.20-9.39 NL: 1.61E+005
T: FTMS + p ESI Full ms2 321.1000@hcd38.33 [50.0000-345.0000]

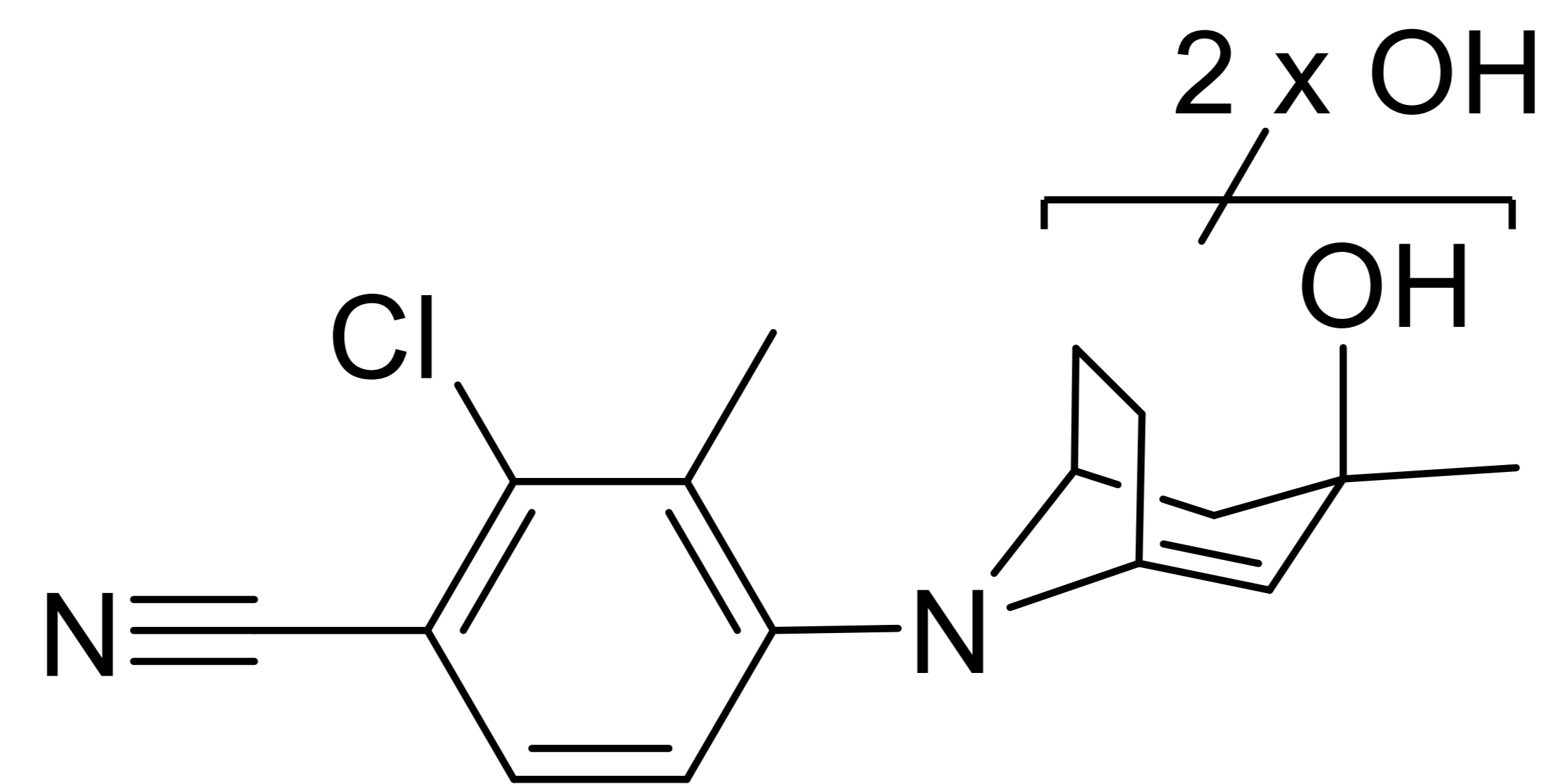


M5b

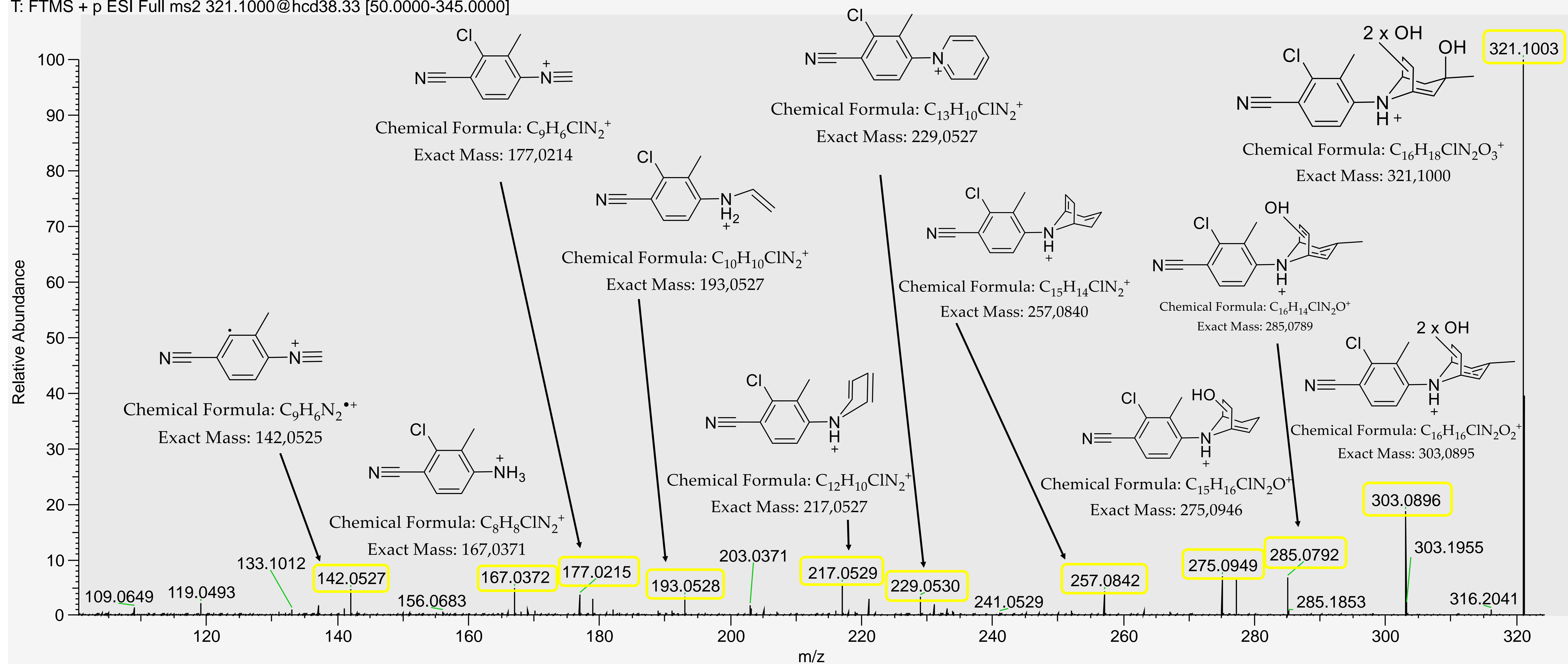
Dihydroxylation + loss of 2H

Retention time: 9.57 min

m/z: 321.1003



MNB_20200730_011 #2010-2030 RT: 9.54-9.63 AV: 21 SB: 34 9.47-9.54 , 9.63-9.70 NL: 1.27E+006
T: FTMS + p ESI Full ms2 321.1000@hcd38.33 [50.0000-345.0000]

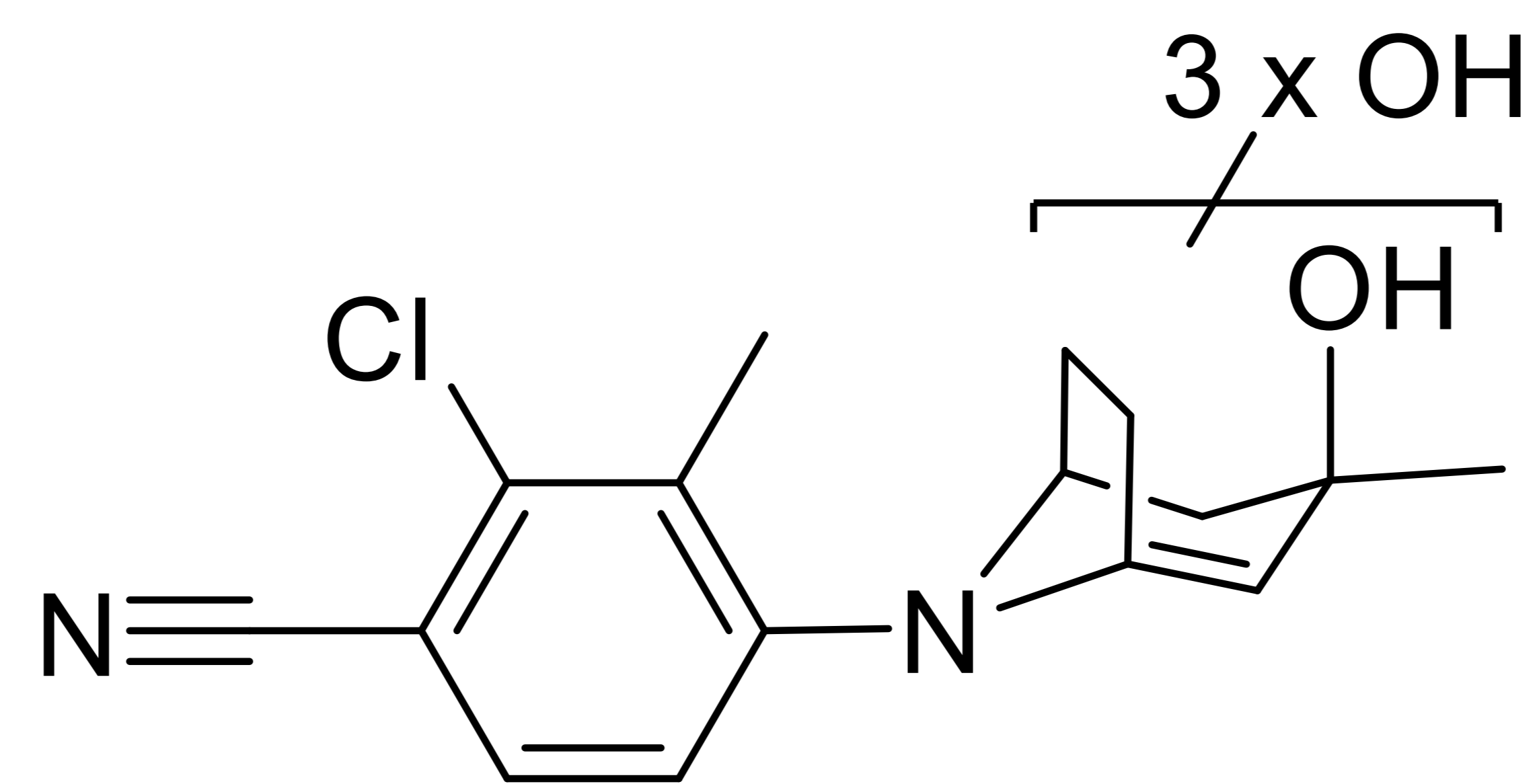


M6a

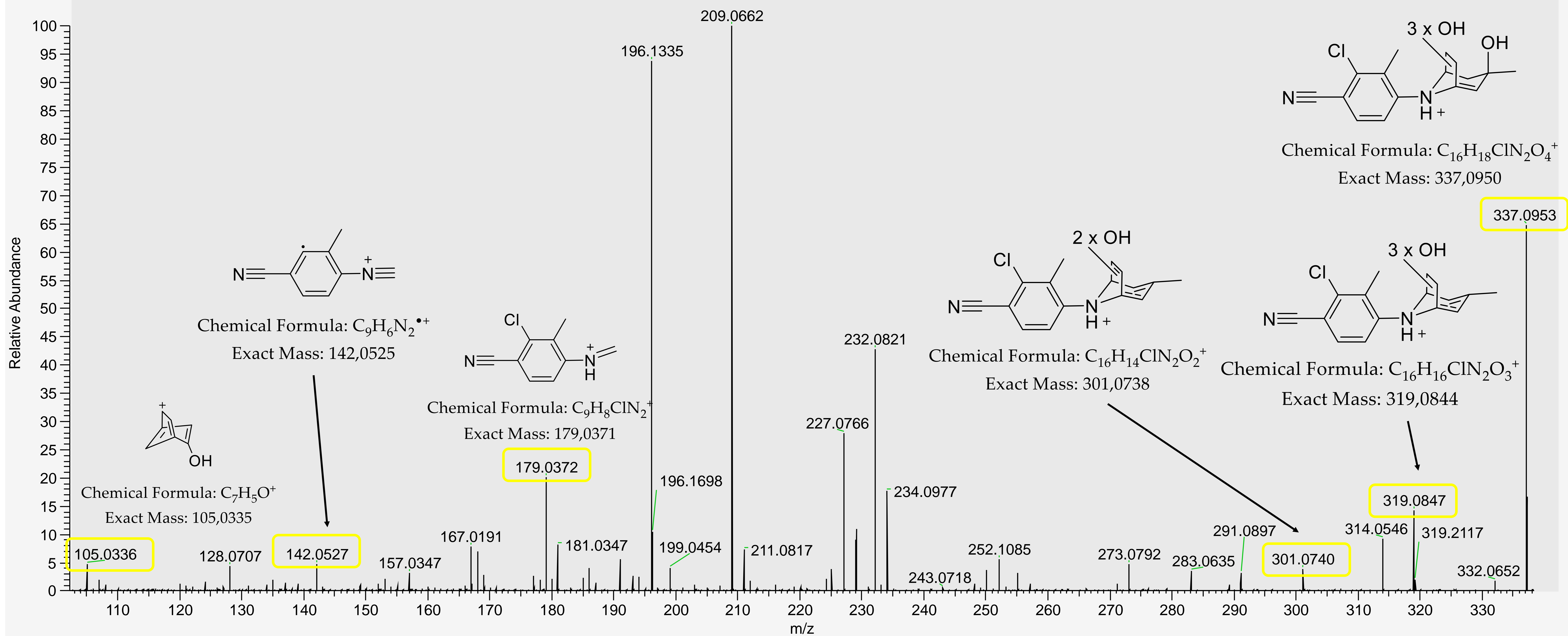
Trihydroxylation + loss of 2H

Retention time: 7.21 min

m/z: 337.0953



MNB_20200730_010 #1486-1503 RT: 7.18-7.26 AV: 18 SB: 20 7.14-7.18 , 7.26-7.30 NL: 2.21E+005
T: FTMS + p ESI Full ms2 337.0950@hcd38.33 [50.0000-360.0000]

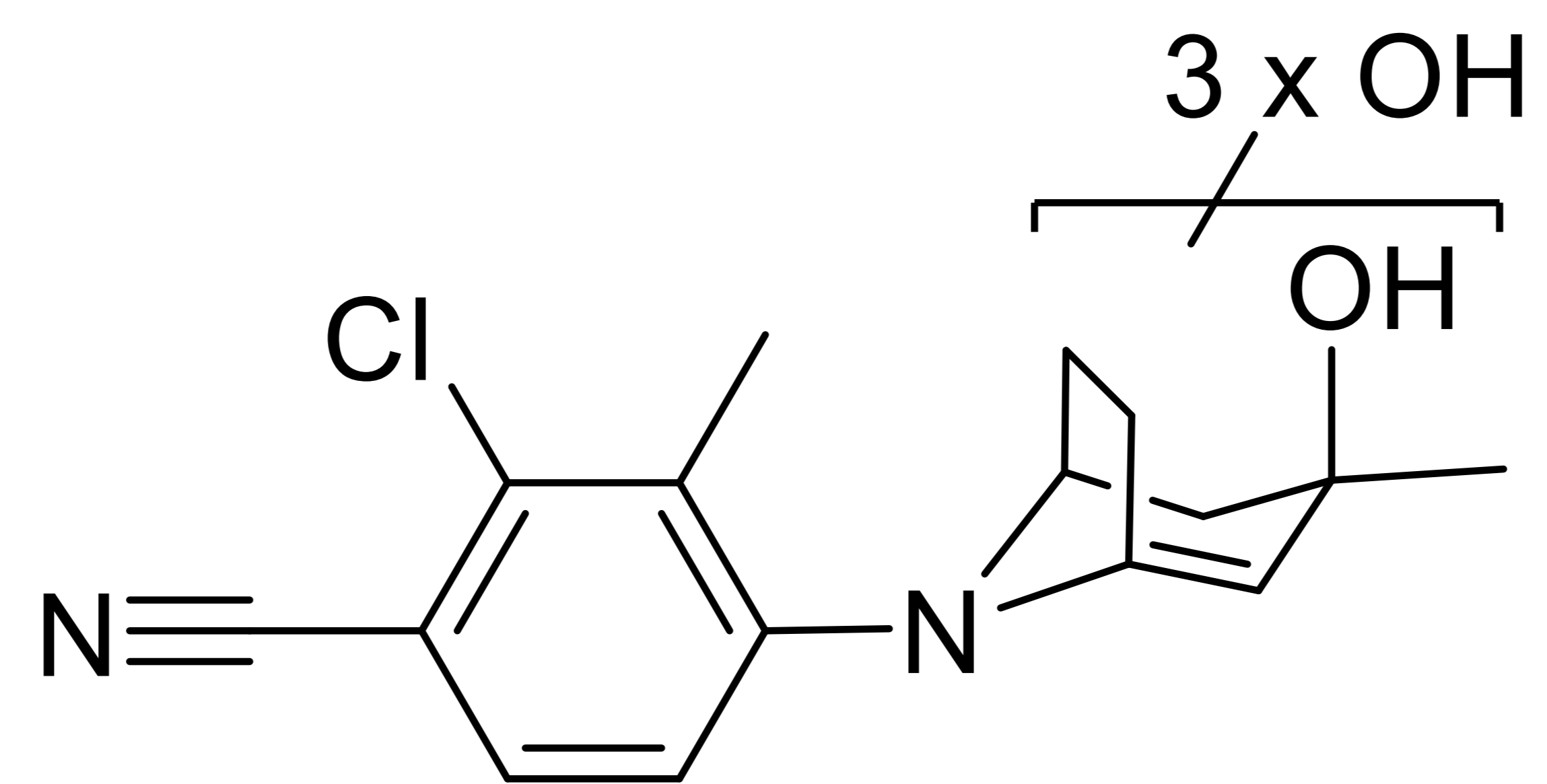


M6b

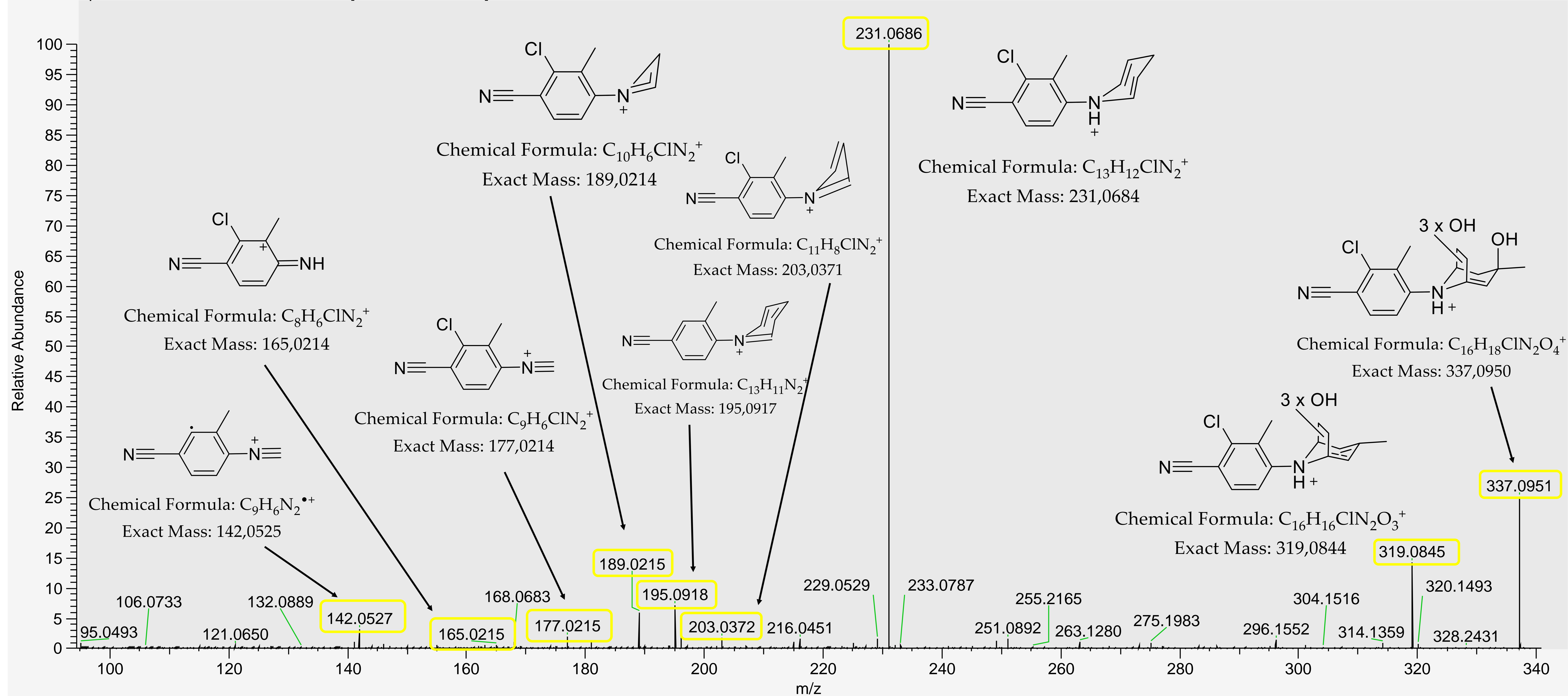
Trihydroxylation + loss of 2H

Retention time: 7.47 min

m/z: 337.0951



MNB_20200730_010 #1548-1562 RT: 7.46-7.52 AV: 15 SB: 17 7.42-7.45 , 7.52-7.56 NL: 8.85E+005
T: FTMS + p ESI Full ms2 337.0950@hcd38.33 [50.0000-360.0000]

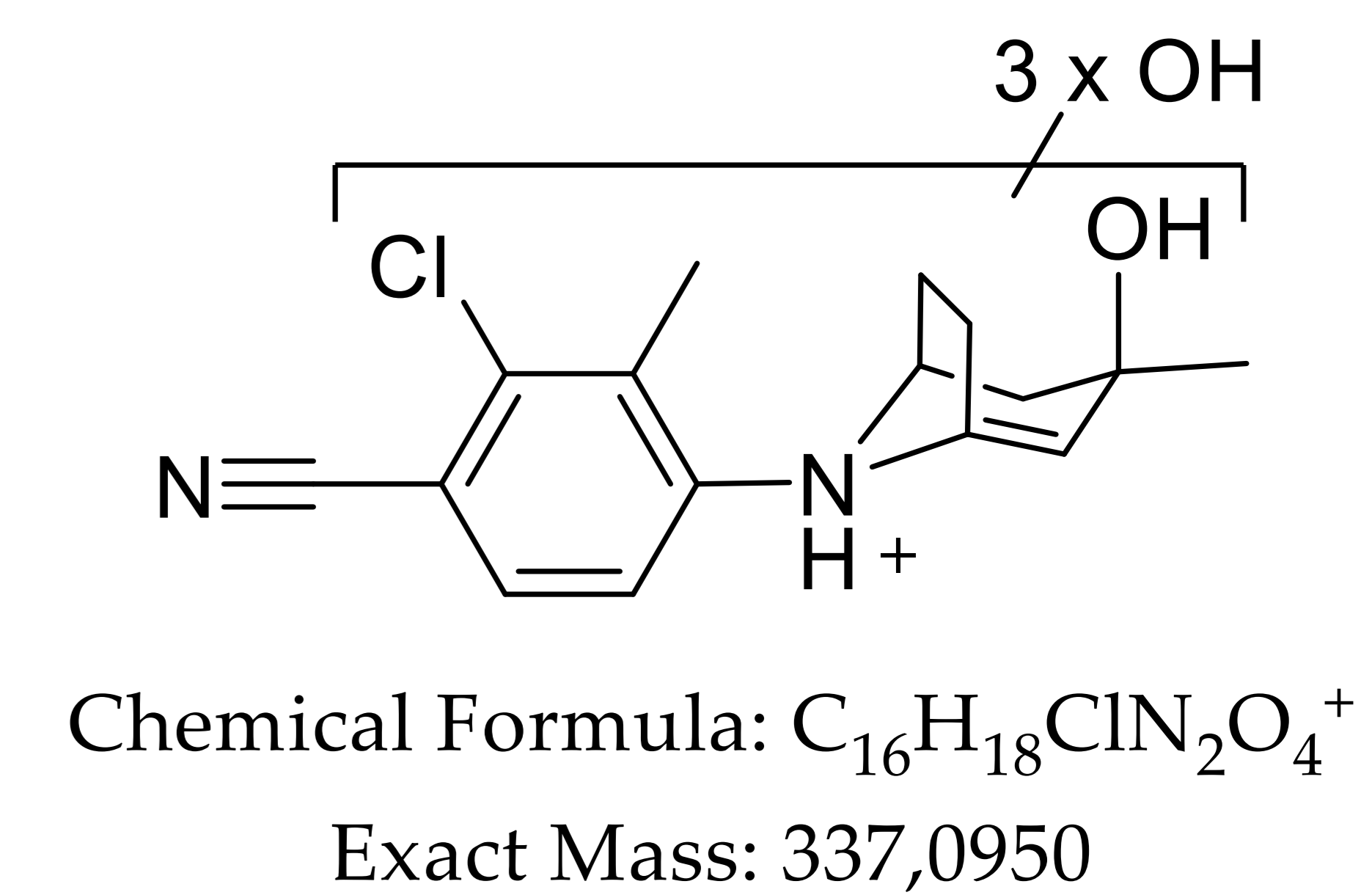
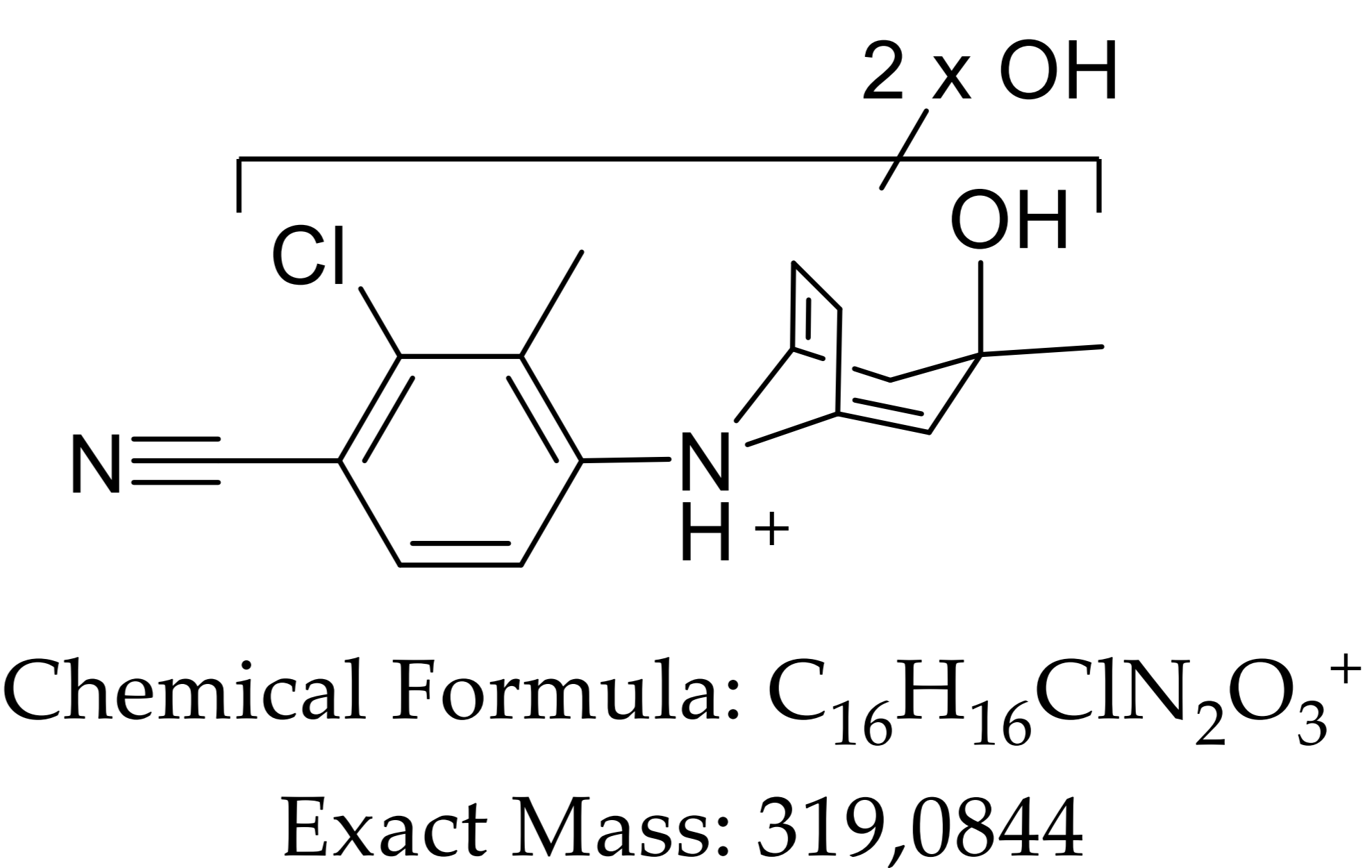
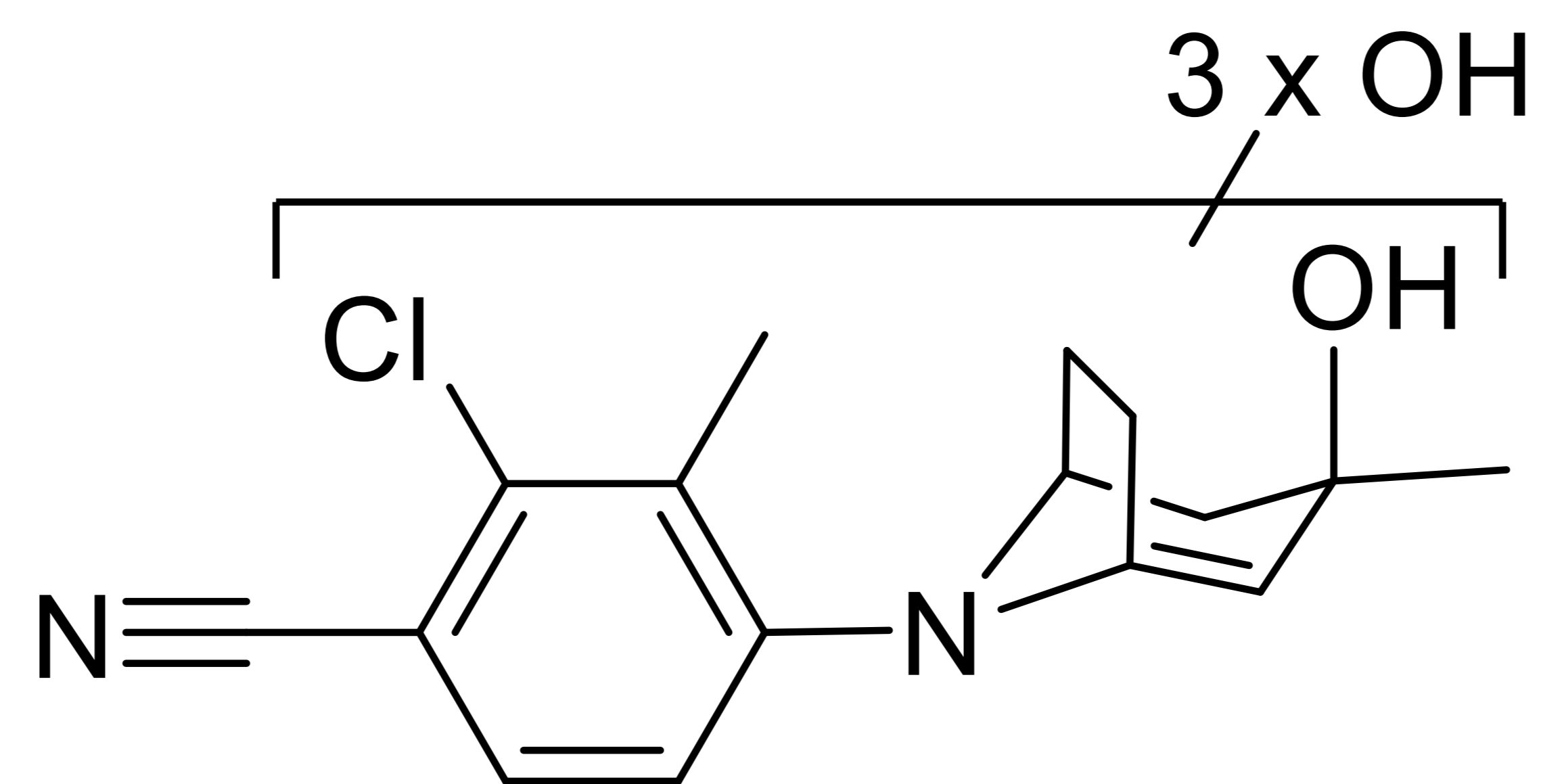


M6c

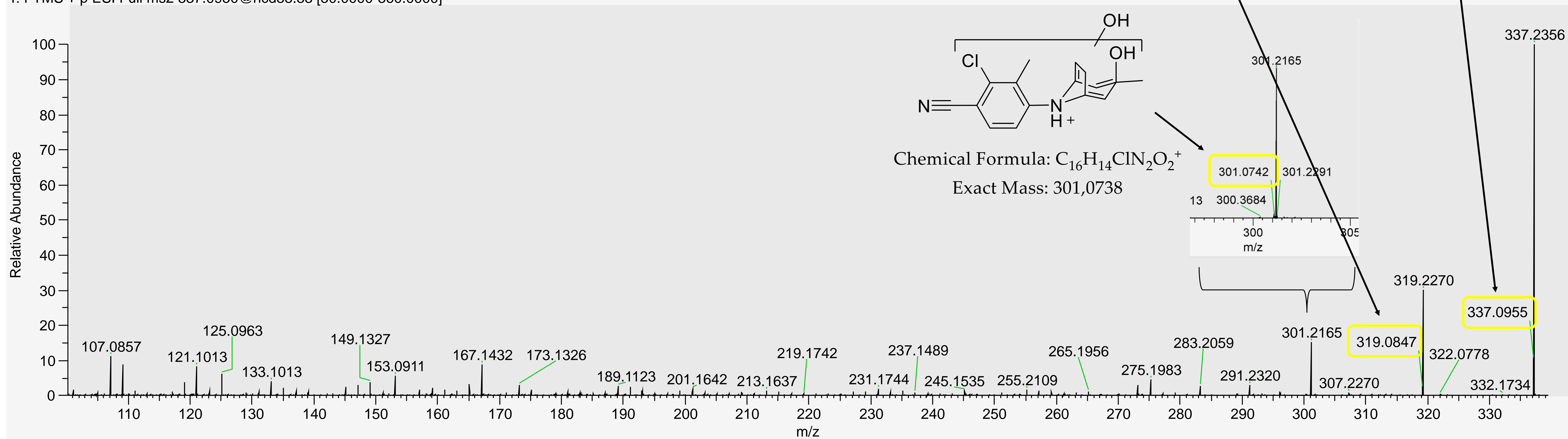
Trihydroxylation + loss of 2H

Retention time: 9.04 min

m/z: 337.0955



MNB_20200730_010 #1896-1910 RT: 9.03-9.09 AV: 15 SB: 14 9.00-9.03 , 9.09-9.12 NL: 1.48E+005
T: FTMS + p ESI Full ms2 337.0950@hcd38.33 [50.0000-360.0000]

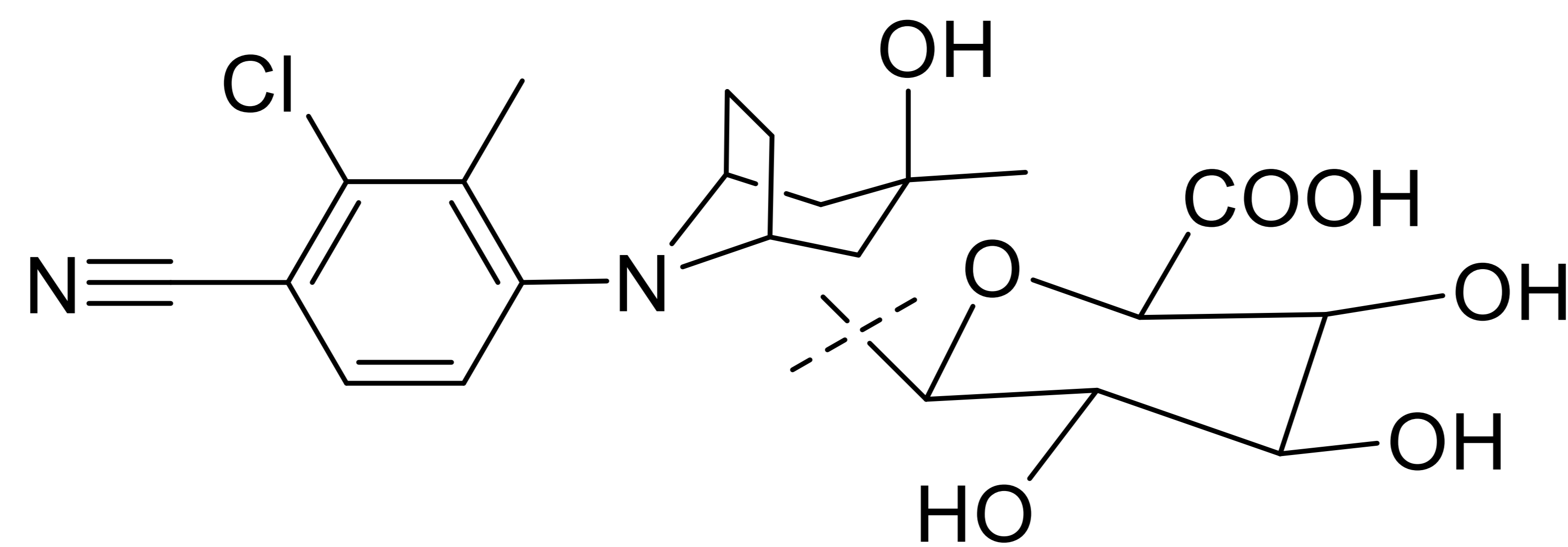


M7

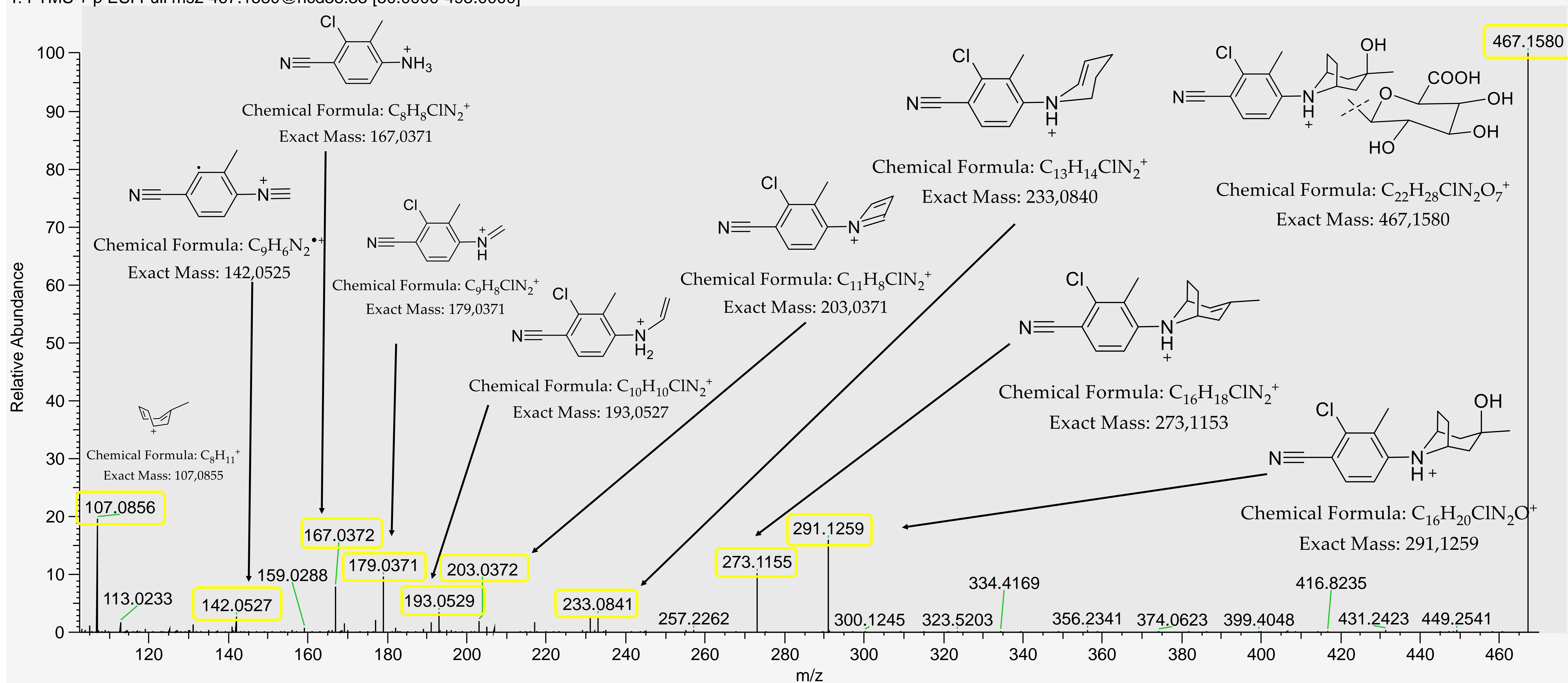
Glucuronidation

Retention time: 8.95 min

m/z: 467.1580



MNB_20200429_05 #1704-1713 RT: 8.93-8.99 AV: 4 SB: 7 8.88-8.93 , 8.99-9.03 NL: 2.73E+005
T: FTMS + p ESI Full ms2 467.1580@hcd38.33 [50.0000-495.0000]

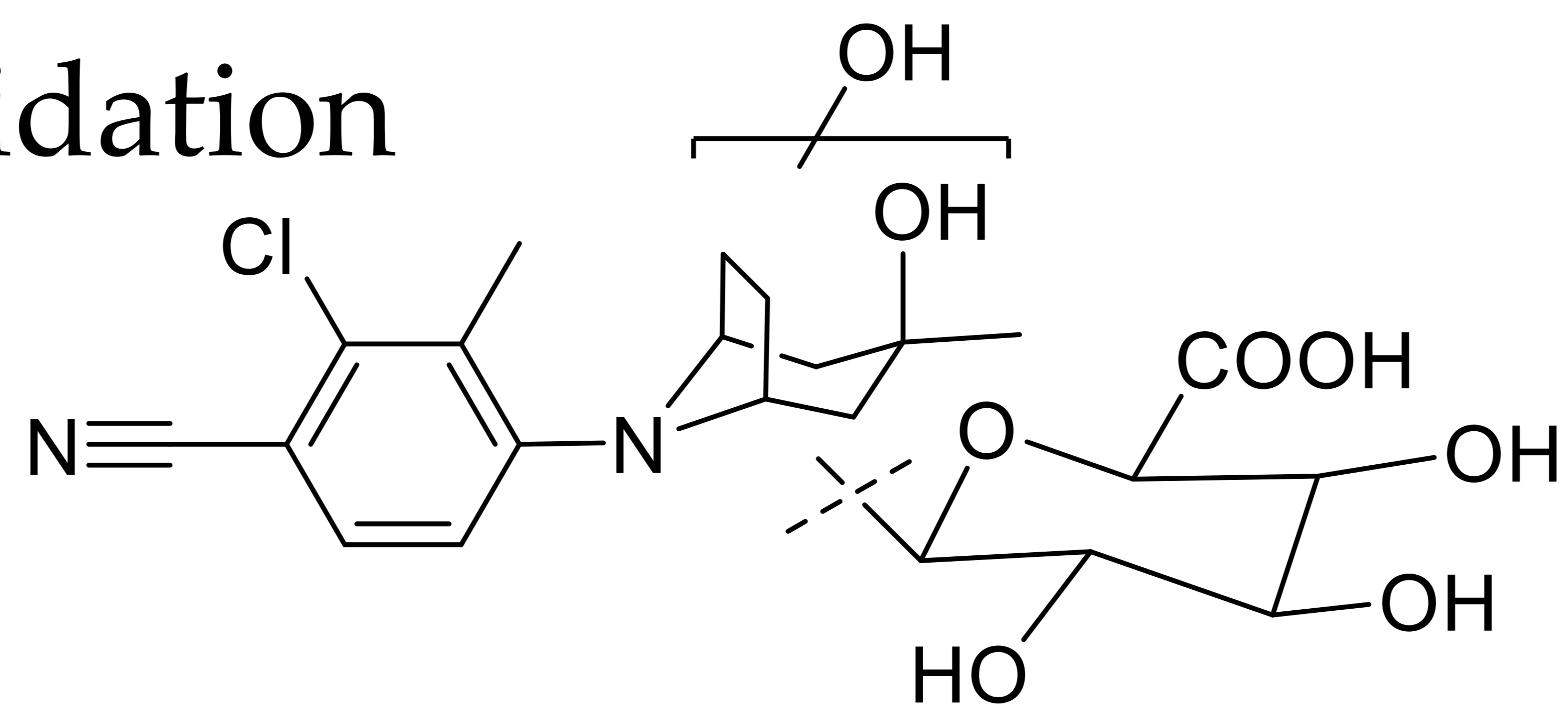


M8a

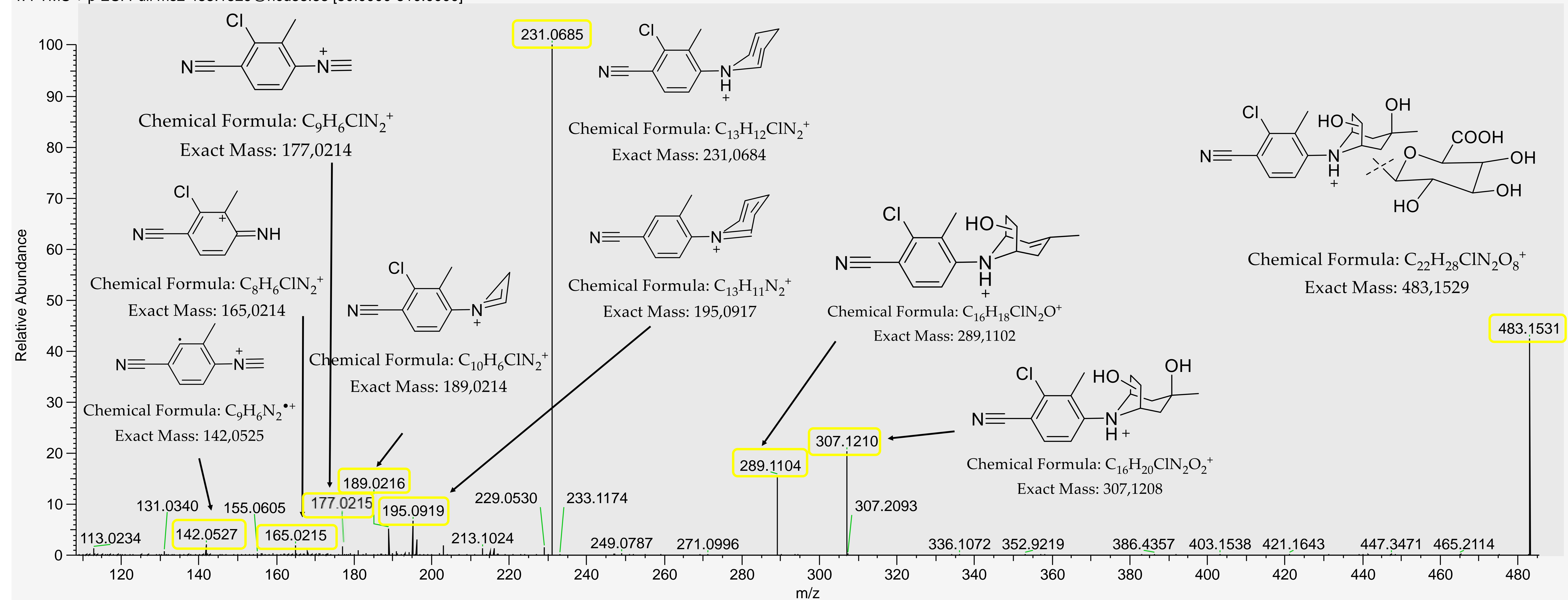
Monohydroxylation + glucuronidation

Retention time: 7.13 min

m/z: 483.1531



MNB_20200429_05 #1408-1421 RT: 7.1-7.18 AV: 3 SB: 5 7.03-7.10 , 7.18-7.22 NL: 7.53E+006
T: FTMS + p ESI Full ms2 483.1529@hcd38.33 [50.0000-510.0000]

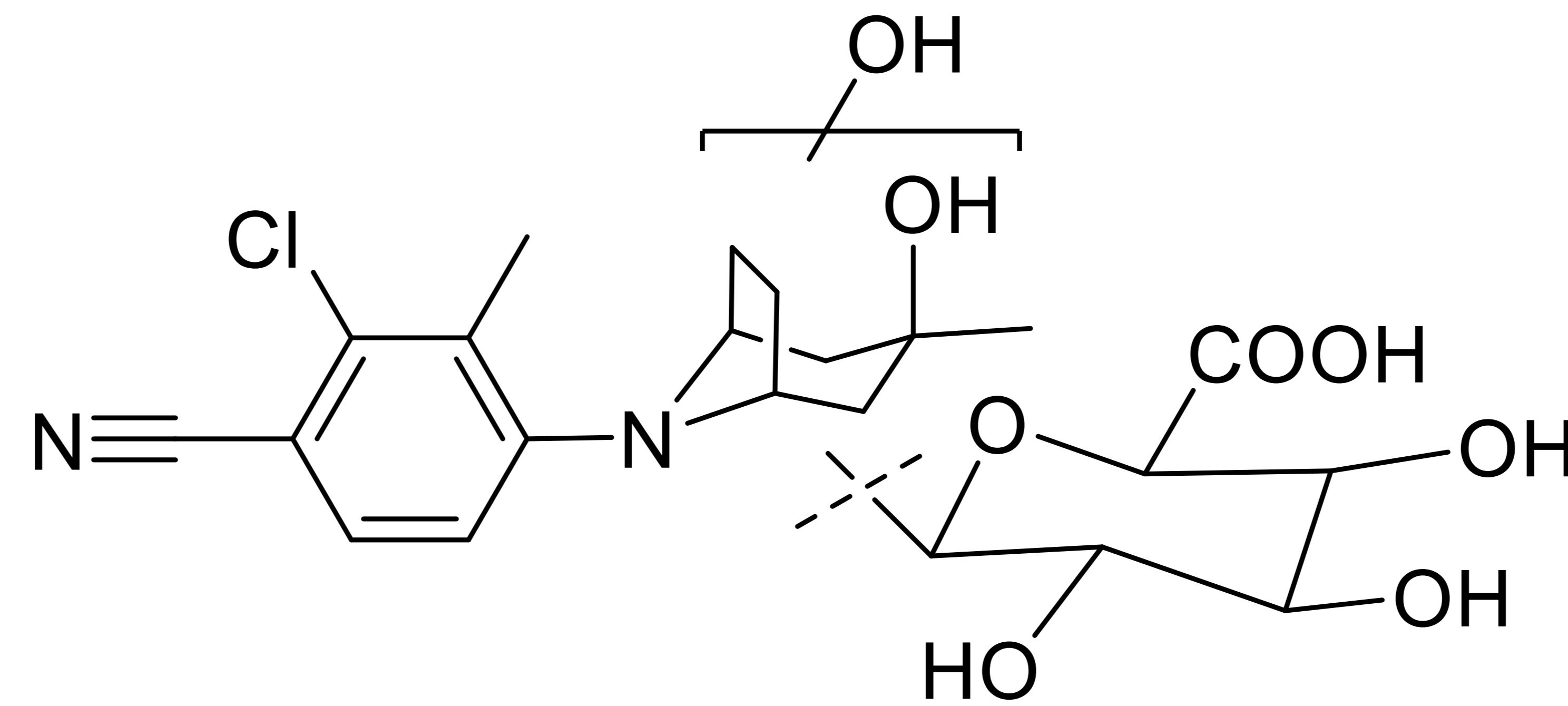


M8b

Monohydroxylation + glucuronidation

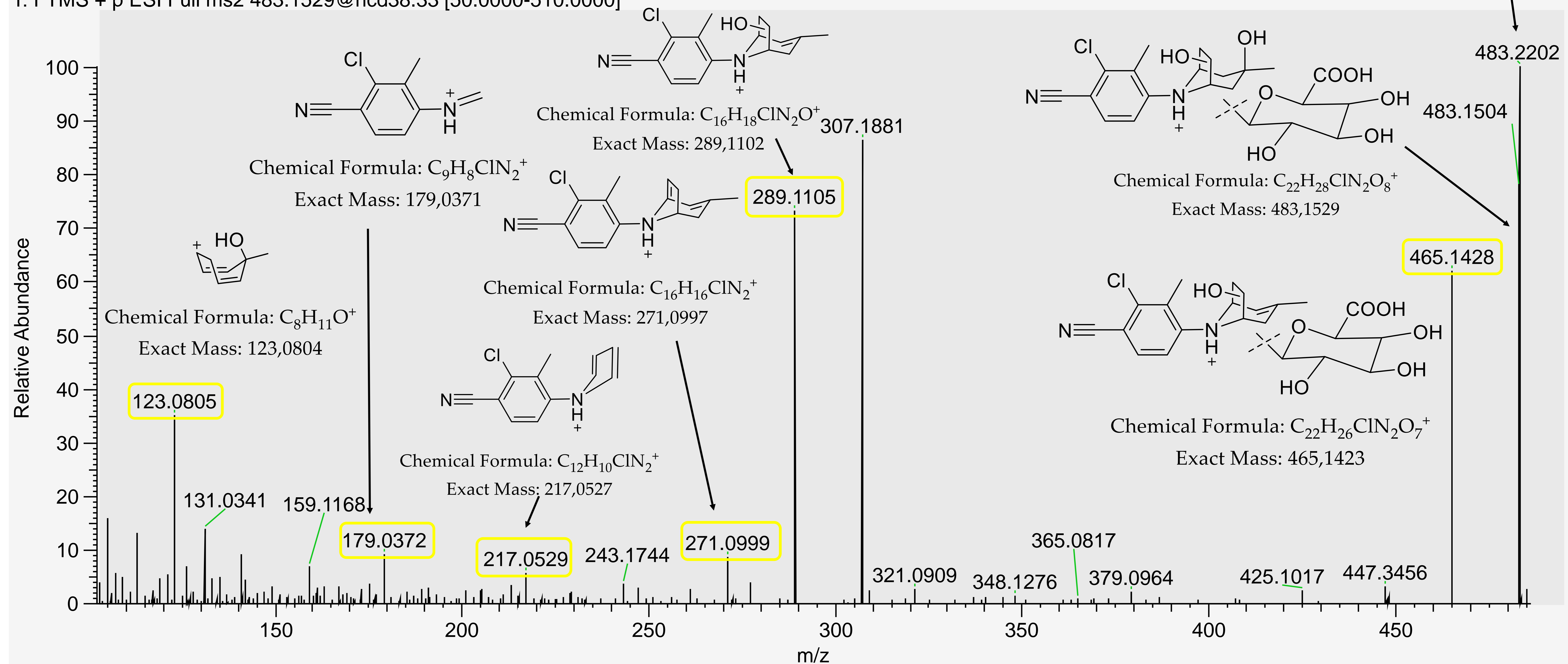
Retention time: 7.61 min

m/z: 483.1527



Very low intensity of remaining m/z 483.1527 and co-eluting with other ions and thereby not visible in spectrum

MNB_20200429_05 #1487-1494 RT: 7.6-7.64 AV: 2 SB: 5 7.53-7.60 , 7.64-7.69 NL: 8.46E+004
T: FTMS + p ESI Full ms2 483.1529@hcd38.33 [50.0000-510.0000]

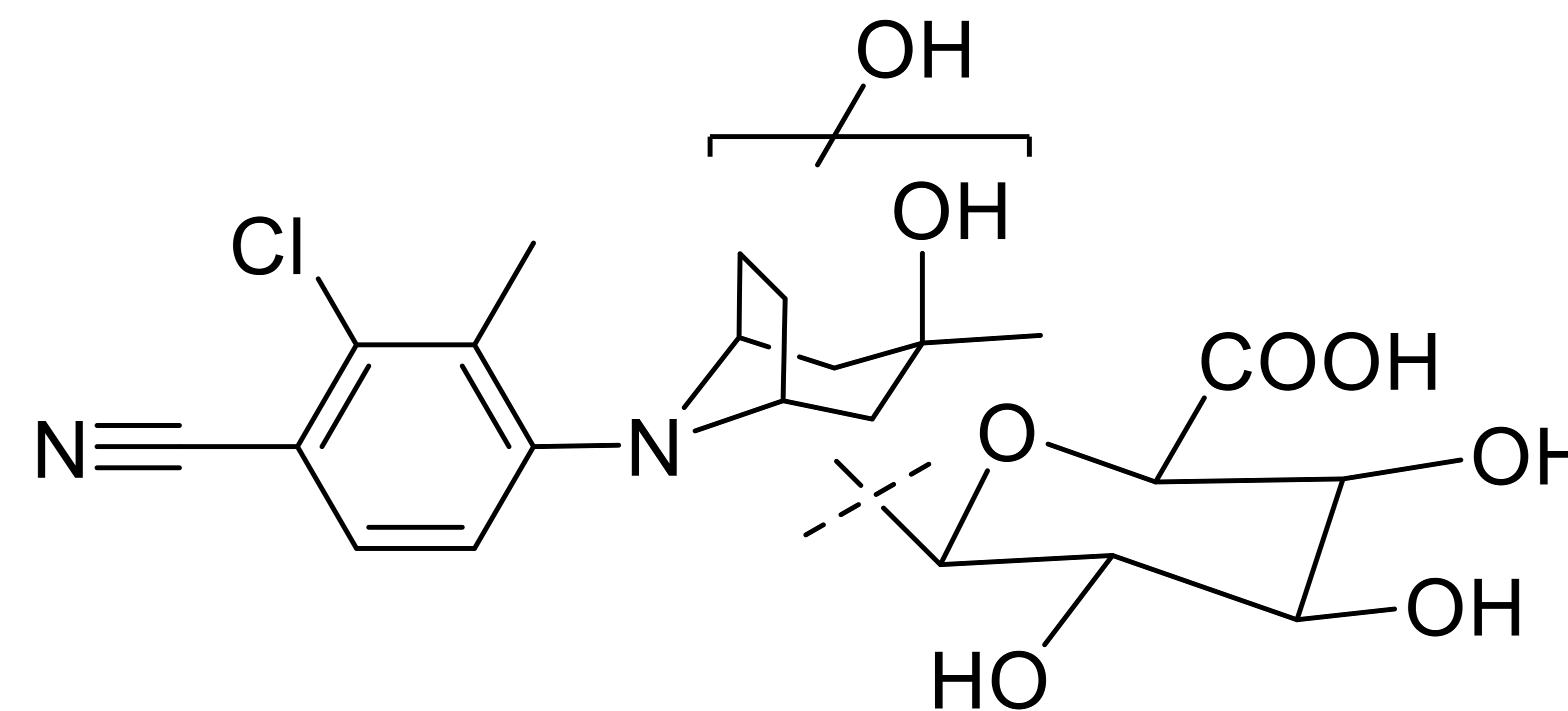


M8c

Monohydroxylation + glucuronidation

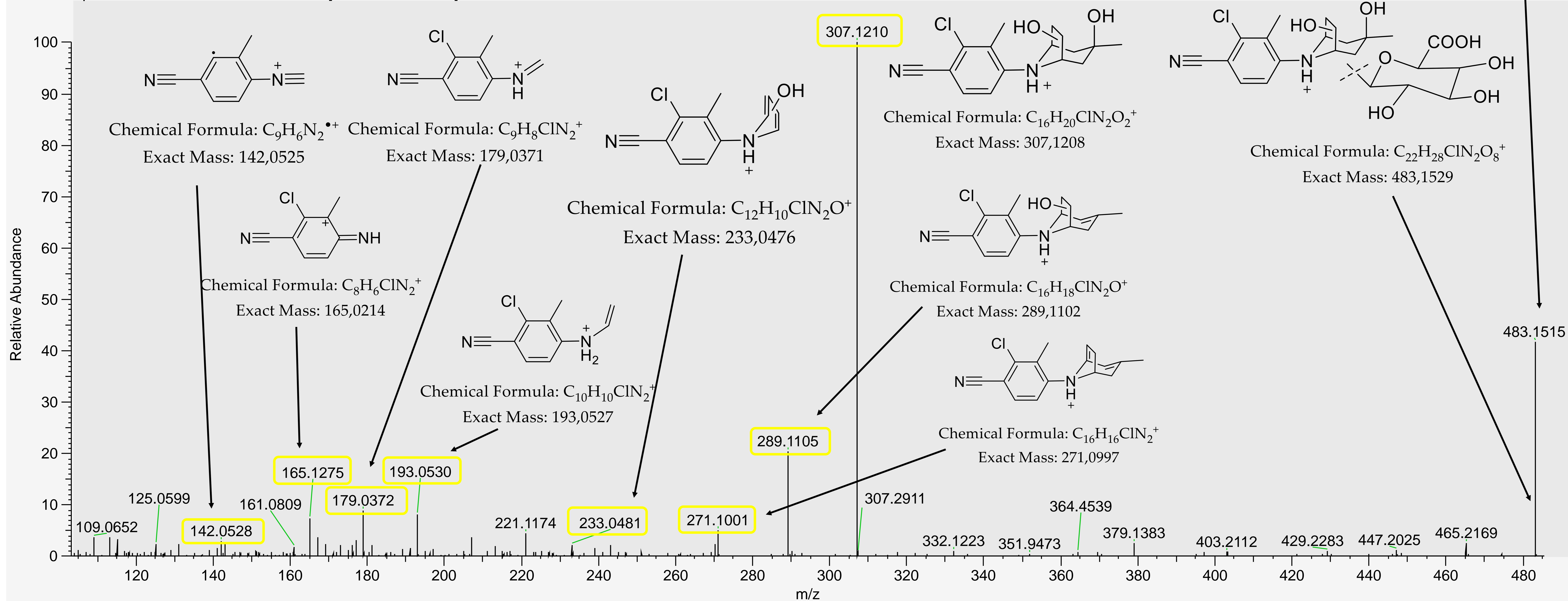
Retention time: 7.85 min

m/z: 483.1524



Very low intensity of remaining m/z 483.1524 and co-eluting with other ions and thereby not visible in spectrum

MNB_20200429_05 #1523-1533 RT: 7.82-7.88 AV: 3 SB: 5 7.76-7.82 , 7.88-7.93 NL: 7.59E+004
T: FTMS + p ESI Full ms2 483.1529@hcd38.33 [50.0000-510.0000]

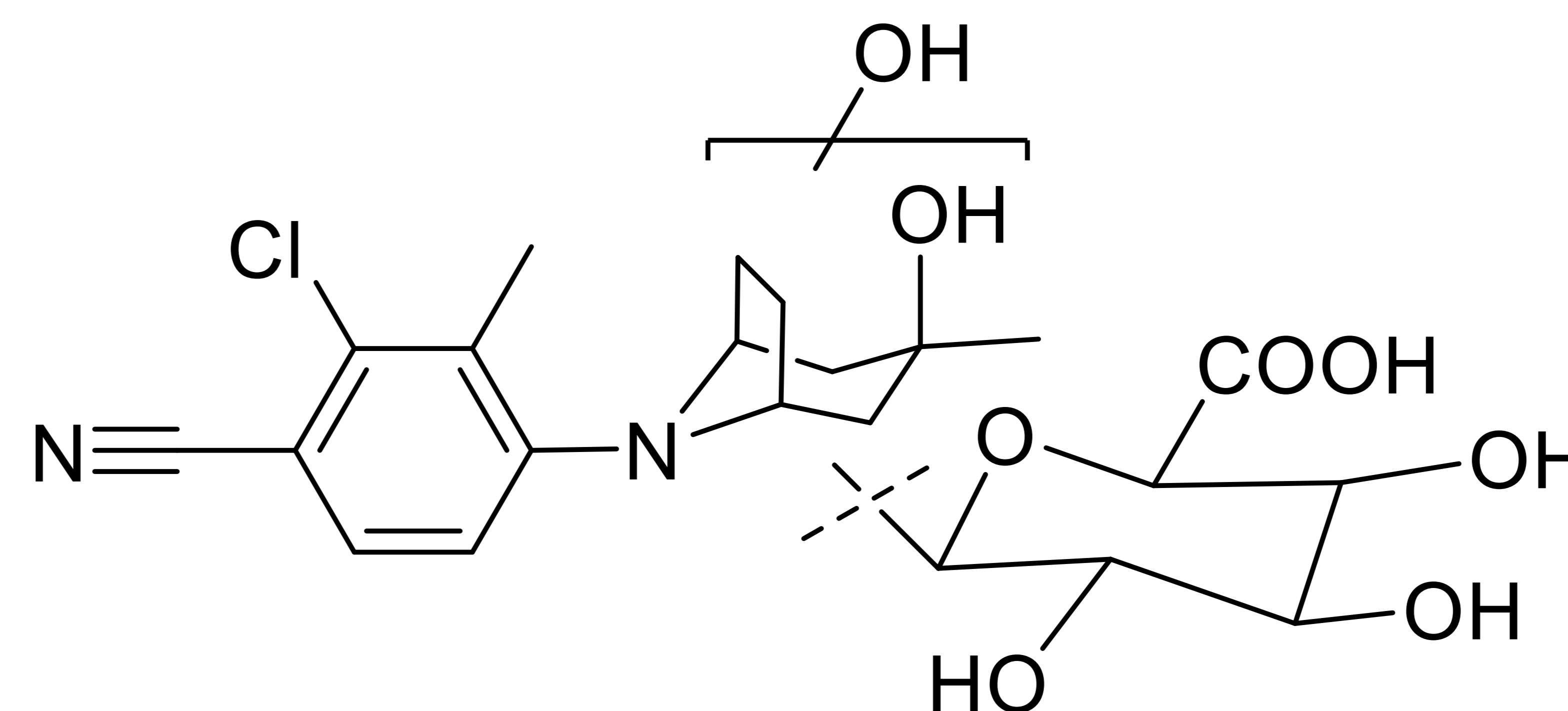


M8d

Monohydroxylation + glucuronidation

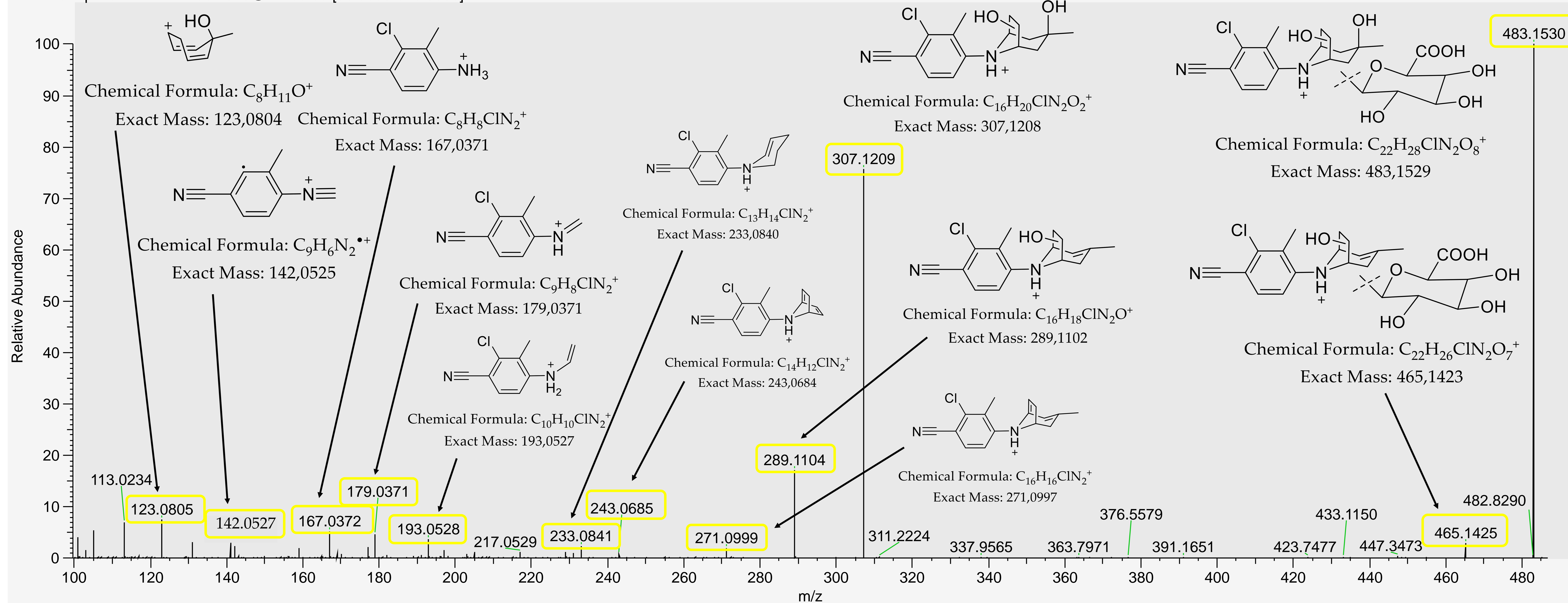
Retention time: 8.00 min

m/z: 483.1530



MNB_20200429_05 #1546-1557 RT: 7.96-8.03 AV: 4 SB: 7 7.90-7.96 , 8.03-8.10 NL: 1.02E+007

T: FTMS + p ESI Full ms2 483.1529@hcd38.33 [50.0000-510.0000]

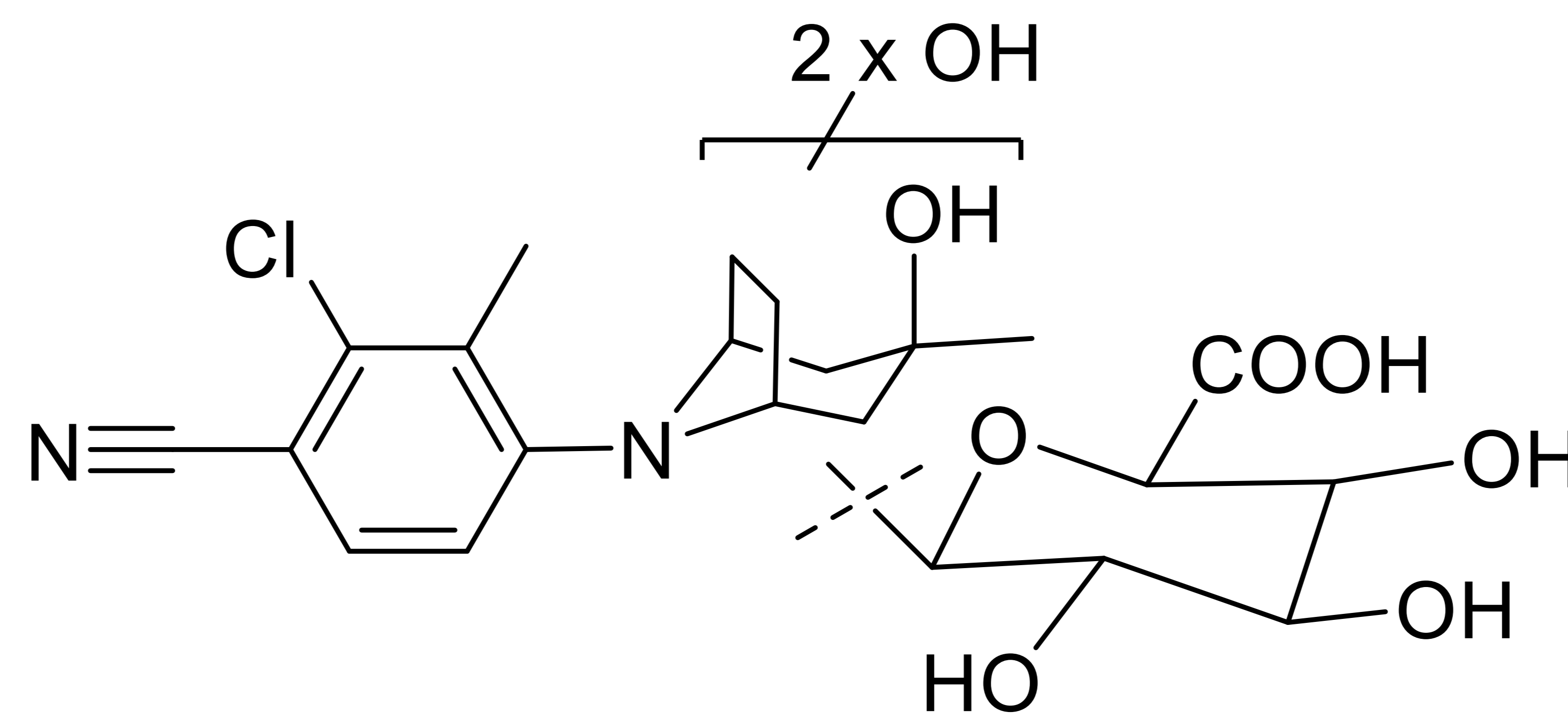


M9a

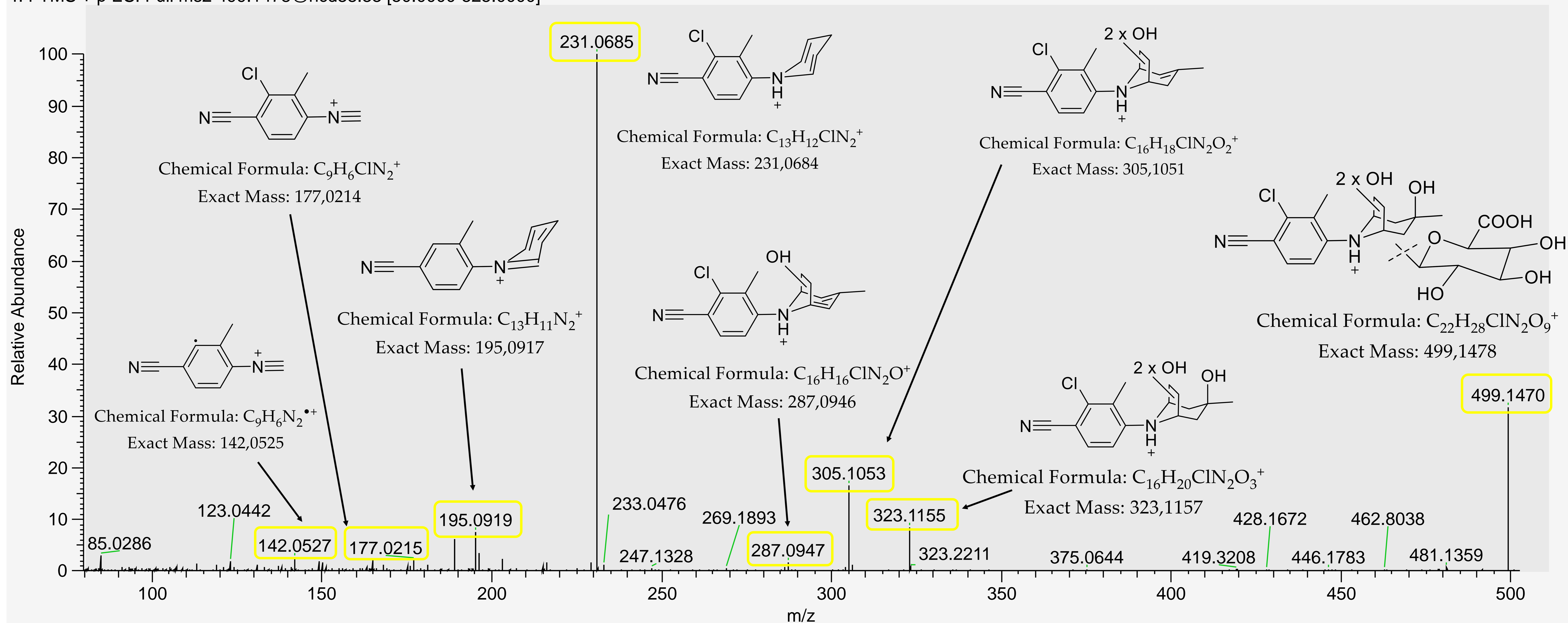
Dihydroxylation + glucuronidation

Retention time: 6.21 min

m/z: 499.1470



MNB_20200429_05 #1265-1275 RT: 6.19-6.25 AV: 4 SB: 7 6.13-6.19 , 6.25-6.31 NL: 4.23E+005
T: FTMS + p ESI Full ms2 499.1478@hcd38.33 [50.0000-525.0000]

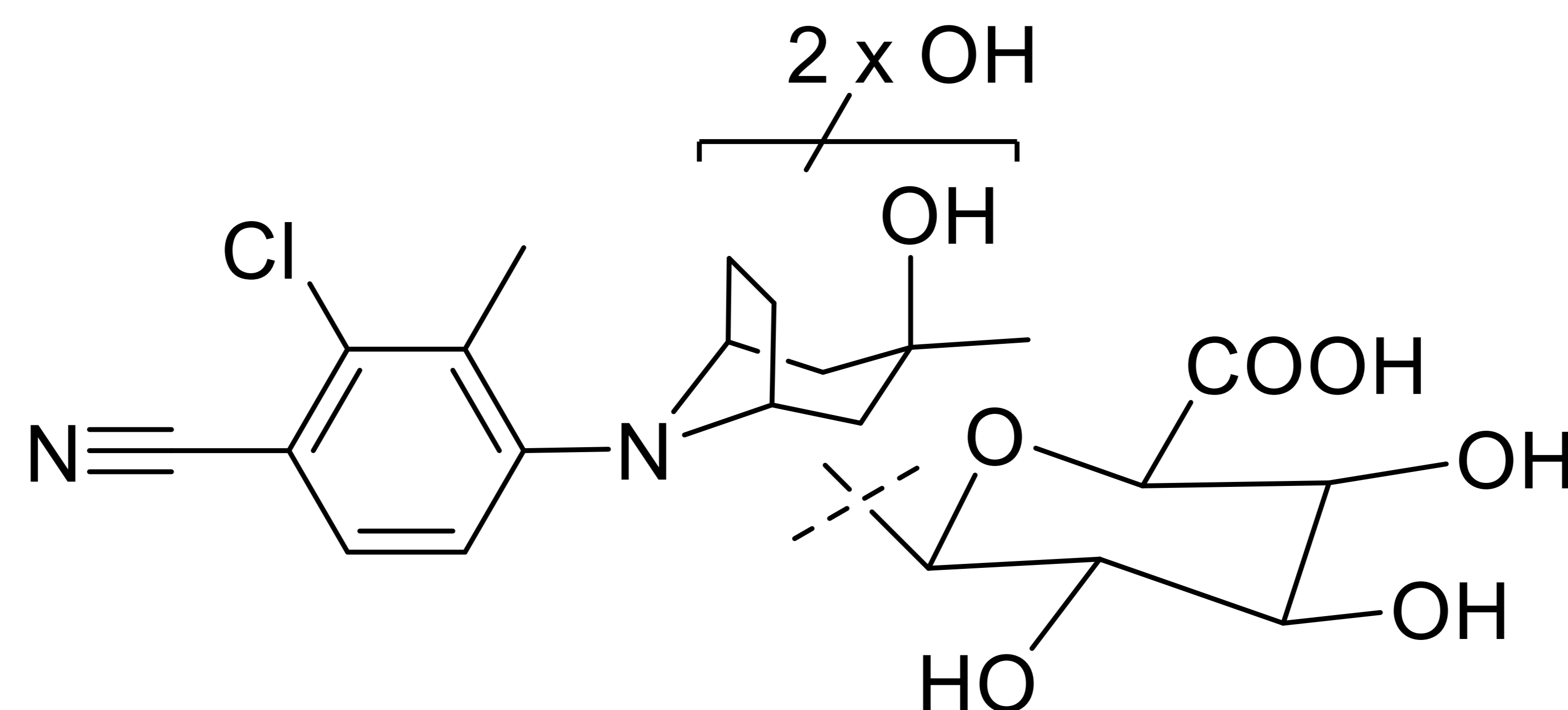


M9b

Dihydroxylation + glucuronidation

Retention time: 6.52 min

m/z: 499.1478



MNB_20200429_05 #1316 RT: 6.51 AV: 1 NL: 1.21E+006
T: FTMS + p ESI Full ms2 499.1478@hcd38.33 [50.0000-525.0000]

