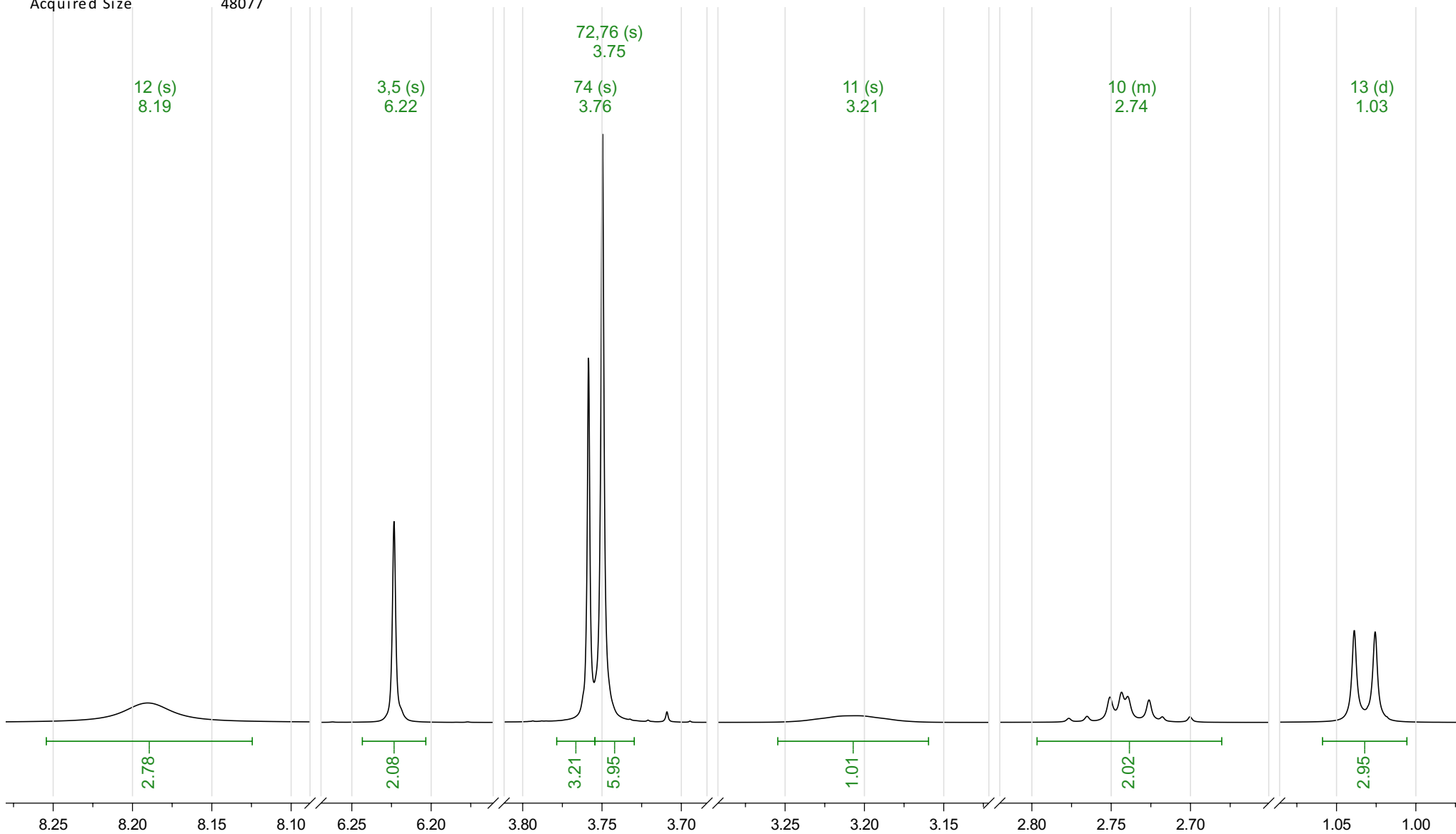
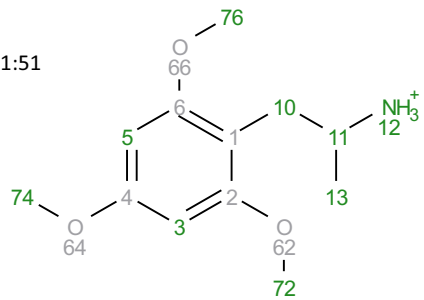
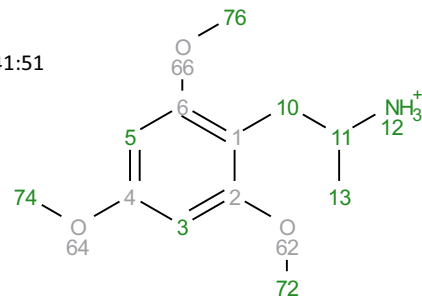


Analyte SB2: TMA-6 H+
 Acquisition Date 2016-12-02T16:41:51
 Solvent dmso
 Temperature 25
 Number of Scans 16
 Relaxation Delay 1
 Spectrometer Frequency 499.66
 Spectral Width 8012.8
 Nucleus 1H
 Acquired Size 48077

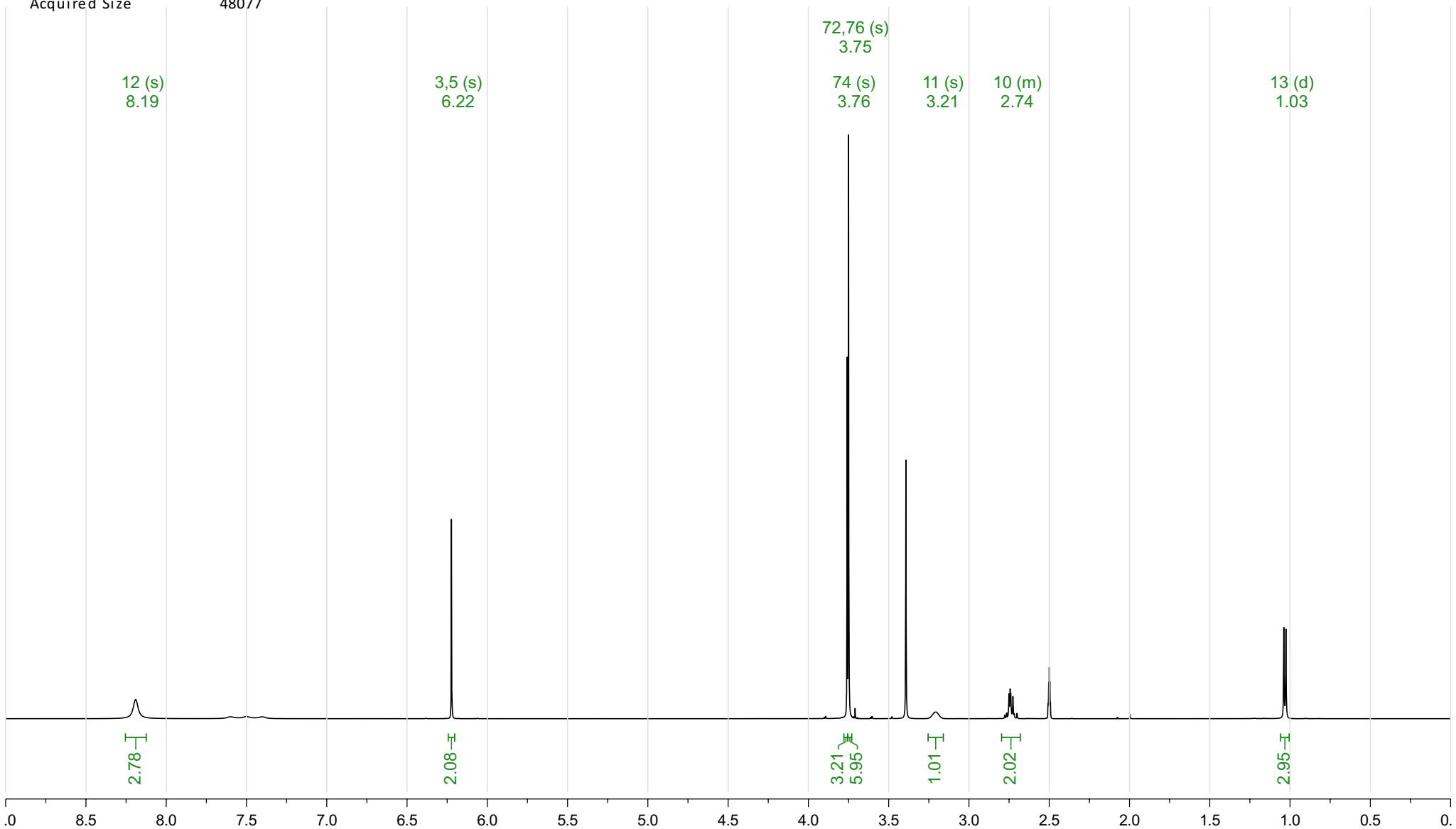
¹H NMR (500 MHz, DMSO-*d*₆) δ 8.19 (s, 3H), 6.22 (s, 2H), 3.76 (s, 3H), 3.75 (s, 6H), 3.21 (s, 1H), 2.80 – 2.68 (m, 2H), 1.03 (d, *J* = 6.6 Hz, 3H).



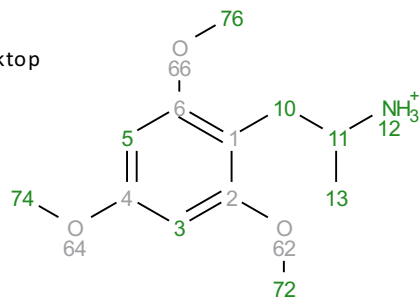
Analyte SB2: TMA-6 H+
 Acquisition Date 2016-12-02T16:41:51
 Solvent dmso
 Temperature 25
 Number of Scans 16
 Relaxation Delay 1
 Spectrometer Frequency 499.66
 Spectral Width 8012.8
 Nucleus 1H
 Acquired Size 48077



¹H NMR (500 MHz, DMSO-*d*₆) δ 8.19 (s, 3H), 6.22 (s, 2H), 3.76 (s, 3H), 3.75 (s, 6H), 3.21 (s, 1H), 2.80 – 2.68 (m, 2H), 1.03 (d, *J* = 6.6 Hz, 3H).



Prediction Prediction for TMA-6 H+
Origin Modgraph NMRPredict Desktop
Solvent DMSO-d6
Algorithm Best
GMMX Cycles 25
Version 18153
Frequency 500.13
Nucleus 1H



¹H NMR (500 MHz, DMSO-*d*₆) δ 8.25 (d, *J* = 5.2 Hz, 3H), 6.20 (s, 2H), 3.82 (s, 3H), 3.77 (s, 6H), 3.54 – 3.44 (m, 1H), 2.84 (dd, *J* = 16.5, 6.3 Hz, 1H), 2.72 (dd, *J* = 16.5, 6.3 Hz, 1H), 1.11 (d, *J* = 5.5 Hz, 3H).

