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**Return of the lysergamides. Part IV: Analytical and pharmacological characterization of lysergic acid morpholide (LSM-775)**

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### Article

**Citation** (please note it is advisable to refer to the publisher's version if you intend to cite from this work)

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## **Return of the lysergamides. Part IV: Analytical and pharmacological characterization of lysergic acid morpholide (LSM-775)**

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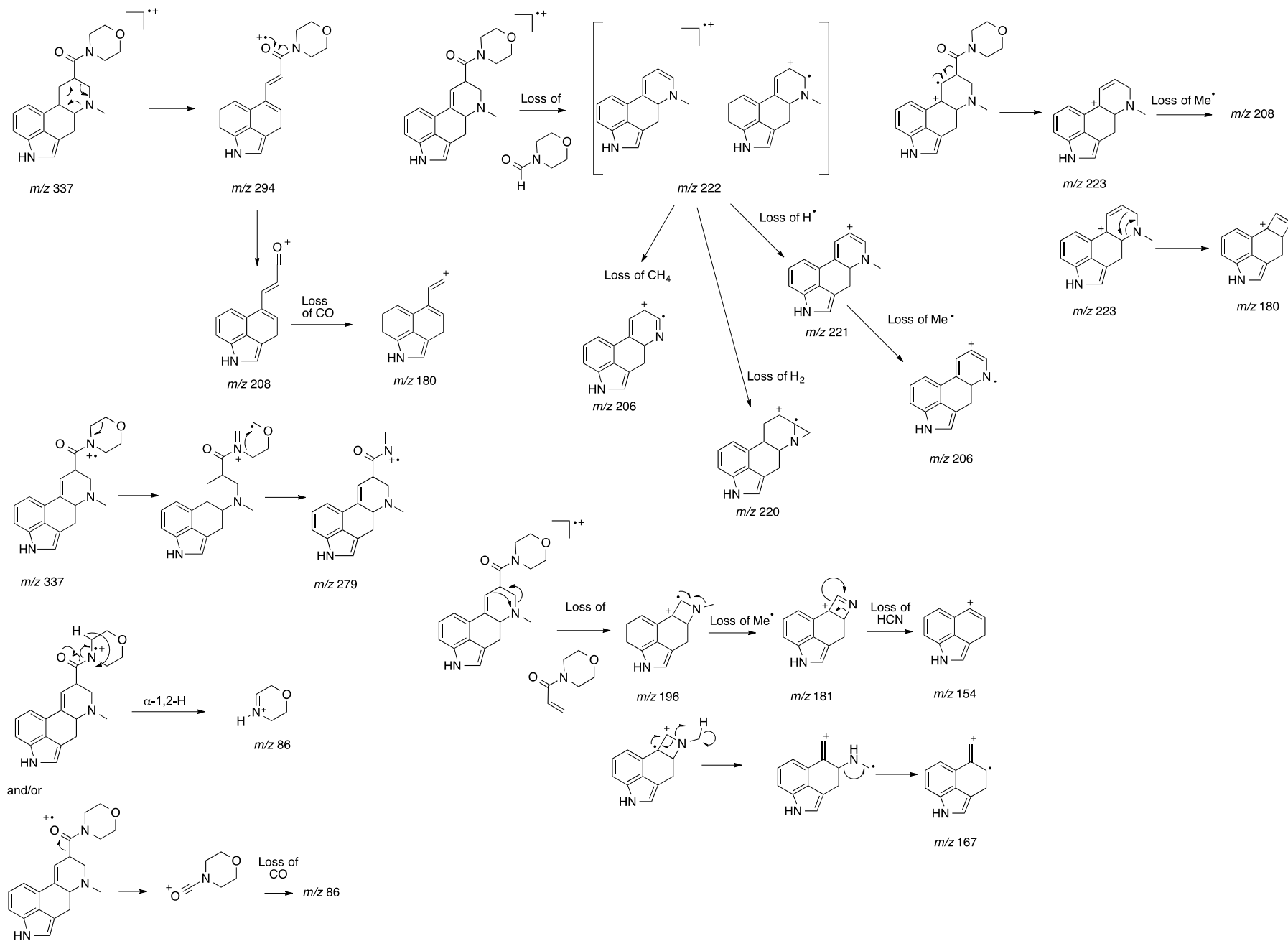
<sup>k</sup> *Department of Psychiatry, University of California San Diego, 9500 Gilman Drive, La Jolla, CA 92093, USA*

\* Correspondence to: Adam L. Halberstadt, Department of Psychiatry, University of California San Diego, 9500 Gilman Dr., La Jolla, CA 92093-0804 USA. E-Mail: ahalberstadt@ucsd.edu

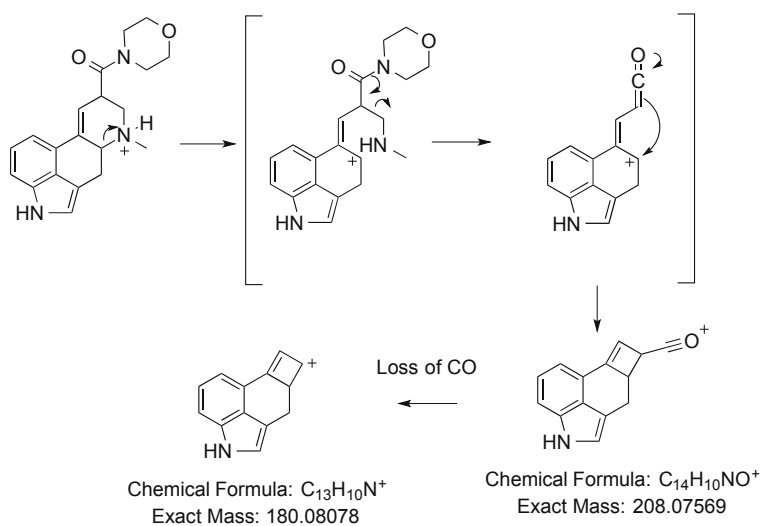
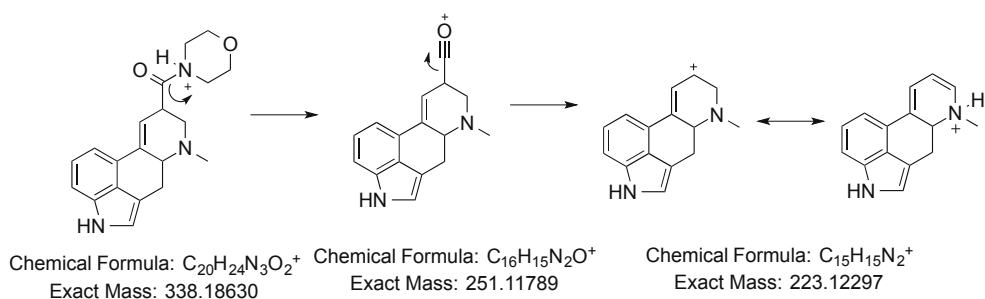
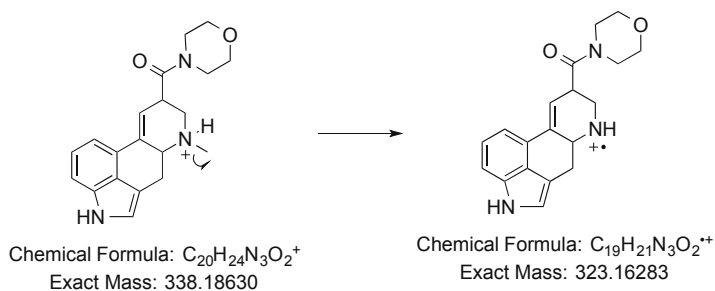
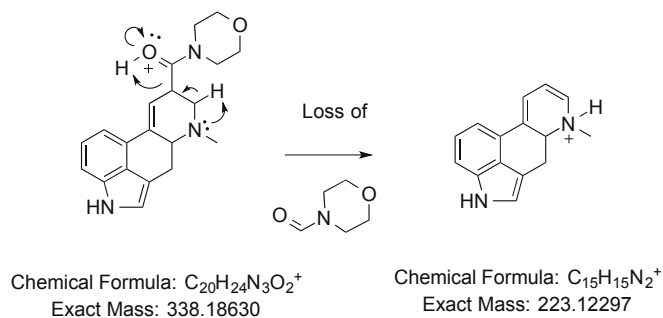
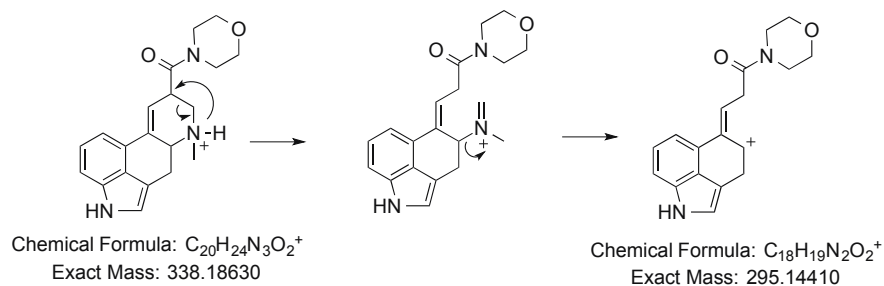


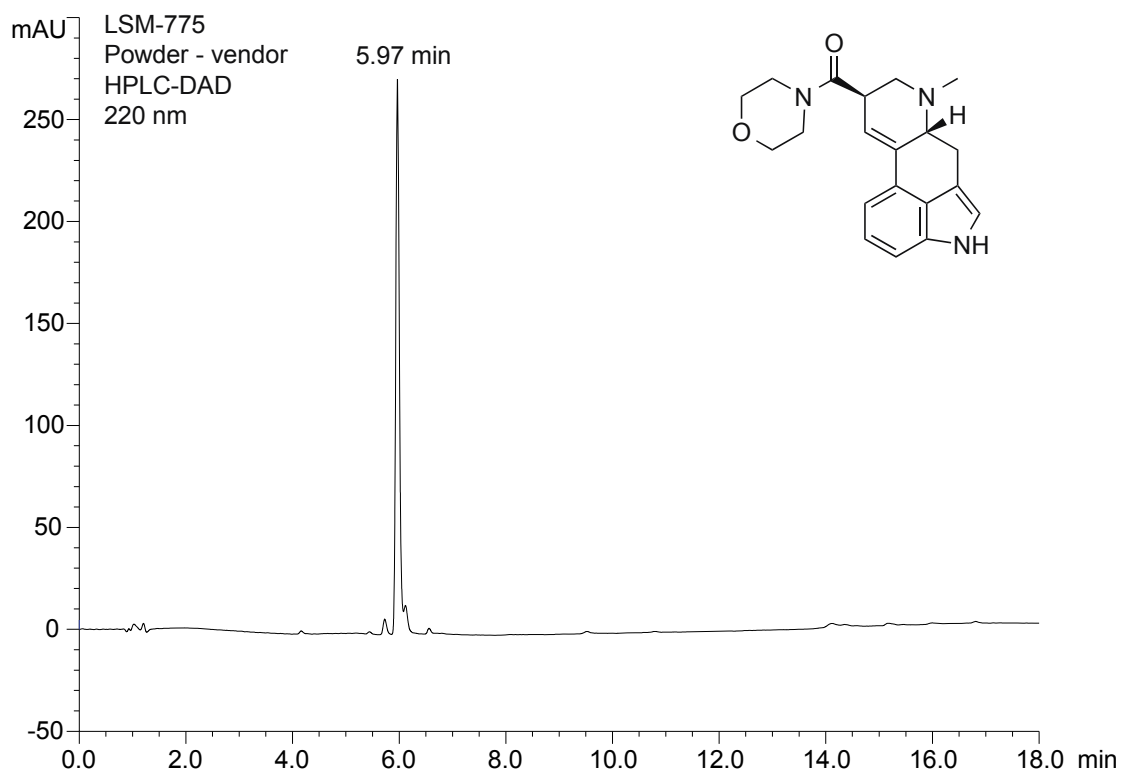
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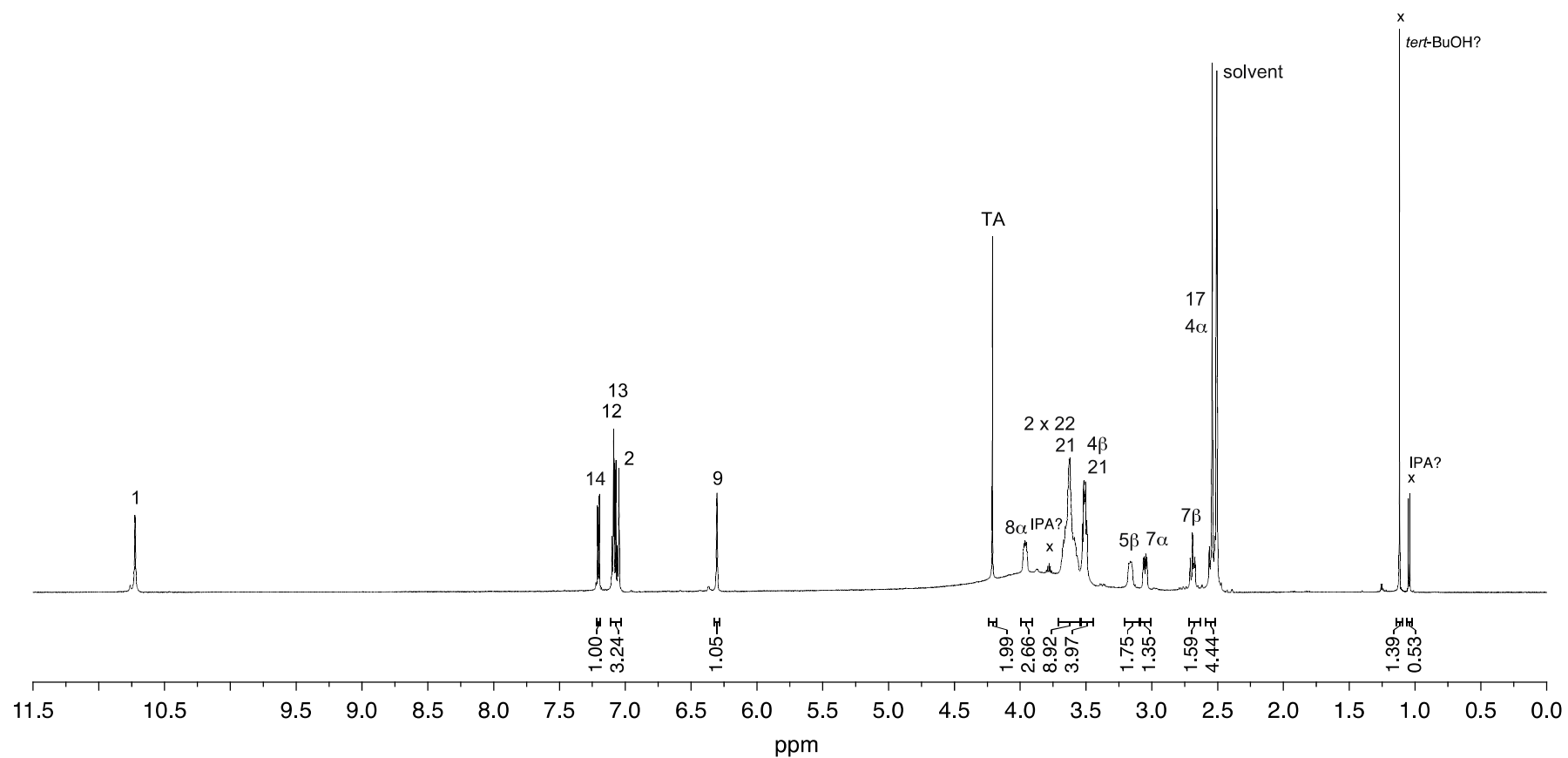
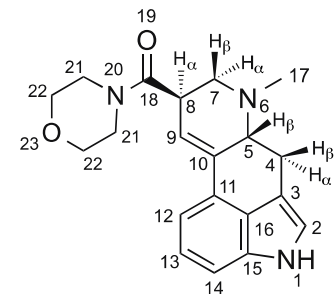
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Drug Testing and Analysis – Brandt *et al.* – Supporting Information

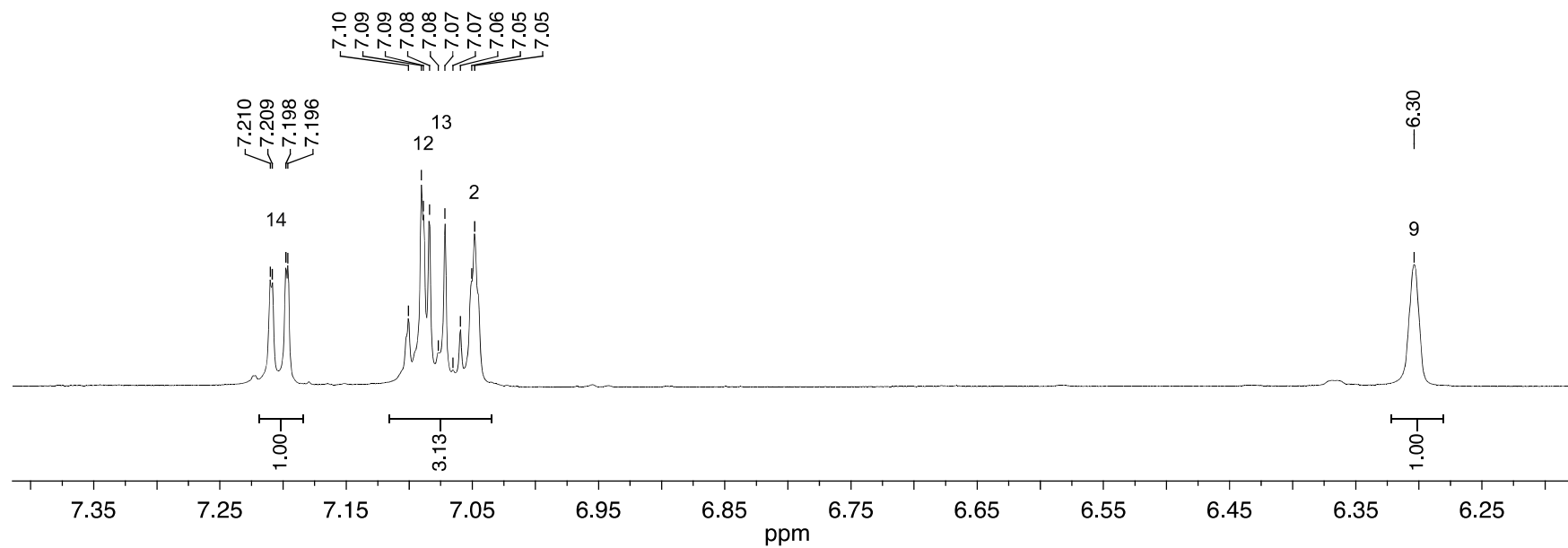
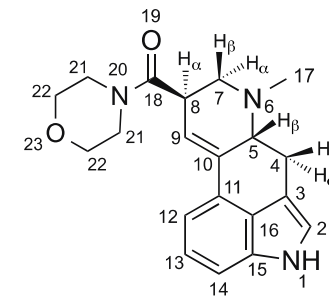
LSM-775 hemitartrate  
<sup>1</sup>H NMR / 600 MHz  
d<sub>6</sub> - DMSO



TA = tartaric acid

Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
<sup>1</sup>H NMR / 600 MHz  
d<sub>6</sub> - DMSO

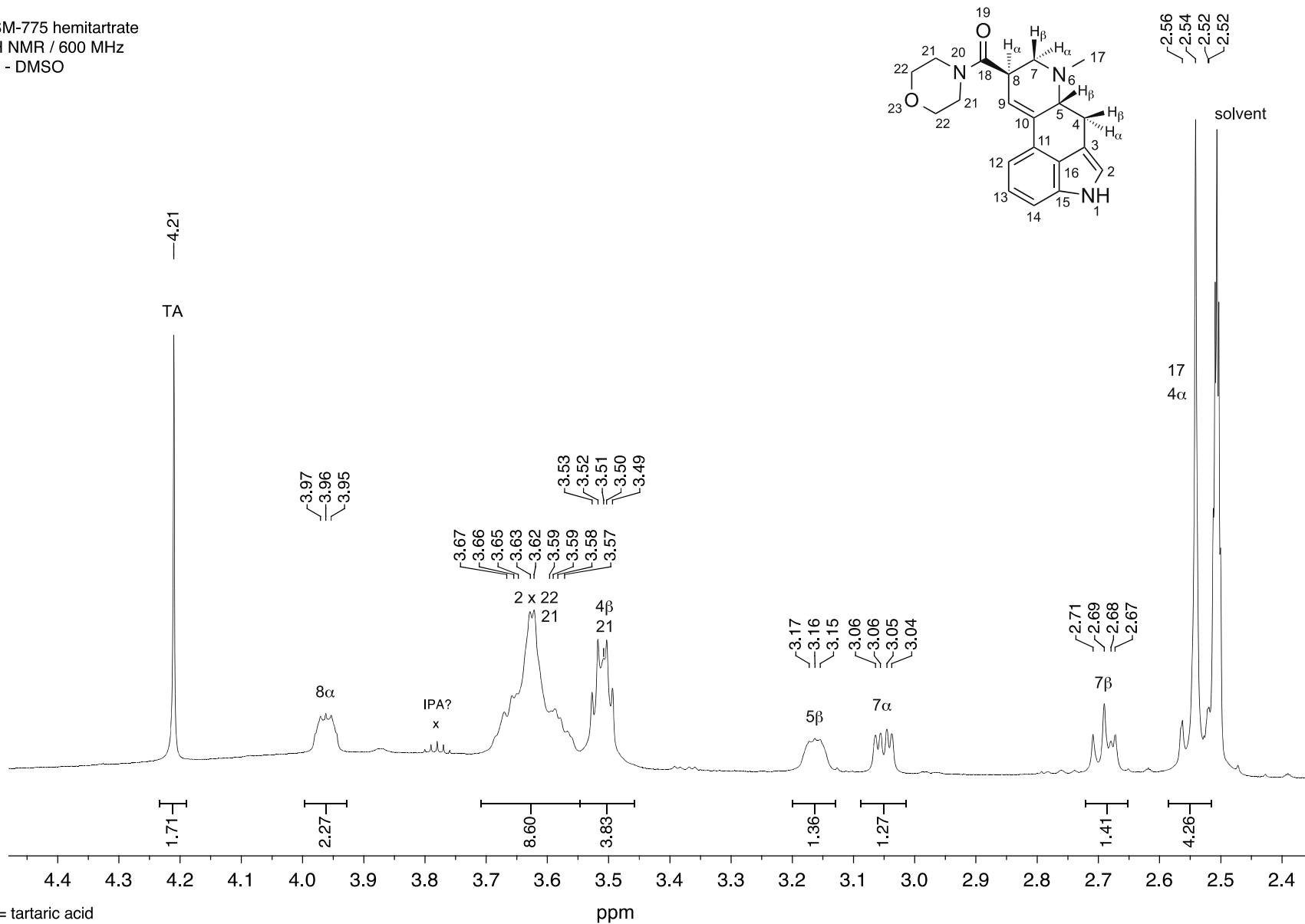


TA = tartaric acid



Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
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d<sub>6</sub> - DMSO

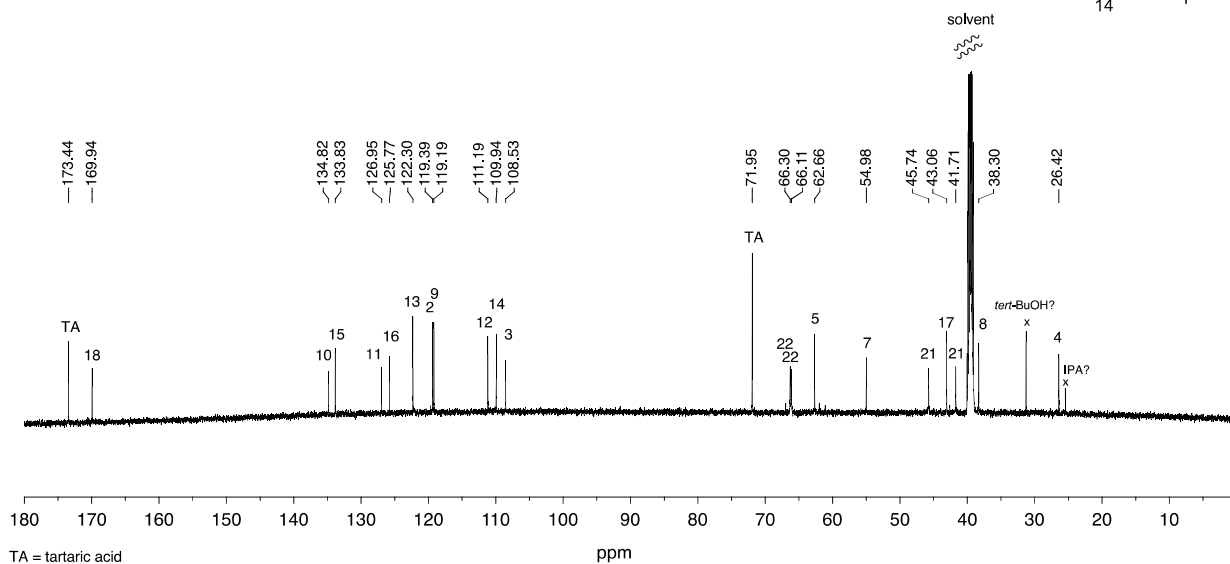
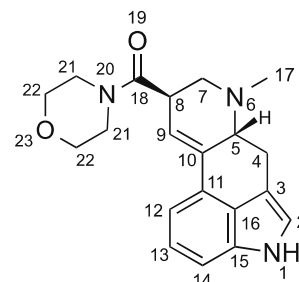


TA = tartaric acid

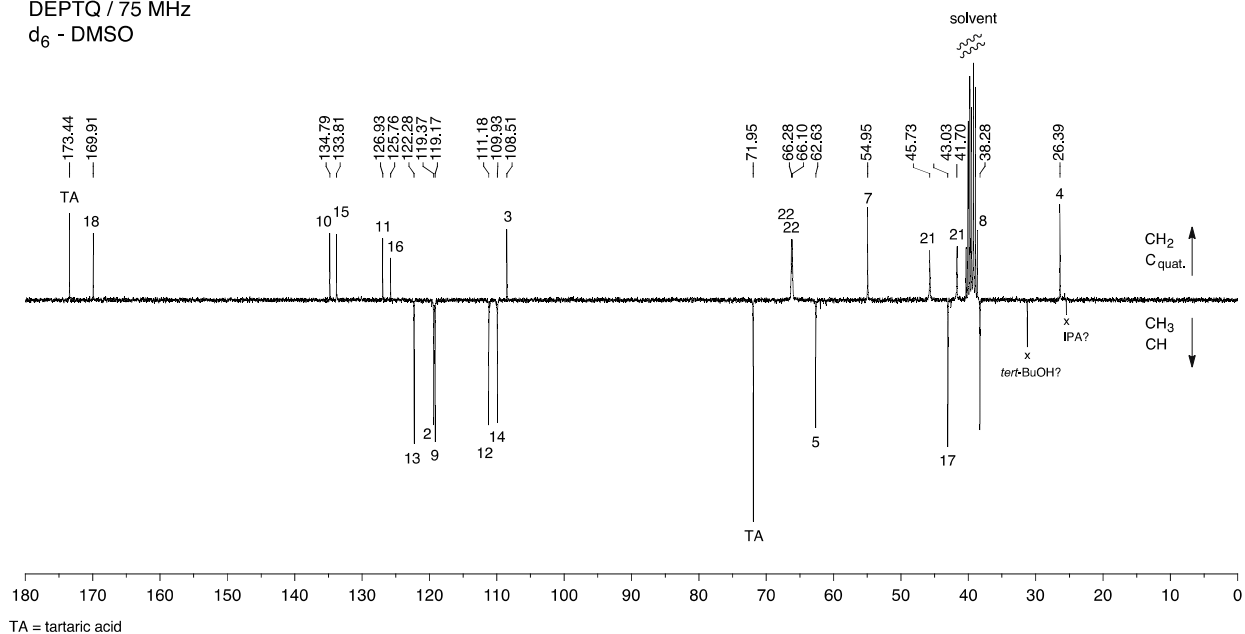
Increased baseline led to inaccurate integration values, particularly between 4.0 and 2.7 ppm

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LSM-775 hemitartrate  
<sup>13</sup>C NMR / 150 MHz  
 d<sub>6</sub> - DMSO

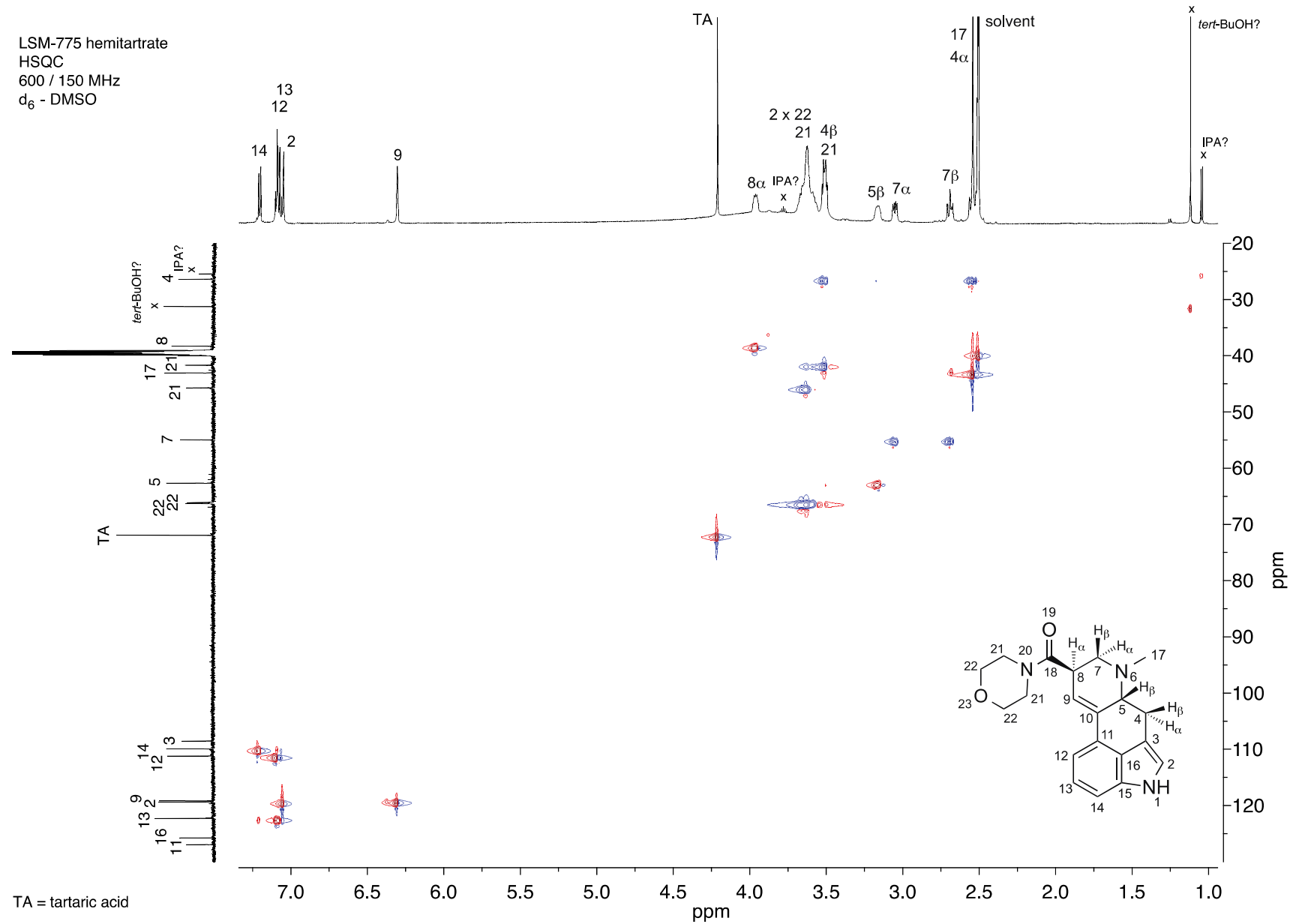


LSM-775 hemitartrate  
 DEPTQ / 75 MHz  
 d<sub>6</sub> - DMSO



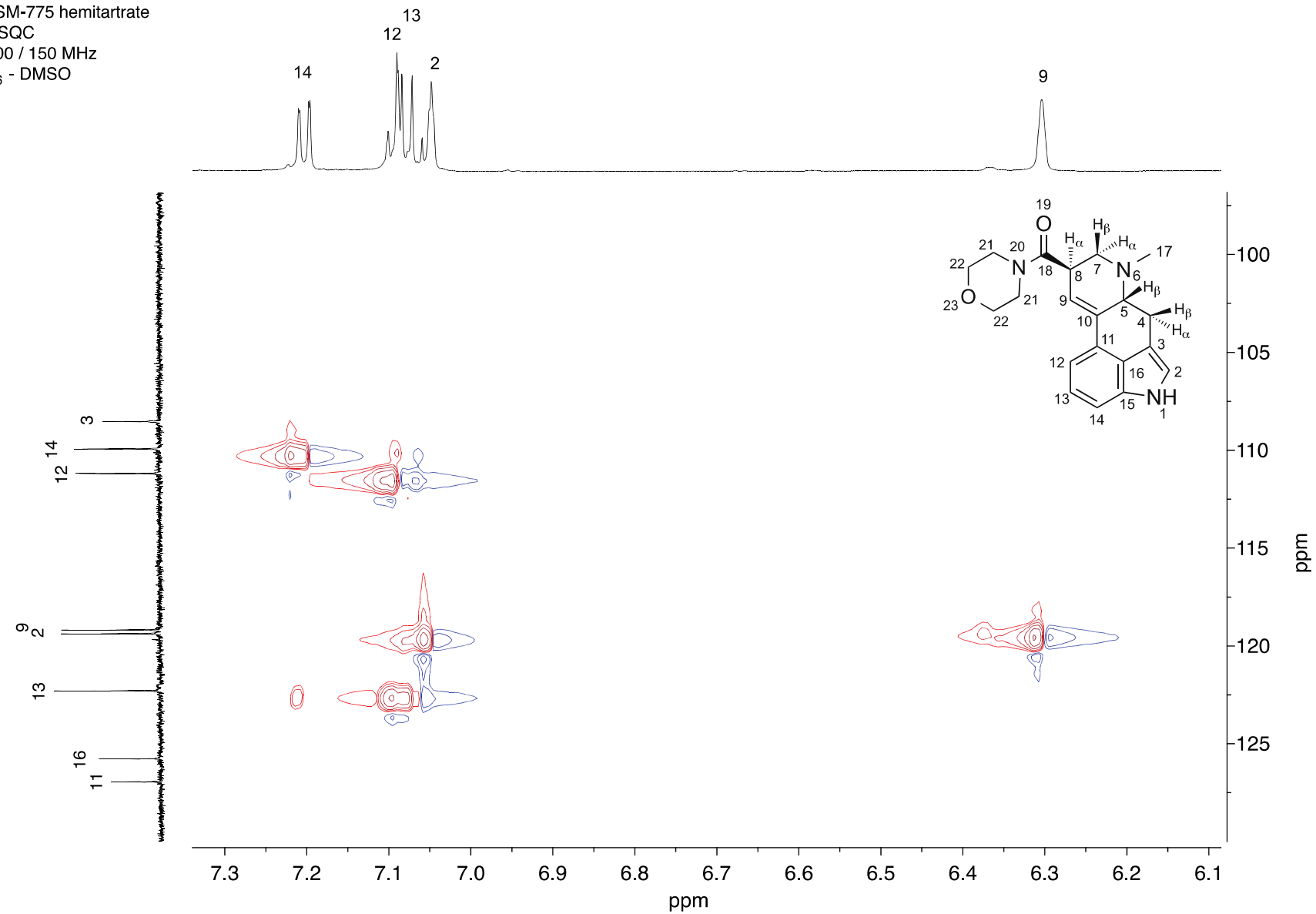
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
 HSQC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO



Drug Testing and Analysis – Brandt *et al.* – Supporting Information

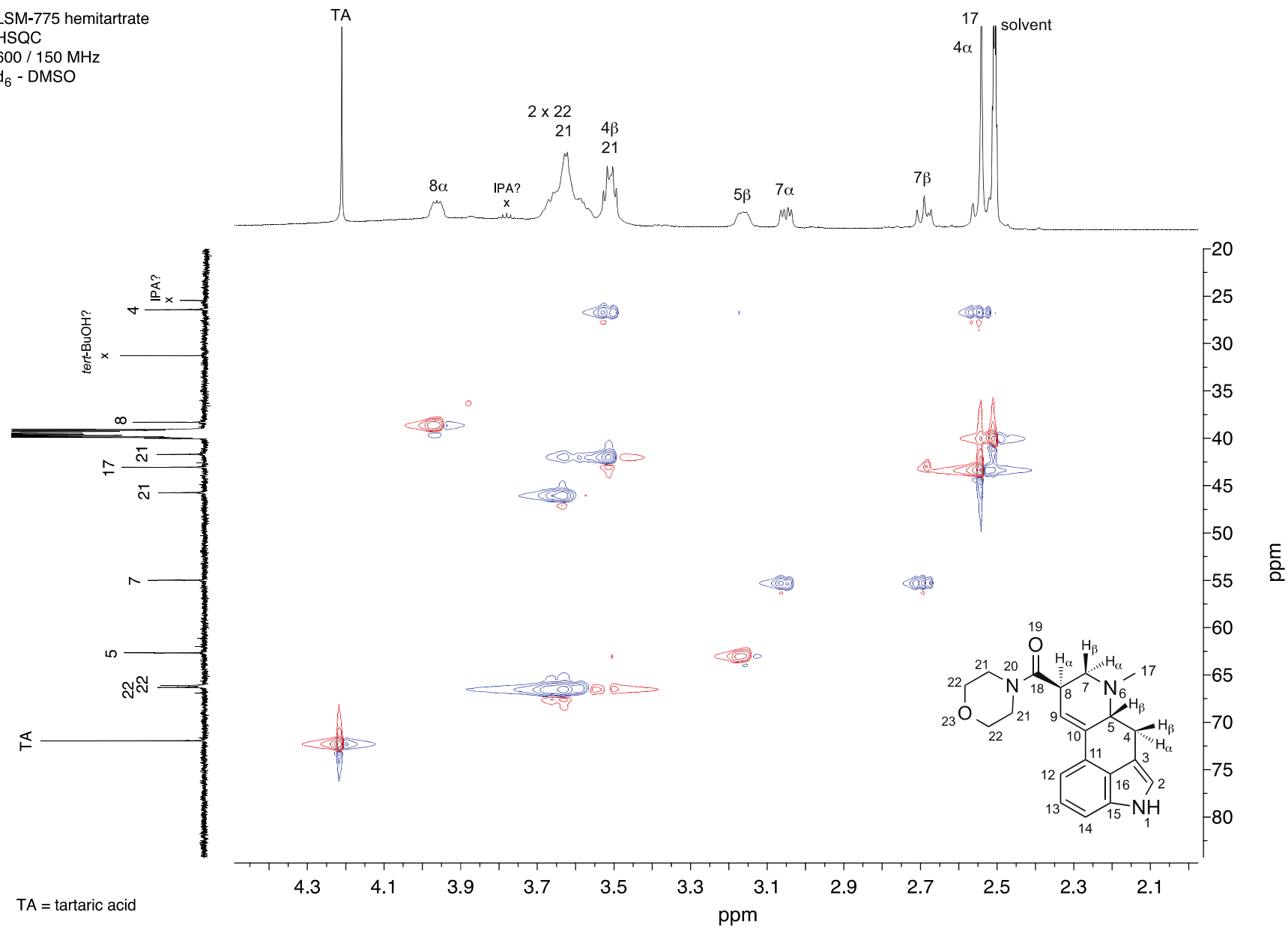
LSM-775 hemitartrate  
HSQC  
600 / 150 MHz  
d<sub>6</sub> - DMSO



TA = tartaric acid

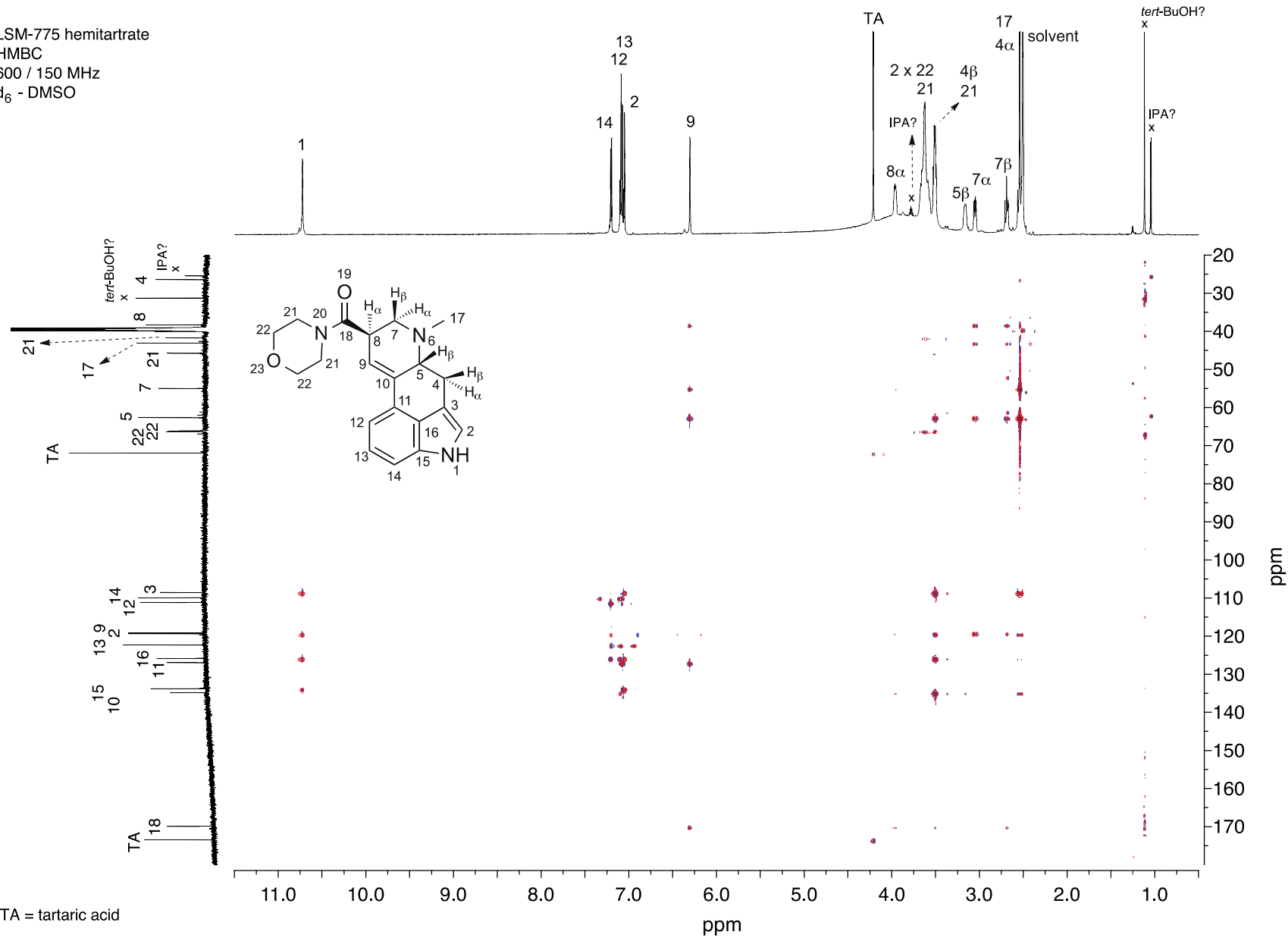
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

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 HSQC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO



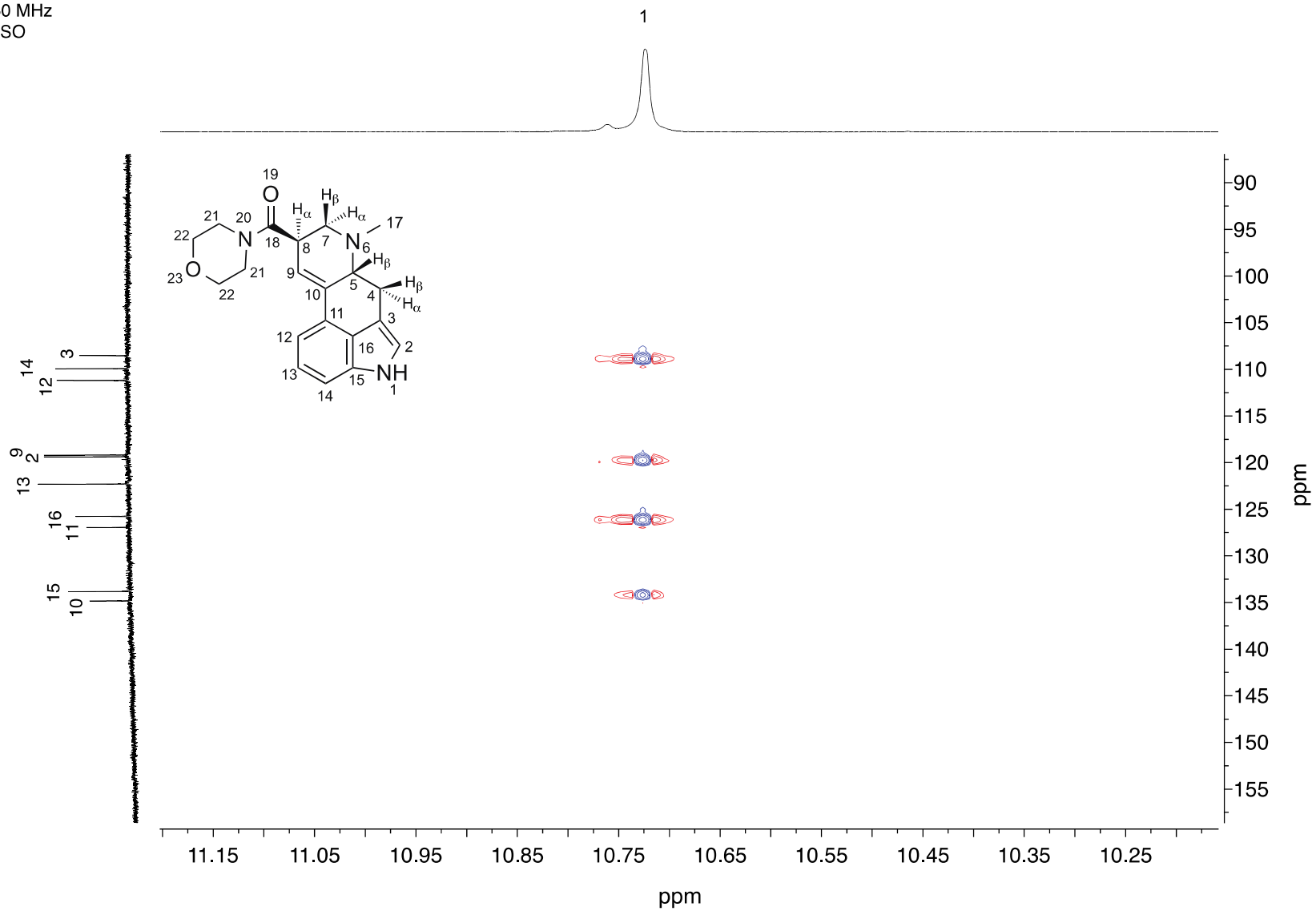
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
 HMBC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO



Drug Testing and Analysis – Brandt *et al.* – Supporting Information

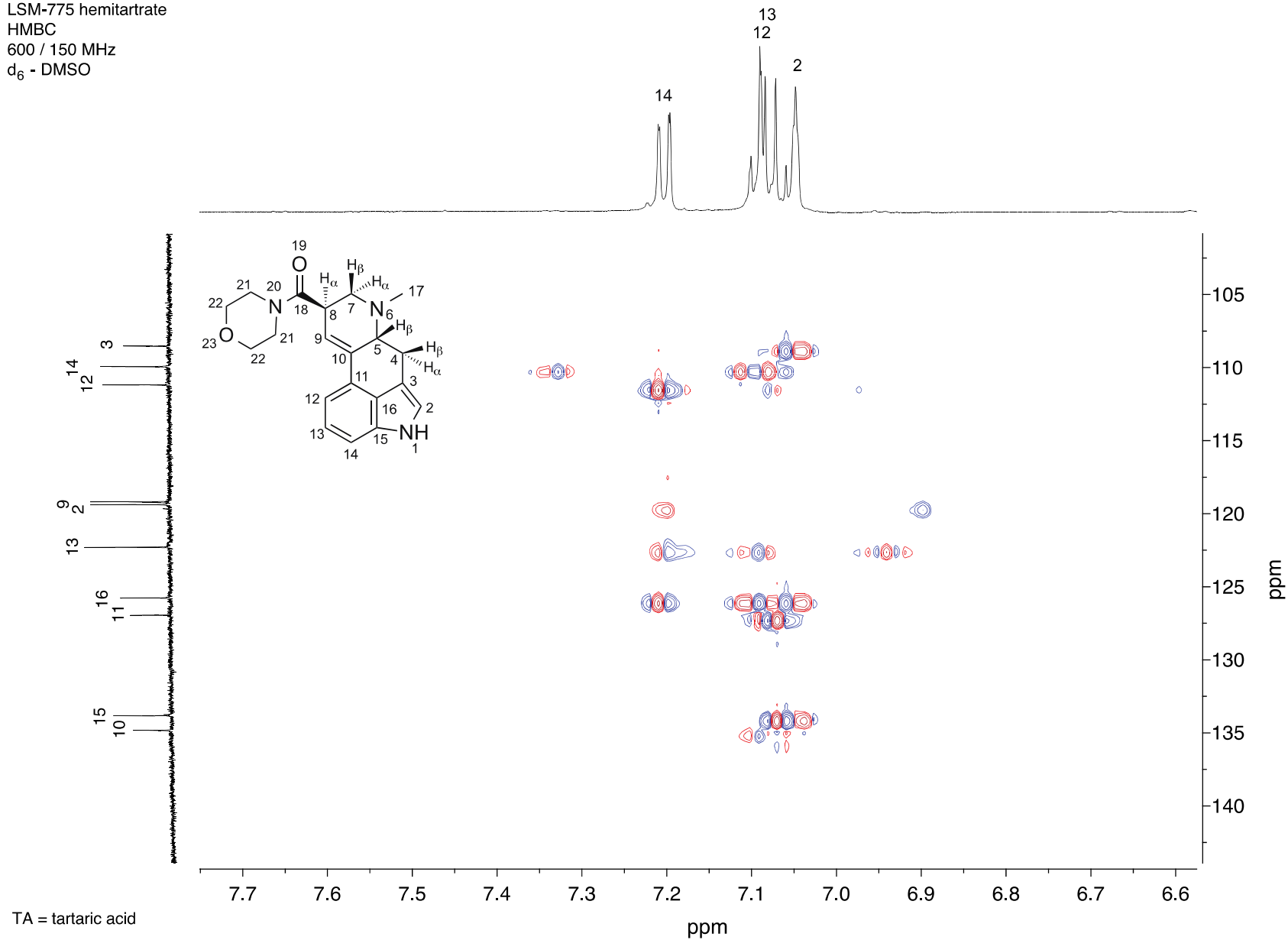
LSM-775 hemitartrate  
 HMBC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO



TA = tartaric acid

Drug Testing and Analysis – Brandt *et al.* – Supporting Information

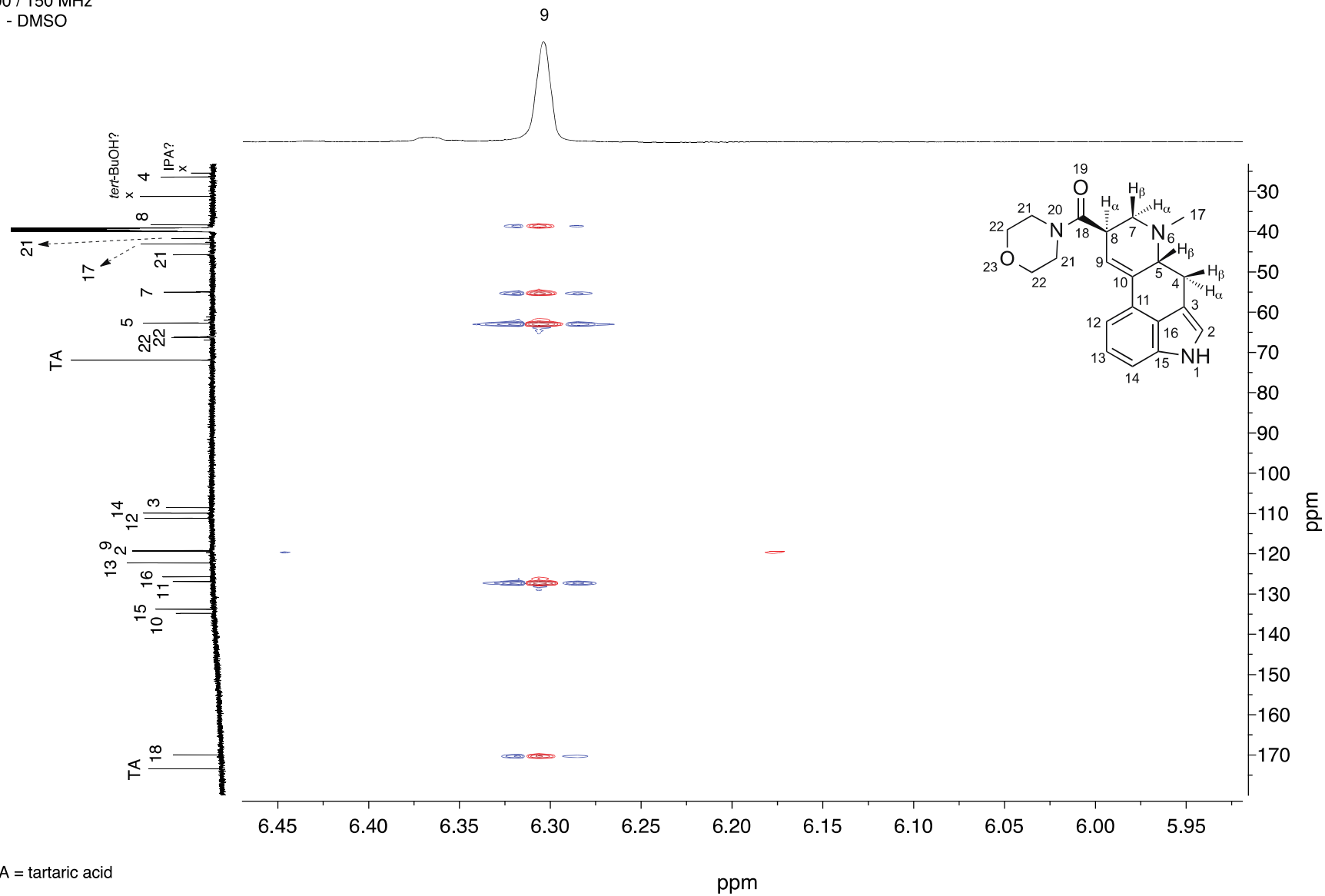
LSM-775 hemitartrate  
 HMBC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO





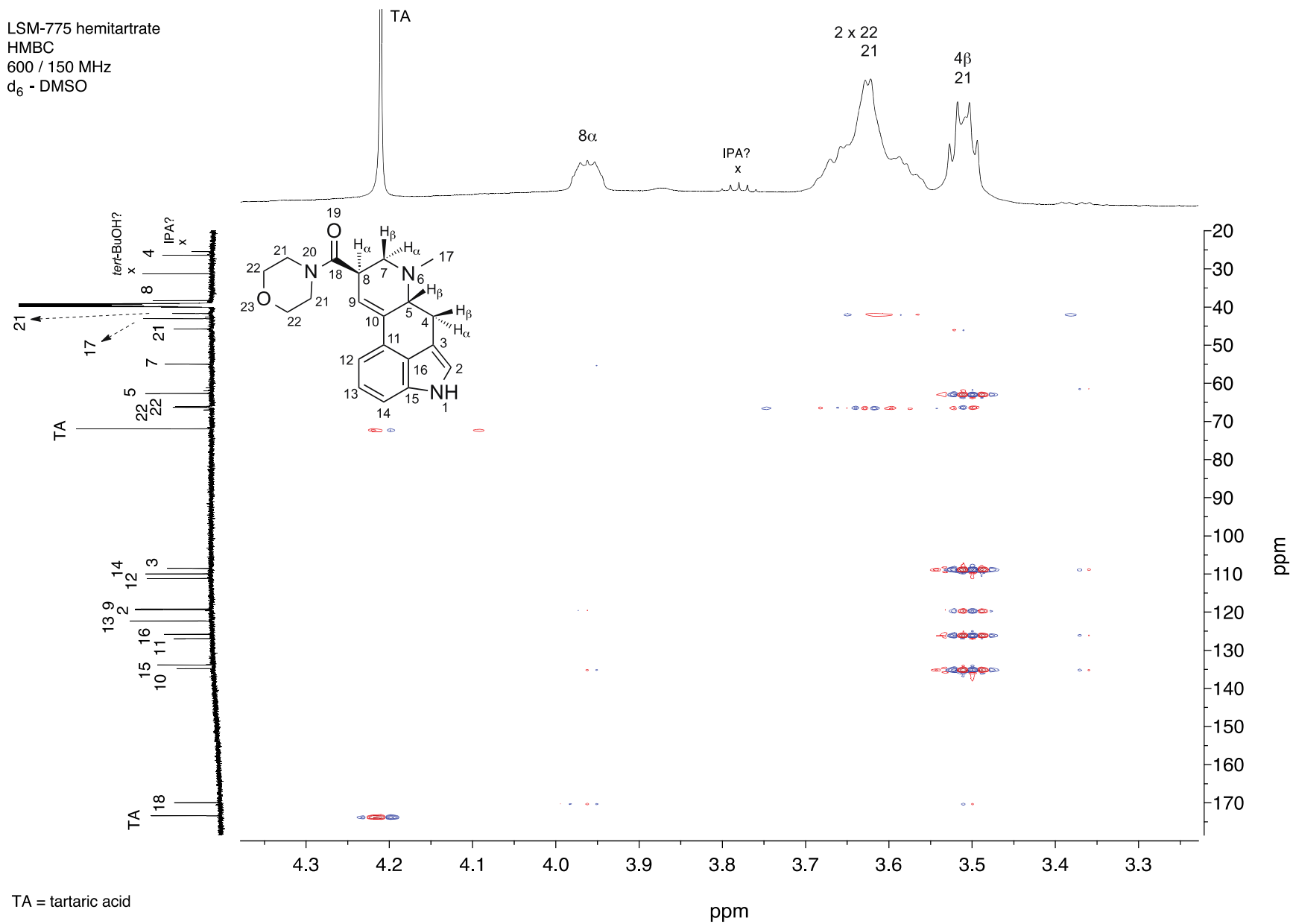
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
 HMBC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO



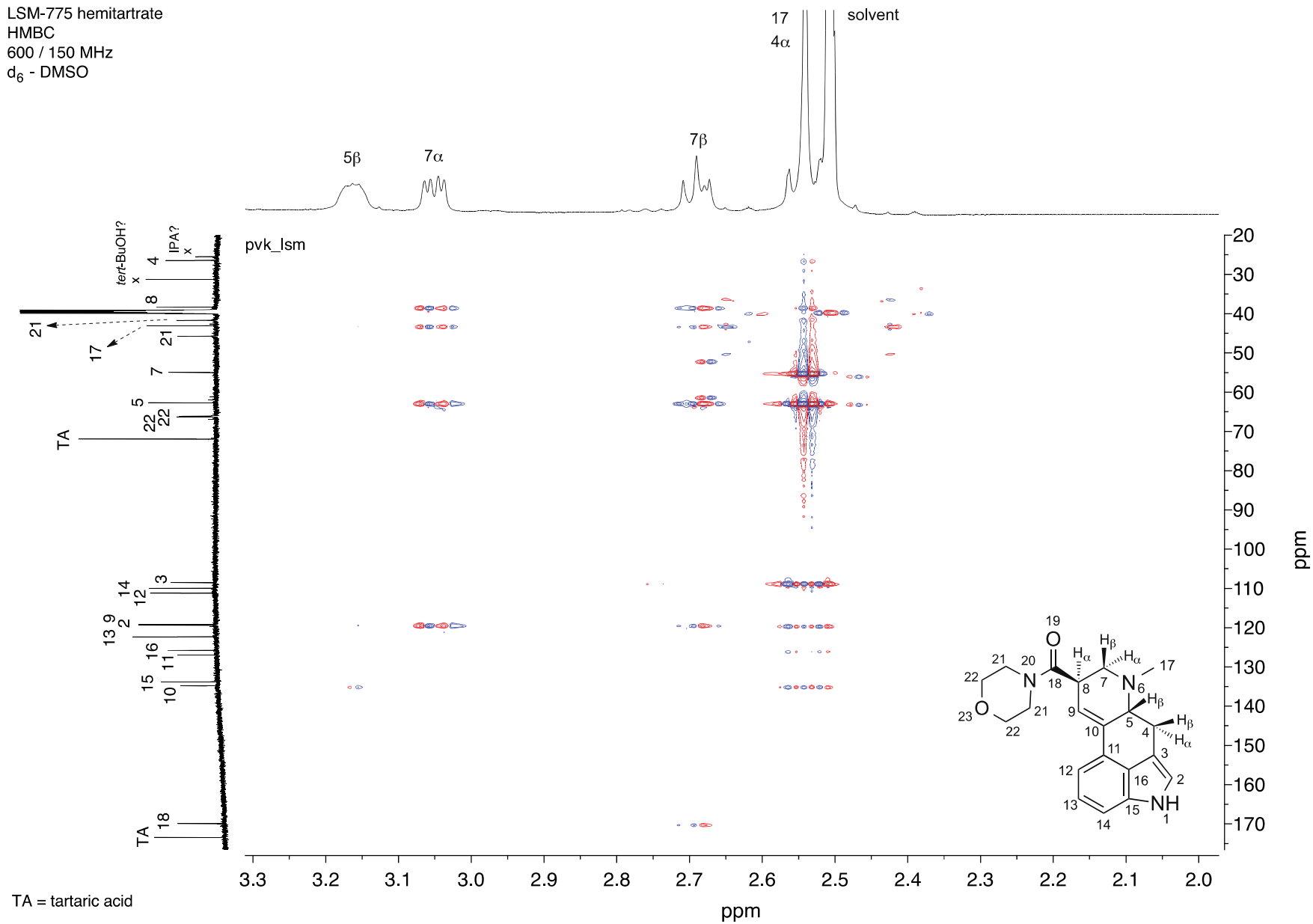
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
 HMBC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO



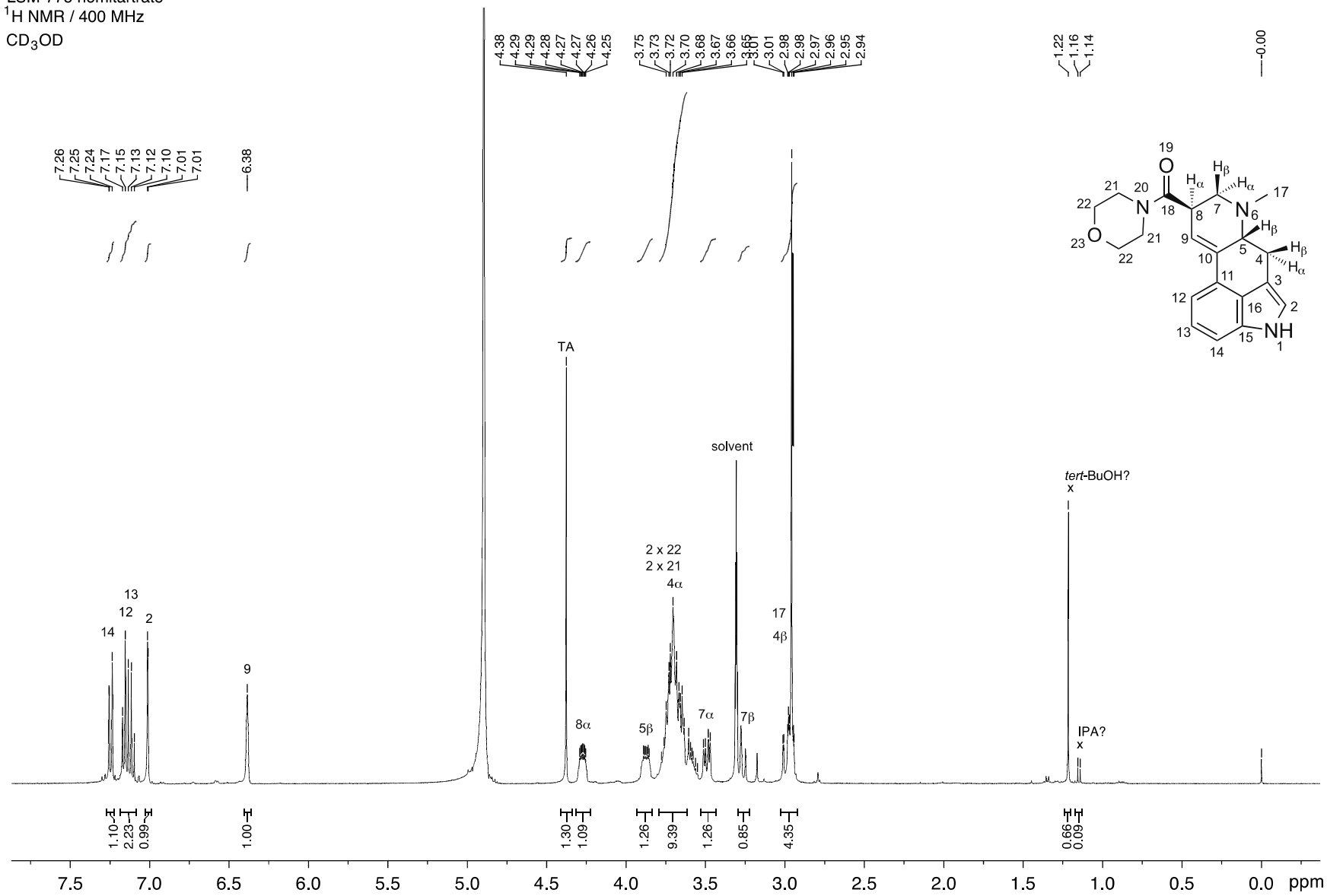
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
 HMBC  
 600 / 150 MHz  
 d<sub>6</sub> - DMSO



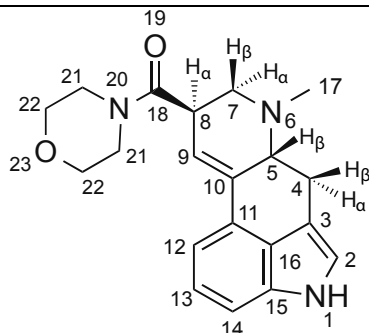
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
<sup>1</sup>H NMR / 400 MHz  
 CD<sub>3</sub>OD



TA = tartaric acid

<sup>1</sup>H and <sup>13</sup>C NMR data for LSM-775 hemitartrate in CD<sub>3</sub>OD at 400 / 100 MHz (20 mg/mL)



| No.             | <sup>13</sup> C [δ / ppm] | <sup>1</sup> H [δ / ppm]   |
|-----------------|---------------------------|--|
| 1               | –                         | –  |
| 2               | 120.86                    | 6.99 (d, <i>J</i> = 1.3 Hz, 1H)  |
| 3               | 108.53                    | –  |
| 4               | 26.32                     | 3.80–3.60 (m, 4β-H, 1H) <sup>a</sup><br>3.00–2.89 (m, 4α-H, 1H) <sup>b</sup> |
| 5               | 64.09                     | 3.90–3.80 (m, 5β-H, 1H)  |
| 6               | –                         | –  |
| 7               | 55.18                     | 3.47 (dd, <i>J</i> = 11.8, 5.1 Hz, 7α-H, 1H)<br>3.27–3.21 (m, 7β-H, 1H)      |
| 8               | 38.66                     | 4.36–4.20 (m, 8α-H, 1H)  |
| 9               | 118.79                    | 6.36 (s, 1H)   |
| 10              | 135.74                    | –  |
| 11              | 127.26                    | –  |
| 12              | 113.22                    | 7.13 (dd, <i>J</i> = 7.4, 0.9 Hz, 1H)  |
| 13              | 124.01                    | 7.09 (t, <i>J</i> = 7.5 Hz, 1H)  |
| 14              | 111.86                    | 7.22 (dd, <i>J</i> = 7.8, 0.9 Hz, 1H)  |
| 15              | 135.60                    | –  |
| 16              | 127.15                    | –  |
| 17              | 42.82                     | 2.94 (s, 3H) <sup>c</sup>  |
| 18              | 171.43                    | –  |
| 19              | –                         | –  |
| 20              | –                         | –  |
| 21              | 47.70                     | 3.80–3.52 (m, 2H) <sup>d, e</sup>  |
| 21              | 43.82                     | 3.80–3.52 (m, 2H) <sup>d, e</sup>  |
| 22              | 68.03                     | 3.80–3.52 (m, 2H) <sup>e, f</sup>  |
| 22              | 67.82                     | 3.80–3.52 (m, 2H) <sup>e, f</sup>  |
| 23              | –                         | –  |
| TA <sup>g</sup> | 74.44                     | 4.35 (s, 1H)   |
| TA <sup>g</sup> | 177.49                    | –  |

<sup>a</sup> Overlapping with 22-CH<sub>2</sub> and 21-CH<sub>2</sub> (4H)

<sup>b</sup> Overlapping with 17-CH<sub>3</sub> (3H)

<sup>c</sup> Overlapping with 4α-H (1H)

<sup>d</sup> Overlapping with 2 x 22-CH<sub>2</sub> (4H)

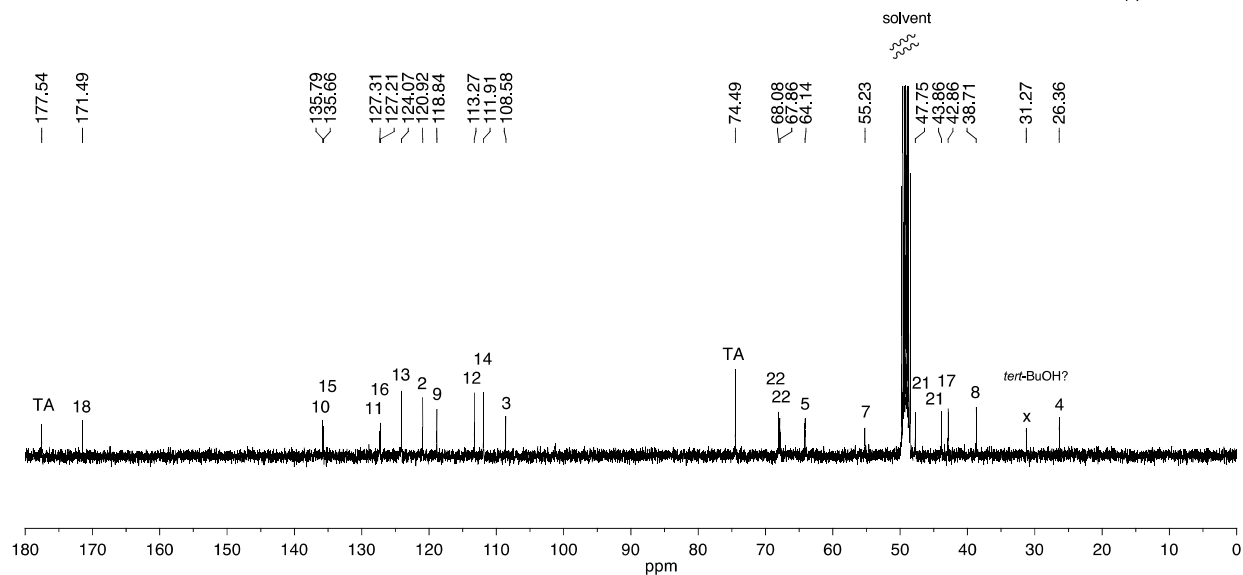
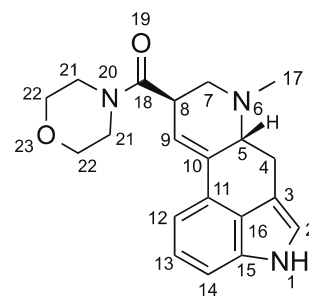
<sup>e</sup> Overlapping with 4β-H (1H)

<sup>f</sup> Overlapping with 1 x 21-CH<sub>2</sub> (2H)

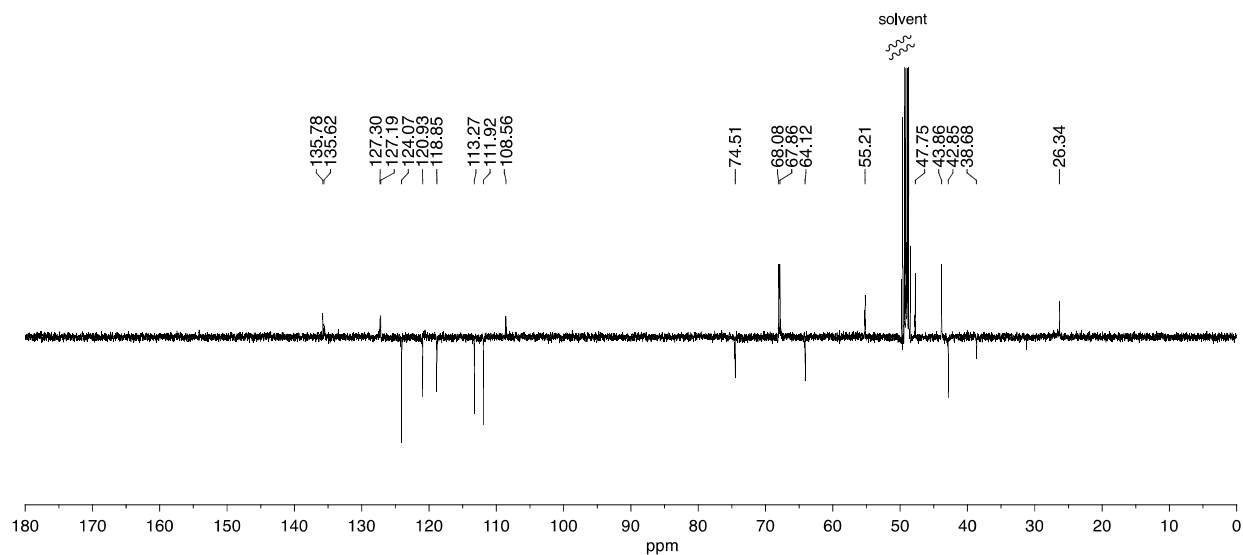
<sup>g</sup> TA: tartaric acid

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LSM-775 hemitartrate  
<sup>13</sup>C NMR / 100 MHz  
 CD<sub>3</sub>OD



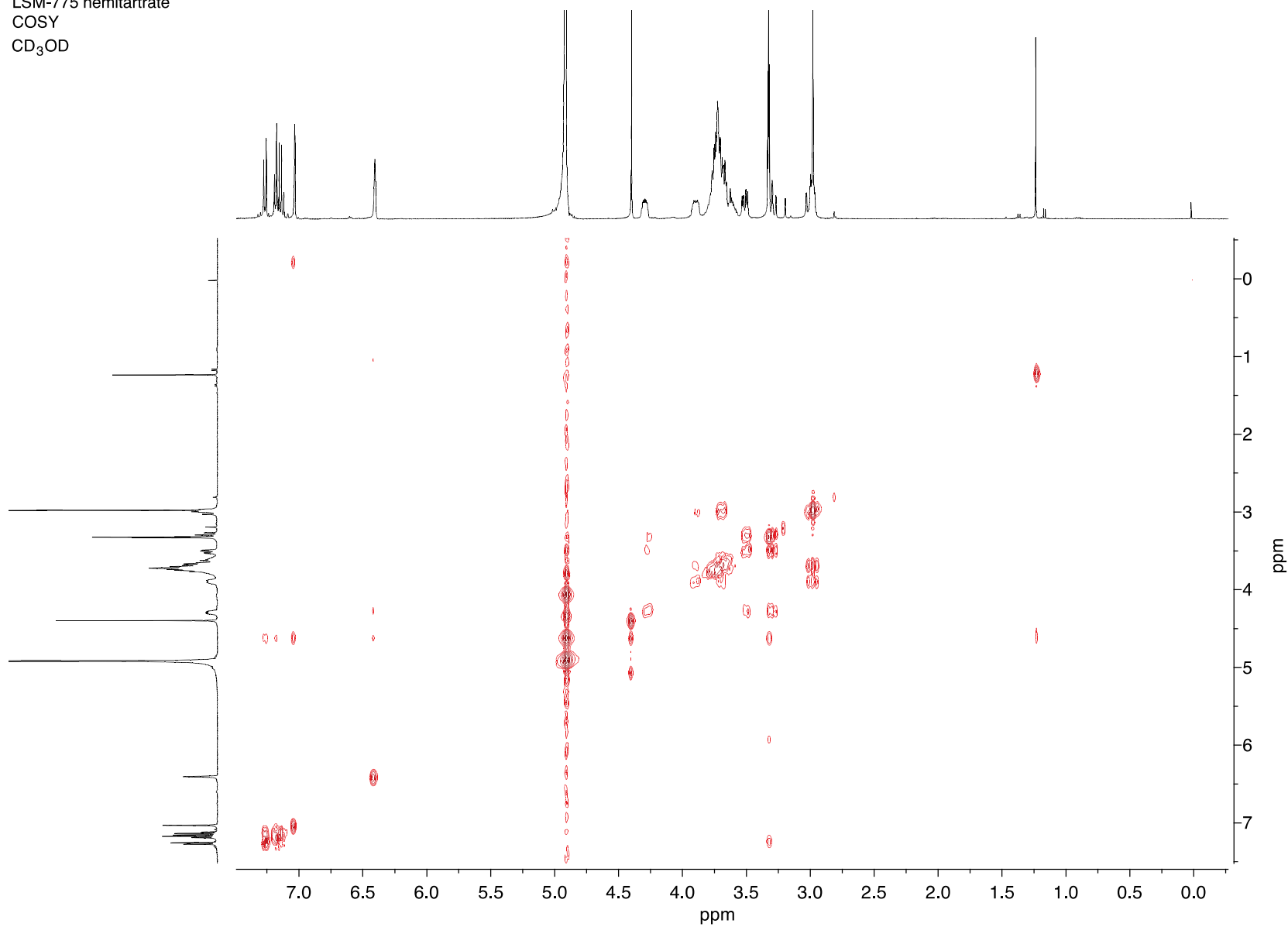
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 PENDANT / 100 MHz  
 CD<sub>3</sub>OD



TA = tartaric acid

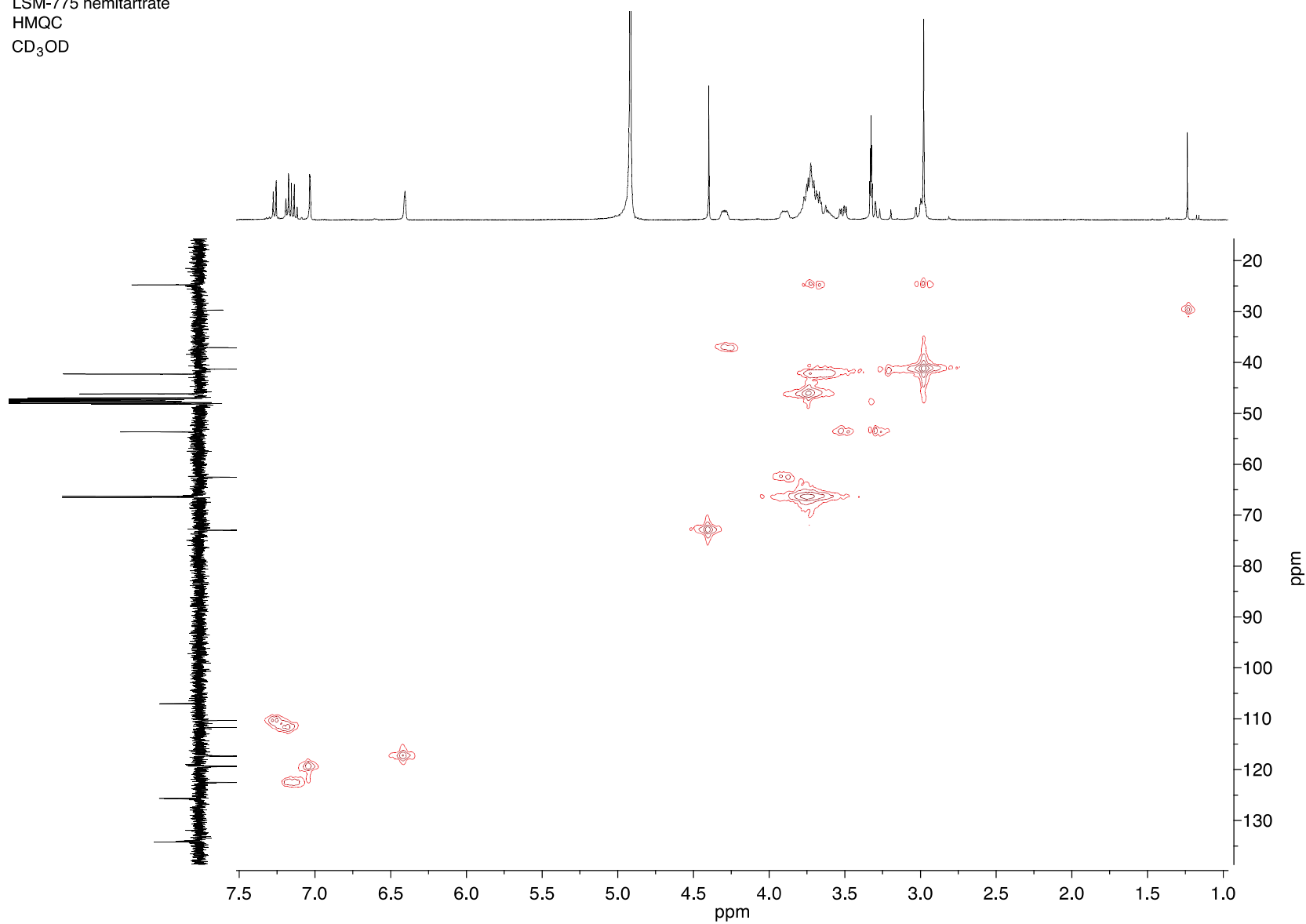
Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
COSY  
CD<sub>3</sub>OD



Drug Testing and Analysis – Brandt *et al.* – Supporting Information

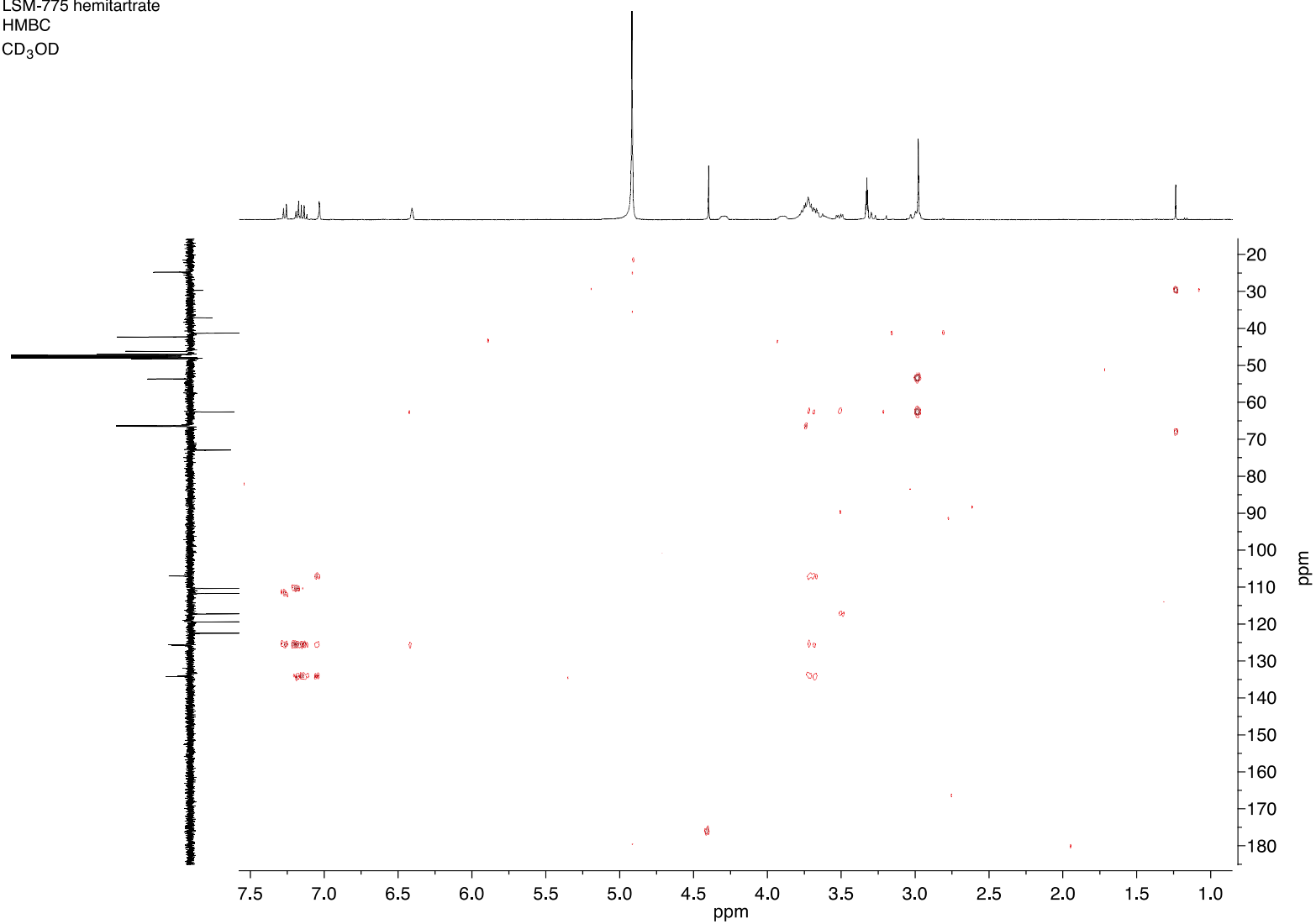
LSM-775 hemitartrate  
HMQC  
CD<sub>3</sub>OD





Drug Testing and Analysis – Brandt *et al.* – Supporting Information

LSM-775 hemitartrate  
HMBC  
CD<sub>3</sub>OD





# Small Molecule X-ray Facility School Of Chemistry

## Structure Report

Filename: TCD173

Submitted by: Pierce Kavanagh

Reference: PKSBLSM

Group: Kavanagh

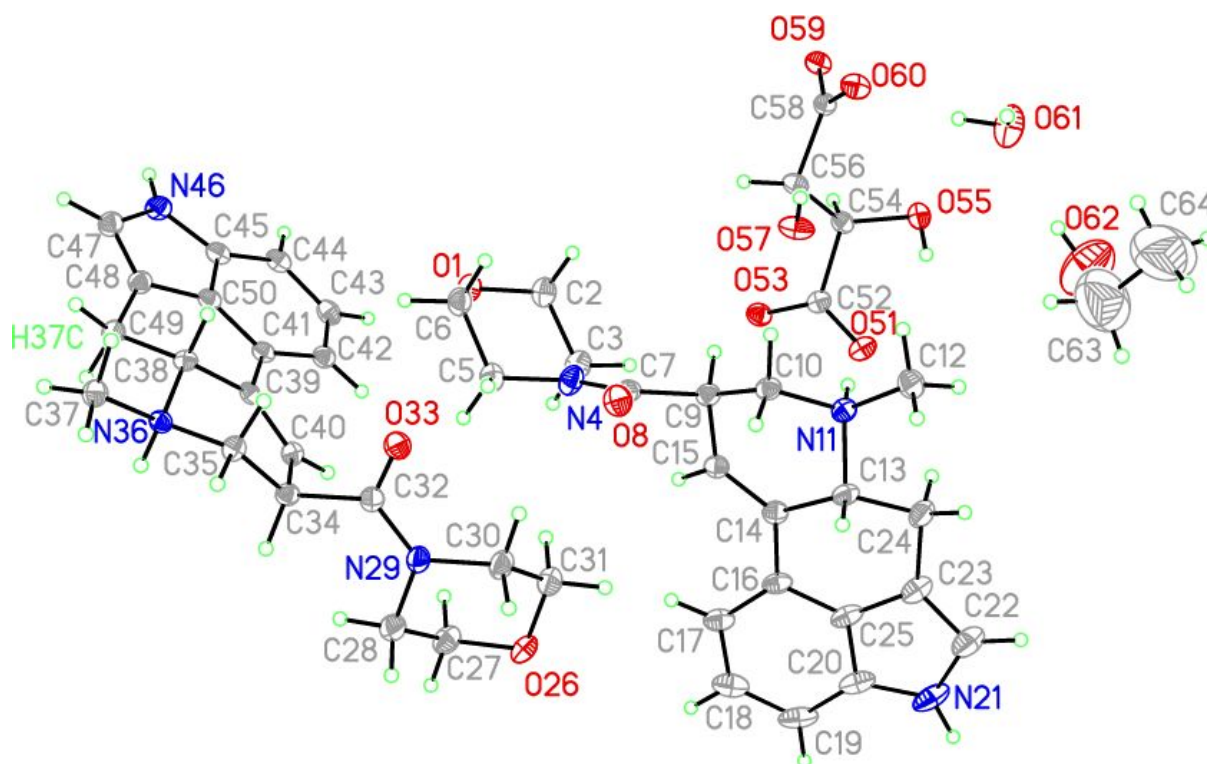


Fig. 1. Asymmetric unit of TCD173 showing both molecules with the tartrate anion as well as solvates H<sub>2</sub>O and EtOH. Displacement ellipsoids shown at 50%.

11/12/14

Brendan Twamley

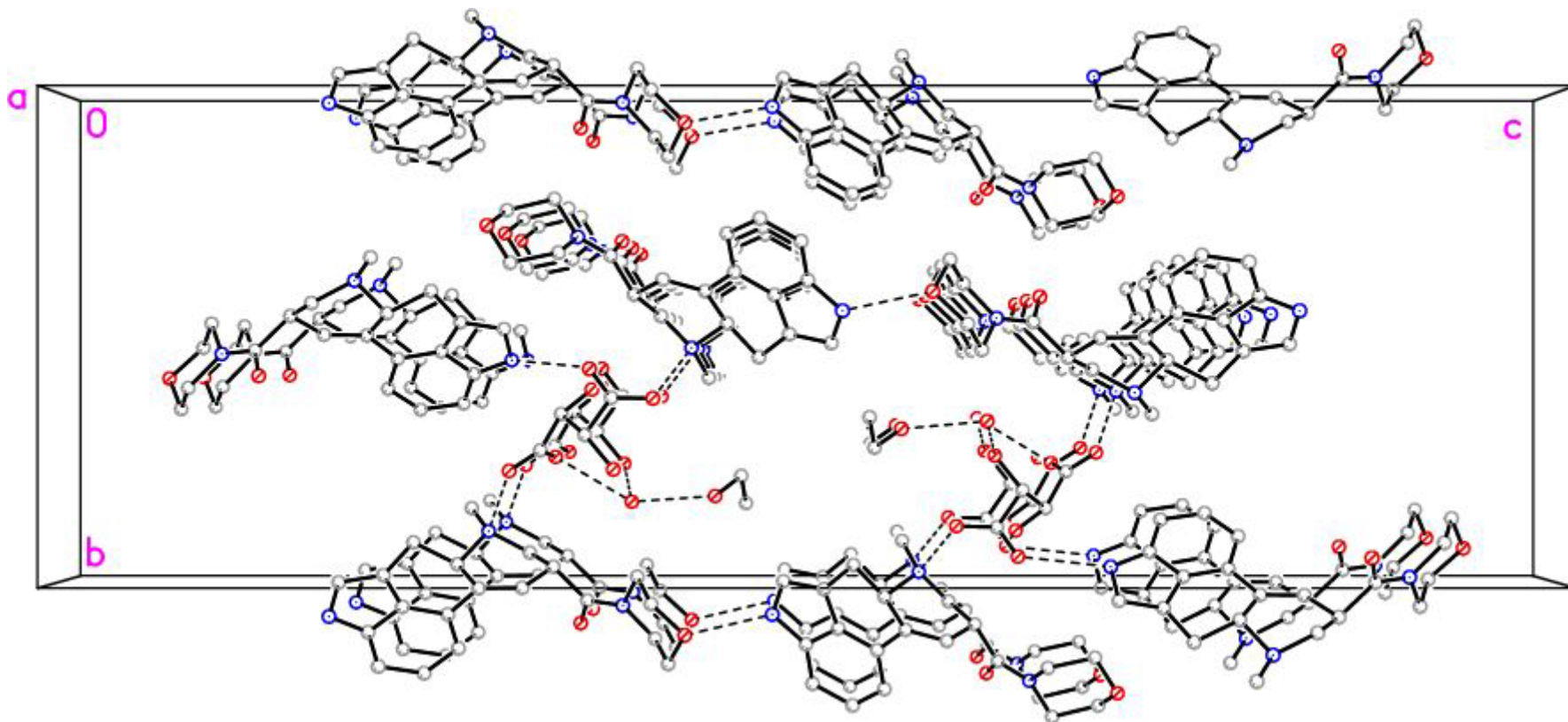


Figure 2. Packing diagram of TCD173 viewed down the a-axis. Hydrogen atoms omitted for clarity.

## Crystal Structure Report for TCD173

A specimen of  $C_{46}H_{60}N_6O_{12}$ , approximate dimensions 0.060 mm x 0.060 mm x 0.210 mm, was used for the X-ray crystallographic analysis. The X-ray intensity data were measured at 100(2)K using an Oxford Cryosystems Cobra low temperature device using a MiTeGen micromount. See Table 1 for collection parameters and exposure time. Bruker APEX software was used to correct for Lorentz and polarization effects.

A total of 5863 frames were collected. The total exposure time was 29.31 hours. The integration of the data using an orthorhombic unit cell yielded a total of 57701 reflections to a maximum  $\theta$  angle of  $68.30^\circ$  (0.83 Å resolution), of which 8027 were independent (average redundancy 7.188, completeness = 99.9%,  $R_{\text{int}} = 4.44\%$ ,  $R_{\text{sig}} = 2.32\%$ ) and 7723 (96.21%) were greater than  $2\sigma(F^2)$ . The final cell constants of  $a = 5.9621(2)$  Å,  $b = 15.5087(6)$  Å,  $c = 47.4811(19)$  Å, volume =  $4390.3(3)$  Å<sup>3</sup>, are based upon the refinement of the XYZ-centroids of reflections above  $20\sigma(I)$ . Data were corrected for absorption effects using the numerical method (SADABS). The calculated minimum and maximum transmission coefficients (based on crystal size) are 0.8490 and 0.9424.

The structure was solved and refined using the Bruker SHELXTL Software Package, using the space group  $P2_12_12_1$ , with  $Z = 4$  for the formula unit,  $C_{46}H_{60}N_6O_{12}$ . The final anisotropic full-matrix least-squares refinement on  $F^2$  with 607 variables converged at  $R1 = 3.71\%$ , for the observed data and  $wR2 = 9.92\%$  for all data. The goodness-of-fit was 1.050. The largest peak in the final difference electron density synthesis was  $0.557 e^-/\text{Å}^3$  and the largest hole was  $-0.395 e^-/\text{Å}^3$  with an RMS deviation of  $0.049 e^-/\text{Å}^3$ . On the basis of the final model, the calculated density was  $1.345 \text{ g/cm}^3$  and  $F(000)$ , 1896  $e^-$ .

**Refinement Note:** All donor hydrogen atoms were located and refined with restraints (SADI). The solvent EtOH molecule C-C distance was also restrained (DFIX). Absolute configuration was established by anomalous-dispersion effects in diffraction measurements on the crystal.

### References:

Bruker APEX v2012.12-0, Bruker AXS Inc., Madison, Wisconsin, USA.

SADABS (2014/3) Bruker AXS Inc., Madison, Wisconsin, USA; Sheldrick, G. M. University of Göttingen, Germany.

SHELXL-2014, (2014), Bruker AXS Inc., Madison, Wisconsin, USA; Sheldrick, G. M. University of Göttingen, Germany.

### Acknowledgement:

Facility funded by PRTL and ERDF.

Table 1: Data collection details for TCD173.

| Axis  | dx/mm  | 2 $\theta$ / $^{\circ}$ | $\omega$ / $^{\circ}$ | $\varphi$ / $^{\circ}$ | $\chi$ / $^{\circ}$ | Width/ $^{\circ}$ | Frames | Time/s | Wavelength/ $\text{\AA}$ | Voltage/kV | Current/mA | Temperature/K |
|-------|--------|-------------------------|-----------------------|------------------------|---------------------|-------------------|--------|--------|--------------------------|------------|------------|---------------|
| Omega | 45.000 | -59.59                  | 184.68                | 360.00                 | 54.74               | 0.60              | 209    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 343.28                | 270.00                 | 64.87               | 0.60              | 215    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Phi   | 45.000 | 104.92                  | 82.49                 | 0.00                   | -23.00              | 0.60              | 600    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | -74.59                  | 169.68                | 270.00                 | 54.74               | 0.60              | 209    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Phi   | 45.000 | 104.92                  | 4.83                  | 180.20                 | 23.00               | 0.60              | 306    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | -39.60                  | 312.65                | 0.00                   | -64.88              | 0.60              | 165    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | -39.60                  | 312.65                | 270.00                 | -64.88              | 0.60              | 165    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Phi   | 45.000 | 89.92                   | 80.25                 | 360.00                 | -57.06              | 0.60              | 600    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 95.82                 | 189.00                 | -54.74              | 0.60              | 206    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Phi   | 45.000 | -36.55                  | 333.12                | 0.00                   | 57.06               | 0.60              | 600    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 95.82                 | 270.00                 | -54.74              | 0.60              | 206    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 343.28                | 27.00                  | 64.88               | 0.60              | 215    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Phi   | 45.000 | 104.92                  | 95.25                 | 0.00                   | -57.06              | 0.60              | 600    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 95.82                 | 108.00                 | -54.74              | 0.60              | 206    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Phi   | 45.000 | -74.92                  | 42.49                 | 360.00                 | -57.06              | 0.60              | 600    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 95.82                 | 216.00                 | -54.74              | 0.60              | 206    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 95.82                 | 81.00                  | -54.74              | 0.60              | 206    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | -26.40                  | 325.84                | 153.00                 | -64.88              | 0.60              | 143    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |
| Omega | 45.000 | 104.59                  | 95.82                 | 135.00                 | -54.74              | 0.60              | 206    | 18.00  | 1.54184                  | 45         | 0.6        | 100           |

Table 2. Crystal data and structure refinement for TCD173.

|                                   |  |          |
|-----------------------------------|--|----------|
| Identification code               | tcd173   |          |
| Empirical formula                 | C <sub>46</sub> H <sub>60</sub> N <sub>6</sub> O <sub>12</sub> |          |
| Formula weight                    | 889.00   |          |
| Temperature                       | 100(2) K   |          |
| Wavelength                        | 1.54178 Å  |          |
| Crystal system                    | Orthorhombic   |          |
| Space group                       | P2 <sub>1</sub> 2 <sub>1</sub> 2 <sub>1</sub>                  |          |
| Unit cell dimensions              | a = 5.9621(2) Å  | α = 90°. |
|                                   | b = 15.5087(6) Å   | β = 90°. |
|                                   | c = 47.4811(19) Å  | γ = 90°. |
| Volume                            | 4390.3(3) Å <sup>3</sup>                                       |          |
| Z                                 | 4  |          |
| Density (calculated)              | 1.345 Mg/m <sup>3</sup>  |          |
| Absorption coefficient            | 0.807 mm <sup>-1</sup>   |          |
| F(000)                            | 1896   |          |
| Crystal size                      | 0.210 x 0.060 x 0.060 mm <sup>3</sup>                          |          |
| Theta range for data collection   | 3.404 to 68.296°.  |          |
| Index ranges                      | -7 ≤ h ≤ 7, -18 ≤ k ≤ 17, -57 ≤ l ≤ 56                         |          |
| Reflections collected             | 57701  |          |
| Independent reflections           | 8027 [R(int) = 0.0444]   |          |
| Completeness to theta = 67.679°   | 99.9 %   |          |
| Absorption correction             | Numerical  |          |
| Max. and min. transmission        | 1.000 and 0.9424   |          |
| Refinement method                 | Full-matrix least-squares on F <sup>2</sup>                    |          |
| Data / restraints / parameters    | 8027 / 10 / 607  |          |
| Goodness-of-fit on F <sup>2</sup> | 1.050  |          |
| Final R indices [I > 2σ(I)]       | R1 = 0.0371, wR2 = 0.0979                                      |          |
| R indices (all data)              | R1 = 0.0387, wR2 = 0.0992                                      |          |
| Absolute structure parameter      | 0.08(4)  |          |
| Largest diff. peak and hole       | 0.557 and -0.395 e.Å <sup>-3</sup>                             |          |

Table 3. Atomic coordinates ( $\times 10^4$ ) and equivalent isotropic displacement parameters ( $\text{\AA}^2 \times 10^3$ ) for TCD173.  $U(\text{eq})$  is defined as one third of the trace of the orthogonalized  $U^{ij}$  tensor.

|       | x        | y        | z        | U(eq) |
|-------|----------|----------|----------|-------|
| C(2)  | 1323(5)  | 6429(2)  | 8112(1)  | 23(1) |
| C(3)  | 1086(5)  | 6623(2)  | 8423(1)  | 24(1) |
| C(5)  | 3978(5)  | 7721(2)  | 8369(1)  | 25(1) |
| C(6)  | 4120(5)  | 7500(2)  | 8061(1)  | 22(1) |
| C(7)  | 4580(4)  | 6612(2)  | 8727(1)  | 19(1) |
| C(9)  | 3663(4)  | 5830(2)  | 8885(1)  | 18(1) |
| C(10) | 5566(4)  | 5415(2)  | 9049(1)  | 18(1) |
| C(12) | 6346(5)  | 4182(2)  | 9358(1)  | 25(1) |
| C(13) | 3382(4)  | 5257(2)  | 9486(1)  | 19(1) |
| C(14) | 1694(4)  | 5881(2)  | 9352(1)  | 18(1) |
| C(15) | 1808(4)  | 6107(2)  | 9080(1)  | 18(1) |
| C(16) | -29(4)   | 6233(2)  | 9543(1)  | 20(1) |
| C(17) | -1369(5) | 6954(2)  | 9501(1)  | 24(1) |
| C(18) | -2928(5) | 7215(2)  | 9708(1)  | 29(1) |
| C(19) | -3193(5) | 6778(2)  | 9963(1)  | 32(1) |
| C(20) | -1854(5) | 6054(2)  | 10006(1) | 28(1) |
| C(22) | -26(6)   | 4861(2)  | 10153(1) | 34(1) |
| C(23) | 772(5)   | 5037(2)  | 9889(1)  | 27(1) |
| C(24) | 2382(5)  | 4591(2)  | 9693(1)  | 25(1) |
| C(25) | -331(5)  | 5788(2)  | 9797(1)  | 24(1) |
| C(27) | 1209(5)  | 9790(2)  | 9010(1)  | 25(1) |
| C(28) | 3271(5)  | 10270(2) | 8909(1)  | 24(1) |
| C(30) | 5484(5)  | 8983(2)  | 9010(1)  | 22(1) |
| C(31) | 3341(5)  | 8553(2)  | 9107(1)  | 26(1) |
| C(32) | 6146(4)  | 9726(2)  | 8563(1)  | 17(1) |
| C(34) | 5447(4)  | 10415(2) | 8347(1)  | 16(1) |
| C(35) | 7495(4)  | 10653(2) | 8173(1)  | 16(1) |
| C(37) | 8721(4)  | 11531(2) | 7774(1)  | 19(1) |
| C(38) | 5612(4)  | 10463(2) | 7720(1)  | 15(1) |
| C(39) | 3693(4)  | 10032(1) | 7882(1)  | 14(1) |
| C(40) | 3621(4)  | 10043(2) | 8164(1)  | 16(1) |
| C(41) | 2025(4)  | 9565(2)  | 7707(1)  | 16(1) |
| C(42) | 468(4)   | 8956(2)  | 7795(1)  | 17(1) |

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|       |           |          |          |        |
|-------|-----------|----------|----------|--------|
| C(43) | -881(4)   | 8522(2)  | 7598(1)  | 20(1)  |
| C(44) | -725(4)   | 8648(2)  | 7308(1)  | 19(1)  |
| C(45) | 857(4)    | 9255(2)  | 7217(1)  | 18(1)  |
| C(47) | 3170(5)   | 10164(2) | 6990(1)  | 21(1)  |
| C(48) | 3557(4)   | 10296(2) | 7271(1)  | 18(1)  |
| C(49) | 4902(5)   | 10914(2) | 7444(1)  | 19(1)  |
| C(50) | 2141(4)   | 9713(2)  | 7415(1)  | 17(1)  |
| C(52) | 871(4)    | 3803(2)  | 8758(1)  | 16(1)  |
| C(54) | 1905(4)   | 3078(2)  | 8583(1)  | 16(1)  |
| C(56) | 3838(4)   | 3469(2)  | 8414(1)  | 17(1)  |
| C(58) | 5120(4)   | 2742(2)  | 8266(1)  | 16(1)  |
| N(4)  | 3248(4)   | 6958(2)  | 8526(1)  | 24(1)  |
| N(11) | 4583(4)   | 4782(1)  | 9253(1)  | 18(1)  |
| N(21) | -1628(5)  | 5475(2)  | 10223(1) | 36(1)  |
| N(29) | 4943(4)   | 9659(1)  | 8804(1)  | 20(1)  |
| N(36) | 6767(3)   | 11102(1) | 7910(1)  | 14(1)  |
| N(46) | 1536(4)   | 9539(1)  | 6957(1)  | 20(1)  |
| O(1)  | 2049(3)   | 7167(1)  | 7957(1)  | 23(1)  |
| O(8)  | 6414(3)   | 6921(1)  | 8789(1)  | 24(1)  |
| O(26) | 1789(3)   | 9168(1)  | 9221(1)  | 24(1)  |
| O(33) | 7685(3)   | 9226(1)  | 8507(1)  | 21(1)  |
| O(51) | 1287(3)   | 3790(1)  | 9021(1)  | 21(1)  |
| O(53) | -234(3)   | 4364(1)  | 8630(1)  | 20(1)  |
| O(55) | 2678(3)   | 2390(1)  | 8759(1)  | 22(1)  |
| O(57) | 5193(3)   | 3952(1)  | 8601(1)  | 23(1)  |
| O(59) | 4066(3)   | 2344(1)  | 8076(1)  | 19(1)  |
| O(60) | 7074(3)   | 2589(1)  | 8347(1)  | 21(1)  |
| O(61) | 8262(4)   | 1668(2)  | 8840(1)  | 37(1)  |
| O(62) | 6805(7)   | 1807(3)  | 9398(1)  | 89(1)  |
| C(63) | 8661(13)  | 2189(5)  | 9564(2)  | 110(2) |
| C(64) | 10528(16) | 1603(6)  | 9591(2)  | 142(4) |

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Table 4. Bond lengths [Å] and angles [°] for TCD173.

|              |          |              |          |
|--------------|----------|--------------|----------|
| C(2)-O(1)    | 1.429(3) | C(17)-C(18)  | 1.413(4) |
| C(2)-C(3)    | 1.512(4) | C(17)-H(17)  | 0.9500   |
| C(2)-H(2A)   | 0.9900   | C(18)-C(19)  | 1.395(5) |
| C(2)-H(2B)   | 0.9900   | C(18)-H(18)  | 0.9500   |
| C(3)-N(4)    | 1.475(4) | C(19)-C(20)  | 1.393(5) |
| C(3)-H(3A)   | 0.9900   | C(19)-H(19)  | 0.9500   |
| C(3)-H(3B)   | 0.9900   | C(20)-N(21)  | 1.373(4) |
| C(5)-N(4)    | 1.466(3) | C(20)-C(25)  | 1.406(4) |
| C(5)-C(6)    | 1.506(4) | C(22)-C(23)  | 1.368(4) |
| C(5)-H(5A)   | 0.9900   | C(22)-N(21)  | 1.389(4) |
| C(5)-H(5B)   | 0.9900   | C(22)-H(22)  | 0.9500   |
| C(6)-O(1)    | 1.426(3) | C(23)-C(25)  | 1.407(4) |
| C(6)-H(6A)   | 0.9900   | C(23)-C(24)  | 1.505(4) |
| C(6)-H(6B)   | 0.9900   | C(24)-H(24A) | 0.9900   |
| C(7)-O(8)    | 1.230(3) | C(24)-H(24B) | 0.9900   |
| C(7)-N(4)    | 1.351(4) | C(27)-O(26)  | 1.436(3) |
| C(7)-C(9)    | 1.529(3) | C(27)-C(28)  | 1.515(4) |
| C(9)-C(15)   | 1.505(3) | C(27)-H(27A) | 0.9900   |
| C(9)-C(10)   | 1.517(4) | C(27)-H(27B) | 0.9900   |
| C(9)-H(9)    | 1.0000   | C(28)-N(29)  | 1.463(3) |
| C(10)-N(11)  | 1.498(3) | C(28)-H(28A) | 0.9900   |
| C(10)-H(10A) | 0.9900   | C(28)-H(28B) | 0.9900   |
| C(10)-H(10B) | 0.9900   | C(30)-N(29)  | 1.469(3) |
| C(12)-N(11)  | 1.490(3) | C(30)-C(31)  | 1.514(4) |
| C(12)-H(12A) | 0.9800   | C(30)-H(30A) | 0.9900   |
| C(12)-H(12B) | 0.9800   | C(30)-H(30B) | 0.9900   |
| C(12)-H(12C) | 0.9800   | C(31)-O(26)  | 1.434(4) |
| C(13)-N(11)  | 1.512(3) | C(31)-H(31A) | 0.9900   |
| C(13)-C(14)  | 1.535(4) | C(31)-H(31B) | 0.9900   |
| C(13)-C(24)  | 1.544(4) | C(32)-O(33)  | 1.231(3) |
| C(13)-H(13)  | 1.0000   | C(32)-N(29)  | 1.354(3) |
| C(14)-C(15)  | 1.338(4) | C(32)-C(34)  | 1.538(3) |
| C(14)-C(16)  | 1.475(4) | C(34)-C(40)  | 1.508(3) |
| C(15)-H(15)  | 0.9500   | C(34)-C(35)  | 1.521(3) |
| C(16)-C(17)  | 1.389(4) | C(34)-H(34)  | 1.0000   |
| C(16)-C(25)  | 1.400(4) | C(35)-N(36)  | 1.494(3) |

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|              |          |                  |          |
|--------------|----------|------------------|----------|
| C(35)-H(35A) | 0.9900   | C(56)-H(56)      | 1.0000   |
| C(35)-H(35B) | 0.9900   | C(58)-O(60)      | 1.250(3) |
| C(37)-N(36)  | 1.488(3) | C(58)-O(59)      | 1.262(3) |
| C(37)-H(37A) | 0.9800   | N(11)-H(11)      | 0.89(2)  |
| C(37)-H(37B) | 0.9800   | N(21)-H(21)      | 0.90(2)  |
| C(37)-H(37C) | 0.9800   | N(36)-H(36)      | 0.90(2)  |
| C(38)-N(36)  | 1.505(3) | N(46)-H(46)      | 0.89(2)  |
| C(38)-C(39)  | 1.531(3) | O(55)-H(55)      | 0.84(3)  |
| C(38)-C(49)  | 1.544(3) | O(57)-H(57)      | 0.85(3)  |
| C(38)-H(38)  | 1.0000   | O(61)-H(61A)     | 0.94(3)  |
| C(39)-C(40)  | 1.341(3) | O(61)-H(61B)     | 0.90(3)  |
| C(39)-C(41)  | 1.485(3) | O(62)-C(63)      | 1.484(8) |
| C(40)-H(40)  | 0.9500   | O(62)-H(62)      | 0.86(3)  |
| C(41)-C(42)  | 1.389(4) | C(63)-C(64)      | 1.443(8) |
| C(41)-C(50)  | 1.405(3) | C(63)-H(63A)     | 0.9900   |
| C(42)-C(43)  | 1.405(4) | C(63)-H(63B)     | 0.9900   |
| C(42)-H(42)  | 0.9500   | C(64)-H(64A)     | 0.9800   |
| C(43)-C(44)  | 1.394(4) | C(64)-H(64B)     | 0.9800   |
| C(43)-H(43)  | 0.9500   | C(64)-H(64C)     | 0.9800   |
| C(44)-C(45)  | 1.401(4) |                  |          |
| C(44)-H(44)  | 0.9500   | O(1)-C(2)-C(3)   | 111.9(2) |
| C(45)-N(46)  | 1.373(3) | O(1)-C(2)-H(2A)  | 109.2    |
| C(45)-C(50)  | 1.404(4) | C(3)-C(2)-H(2A)  | 109.2    |
| C(47)-C(48)  | 1.368(4) | O(1)-C(2)-H(2B)  | 109.2    |
| C(47)-N(46)  | 1.383(4) | C(3)-C(2)-H(2B)  | 109.2    |
| C(47)-H(47)  | 0.9500   | H(2A)-C(2)-H(2B) | 107.9    |
| C(48)-C(50)  | 1.413(4) | N(4)-C(3)-C(2)   | 108.2(2) |
| C(48)-C(49)  | 1.497(3) | N(4)-C(3)-H(3A)  | 110.0    |
| C(49)-H(49A) | 0.9900   | C(2)-C(3)-H(3A)  | 110.0    |
| C(49)-H(49B) | 0.9900   | N(4)-C(3)-H(3B)  | 110.0    |
| C(52)-O(53)  | 1.249(3) | C(2)-C(3)-H(3B)  | 110.0    |
| C(52)-O(51)  | 1.270(3) | H(3A)-C(3)-H(3B) | 108.4    |
| C(52)-C(54)  | 1.531(3) | N(4)-C(5)-C(6)   | 109.2(2) |
| C(54)-O(55)  | 1.432(3) | N(4)-C(5)-H(5A)  | 109.8    |
| C(54)-C(56)  | 1.529(3) | C(6)-C(5)-H(5A)  | 109.8    |
| C(54)-H(54)  | 1.0000   | N(4)-C(5)-H(5B)  | 109.8    |
| C(56)-O(57)  | 1.416(3) | C(6)-C(5)-H(5B)  | 109.8    |
| C(56)-C(58)  | 1.531(3) | H(5A)-C(5)-H(5B) | 108.3    |

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|                     |            |                     |          |
|---------------------|------------|---------------------|----------|
| O(1)-C(6)-C(5)      | 111.7(2)   | C(9)-C(15)-H(15)    | 118.1    |
| O(1)-C(6)-H(6A)     | 109.3      | C(17)-C(16)-C(25)   | 116.5(2) |
| C(5)-C(6)-H(6A)     | 109.3      | C(17)-C(16)-C(14)   | 127.6(2) |
| O(1)-C(6)-H(6B)     | 109.3      | C(25)-C(16)-C(14)   | 115.9(2) |
| C(5)-C(6)-H(6B)     | 109.3      | C(16)-C(17)-C(18)   | 120.6(3) |
| H(6A)-C(6)-H(6B)    | 107.9      | C(16)-C(17)-H(17)   | 119.7    |
| O(8)-C(7)-N(4)      | 122.4(2)   | C(18)-C(17)-H(17)   | 119.7    |
| O(8)-C(7)-C(9)      | 120.6(2)   | C(19)-C(18)-C(17)   | 122.7(3) |
| N(4)-C(7)-C(9)      | 116.9(2)   | C(19)-C(18)-H(18)   | 118.7    |
| C(15)-C(9)-C(10)    | 110.9(2)   | C(17)-C(18)-H(18)   | 118.7    |
| C(15)-C(9)-C(7)     | 109.8(2)   | C(20)-C(19)-C(18)   | 116.9(3) |
| C(10)-C(9)-C(7)     | 108.7(2)   | C(20)-C(19)-H(19)   | 121.6    |
| C(15)-C(9)-H(9)     | 109.1      | C(18)-C(19)-H(19)   | 121.6    |
| C(10)-C(9)-H(9)     | 109.1      | N(21)-C(20)-C(19)   | 133.8(3) |
| C(7)-C(9)-H(9)      | 109.1      | N(21)-C(20)-C(25)   | 105.9(3) |
| N(11)-C(10)-C(9)    | 108.4(2)   | C(19)-C(20)-C(25)   | 120.3(3) |
| N(11)-C(10)-H(10A)  | 110.0      | C(23)-C(22)-N(21)   | 108.7(3) |
| C(9)-C(10)-H(10A)   | 110.0      | C(23)-C(22)-H(22)   | 125.7    |
| N(11)-C(10)-H(10B)  | 110.0      | N(21)-C(22)-H(22)   | 125.7    |
| C(9)-C(10)-H(10B)   | 110.0      | C(22)-C(23)-C(25)   | 106.7(3) |
| H(10A)-C(10)-H(10B) | 108.4      | C(22)-C(23)-C(24)   | 134.1(3) |
| N(11)-C(12)-H(12A)  | 109.5      | C(25)-C(23)-C(24)   | 119.1(2) |
| N(11)-C(12)-H(12B)  | 109.5      | C(23)-C(24)-C(13)   | 109.4(2) |
| H(12A)-C(12)-H(12B) | 109.5      | C(23)-C(24)-H(24A)  | 109.8    |
| N(11)-C(12)-H(12C)  | 109.5      | C(13)-C(24)-H(24A)  | 109.8    |
| H(12A)-C(12)-H(12C) | 109.5      | C(23)-C(24)-H(24B)  | 109.8    |
| H(12B)-C(12)-H(12C) | 109.5      | C(13)-C(24)-H(24B)  | 109.8    |
| N(11)-C(13)-C(14)   | 108.25(19) | H(24A)-C(24)-H(24B) | 108.2    |
| N(11)-C(13)-C(24)   | 108.9(2)   | C(16)-C(25)-C(20)   | 123.1(3) |
| C(14)-C(13)-C(24)   | 115.6(2)   | C(16)-C(25)-C(23)   | 128.0(3) |
| N(11)-C(13)-H(13)   | 107.9      | C(20)-C(25)-C(23)   | 108.9(3) |
| C(14)-C(13)-H(13)   | 107.9      | O(26)-C(27)-C(28)   | 110.9(2) |
| C(24)-C(13)-H(13)   | 107.9      | O(26)-C(27)-H(27A)  | 109.5    |
| C(15)-C(14)-C(16)   | 122.1(2)   | C(28)-C(27)-H(27A)  | 109.5    |
| C(15)-C(14)-C(13)   | 122.1(2)   | O(26)-C(27)-H(27B)  | 109.5    |
| C(16)-C(14)-C(13)   | 115.8(2)   | C(28)-C(27)-H(27B)  | 109.5    |
| C(14)-C(15)-C(9)    | 123.7(2)   | H(27A)-C(27)-H(27B) | 108.1    |
| C(14)-C(15)-H(15)   | 118.1      | N(29)-C(28)-C(27)   | 110.0(2) |

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|                     |            |                    |            |
|---------------------|------------|--------------------|------------|
| N(29)-C(28)-H(28A)  | 109.7      | N(36)-C(38)-C(39)  | 109.23(19) |
| C(27)-C(28)-H(28A)  | 109.7      | N(36)-C(38)-C(49)  | 109.59(19) |
| N(29)-C(28)-H(28B)  | 109.7      | C(39)-C(38)-C(49)  | 114.7(2)   |
| C(27)-C(28)-H(28B)  | 109.7      | N(36)-C(38)-H(38)  | 107.7      |
| H(28A)-C(28)-H(28B) | 108.2      | C(39)-C(38)-H(38)  | 107.7      |
| N(29)-C(30)-C(31)   | 109.4(2)   | C(49)-C(38)-H(38)  | 107.7      |
| N(29)-C(30)-H(30A)  | 109.8      | C(40)-C(39)-C(41)  | 123.0(2)   |
| C(31)-C(30)-H(30A)  | 109.8      | C(40)-C(39)-C(38)  | 121.3(2)   |
| N(29)-C(30)-H(30B)  | 109.8      | C(41)-C(39)-C(38)  | 115.7(2)   |
| C(31)-C(30)-H(30B)  | 109.8      | C(39)-C(40)-C(34)  | 124.0(2)   |
| H(30A)-C(30)-H(30B) | 108.2      | C(39)-C(40)-H(40)  | 118.0      |
| O(26)-C(31)-C(30)   | 111.6(2)   | C(34)-C(40)-H(40)  | 118.0      |
| O(26)-C(31)-H(31A)  | 109.3      | C(42)-C(41)-C(50)  | 116.2(2)   |
| C(30)-C(31)-H(31A)  | 109.3      | C(42)-C(41)-C(39)  | 127.6(2)   |
| O(26)-C(31)-H(31B)  | 109.3      | C(50)-C(41)-C(39)  | 116.0(2)   |
| C(30)-C(31)-H(31B)  | 109.3      | C(41)-C(42)-C(43)  | 120.5(2)   |
| H(31A)-C(31)-H(31B) | 108.0      | C(41)-C(42)-H(42)  | 119.8      |
| O(33)-C(32)-N(29)   | 122.0(2)   | C(43)-C(42)-H(42)  | 119.8      |
| O(33)-C(32)-C(34)   | 119.6(2)   | C(44)-C(43)-C(42)  | 123.6(2)   |
| N(29)-C(32)-C(34)   | 118.2(2)   | C(44)-C(43)-H(43)  | 118.2      |
| C(40)-C(34)-C(35)   | 111.0(2)   | C(42)-C(43)-H(43)  | 118.2      |
| C(40)-C(34)-C(32)   | 108.35(19) | C(43)-C(44)-C(45)  | 116.2(2)   |
| C(35)-C(34)-C(32)   | 108.3(2)   | C(43)-C(44)-H(44)  | 121.9      |
| C(40)-C(34)-H(34)   | 109.7      | C(45)-C(44)-H(44)  | 121.9      |
| C(35)-C(34)-H(34)   | 109.7      | N(46)-C(45)-C(44)  | 133.6(2)   |
| C(32)-C(34)-H(34)   | 109.7      | N(46)-C(45)-C(50)  | 106.2(2)   |
| N(36)-C(35)-C(34)   | 109.6(2)   | C(44)-C(45)-C(50)  | 120.2(2)   |
| N(36)-C(35)-H(35A)  | 109.8      | C(48)-C(47)-N(46)  | 109.5(2)   |
| C(34)-C(35)-H(35A)  | 109.8      | C(48)-C(47)-H(47)  | 125.3      |
| N(36)-C(35)-H(35B)  | 109.8      | N(46)-C(47)-H(47)  | 125.3      |
| C(34)-C(35)-H(35B)  | 109.8      | C(47)-C(48)-C(50)  | 106.0(2)   |
| H(35A)-C(35)-H(35B) | 108.2      | C(47)-C(48)-C(49)  | 136.2(2)   |
| N(36)-C(37)-H(37A)  | 109.5      | C(50)-C(48)-C(49)  | 117.6(2)   |
| N(36)-C(37)-H(37B)  | 109.5      | C(48)-C(49)-C(38)  | 108.9(2)   |
| H(37A)-C(37)-H(37B) | 109.5      | C(48)-C(49)-H(49A) | 109.9      |
| N(36)-C(37)-H(37C)  | 109.5      | C(38)-C(49)-H(49A) | 109.9      |
| H(37A)-C(37)-H(37C) | 109.5      | C(48)-C(49)-H(49B) | 109.9      |
| H(37B)-C(37)-H(37C) | 109.5      | C(38)-C(49)-H(49B) | 109.9      |

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|                     |            |                     |            |
|---------------------|------------|---------------------|------------|
| H(49A)-C(49)-H(49B) | 108.3      | C(20)-N(21)-H(21)   | 126(3)     |
| C(45)-C(50)-C(41)   | 123.3(2)   | C(22)-N(21)-H(21)   | 124(3)     |
| C(45)-C(50)-C(48)   | 109.0(2)   | C(32)-N(29)-C(28)   | 126.7(2)   |
| C(41)-C(50)-C(48)   | 127.7(2)   | C(32)-N(29)-C(30)   | 120.0(2)   |
| O(53)-C(52)-O(51)   | 126.3(2)   | C(28)-N(29)-C(30)   | 112.7(2)   |
| O(53)-C(52)-C(54)   | 117.3(2)   | C(37)-N(36)-C(35)   | 110.07(19) |
| O(51)-C(52)-C(54)   | 116.4(2)   | C(37)-N(36)-C(38)   | 113.16(18) |
| O(55)-C(54)-C(56)   | 111.1(2)   | C(35)-N(36)-C(38)   | 109.05(18) |
| O(55)-C(54)-C(52)   | 110.90(19) | C(37)-N(36)-H(36)   | 106(3)     |
| C(56)-C(54)-C(52)   | 107.3(2)   | C(35)-N(36)-H(36)   | 106(3)     |
| O(55)-C(54)-H(54)   | 109.2      | C(38)-N(36)-H(36)   | 112(3)     |
| C(56)-C(54)-H(54)   | 109.2      | C(45)-N(46)-C(47)   | 109.3(2)   |
| C(52)-C(54)-H(54)   | 109.2      | C(45)-N(46)-H(46)   | 127(3)     |
| O(57)-C(56)-C(54)   | 108.1(2)   | C(47)-N(46)-H(46)   | 124(3)     |
| O(57)-C(56)-C(58)   | 113.1(2)   | C(6)-O(1)-C(2)      | 111.96(19) |
| C(54)-C(56)-C(58)   | 108.9(2)   | C(31)-O(26)-C(27)   | 109.79(19) |
| O(57)-C(56)-H(56)   | 108.9      | C(54)-O(55)-H(55)   | 101(2)     |
| C(54)-C(56)-H(56)   | 108.9      | C(56)-O(57)-H(57)   | 108(2)     |
| C(58)-C(56)-H(56)   | 108.9      | H(61A)-O(61)-H(61B) | 106.7      |
| O(60)-C(58)-O(59)   | 126.3(2)   | C(63)-O(62)-H(62)   | 110.9      |
| O(60)-C(58)-C(56)   | 117.7(2)   | C(64)-C(63)-O(62)   | 111.8(6)   |
| O(59)-C(58)-C(56)   | 116.1(2)   | C(64)-C(63)-H(63A)  | 109.3      |
| C(7)-N(4)-C(5)      | 120.4(2)   | O(62)-C(63)-H(63A)  | 109.3      |
| C(7)-N(4)-C(3)      | 127.5(2)   | C(64)-C(63)-H(63B)  | 109.3      |
| C(5)-N(4)-C(3)      | 112.0(2)   | O(62)-C(63)-H(63B)  | 109.3      |
| C(12)-N(11)-C(10)   | 110.5(2)   | H(63A)-C(63)-H(63B) | 107.9      |
| C(12)-N(11)-C(13)   | 113.09(19) | C(63)-C(64)-H(64A)  | 109.5      |
| C(10)-N(11)-C(13)   | 109.92(19) | C(63)-C(64)-H(64B)  | 109.5      |
| C(12)-N(11)-H(11)   | 106(2)     | H(64A)-C(64)-H(64B) | 109.5      |
| C(10)-N(11)-H(11)   | 108(2)     | C(63)-C(64)-H(64C)  | 109.5      |
| C(13)-N(11)-H(11)   | 109(2)     | H(64A)-C(64)-H(64C) | 109.5      |
| C(20)-N(21)-C(22)   | 109.7(2)   | H(64B)-C(64)-H(64C) | 109.5      |

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Table 5. Anisotropic displacement parameters ( $\text{\AA}^2 \times 10^3$ ) for TCD173. The anisotropic displacement factor exponent takes the form:  $-2\pi^2 [ h^2 a^{*2} U^{11} + \dots + 2 h k a^* b^* U^{12} ]$

|       | $U^{11}$ | $U^{22}$ | $U^{33}$ | $U^{23}$ | $U^{13}$ | $U^{12}$ |
|-------|----------|----------|----------|----------|----------|----------|
| C(2)  | 21(1)    | 24(1)    | 25(1)    | 4(1)     | -3(1)    | -3(1)    |
| C(3)  | 21(1)    | 28(1)    | 24(1)    | 3(1)     | -2(1)    | -4(1)    |
| C(5)  | 30(2)    | 20(1)    | 24(1)    | 3(1)     | 1(1)     | -2(1)    |
| C(6)  | 25(1)    | 20(1)    | 22(1)    | 3(1)     | 0(1)     | -3(1)    |
| C(7)  | 21(1)    | 18(1)    | 18(1)    | -2(1)    | 3(1)     | -1(1)    |
| C(9)  | 19(1)    | 18(1)    | 16(1)    | -2(1)    | 0(1)     | -2(1)    |
| C(10) | 17(1)    | 19(1)    | 17(1)    | -2(1)    | 3(1)     | -3(1)    |
| C(12) | 23(1)    | 25(1)    | 26(1)    | 2(1)     | -1(1)    | 6(1)     |
| C(13) | 19(1)    | 25(1)    | 14(1)    | -4(1)    | -1(1)    | 2(1)     |
| C(14) | 18(1)    | 20(1)    | 18(1)    | -3(1)    | -2(1)    | -2(1)    |
| C(15) | 17(1)    | 18(1)    | 18(1)    | -2(1)    | 0(1)     | -1(1)    |
| C(16) | 17(1)    | 27(1)    | 17(1)    | -7(1)    | -1(1)    | -2(1)    |
| C(17) | 21(1)    | 30(1)    | 21(1)    | -8(1)    | -2(1)    | 1(1)     |
| C(18) | 20(1)    | 35(2)    | 33(2)    | -15(1)   | -2(1)    | 3(1)     |
| C(19) | 22(1)    | 47(2)    | 26(1)    | -18(1)   | 4(1)     | 0(1)     |
| C(20) | 23(1)    | 42(2)    | 19(1)    | -11(1)   | 4(1)     | -5(1)    |
| C(22) | 34(2)    | 49(2)    | 21(1)    | 3(1)     | 4(1)     | 2(2)     |
| C(23) | 24(1)    | 40(2)    | 17(1)    | -1(1)    | 1(1)     | -1(1)    |
| C(24) | 26(1)    | 32(1)    | 18(1)    | 5(1)     | 3(1)     | 4(1)     |
| C(25) | 20(1)    | 36(2)    | 15(1)    | -6(1)    | 0(1)     | -3(1)    |
| C(27) | 24(1)    | 32(1)    | 19(1)    | 4(1)     | -1(1)    | 3(1)     |
| C(28) | 27(2)    | 24(1)    | 22(1)    | 2(1)     | 4(1)     | 4(1)     |
| C(30) | 25(1)    | 25(1)    | 17(1)    | 5(1)     | -2(1)    | 2(1)     |
| C(31) | 34(2)    | 24(1)    | 20(1)    | 4(1)     | -1(1)    | -1(1)    |
| C(32) | 18(1)    | 17(1)    | 15(1)    | -1(1)    | -5(1)    | -3(1)    |
| C(34) | 16(1)    | 17(1)    | 15(1)    | -1(1)    | -1(1)    | 0(1)     |
| C(35) | 15(1)    | 18(1)    | 15(1)    | 1(1)     | -2(1)    | 0(1)     |
| C(37) | 18(1)    | 19(1)    | 21(1)    | 0(1)     | 2(1)     | -3(1)    |
| C(38) | 15(1)    | 16(1)    | 15(1)    | -3(1)    | 0(1)     | 0(1)     |
| C(39) | 13(1)    | 11(1)    | 18(1)    | 1(1)     | -2(1)    | 2(1)     |
| C(40) | 15(1)    | 16(1)    | 17(1)    | 1(1)     | -1(1)    | 0(1)     |
| C(41) | 16(1)    | 14(1)    | 17(1)    | -2(1)    | -2(1)    | 2(1)     |
| C(42) | 17(1)    | 17(1)    | 18(1)    | 0(1)     | -1(1)    | 2(1)     |

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|       |        |        |        |        |        |        |
|-------|--------|--------|--------|--------|--------|--------|
| C(43) | 16(1)  | 16(1)  | 27(1)  | -1(1)  | -2(1)  | 1(1)   |
| C(44) | 17(1)  | 17(1)  | 22(1)  | -4(1)  | -5(1)  | 1(1)   |
| C(45) | 19(1)  | 16(1)  | 19(1)  | -4(1)  | -4(1)  | 3(1)   |
| C(47) | 22(1)  | 22(1)  | 18(1)  | 0(1)   | -2(1)  | -1(1)  |
| C(48) | 18(1)  | 19(1)  | 18(1)  | 0(1)   | 0(1)   | 2(1)   |
| C(49) | 20(1)  | 20(1)  | 17(1)  | 1(1)   | -1(1)  | -3(1)  |
| C(50) | 17(1)  | 15(1)  | 18(1)  | -1(1)  | -1(1)  | 3(1)   |
| C(52) | 13(1)  | 19(1)  | 17(1)  | -2(1)  | 1(1)   | -4(1)  |
| C(54) | 15(1)  | 16(1)  | 19(1)  | -1(1)  | -2(1)  | 1(1)   |
| C(56) | 15(1)  | 16(1)  | 20(1)  | -5(1)  | -1(1)  | -1(1)  |
| C(58) | 16(1)  | 16(1)  | 16(1)  | 0(1)   | 1(1)   | 1(1)   |
| N(4)  | 27(1)  | 25(1)  | 19(1)  | 4(1)   | -1(1)  | -9(1)  |
| N(11) | 16(1)  | 20(1)  | 16(1)  | 0(1)   | 0(1)   | -1(1)  |
| N(21) | 35(2)  | 57(2)  | 16(1)  | -3(1)  | 8(1)   | 0(1)   |
| N(29) | 22(1)  | 19(1)  | 18(1)  | 3(1)   | 0(1)   | 3(1)   |
| N(36) | 14(1)  | 14(1)  | 16(1)  | -1(1)  | 0(1)   | -1(1)  |
| N(46) | 22(1)  | 23(1)  | 16(1)  | -3(1)  | -4(1)  | -1(1)  |
| O(1)  | 23(1)  | 23(1)  | 21(1)  | 4(1)   | -1(1)  | -1(1)  |
| O(8)  | 22(1)  | 22(1)  | 28(1)  | 1(1)   | 0(1)   | -5(1)  |
| O(26) | 26(1)  | 31(1)  | 14(1)  | 3(1)   | 1(1)   | -1(1)  |
| O(33) | 21(1)  | 21(1)  | 21(1)  | 1(1)   | -1(1)  | 4(1)   |
| O(51) | 21(1)  | 27(1)  | 16(1)  | -2(1)  | 1(1)   | -3(1)  |
| O(53) | 18(1)  | 21(1)  | 21(1)  | -2(1)  | -1(1)  | 2(1)   |
| O(55) | 25(1)  | 19(1)  | 22(1)  | 3(1)   | -1(1)  | 2(1)   |
| O(57) | 14(1)  | 26(1)  | 30(1)  | -13(1) | 1(1)   | -4(1)  |
| O(59) | 17(1)  | 20(1)  | 20(1)  | -4(1)  | -4(1)  | 3(1)   |
| O(60) | 14(1)  | 24(1)  | 24(1)  | -5(1)  | -3(1)  | 2(1)   |
| O(61) | 34(1)  | 41(1)  | 36(1)  | 13(1)  | -8(1)  | -9(1)  |
| O(62) | 91(3)  | 112(3) | 66(2)  | 13(2)  | 12(2)  | -28(3) |
| C(63) | 121(6) | 97(5)  | 112(5) | -1(4)  | -34(5) | 29(5)  |
| C(64) | 159(8) | 160(7) | 107(5) | -21(5) | -19(5) | 94(7)  |

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Table 6. Hydrogen coordinates ( $\times 10^4$ ) and isotropic displacement parameters ( $\text{\AA}^2 \times 10^{-3}$ ) for TCD173.

|        | x     | y     | z     | U(eq) |
|--------|-------|-------|-------|-------|
| H(2A)  | 2419  | 5956  | 8087  | 28    |
| H(2B)  | -139  | 6231  | 8038  | 28    |
| H(3A)  | -111  | 7056  | 8453  | 29    |
| H(3B)  | 677   | 6092  | 8527  | 29    |
| H(5A)  | 5464  | 7913  | 8438  | 30    |
| H(5B)  | 2897  | 8198  | 8397  | 30    |
| H(6A)  | 4528  | 8022  | 7953  | 27    |
| H(6B)  | 5316  | 7066  | 8032  | 27    |
| H(9)   | 3056  | 5404  | 8747  | 21    |
| H(10A) | 6597  | 5116  | 8918  | 21    |
| H(10B) | 6426  | 5861  | 9151  | 21    |
| H(12A) | 7244  | 3973  | 9199  | 38    |
| H(12B) | 5634  | 3692  | 9453  | 38    |
| H(12C) | 7319  | 4487  | 9491  | 38    |
| H(13)  | 4518  | 5607  | 9591  | 23    |
| H(15)  | 653   | 6463  | 9006  | 21    |
| H(17)  | -1237 | 7275  | 9331  | 29    |
| H(18)  | -3832 | 7707  | 9673  | 35    |
| H(19)  | -4238 | 6966  | 10101 | 38    |
| H(22)  | 440   | 4396  | 10269 | 41    |
| H(24A) | 1589  | 4135  | 9587  | 30    |
| H(24B) | 3601  | 4318  | 9803  | 30    |
| H(27A) | 116   | 10206 | 9088  | 30    |
| H(27B) | 494   | 9494  | 8848  | 30    |
| H(28A) | 2849  | 10676 | 8757  | 29    |
| H(28B) | 3913  | 10609 | 9066  | 29    |
| H(30A) | 6274  | 9239  | 9173  | 27    |
| H(30B) | 6486  | 8550  | 8923  | 27    |
| H(31A) | 2635  | 8252  | 8946  | 31    |
| H(31B) | 3701  | 8117  | 9253  | 31    |
| H(34)  | 4874  | 10937 | 8448  | 19    |
| H(35A) | 8491  | 11035 | 8283  | 19    |



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|        |           |           |          |        |
|--------|-----------|-----------|----------|--------|
| H(35B) | 8345      | 10125     | 8124     | 19     |
| H(37A) | 9548      | 11864     | 7915     | 29     |
| H(37B) | 8190      | 11918     | 7625     | 29     |
| H(37C) | 9712      | 11093     | 7692     | 29     |
| H(38)  | 6726      | 10005     | 7671     | 18     |
| H(40)  | 2342      | 9803      | 8254     | 19     |
| H(42)  | 313       | 8832      | 7990     | 21     |
| H(43)  | -1957     | 8122      | 7666     | 24     |
| H(44)  | -1640     | 8338      | 7179     | 22     |
| H(47)  | 3910      | 10456     | 6841     | 25     |
| H(49A) | 3994      | 11431     | 7488     | 23     |
| H(49B) | 6249      | 11099     | 7338     | 23     |
| H(54)  | 749       | 2849      | 8449     | 20     |
| H(56)  | 3206      | 3866      | 8268     | 20     |
| H(11)  | 3590(50)  | 4460(20)  | 9160(7)  | 39(10) |
| H(21)  | -2380(70) | 5490(30)  | 10387(6) | 58(12) |
| H(36)  | 5850(60)  | 11530(20) | 7965(8)  | 55(12) |
| H(46)  | 1000(60)  | 9370(20)  | 6790(6)  | 42(10) |
| H(55)  | 2990(60)  | 2660(20)  | 8908(6)  | 33     |
| H(57)  | 6550(50)  | 3860(20)  | 8560(7)  | 35     |
| H(61A) | 7610(20)  | 1970(11)  | 8689(5)  | 59(13) |
| H(61B) | 9710(50)  | 1838(6)   | 8850(1)  | 73(15) |
| H(62)  | 7108(13)  | 1816(3)   | 9222(6)  | 134    |
| H(63A) | 9175      | 2725      | 9472     | 132    |
| H(63B) | 8101      | 2341      | 9754     | 132    |
| H(64A) | 9968      | 1021      | 9630     | 213    |
| H(64B) | 11494     | 1791      | 9747     | 213    |
| H(64C) | 11392     | 1600      | 9416     | 213    |

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Table 7. Torsion angles [°] for TCD173.

|                         |            |                         |            |
|-------------------------|------------|-------------------------|------------|
| O(1)-C(2)-C(3)-N(4)     | -56.0(3)   | N(21)-C(20)-C(25)-C(16) | -178.5(3)  |
| N(4)-C(5)-C(6)-O(1)     | 55.7(3)    | C(19)-C(20)-C(25)-C(16) | 1.6(4)     |
| O(8)-C(7)-C(9)-C(15)    | 105.9(3)   | N(21)-C(20)-C(25)-C(23) | 1.1(3)     |
| N(4)-C(7)-C(9)-C(15)    | -71.5(3)   | C(19)-C(20)-C(25)-C(23) | -178.8(3)  |
| O(8)-C(7)-C(9)-C(10)    | -15.6(3)   | C(22)-C(23)-C(25)-C(16) | 178.0(3)   |
| N(4)-C(7)-C(9)-C(10)    | 167.0(2)   | C(24)-C(23)-C(25)-C(16) | -4.5(5)    |
| C(15)-C(9)-C(10)-N(11)  | 46.5(3)    | C(22)-C(23)-C(25)-C(20) | -1.5(3)    |
| C(7)-C(9)-C(10)-N(11)   | 167.32(19) | C(24)-C(23)-C(25)-C(20) | 175.9(3)   |
| N(11)-C(13)-C(14)-C(15) | -16.7(3)   | O(26)-C(27)-C(28)-N(29) | 56.5(3)    |
| C(24)-C(13)-C(14)-C(15) | -139.1(3)  | N(29)-C(30)-C(31)-O(26) | -56.2(3)   |
| N(11)-C(13)-C(14)-C(16) | 164.7(2)   | O(33)-C(32)-C(34)-C(40) | 91.1(3)    |
| C(24)-C(13)-C(14)-C(16) | 42.3(3)    | N(29)-C(32)-C(34)-C(40) | -85.0(3)   |
| C(16)-C(14)-C(15)-C(9)  | 174.0(2)   | O(33)-C(32)-C(34)-C(35) | -29.3(3)   |
| C(13)-C(14)-C(15)-C(9)  | -4.5(4)    | N(29)-C(32)-C(34)-C(35) | 154.6(2)   |
| C(10)-C(9)-C(15)-C(14)  | -10.8(3)   | C(40)-C(34)-C(35)-N(36) | 43.9(3)    |
| C(7)-C(9)-C(15)-C(14)   | -130.9(3)  | C(32)-C(34)-C(35)-N(36) | 162.72(18) |
| C(15)-C(14)-C(16)-C(17) | -16.3(4)   | N(36)-C(38)-C(39)-C(40) | -19.0(3)   |
| C(13)-C(14)-C(16)-C(17) | 162.2(3)   | C(49)-C(38)-C(39)-C(40) | -142.5(2)  |
| C(15)-C(14)-C(16)-C(25) | 164.1(3)   | N(36)-C(38)-C(39)-C(41) | 164.38(19) |
| C(13)-C(14)-C(16)-C(25) | -17.3(3)   | C(49)-C(38)-C(39)-C(41) | 40.9(3)    |
| C(25)-C(16)-C(17)-C(18) | 0.5(4)     | C(41)-C(39)-C(40)-C(34) | 171.9(2)   |
| C(14)-C(16)-C(17)-C(18) | -179.0(3)  | C(38)-C(39)-C(40)-C(34) | -4.4(4)    |
| C(16)-C(17)-C(18)-C(19) | 0.6(4)     | C(35)-C(34)-C(40)-C(39) | -8.1(3)    |
| C(17)-C(18)-C(19)-C(20) | -0.7(4)    | C(32)-C(34)-C(40)-C(39) | -126.9(2)  |
| C(18)-C(19)-C(20)-N(21) | 179.7(3)   | C(40)-C(39)-C(41)-C(42) | -13.4(4)   |
| C(18)-C(19)-C(20)-C(25) | -0.3(4)    | C(38)-C(39)-C(41)-C(42) | 163.1(2)   |
| N(21)-C(22)-C(23)-C(25) | 1.3(4)     | C(40)-C(39)-C(41)-C(50) | 171.4(2)   |
| N(21)-C(22)-C(23)-C(24) | -175.6(3)  | C(38)-C(39)-C(41)-C(50) | -12.1(3)   |
| C(22)-C(23)-C(24)-C(13) | -156.1(3)  | C(50)-C(41)-C(42)-C(43) | 0.3(3)     |
| C(25)-C(23)-C(24)-C(13) | 27.4(4)    | C(39)-C(41)-C(42)-C(43) | -175.0(2)  |
| N(11)-C(13)-C(24)-C(23) | -167.7(2)  | C(41)-C(42)-C(43)-C(44) | 1.7(4)     |
| C(14)-C(13)-C(24)-C(23) | -45.6(3)   | C(42)-C(43)-C(44)-C(45) | -1.1(4)    |
| C(17)-C(16)-C(25)-C(20) | -1.6(4)    | C(43)-C(44)-C(45)-N(46) | 178.1(3)   |
| C(14)-C(16)-C(25)-C(20) | 178.0(2)   | C(43)-C(44)-C(45)-C(50) | -1.5(4)    |
| C(17)-C(16)-C(25)-C(23) | 178.9(3)   | N(46)-C(47)-C(48)-C(50) | 1.4(3)     |
| C(14)-C(16)-C(25)-C(23) | -1.5(4)    | N(46)-C(47)-C(48)-C(49) | -173.2(3)  |

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|                         |           |                         |             |
|-------------------------|-----------|-------------------------|-------------|
| C(47)-C(48)-C(49)-C(38) | -150.8(3) | C(6)-C(5)-N(4)-C(3)     | -56.9(3)    |
| C(50)-C(48)-C(49)-C(38) | 35.1(3)   | C(2)-C(3)-N(4)-C(7)     | -118.7(3)   |
| N(36)-C(38)-C(49)-C(48) | -174.4(2) | C(2)-C(3)-N(4)-C(5)     | 56.8(3)     |
| C(39)-C(38)-C(49)-C(48) | -51.1(3)  | C(9)-C(10)-N(11)-C(12)  | 163.7(2)    |
| N(46)-C(45)-C(50)-C(41) | -176.1(2) | C(9)-C(10)-N(11)-C(13)  | -70.8(2)    |
| C(44)-C(45)-C(50)-C(41) | 3.6(4)    | C(14)-C(13)-N(11)-C(12) | 177.5(2)    |
| N(46)-C(45)-C(50)-C(48) | 1.8(3)    | C(24)-C(13)-N(11)-C(12) | -56.1(3)    |
| C(44)-C(45)-C(50)-C(48) | -178.6(2) | C(14)-C(13)-N(11)-C(10) | 53.5(3)     |
| C(42)-C(41)-C(50)-C(45) | -2.9(4)   | C(24)-C(13)-N(11)-C(10) | 179.9(2)    |
| C(39)-C(41)-C(50)-C(45) | 172.9(2)  | C(19)-C(20)-N(21)-C(22) | 179.7(3)    |
| C(42)-C(41)-C(50)-C(48) | 179.7(2)  | C(25)-C(20)-N(21)-C(22) | -0.3(3)     |
| C(39)-C(41)-C(50)-C(48) | -4.5(4)   | C(23)-C(22)-N(21)-C(20) | -0.7(4)     |
| C(47)-C(48)-C(50)-C(45) | -2.0(3)   | O(33)-C(32)-N(29)-C(28) | 172.2(3)    |
| C(49)-C(48)-C(50)-C(45) | 173.8(2)  | C(34)-C(32)-N(29)-C(28) | -11.8(4)    |
| C(47)-C(48)-C(50)-C(41) | 175.8(2)  | O(33)-C(32)-N(29)-C(30) | 1.1(4)      |
| C(49)-C(48)-C(50)-C(41) | -8.5(4)   | C(34)-C(32)-N(29)-C(30) | 177.0(2)    |
| O(53)-C(52)-C(54)-O(55) | -167.1(2) | C(27)-C(28)-N(29)-C(32) | 134.7(3)    |
| O(51)-C(52)-C(54)-O(55) | 15.1(3)   | C(27)-C(28)-N(29)-C(30) | -53.6(3)    |
| O(53)-C(52)-C(54)-C(56) | 71.4(3)   | C(31)-C(30)-N(29)-C(32) | -134.6(2)   |
| O(51)-C(52)-C(54)-C(56) | -106.4(2) | C(31)-C(30)-N(29)-C(28) | 53.1(3)     |
| O(55)-C(54)-C(56)-O(57) | -72.3(2)  | C(34)-C(35)-N(36)-C(37) | 165.7(2)    |
| C(52)-C(54)-C(56)-O(57) | 49.1(2)   | C(34)-C(35)-N(36)-C(38) | -69.6(2)    |
| O(55)-C(54)-C(56)-C(58) | 50.9(3)   | C(39)-C(38)-N(36)-C(37) | 177.78(19)  |
| C(52)-C(54)-C(56)-C(58) | 172.3(2)  | C(49)-C(38)-N(36)-C(37) | -55.8(3)    |
| O(57)-C(56)-C(58)-O(60) | 8.1(3)    | C(39)-C(38)-N(36)-C(35) | 54.9(2)     |
| C(54)-C(56)-C(58)-O(60) | -112.1(2) | C(49)-C(38)-N(36)-C(35) | -178.63(19) |
| O(57)-C(56)-C(58)-O(59) | -173.1(2) | C(44)-C(45)-N(46)-C(47) | 179.5(3)    |
| C(54)-C(56)-C(58)-O(59) | 66.7(3)   | C(50)-C(45)-N(46)-C(47) | -0.9(3)     |
| O(8)-C(7)-N(4)-C(5)     | 1.3(4)    | C(48)-C(47)-N(46)-C(45) | -0.4(3)     |
| C(9)-C(7)-N(4)-C(5)     | 178.6(2)  | C(5)-C(6)-O(1)-C(2)     | -56.6(3)    |
| O(8)-C(7)-N(4)-C(3)     | 176.5(2)  | C(3)-C(2)-O(1)-C(6)     | 57.2(3)     |
| C(9)-C(7)-N(4)-C(3)     | -6.2(4)   | C(30)-C(31)-O(26)-C(27) | 60.5(3)     |
| C(6)-C(5)-N(4)-C(7)     | 119.0(3)  | C(28)-C(27)-O(26)-C(31) | -60.2(3)    |

Table 8. Hydrogen bonds for TCD173 [ $\text{\AA}$  and  $^\circ$ ].

| D-H...A                | d(D-H)  | d(H...A) | d(D...A) | $\angle(\text{DHA})$ |
|------------------------|---------|----------|----------|----------------------|
| C(5)-H(5A)...O(33)     | 0.99    | 2.45     | 3.280(3) | 141                  |
| C(9)-H(9)...O(53)      | 1.00    | 2.60     | 3.469(3) | 145                  |
| C(10)-H(10A)...O(57)   | 0.99    | 2.49     | 3.117(3) | 121                  |
| C(12)-H(12A)...O(51)#1 | 0.98    | 2.57     | 3.407(3) | 143                  |
| C(22)-H(22)...O(62)#2  | 0.95    | 2.58     | 3.527(5) | 177                  |
| C(27)-H(27B)...O(33)#3 | 0.99    | 2.37     | 3.299(3) | 156                  |
| C(30)-H(30B)...O(8)    | 0.99    | 2.60     | 3.411(3) | 139                  |
| C(35)-H(35A)...O(60)#4 | 0.99    | 2.57     | 3.125(3) | 115                  |
| C(44)-H(44)...O(59)#5  | 0.95    | 2.44     | 3.373(3) | 169                  |
| C(49)-H(49B)...O(1)#6  | 0.99    | 2.40     | 3.273(3) | 147                  |
| C(54)-H(54)...O(60)#3  | 1.00    | 2.28     | 3.181(3) | 149                  |
| N(11)-H(11)...O(51)    | 0.89(2) | 1.84(2)  | 2.729(3) | 171(3)               |
| N(21)-H(21)...O(26)#7  | 0.90(2) | 2.00(3)  | 2.859(3) | 160(4)               |
| N(36)-H(36)...O(59)#4  | 0.90(2) | 1.73(2)  | 2.631(3) | 179(4)               |
| N(36)-H(36)...O(60)#4  | 0.90(2) | 2.55(4)  | 3.109(3) | 120(3)               |
| N(46)-H(46)...O(53)#5  | 0.89(2) | 2.05(2)  | 2.908(3) | 161(3)               |
| O(55)-H(55)...O(51)    | 0.84(3) | 2.09(3)  | 2.634(3) | 122(3)               |
| O(57)-H(57)...O(53)#1  | 0.85(3) | 2.10(3)  | 2.804(3) | 141(3)               |
| O(57)-H(57)...O(60)    | 0.85(3) | 2.23(4)  | 2.679(3) | 113(3)               |
| O(61)-H(61A)...O(60)   | 0.94    | 1.91     | 2.831(3) | 165                  |
| O(61)-H(61B)...O(55)#1 | 0.90    | 2.01     | 2.887(3) | 162                  |
| O(62)-H(62)...O(61)    | 0.86    | 1.95     | 2.795(4) | 169                  |

Symmetry transformations used to generate equivalent atoms:

#1  $x+1, y, z$  #2  $x-1/2, -y+1/2, -z+2$  #3  $x-1, y, z$

#4  $x, y+1, z$  #5  $-x, y+1/2, -z+3/2$  #6  $-x+1, y+1/2, -z+3/2$

#7  $x-1/2, -y+3/2, -z+2$