

**Supplementary Information for**  
**“Highly fluorescent donor–acceptor purines”**

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**Experimental**

**General.** Reagents and solvents were purchased from Aldrich, Fluka, or Acros and used without further purification unless otherwise specified. THF, ether, CH<sub>2</sub>Cl<sub>2</sub>, and DMF were degassed in 20 L drums and passed through two sequential purification columns (activated alumina; molecular sieves for DMF) under a positive argon atmosphere. Flash chromatography (FC) was performed on Purasil SiO<sub>2</sub>-60, 230–400 mesh from Whatman. Thin layer chromatography (TLC) was performed on SiO<sub>2</sub>-60 F<sub>254</sub> aluminum plates from EMD Chemicals with visualization by UV light or staining (e.g. KMnO<sub>4</sub>). Melting points (m.p.) were determined on a Mel-temp electrothermal melting point apparatus and are uncorrected. <sup>1</sup>H (300) and <sup>13</sup>C NMR (75 MHz) spectra were recorded on Varian Mercury 300, Gemini 300, and VXR 300 spectrometers. Chemical shifts (δ) are given in parts per million (ppm) relative to residual protonated solvent (CHCl<sub>3</sub>: δ<sub>H</sub> 7.26 ppm, δ<sub>C</sub> 77.00 ppm). Abbreviations used are s (singlet), d (doublet), t (triplet), q (quartet), quin (quintet), b (broad), and m (multiplet). UV-Visible absorption spectra were obtained on a Perkin-Elmer Lambda 25 dual beam absorption spectrometer and a Cary 100 Bio spectrophotometer using 1 cm quartz cells. Steady-state fluorescence emission spectra were recorded on a SPEX Fluoromax spectrophotometer. Fluorescence lifetime measurements were obtained on a Photon Technology International, Inc. Time Master Fluorescence Lifetime system. High resolution mass spectra (HRMS) were obtained by the University of Florida Mass Spectrometry Services using a Finnigan MAT95Q Hybrid

Sector spectrometer. Purines **3a**<sup>1</sup>, **3d**<sup>2</sup>, and **4a**<sup>3</sup> were obtained as described in the literature with some modification.

## Synthesis of Compounds

### **General Method A** (*C(8) bromination of purines; conversion of 3 to 4*)<sup>4</sup>

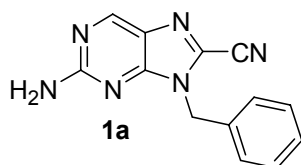
To a solution (6 mM) of starting material in CH<sub>2</sub>Cl<sub>2</sub> was added bromine (33 equiv.) dropwise and the mixture was stirred at room temperature for 1 h to 24 h. The reaction mixture was poured into 10% aqueous Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub>. The organic layer was separated and washed with brine, dried over MgSO<sub>4</sub>, and concentrated under reduced pressure. The residue was purified by silica gel column chromatography (1% MeOH/CH<sub>2</sub>Cl<sub>2</sub>) to give the C(8) brominated product.

### **General Method B** (*coupling of bromopurines with zinc cyanide; conversion of 4 to 1*)<sup>5</sup>

To a solution (0.16 M) of bromopurine **4** in NMP was added tetrakis(triphenylphosphine)palladium(0) (20–40 mol %), which was purified by washing with benzene and methanol. After stirring at 90 °C for 20 min a mixture of zinc cyanide (2 equiv.) suspended in NMP (0.90 mM) was added dropwise to the reaction (10 drops every 30 min for 4 h). The reaction was stirred at 90 °C overnight and then cooled and treated with aqueous ammonia (2 N, 15 mL). The mixture was extracted with ethyl acetate (3 x 100 mL). The combined organic extracts were washed with 2 N NH<sub>4</sub>OH and brine, dried over MgSO<sub>4</sub>, and concentrated under reduced pressure. The residue was purified by silica gel column chromatography (0.5% MeOH/CH<sub>2</sub>Cl<sub>2</sub>).

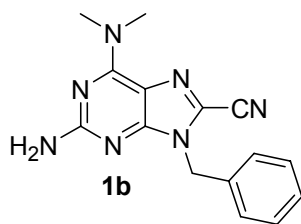
**General Method C** (*methanolysis of C(8) cyanopurines; conversion of 1 to 2*)<sup>6</sup>

A solution of methanolic ammonia (7 N) and cyanopurine **1** was stirred at room temperature overnight. The residue, after evaporation, was stirred in methanol while 1 N aqueous HCl was added dropwise. The solution was stirred for 2 h at room temperature, neutralized with Dowex 1 (OH<sup>-</sup>) resin, filtered, and concentrated under reduced pressure. The crude product was purified by silica gel column chromatography (1% MeOH/CH<sub>2</sub>Cl<sub>2</sub>).



**2-Amino-8-cyano-9-benzylpurine (1a)**

Compound **4a** (0.10 g, 0.33 mmol) was reacted under the conditions of general method **B** to yield a white solid (0.065 g, 80%): m.p. 139–142 °C. <sup>1</sup>H NMR (300 MHz, DMSO-*d*<sub>6</sub>) δ 5.40 (s, 2H), 7.26 (m, 7H), 8.88 (s, 1H). <sup>13</sup>C NMR (75 MHz, DMSO-*d*<sub>6</sub>) δ 46.62, 112.17, 124.40, 126.58, 127.87, 128.87, 128.90, 129.67, 136.11. HRMS calcd for C<sub>13</sub>H<sub>10</sub>N<sub>6</sub> [M<sup>+</sup>]: 250.0967, found: 250.0973.

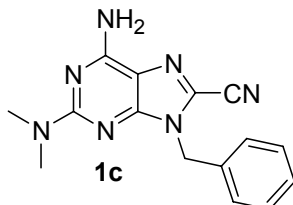


**2-Amino-6-dimethylamino-8-cyano-9-benzylpurine (1b)**

Compound **4b** (0.20 g, 0.58 mmol) was reacted under the conditions of general method **B** to yield a white solid (0.15 g, 91%): m.p. 175–176°C. <sup>1</sup>H NMR (300 MHz, DMSO-*d*<sub>6</sub>) δ 3.40 (bs, 6H), 5.33 (s, 2H), 6.44 (s, 2H), 7.33 (m, 5H). <sup>13</sup>C NMR (75 MHz, DMSO-*d*<sub>6</sub>) δ

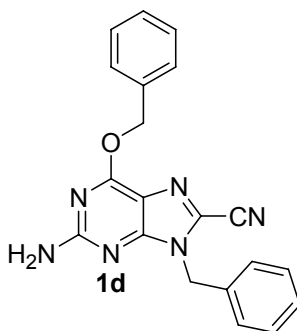
47.56, 112.08, 114.91, 117.72, 126.90, 127.90, 128.78, 135.82, 152.77, 154.86, 161.50.

HRMS calcd for  $C_{15}H_{16}N_7$   $[M+H]^+$ : 294.1467, found: 294.1479.



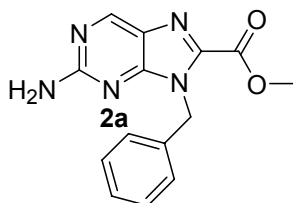
### 2-Dimethylamino-6-amino-8-cyano-9-benzylpurine (1c)

Compound **4c** (0.10 g, 0.29 mmol) was reacted under the conditions of general method **B** to yield a white solid (85 mg, 77 %): m.p. 205–209 ° C.  $^1H$  NMR (300 MHz, DMSO- $d_6$ )  $\delta$  3.12 (s, 6H), 5.33 (s, 2H), 7.34 (m, 7H).  $^{13}C$  NMR (75 MHz, DMSO- $d_6$ )  $\delta$  36.90, 46.02, 112.25, 113.92, 118.50, 127.67, 128.10, 135.86, 136.00, 151.49, 156.28, 161.02. HRMS calcd for  $C_{15}H_{16}N_7$   $[M+H]^+$ : 294.1467, found: 294.1492.



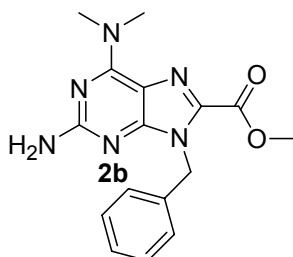
### 2-Amino-6-benzyloxy-8-cyano-9-benzylpurine (1d)

Compound **4d** (0.30 g, 0.73 mmol) was reacted under the conditions of general method **B** to yield a white solid (0.14 g, 54%): m.p. 246–248 ° C.  $^1H$  NMR (300 MHz, DMSO- $d_6$ )  $\delta$  5.37 (s, 2H), 5.52 (s, 2H), 7.14, (s, 2H) 7.38 (m, 10H).  $^{13}C$  NMR (75 MHz, DMSO- $d_6$ )  $\delta$  46.22, 67.61, 111.58, 114.62, 121.17, 127.01, 128.07, 128.25, 128.42, 128.64, 128.88, 135.48, 135.89, 153.69, 161.10, 161.79. HRMS calcd for  $C_{20}H_{17}N_6O$   $[M+H]^+$ : 357.1464, found: 357.1432.



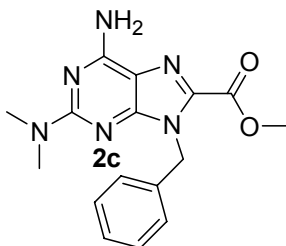
**2-Amino-8-methylester-9-benzylpurine (2a)**

Compound **1a** (0.10 g, 0.40 mmol) was reacted under the conditions of general method **C** to yield a white solid (0.068 g, 60%): m.p. 226–228 ° C.  $^1\text{H}$  NMR (300 MHz,  $\text{CDCl}_3$ )  $\delta$  3.97 (s, 3H), 5.27 (s, 2H), 5.74 (s, 2H), 7.29 (m, 5H), 8.84 (s, 1H).  $^{13}\text{C}$  NMR (75 MHz,  $\text{CDCl}_3$ )  $\delta$  47.05, 53.22, 127.18, 127.92, 128.13, 128.87, 136.71, 151.75, 153.26, 159.86, 161.43. HRMS calcd for  $\text{C}_{14}\text{H}_{13}\text{N}_5\text{O}_2$   $[\text{M}+\text{H}]^+$ : 283.1069, found: 283.1086.



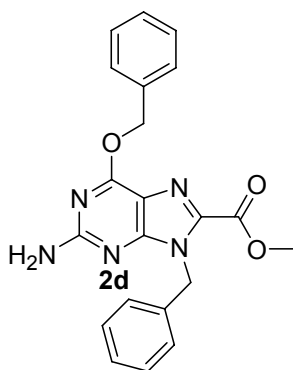
**2-Amino-6-dimethylamino-8-methylester-9-benzylpurine (2b)**

Compound **1b** (0.062 g, 0.21 mmol) was reacted under the general method **C** to yield a white solid (0.054 g, 86%): m.p. 175–180 ° C.  $^1\text{H}$  NMR (300 MHz,  $\text{CDCl}_3$ )  $\delta$  3.51 (bs, 6H) 3.91 (s, 3H) 4.77 (s, 2H) 5.70 (s, 2H) 7.26 (m, 5H).  $^{13}\text{C}$  NMR (75 MHz,  $\text{DMSO}-d_6$ )  $\delta$  37.00, 46.12, 52.02, 114.23, 126.49, 127.12, 128.41, 132.89, 137.89, 137.60, 154.38, 155.46, 159.28, 161.02. HRMS calcd for  $\text{C}_{16}\text{H}_{19}\text{N}_6\text{O}_2$   $[\text{M}+\text{H}]^+$ : 327.1569, found: 327.1583.



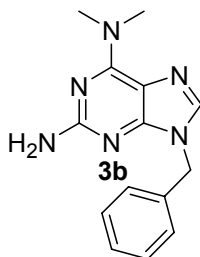
**2-Dimethylamino-6-amino-8-methylester-9-benzylpurine (2c)**

Compound **1c** (0.10 g, 0.34 mmol) was reacted under the conditions of general method **C** to yield a white solid (0.067 g, 60%): m.p. 250–252 °C.  $^1\text{H}$  NMR (300 MHz,  $\text{DMSO}-d_6$ )  $\delta$  3.10 (s, 6H), 3.80 (s, 3H), 5.55 (s, 2H), 7.26 (m, 7H).  $^{13}\text{C}$  NMR (75 MHz,  $\text{DMSO}-d_6$ )  $\delta$  36.91, 46.25, 51.94, 113.12, 127.30, 128.39, 133.75, 137.67, 153.04, 156.83, 159.39, 160.58. HRMS calcd For  $\text{C}_{16}\text{H}_{18}\text{N}_6\text{O}_2$  [ $\text{M}^+$ ]: 326.1491, found: 326.1488.



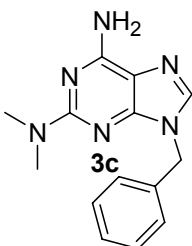
**2-Amino-6-benzyloxy-8-methylester-9-benzylpurine (2d)**

Compound **1d** (0.037 g, 0.10 mmol) was reacted under the conditions of general method **C** to yield a white solid (0.038 g, 95%): m.p. 180–185 °C.  $^1\text{H}$  NMR (300 MHz,  $\text{CDCl}_3$ )  $\delta$  3.90 (s, 3H), 5.03 (s, 2H), 5.53 (s, 2H), 5.70 (s, 2H), 7.31 (m, 10H).  $^{13}\text{C}$  NMR (75 MHz,  $\text{CDCl}_3$ )  $\delta$  47.20, 52.64, 68.46, 115.51, 127.52, 127.69, 128.18, 128.37, 128.53, 128.65, 135.90, 136.74, 137.13, 155.26, 159.70, 160.53, 162.56. HRMS calcd for  $\text{C}_{21}\text{H}_{19}\text{N}_5\text{O}_3$  [ $\text{M}+\text{H}$ ] $^+$ : 390.1566, found: 390.1544.



### 2-Amino-6-dimethylamino-9-benzylpurine (3b)

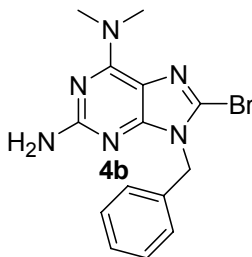
Dimethylamine (4.0 mL, 33% in abs ethanol) was added to 2-amino-6-chloro-9-benzylpurine<sup>7</sup> (1.0 g, 3.9 mmol) in abs ethanol (40 mL) and the solution was heated to 60 °C for 20 h. The reaction mixture was evaporated to dryness and triturated with deionized H<sub>2</sub>O. The white crystals (0.95 g, 92%) were then collected by vacuum filtration and dried under high vacuum: m.p. 170–173 °C. <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>) δ 3.46 (s, 6H), 4.61 (s, 2H), 5.21 (s, 2H), 7.27 (m, 5H), 7.42 (s, 1H). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>) δ 38.22, 46.39, 114.98, 127.49, 127.93, 128.82, 135.87, 136.36, 152.91, 155.39, 159.40. HRMS Calcd for C<sub>14</sub>H<sub>17</sub>N<sub>5</sub> [M+H]<sup>+</sup>: 269.1509, found: 269.1516.



### 2-Dimethylamino-6-amino-9-benzylpurine (3c)

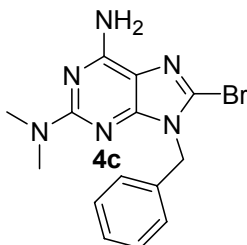
Dimethylamine (10 mL, 33% in abs ethanol) was added to 2-chloro-6-amino-9-benzyl purine (0.30 g, 1.2 mmol) in a sealed tube. The reaction was stirred at 100 °C for 20 h, and then concentrated under reduced pressure. The crude solid was purified by silica gel column chromatography (1% MeOH/CH<sub>2</sub>Cl<sub>2</sub>) to afford a white solid (0.31 g, 99%): m.p. 168–171 °C. <sup>1</sup>H NMR (300 MHz, DMSO-*d*<sub>6</sub>) δ 3.07 (s, 6H), 5.19 (s, 2H), 6.73 (s, 2H), 7.34 (m, 5H), 7.82 (s, 1H). <sup>13</sup>C NMR (75 MHz, DMSO-*d*<sub>6</sub>) δ 37.00, 45.64, 112.39,

127.50, 127.77, 128.48, 137.35, 137.56, 151.60, 155.54, 159.48. HRMS calcd for  $C_{14}H_{17}N_6$   $[M+H]^+$ : 269.1509, found: 269.1515.



**2-Amino-6-dimethylamino-8-bromo-9-benzylpurine (4b)**

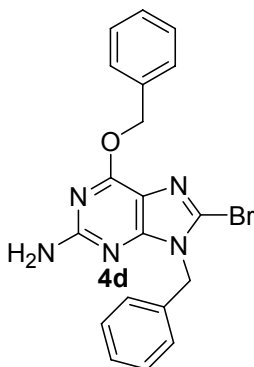
Compound **3b** (0.50 g, 1.8 mmol) was reacted under the conditions of general method A to yield a white solid (0.52 g, 81%): m.p. 168–169 °C.  $^1H$  NMR (300 MHz,  $CDCl_3$ )  $\delta$  3.42 (s, 6H), 4.69 (s, 2H), 5.24 (s, 2H), 7.28 (m, 5H).  $^{13}C$  NMR (75 MHz,  $CDCl_3$ )  $\delta$  38.27, 46.69, 115.54, 121.10, 127.40, 127.77, 128.63, 135.87, 154.13, 154.21, 159.19. HRMS calcd for  $C_{14}H_{15}BrN_6$   $[M+H]^+$ : 347.0620, found: 347.0618.



**2-Dimethylamino-6-amino-8-bromo-9-benzylpurine (4c)**

Compound **3c** (0.59 g, 2.2 mmol) was reacted under the conditions of general method A to yield a white solid (0.52 g, 68%): m.p. 195–197 °C.  $^1H$  NMR (300 MHz,  $CDCl_3$ )  $\delta$  3.15 (s, 6H), 5.21 (bs, 2H), 5.23 (s, 2H), 7.29 (m, 5H).  $^{13}C$  NMR (75 MHz,  $CDCl_3$ )  $\delta$  37.33, 46.11, 46.90, 113.31, 122.36, 127.90, 128.06, 128.61, 136.03, 153.87, 159.77. HRMS calcd for  $C_{14}H_{16}BrN_6$   $[M+H]^+$ : 347.0614, found: 347.0626.





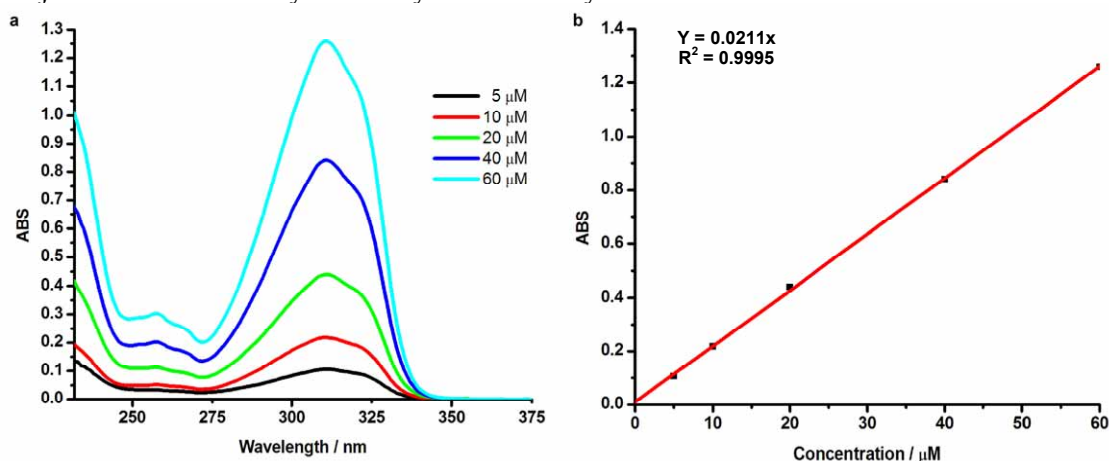
### 2-Amino-6-benzyloxy-8-bromo-9-benzylpurine (4d)

Compound **3d** (0.38 g, 0.93 mmol) was reacted under the conditions of general method **A** to yield a white solid (0.44 g, 77%): m.p. 164–166 °C.  $^1\text{H}$  NMR (300 MHz,  $\text{CDCl}_3$ )  $\delta$  7.50 (m, 2H), 7.30 (m, 8H), 5.53 (s, 2H), 5.26 (s, 2H), 4.90 (s, 2H).  $^{13}\text{C}$  NMR (75 MHz,  $\text{CDCl}_3$ )  $\delta$  47.11, 68.25, 115.85, 125.56, 127.50, 128.03, 128.07, 128.39, 128.75, 135.37, 136.20, 155.29, 159.17, 159.85. HRMS calcd for  $\text{C}_{19}\text{H}_{17}\text{BrN}_6\text{O}$   $[\text{M}+\text{H}]^+$ : 410.0616, found: 410.0627.

## Optical Studies

### UV-Visible Data

The absorbance was measured for 5, 10, 20, 40, and 60  $\mu\text{M}$  solutions on the Cary 100 Bio UV-Visible spectrophotometer (thermostatted at 25 °C) using dried and degassed methylene chloride ( $\text{CH}_2\text{Cl}_2$ ). The absorbance intensity at  $\lambda_{\text{max}}$  was then plotted against the concentration to confirm, by linearity, that the compounds followed Beer's law. Once confirmed, molar extinction coefficients ( $\epsilon$ ) were determined from the linear plot for each compound (where  $A = \epsilon bc$ ). An example of this procedure is shown below for **1d**.



The slope of the line is then multiplied by  $10^6$  (this accounts for the concentration) to generate  $\epsilon$ . This is then displayed as  $\log \epsilon$ .

## Fluorescence Data

### Fluorescence Lifetime Measurements

Fluorescence lifetime measurements were performed on a Photon Technology International, Inc. Time Master Fluorescence Lifetime system. Excitation was performed via a nitrogen laser at 337 nm (pulse width approximately 800 ps), and a quartz fiber optic was used to convey light to the sample chamber. Samples for fluorescence lifetime measurements of compounds **1a–d** and **2a–d** were prepared by making  $\sim 5 \mu\text{M}$  solutions of each compound in dried and degassed methylene chloride. Each compound was excited at 337 nm and the monochromator was set to the corresponding  $\lambda_{\text{em(max)}}$  for each compound. After the measurement of each compound, a scattering agent (Ludox<sup>®</sup> AM-30 colloidal silica, 30% wt. % suspension in water) was measured under the same parameters as the compound in order to find the instrument response time. The instrument response time along with the data collected for each sample was placed into the Time Master software to generate the fluorescence lifetime data.<sup>8,9,10</sup>

**Table 1: Lifetime Values and Parameters**

| <b>Compound</b> | <b>Lifetime of the<br/>excited state<br/>[ns]<br/>(<math>\lambda_{\text{ex}} = 373 \text{ nm}</math>)</b> | <b><math>\chi^2</math></b> | <b>DW</b> | <b>Z</b> |
|-----------------|-----------------------------------------------------------------------------------------------------------|----------------------------|-----------|----------|
| <b>1a</b>       | 0.51 $\pm$ 0.23                                                                                           | 1.049                      | 0.727     | -1.454   |
| <b>1b</b>       | 1.59 $\pm$ 0.25                                                                                           | 1.158                      | 1.158     | -1.577   |
| <b>1c</b>       | 3.15 $\pm$ 0.25                                                                                           | 0.885                      | 0.870     | -0.980   |
| <b>1d</b>       | 3.05 $\pm$ 0.16                                                                                           | 1.041                      | 0.463     | -2.534   |
| <b>2a</b>       | 1.75 $\pm$ 0.18                                                                                           | 1.333                      | 0.937     | -2.336   |
| <b>2b</b>       | 3.06 $\pm$ 0.27                                                                                           | 1.219                      | 0.398     | -1.911   |
| <b>2c</b>       | 3.25 $\pm$ 0.08                                                                                           | 0.968                      | 1.451     | -1.714   |
| <b>2d</b>       | 2.54 $\pm$ 0.07                                                                                           | 1.092                      | 2.414     | -0.564   |

Deviations from the best fit are characterized by the reduced chi-square statistic,  $\chi^2$ . Good results typically produce  $\chi^2$ 's of 0.9 to 1.2. Most of the  $\chi^2$  values displayed in the table above are within the range indicated, those that are not are very close. The Durbin-Watson (DW) parameter was introduced to test for correlations. The fit is likely satisfactory if the value of DW is greater than 1.7 for single exponential fits. Most of the values are well below the satisfactory value of DW, however, the correlation is looking for structured patterns around the baseline, and can be sensitive to radio-frequency (RF) noise. The runs test determines the number of positive and negative groups of the residuals. A value of  $-1.96 < Z$  indicates a satisfactory fit at the 95% confidence level. The general structure of each analysis program is the same so there is considerable similarity in running the programs.

### Fluorescence Quantum Yield Measurements

Fluorescence quantum yield measurements were performed by collecting correlating absorption and steady-state emission spectra on a Perkin-Elmer Lambda 25 dual beam

absorption spectrometer and a SPEX Fluoromax spectrophotometer, respectively. The unknowns and standards were prepared with dried and degassed solvent at or below a 10  $\mu\text{M}$  concentration and absorbance below an intensity of 0.1. The equation used to calculate the quantum yield values follows,

$$\Phi_{\text{F(x)}} = (A_{\text{s}}/A_{\text{x}})(F_{\text{x}}/F_{\text{s}})(\eta_{\text{x}}/\eta_{\text{s}})^2 \Phi_{\text{F(s)}}$$

where  $\Phi_{\text{F}}$  is the fluorescence quantum yield,  $A$  is the absorbance at the excitation wavelength,  $F$  is the area under the emission curve, and  $\eta$  is the refractive index of the solvent used. Subscripts s and x refer to the standard and unknown, respectively. Quantum yields were calculated using several different solutions of three different standards (shown in the table below).

**Table 2: Fluorescence Quantum Yield ( $\Phi_{\text{F}}$ ) Data in  $\text{CH}_2\text{Cl}_2$  Calculated Using Three Different Standards**

| Standards <sup>a</sup> | 1a    | 1b    | 1c    | 1d    | 2a    | 2b    | 2c  | 2d    |
|------------------------|-------|-------|-------|-------|-------|-------|-----|-------|
| DPA 1                  | 0.194 | 0.302 | 0.976 | 0.820 | 0.419 | > 1   | > 1 | 0.986 |
| DPA 2                  | 0.148 | 0.320 | –     | 0.871 | 0.445 | > 1   | > 1 | 0.749 |
| Q3                     | 0.176 | 0.299 | 0.886 | 0.744 | 0.380 | 0.939 | > 1 | 0.894 |
| Q4                     | 0.197 | 0.300 | 0.993 | 0.834 | 0.426 | > 1   | > 1 | > 1   |
| Q5                     | 0.193 | 0.301 | 0.972 | 0.816 | 0.417 | > 1   | > 1 | 0.981 |
| Q6                     | 0.193 | 0.302 | 0.972 | 0.817 | 0.417 | > 1   | > 1 | 0.982 |
| Q7                     | 0.193 | 0.306 | 0.969 | 0.814 | 0.416 | > 1   | > 1 | 0.979 |
| A1                     | 0.193 | 0.273 | 0.973 | 0.816 | 0.418 | > 1   | > 1 | 0.982 |
| A3                     | 0.194 | 0.307 | 0.976 | 0.819 | 0.419 | > 1   | > 1 | 0.986 |
| A4                     | 0.194 | 0.300 | 0.977 | –     | 0.420 | > 1   | > 1 | 0.987 |
| A5                     | 0.197 | 0.300 | 0.991 | 0.832 | 0.426 | > 1   | > 1 | > 1   |

<sup>a</sup> DPA = 9,10-diphenylanthracene in cyclohexane ( $\Phi_F = 0.90$ );<sup>11</sup> Q = quinine sulfate in 0.05 M H<sub>2</sub>SO<sub>4</sub> ( $\Phi_F = 0.546$ );<sup>10,12</sup> A = anthracene in ethanol. The numbers after the standards indicate different solutions.

**Table 3: Absorption and Emission Data for Donor-Acceptor Purines in Methanol (10<sup>-6</sup> M)**

| Compound  | $\lambda_{(\max)}$ | $\log \epsilon$ | $\lambda_{\text{em}(\max)}$<br>excited at 320 nm |
|-----------|--------------------|-----------------|--------------------------------------------------|
| <b>1a</b> | 222, 318           | 4.4, 3.2        | 399                                              |
| <b>1b</b> | 248, 327           | 4.2, 4.2        | 382                                              |
| <b>1c</b> | 291, 338           | 4.2, 4.2        | 412                                              |
| <b>1d</b> | 208, 318           | 4.3, 4.1        | 383                                              |
| <b>2a</b> | 248, 335           | 3.9, 4.1        | 409                                              |
| <b>2b</b> | 249, 331           | 4.2, 4.2        | 419                                              |
| <b>2c</b> | 226, 345           | 4.2, 4.1        | 436                                              |
| <b>2d</b> | 209, 321           | 4.6, 4.3        | 405                                              |

## X-ray Crystallography

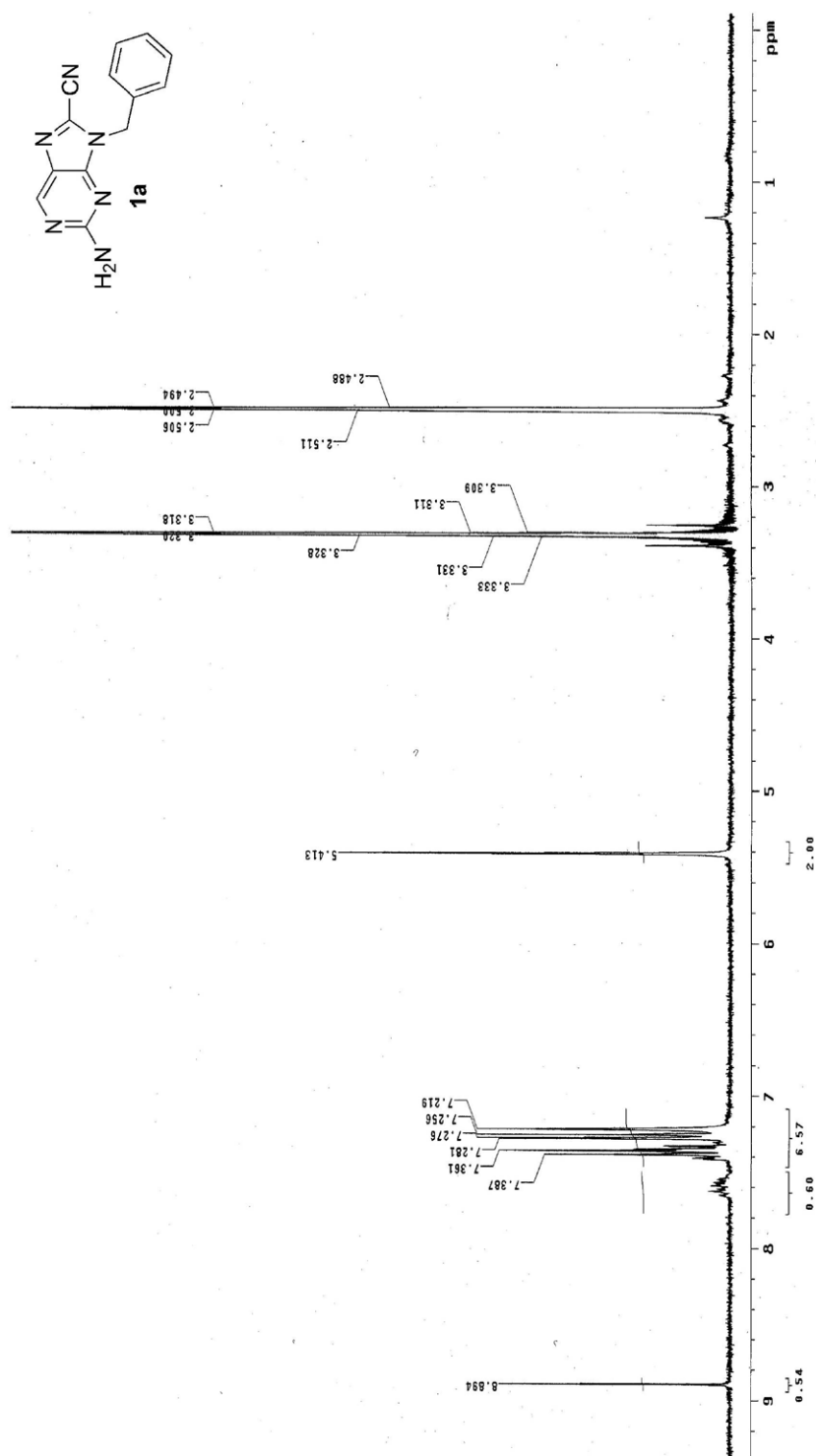
**General:** Data were collected at 173 K on a Siemens SMART PLATFORM equipped with a CCD area detector and a graphite monochromator utilizing MoK $\alpha$  radiation ( $\lambda = 0.71073$  Å). Cell parameters were refined using up to 8192 reflections. A full sphere of data (1850 frames) was collected using the  $\omega$ -scan method (0.3° frame width). The first 50 frames were re-measured at the end of data collection to monitor instrument and crystal stability (maximum correction on  $I$  was  $< 1$  %). Absorption corrections by integration were applied based on measured indexed crystal faces.

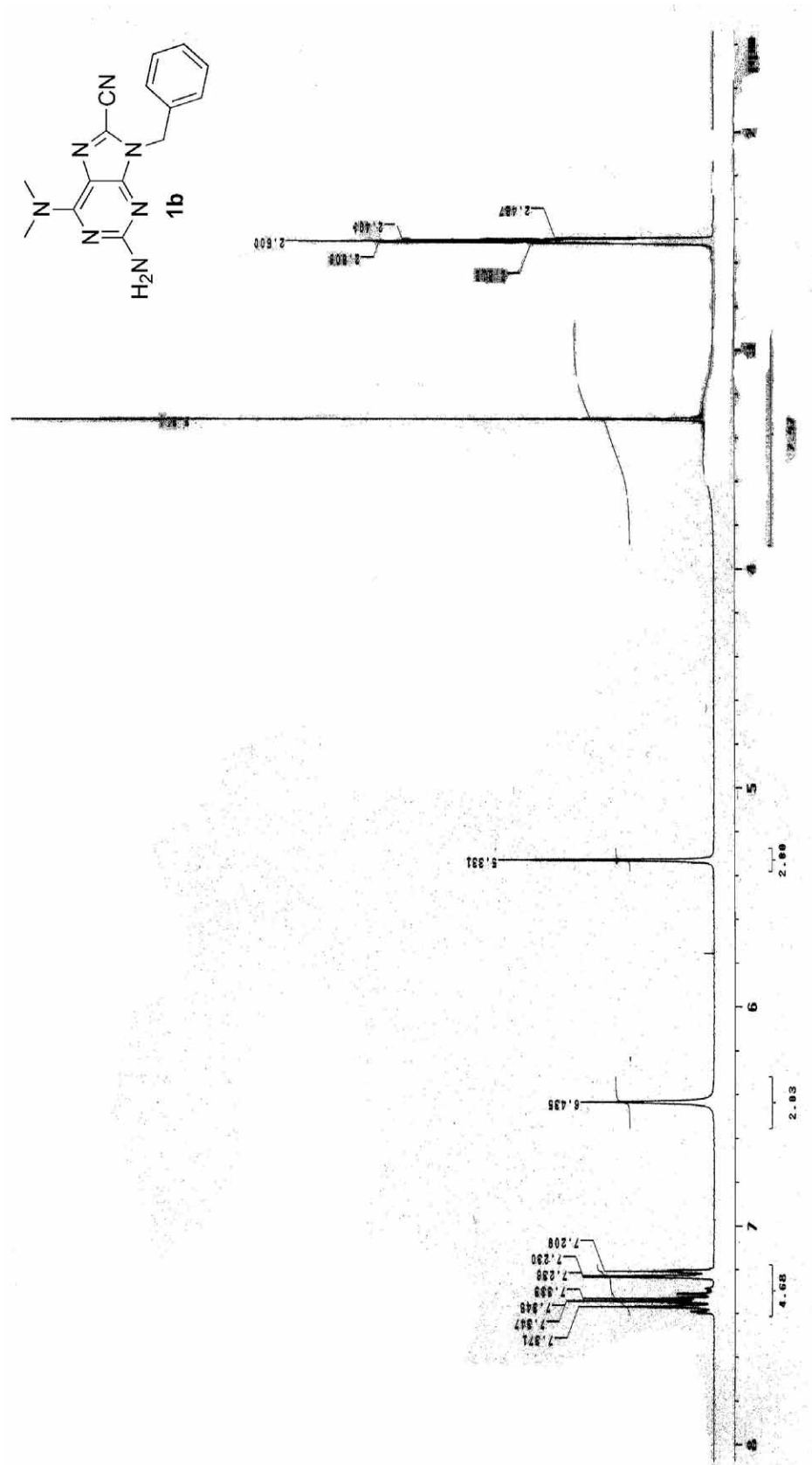
The structures were solved by the Direct Methods in *SHELXTL6*,<sup>13</sup> and refined using full-matrix least squares. The non-H atoms were treated anisotropically, whereas the hydrogen atoms were calculated in ideal positions and were riding on their respective carbon atoms.

**Compound 1b:** The methyl H atoms on C(14) and C(15) are disordered and each set was refined in two parts and in a riding model with their site occupation factors fixed at 50%. A total of 208 parameters were refined in the final cycle of refinement using 4656 reflections with  $I > 2\sigma(I)$  to yield  $R_1$  and  $wR_2$  of 3.83% and 10.27%, respectively. Refinement was done using  $F^2$ .

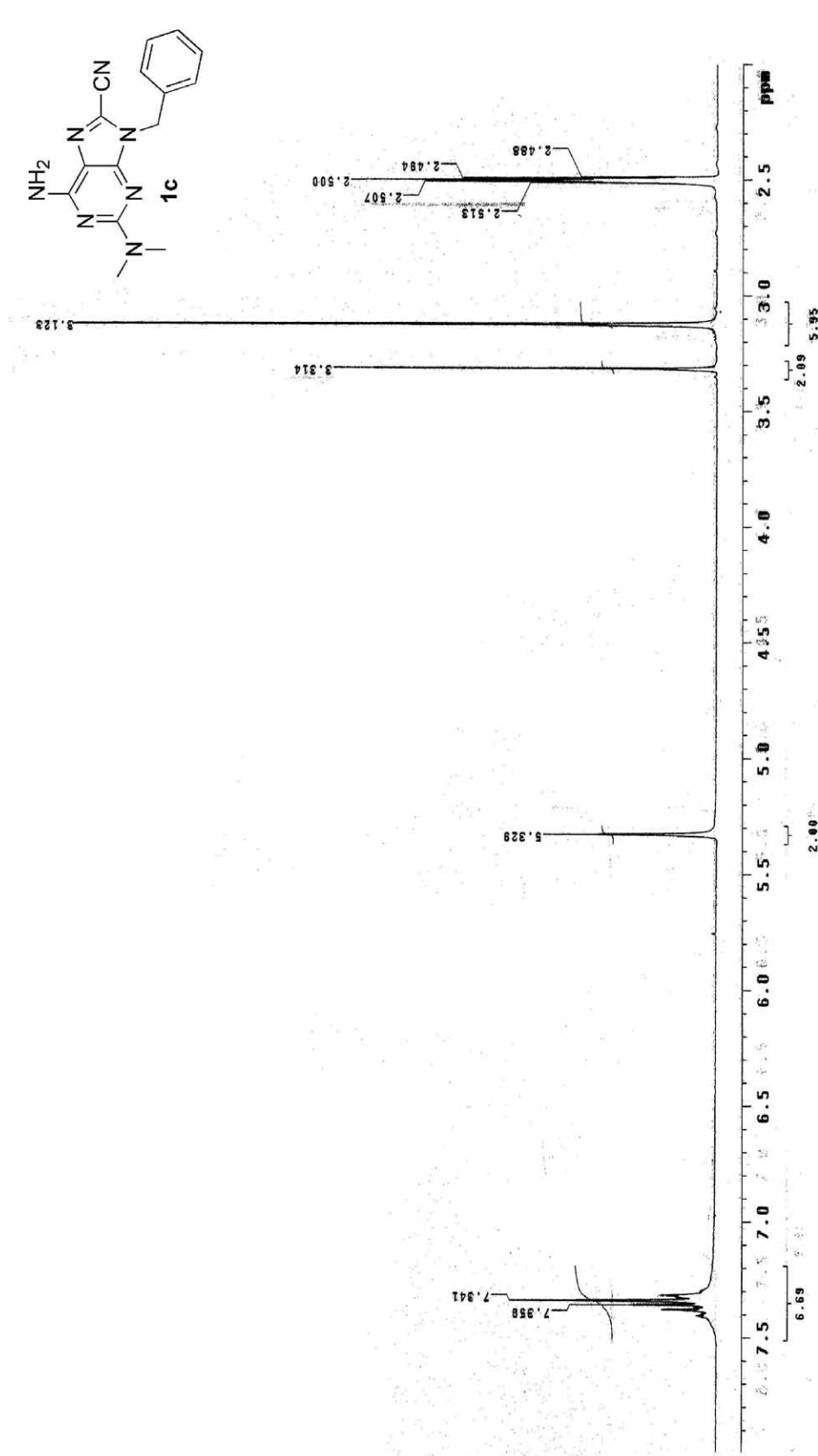
**Compound 1c:** A total of 207 parameters were refined in the final cycle of refinement using 3304 reflections with  $I > 2\sigma(I)$  to yield  $R_1$  and  $wR_2$  of 4.75% and 12.0%, respectively. Refinement was done using  $F^2$ .

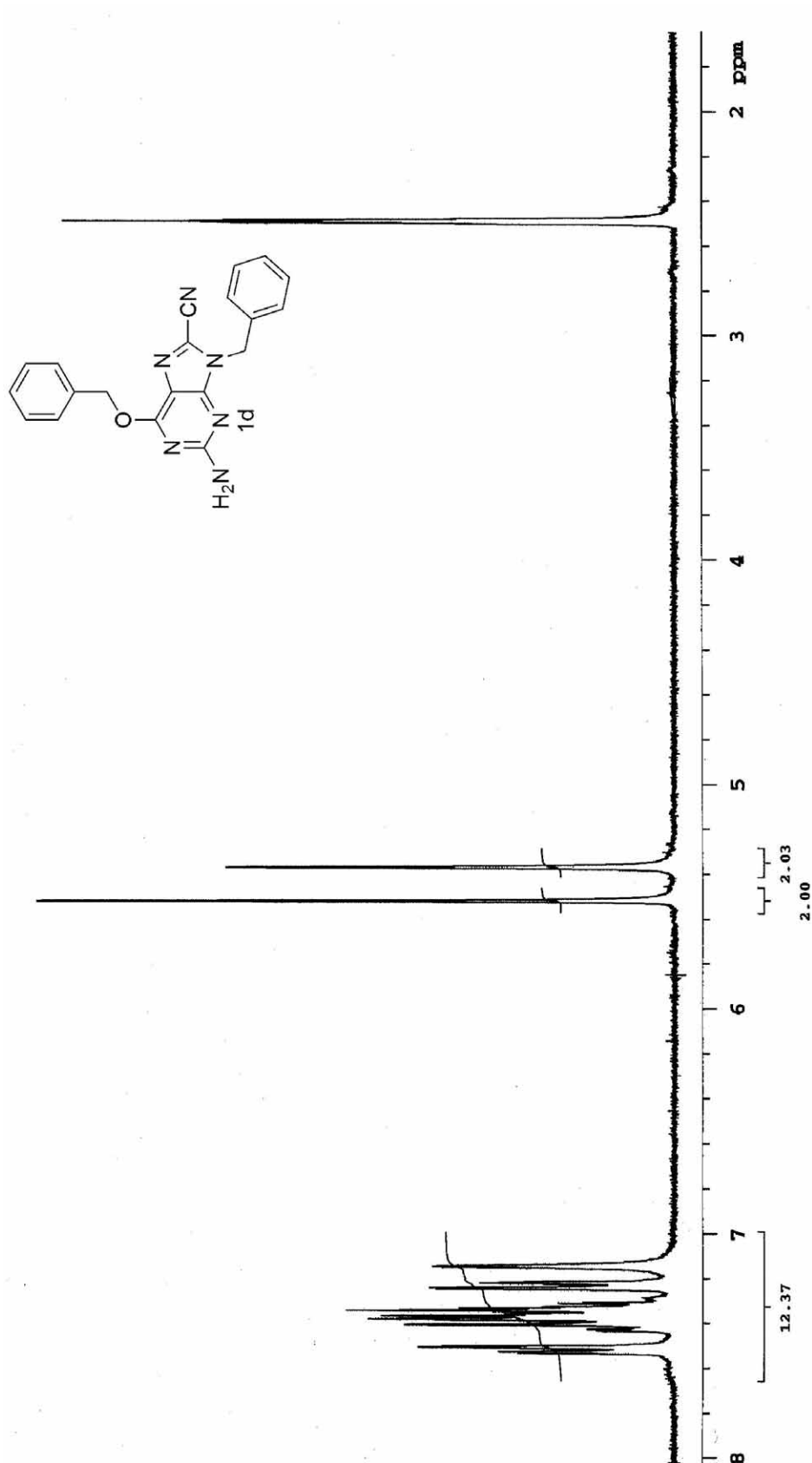
### <sup>1</sup>H NMR Spectra

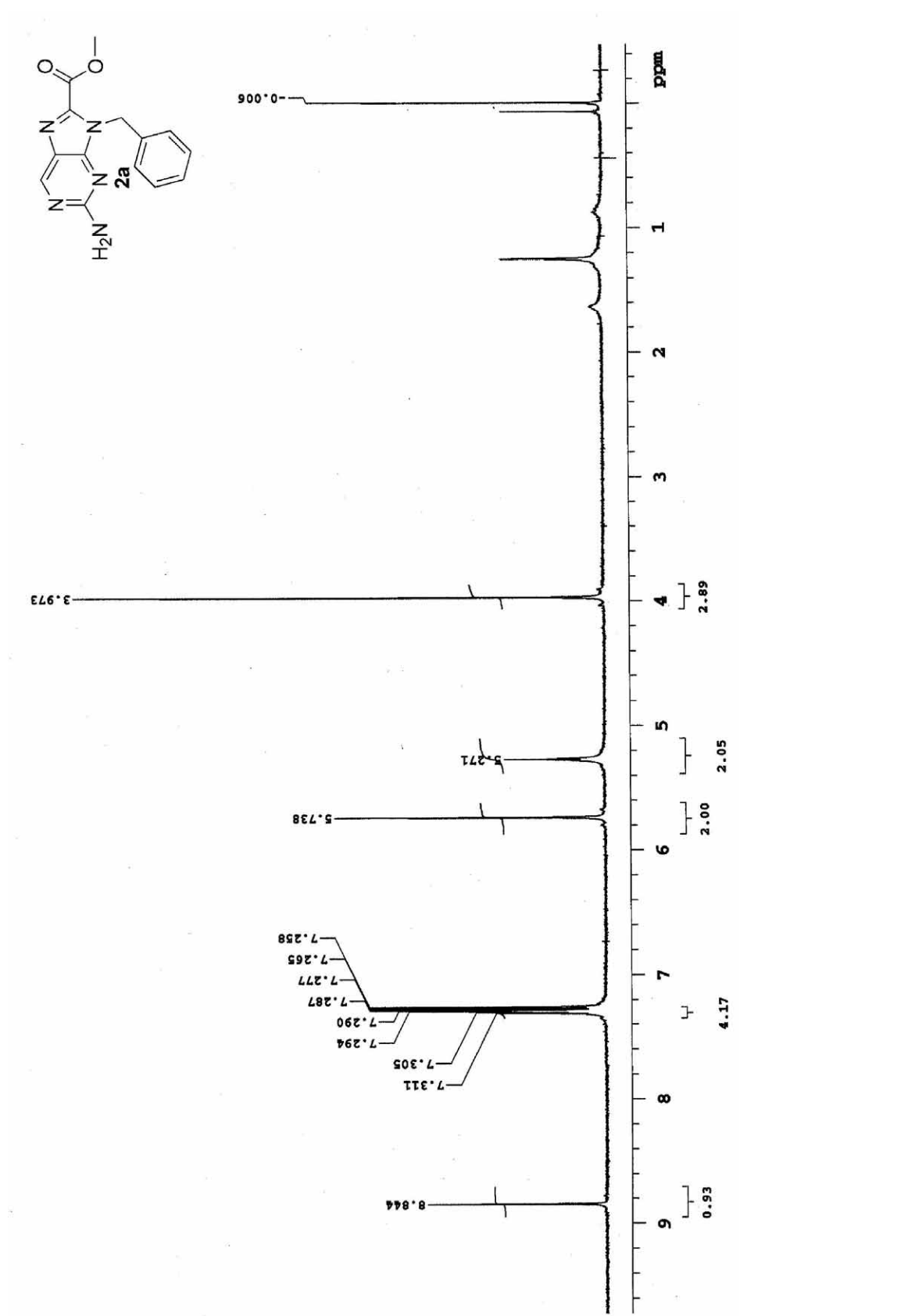


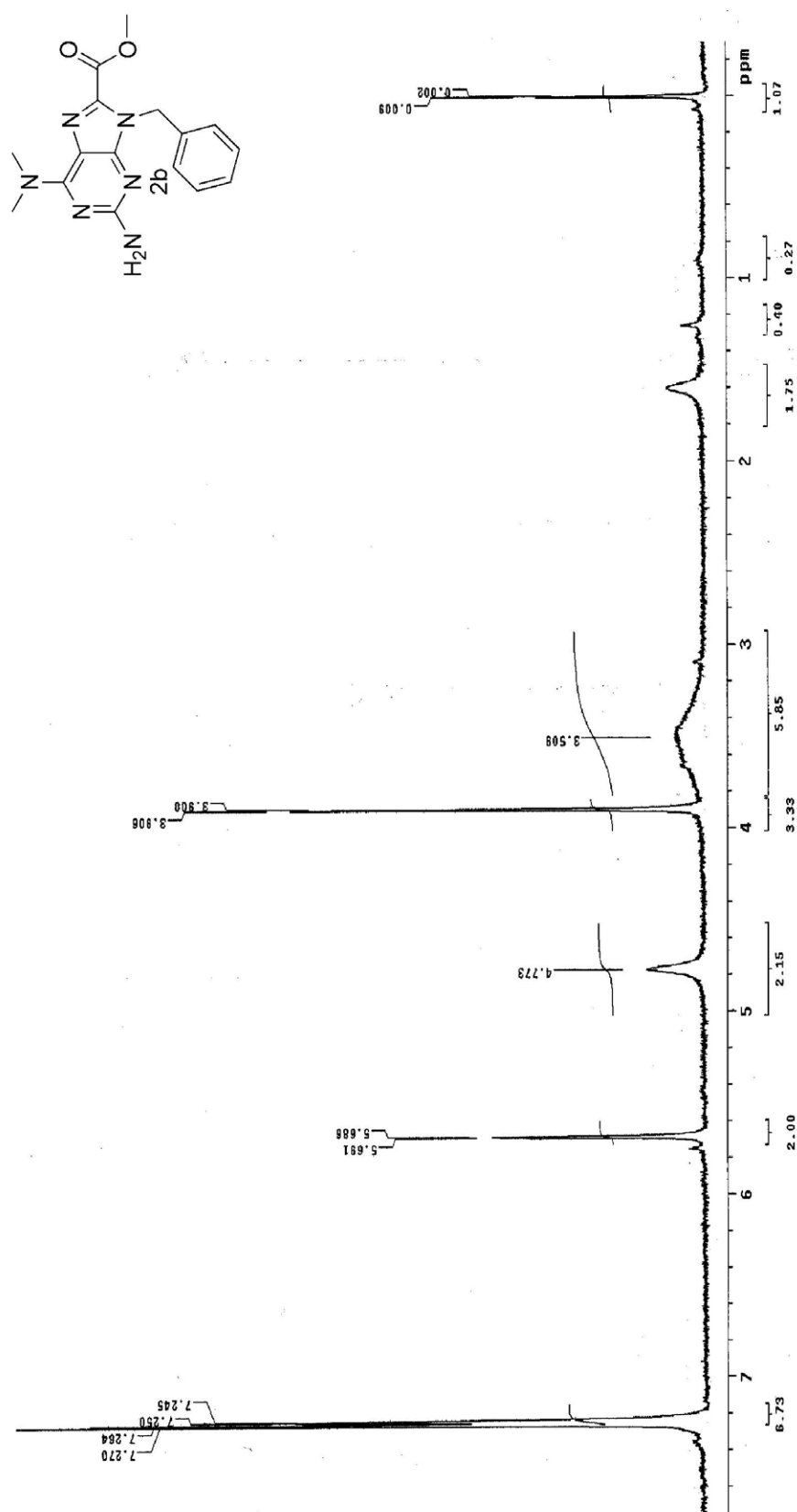


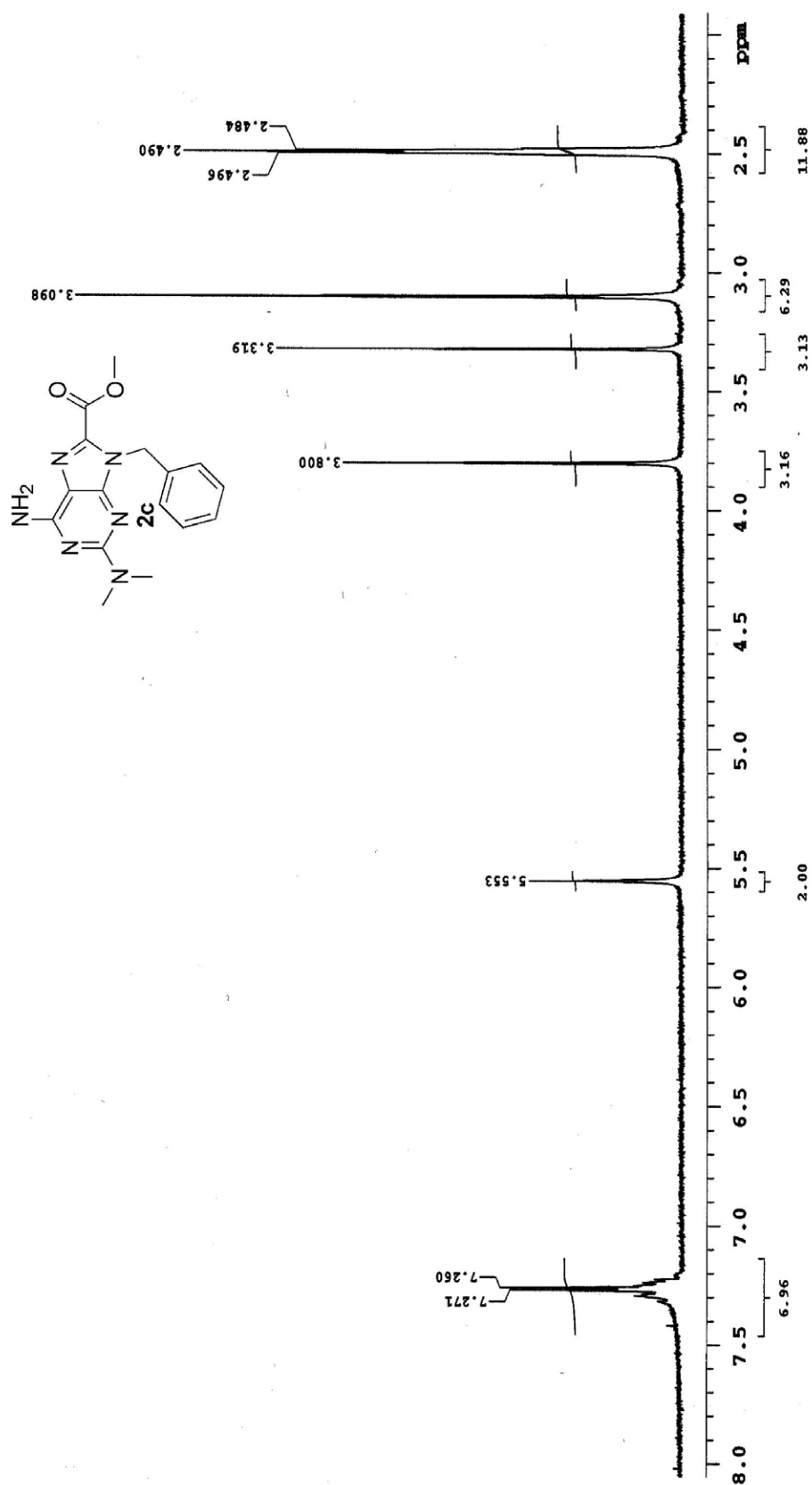


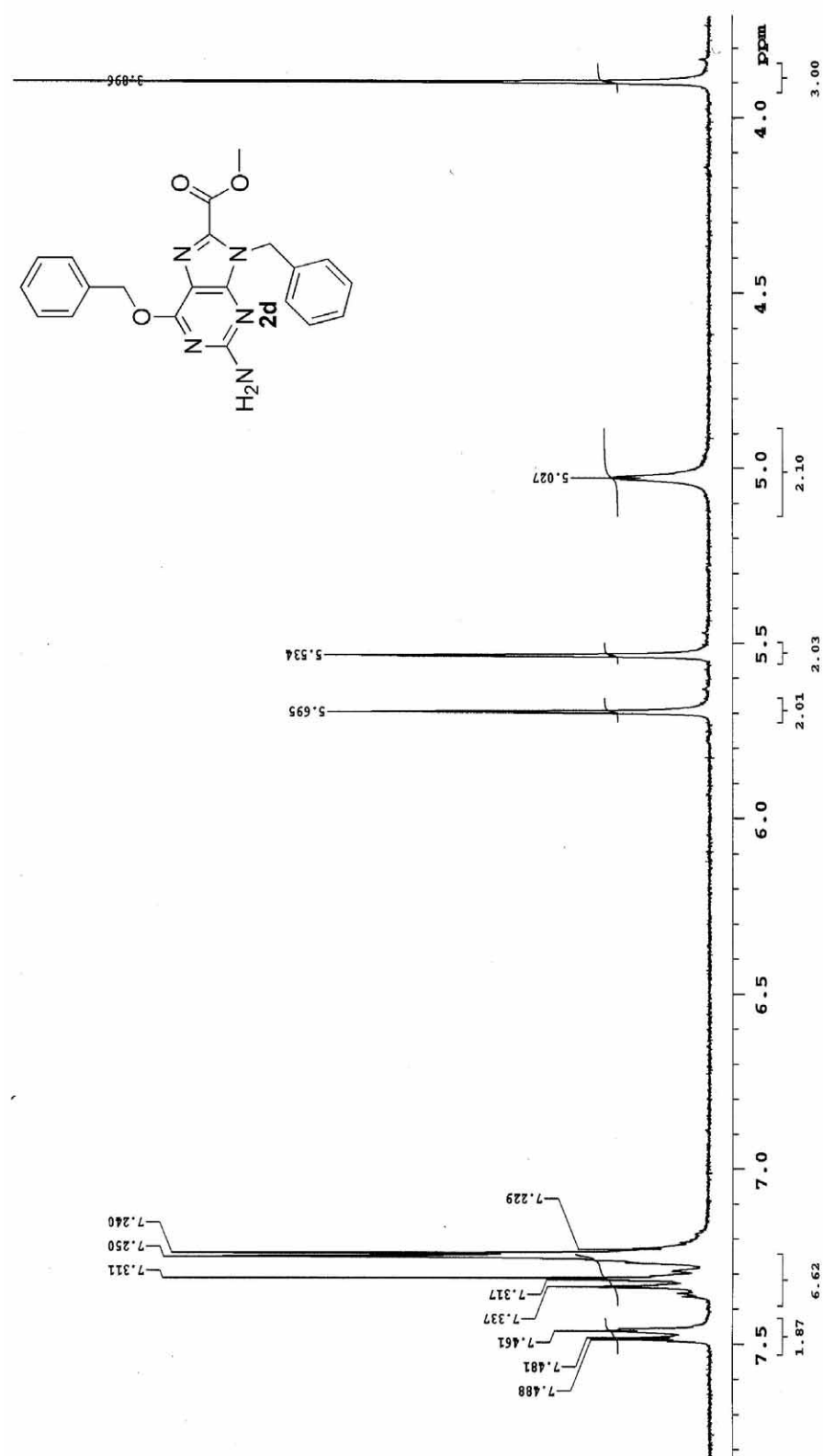


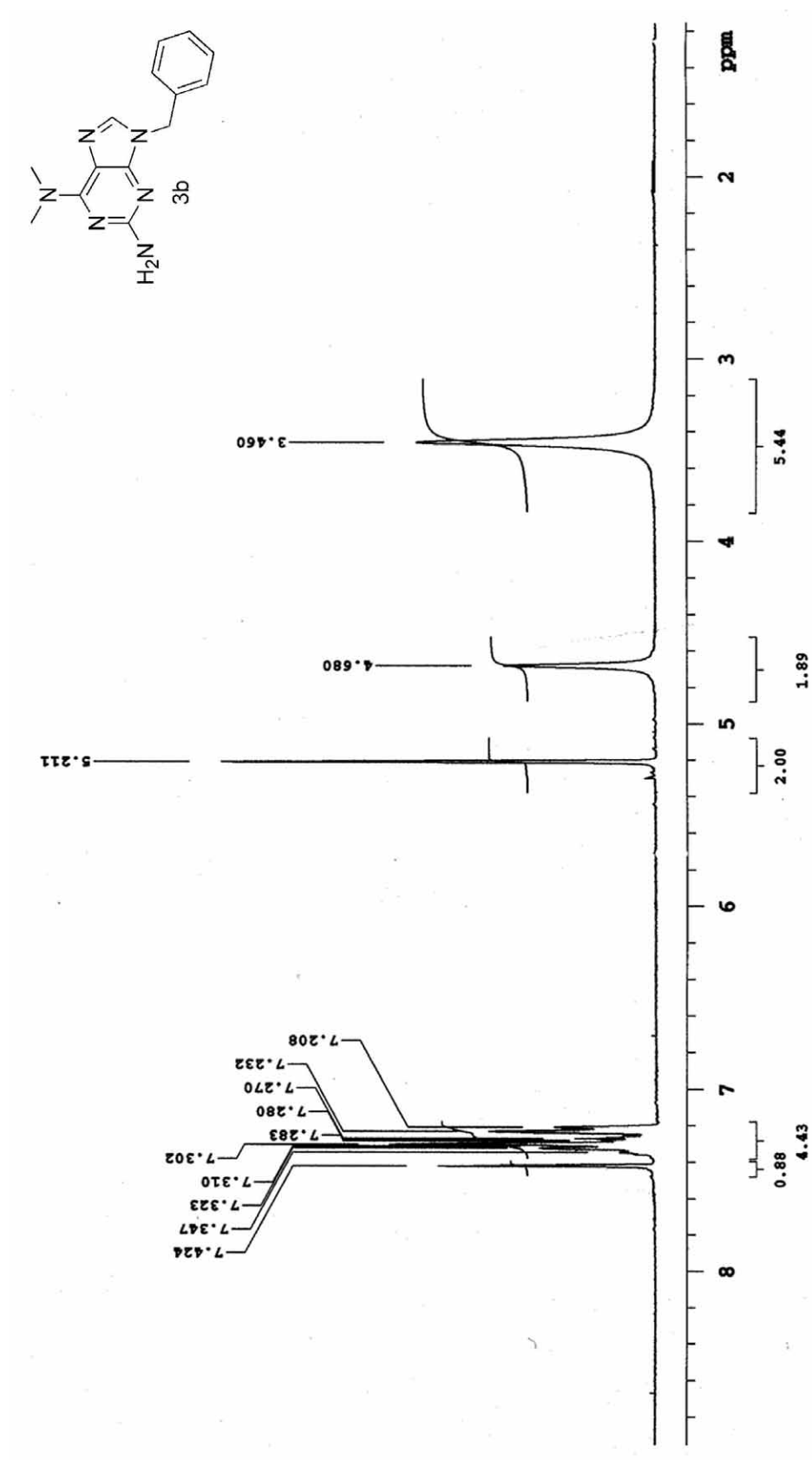


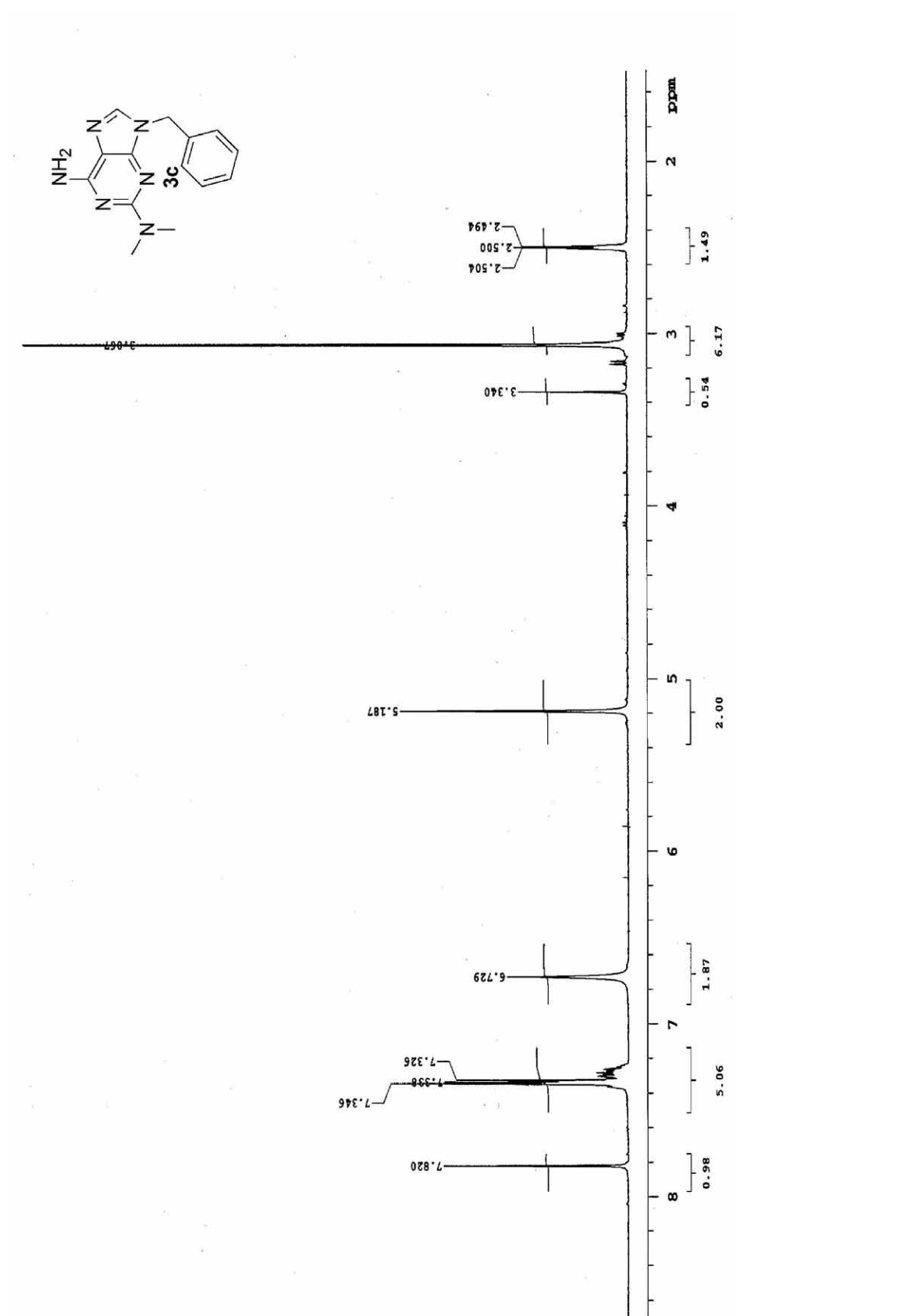




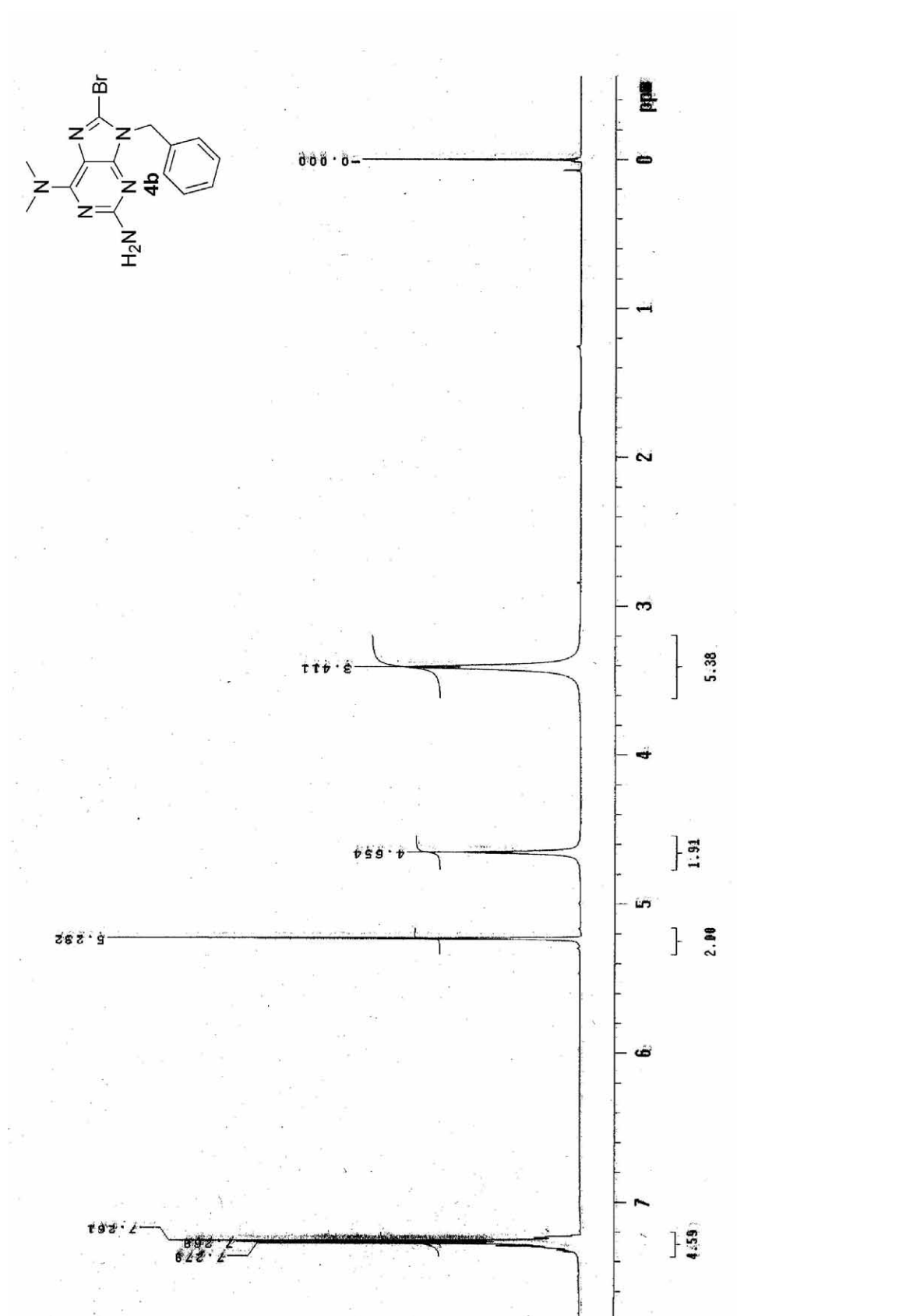


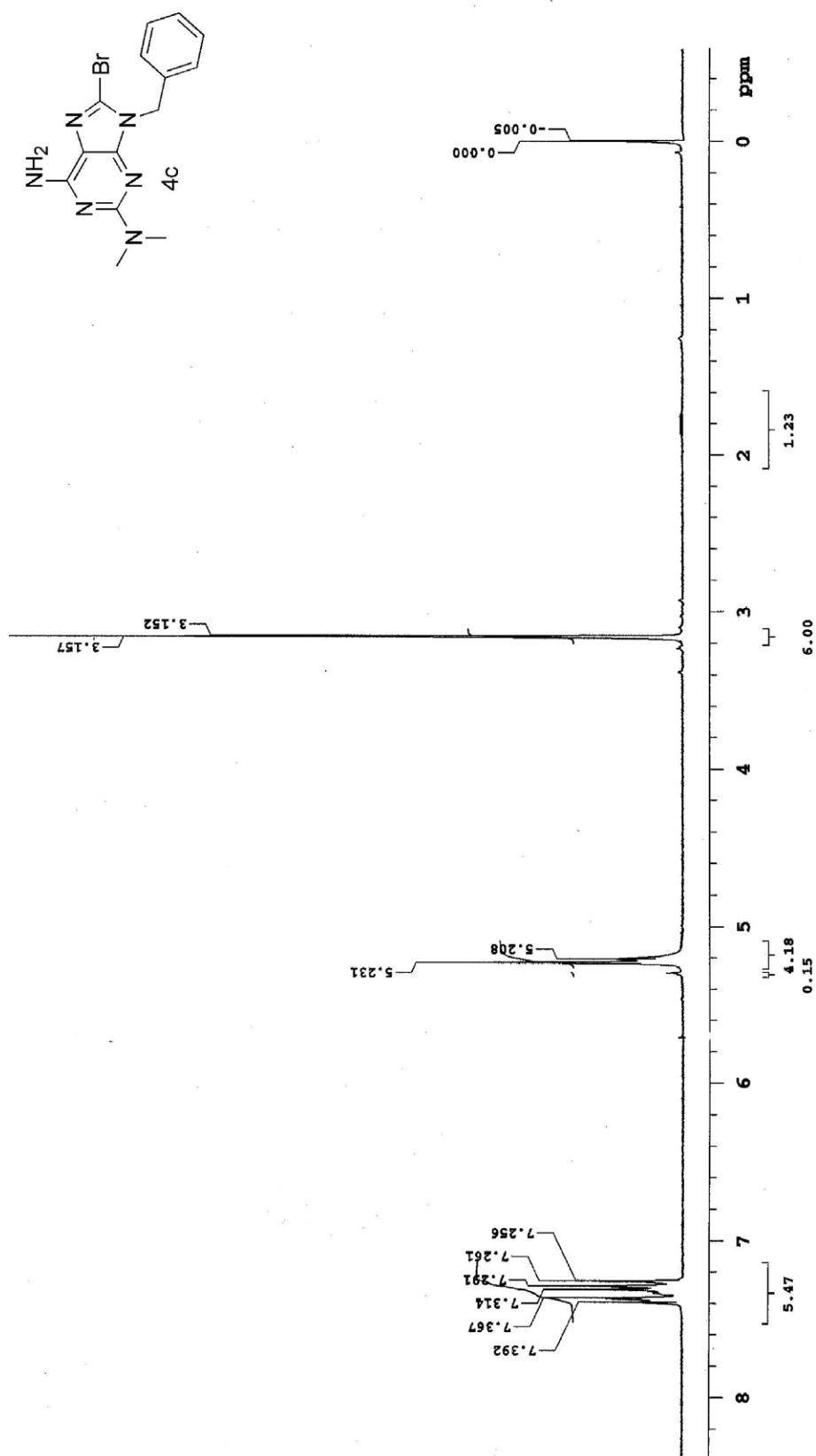


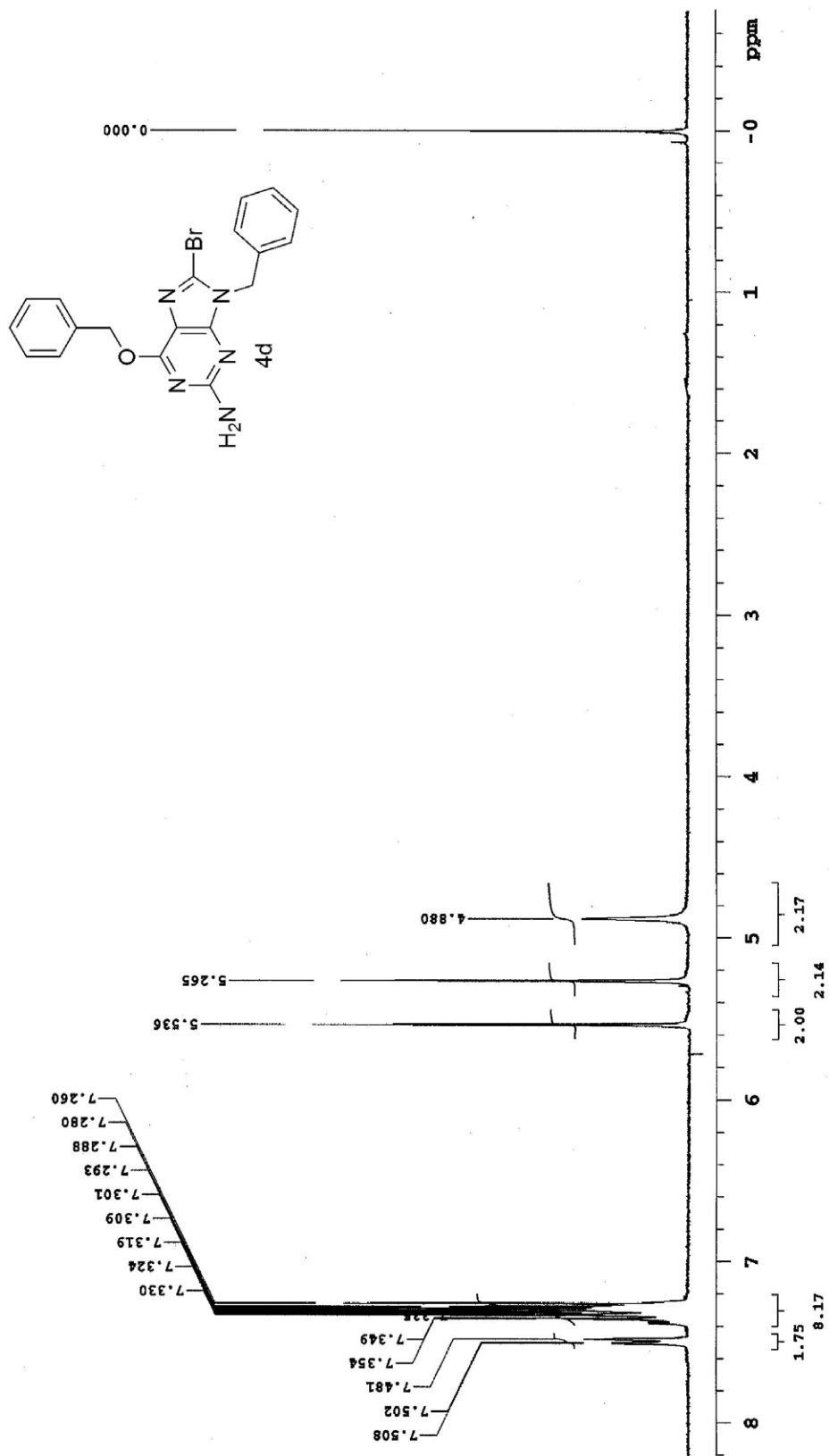




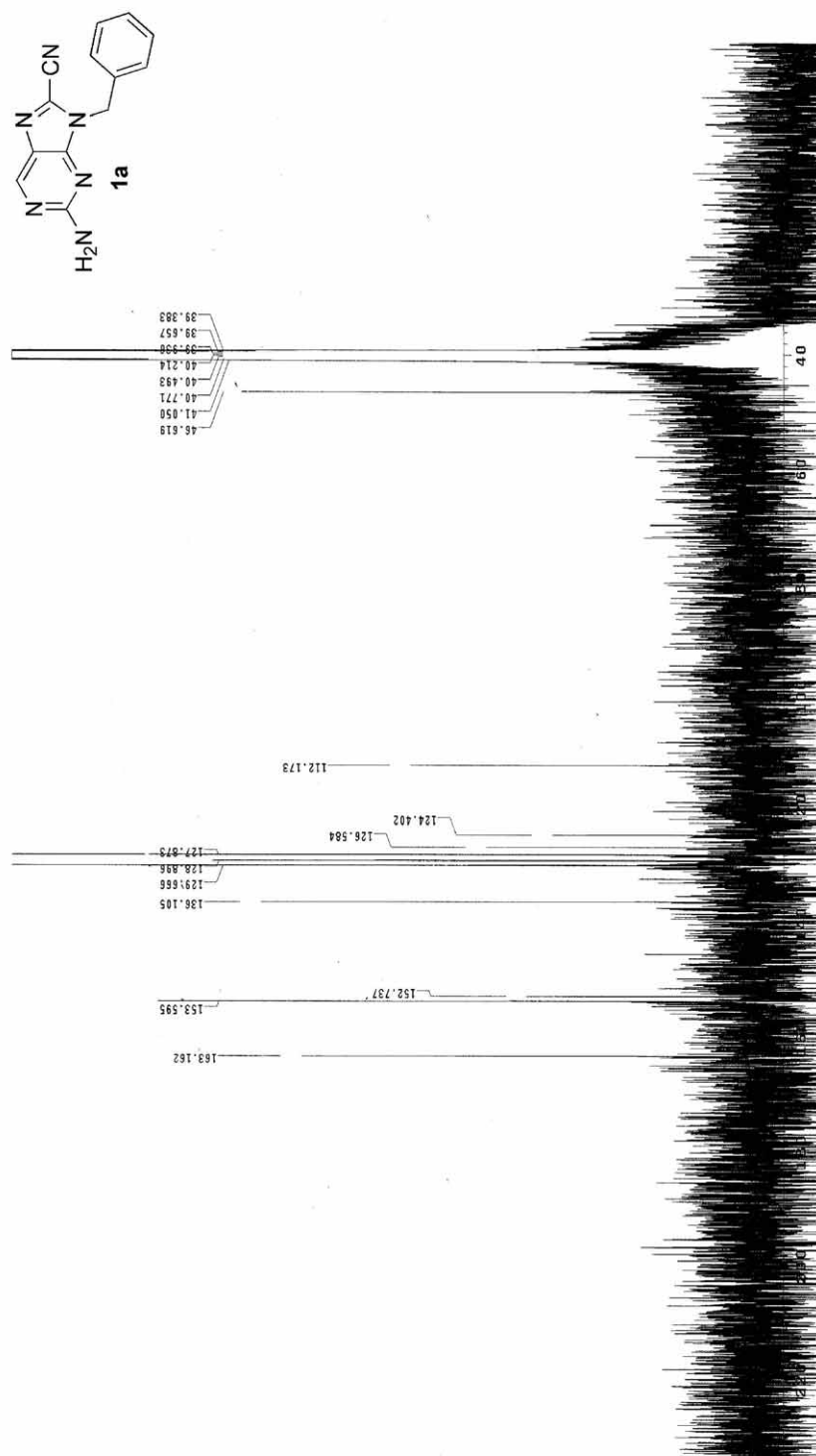


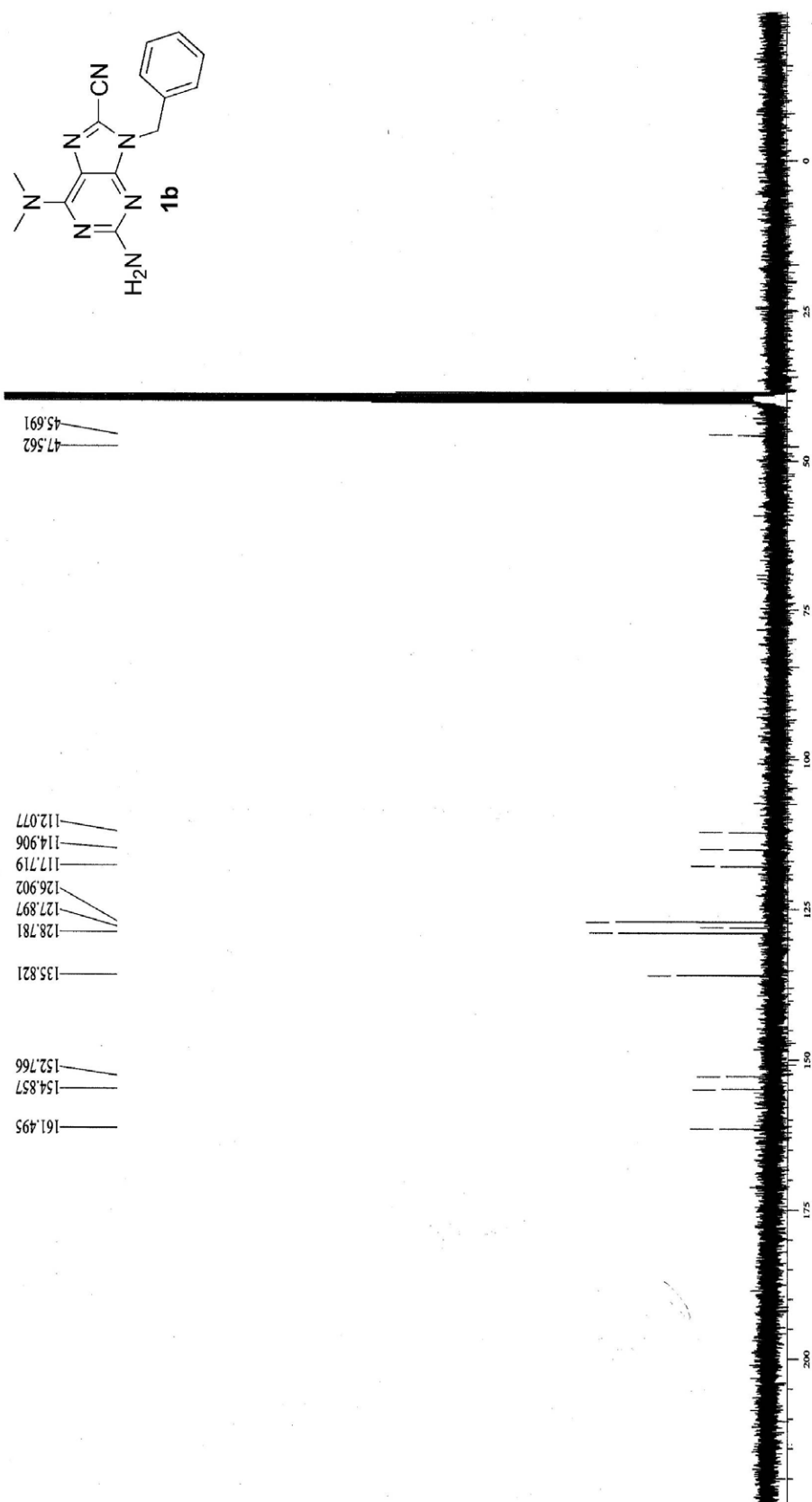


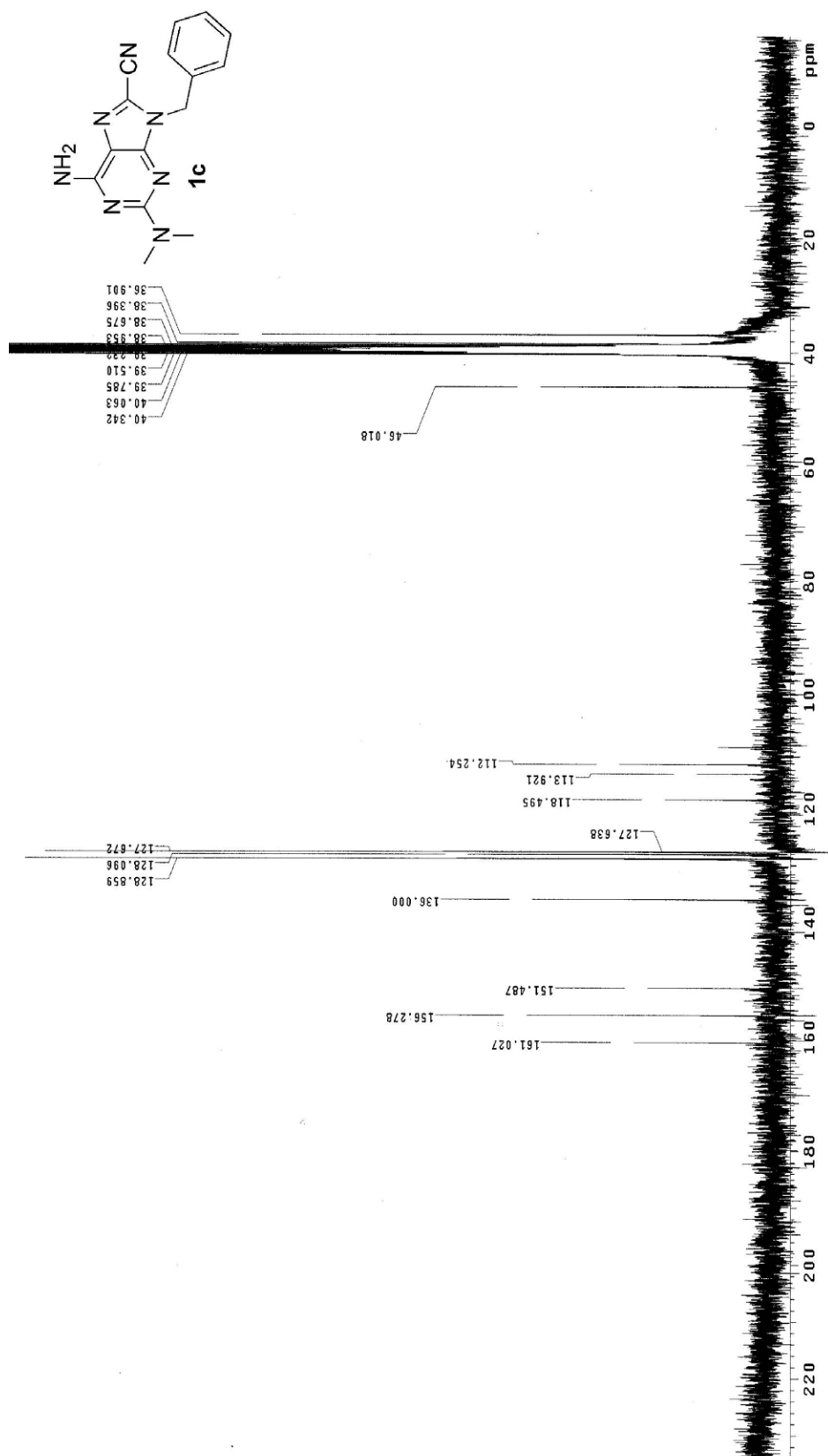


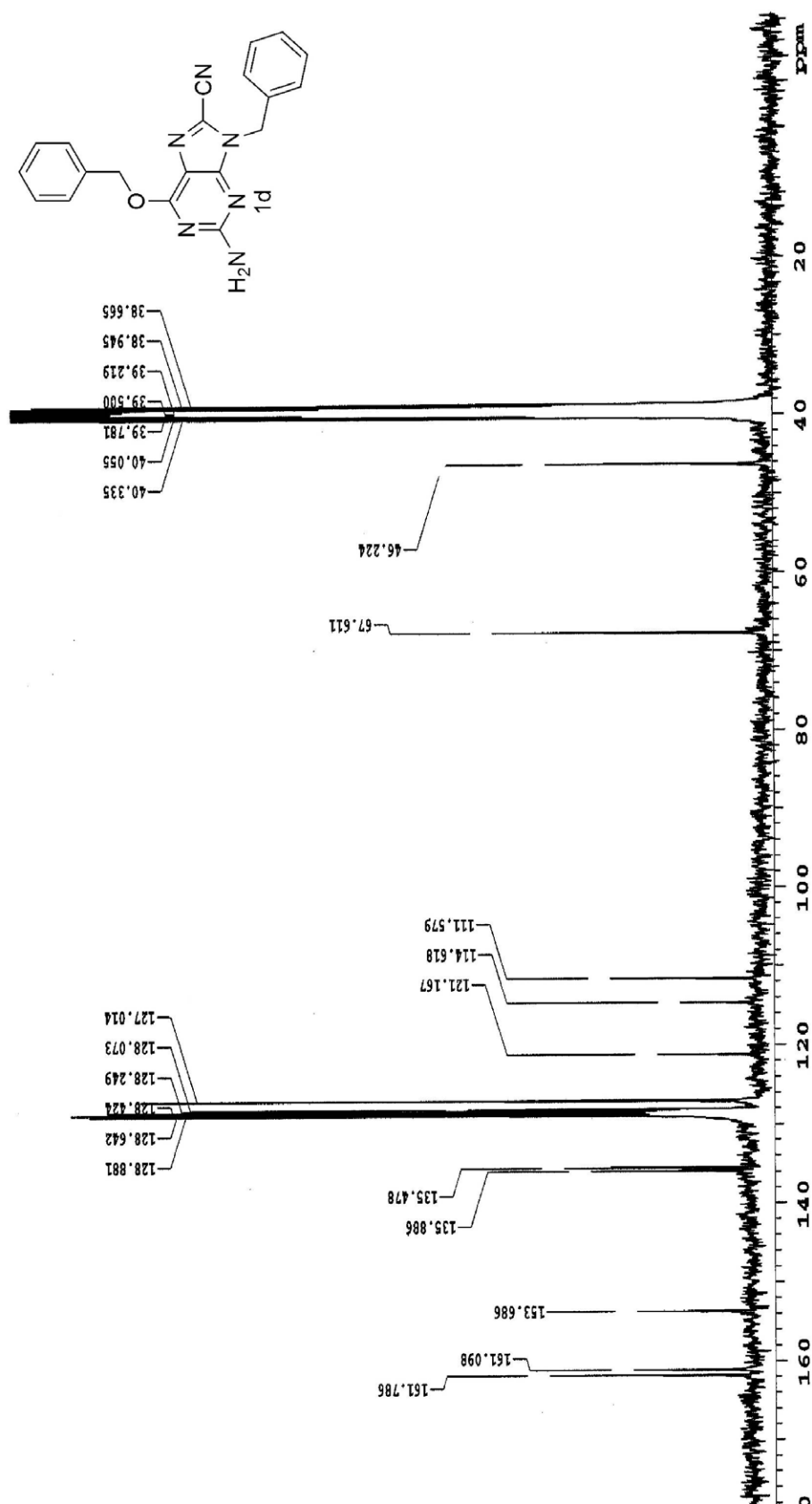


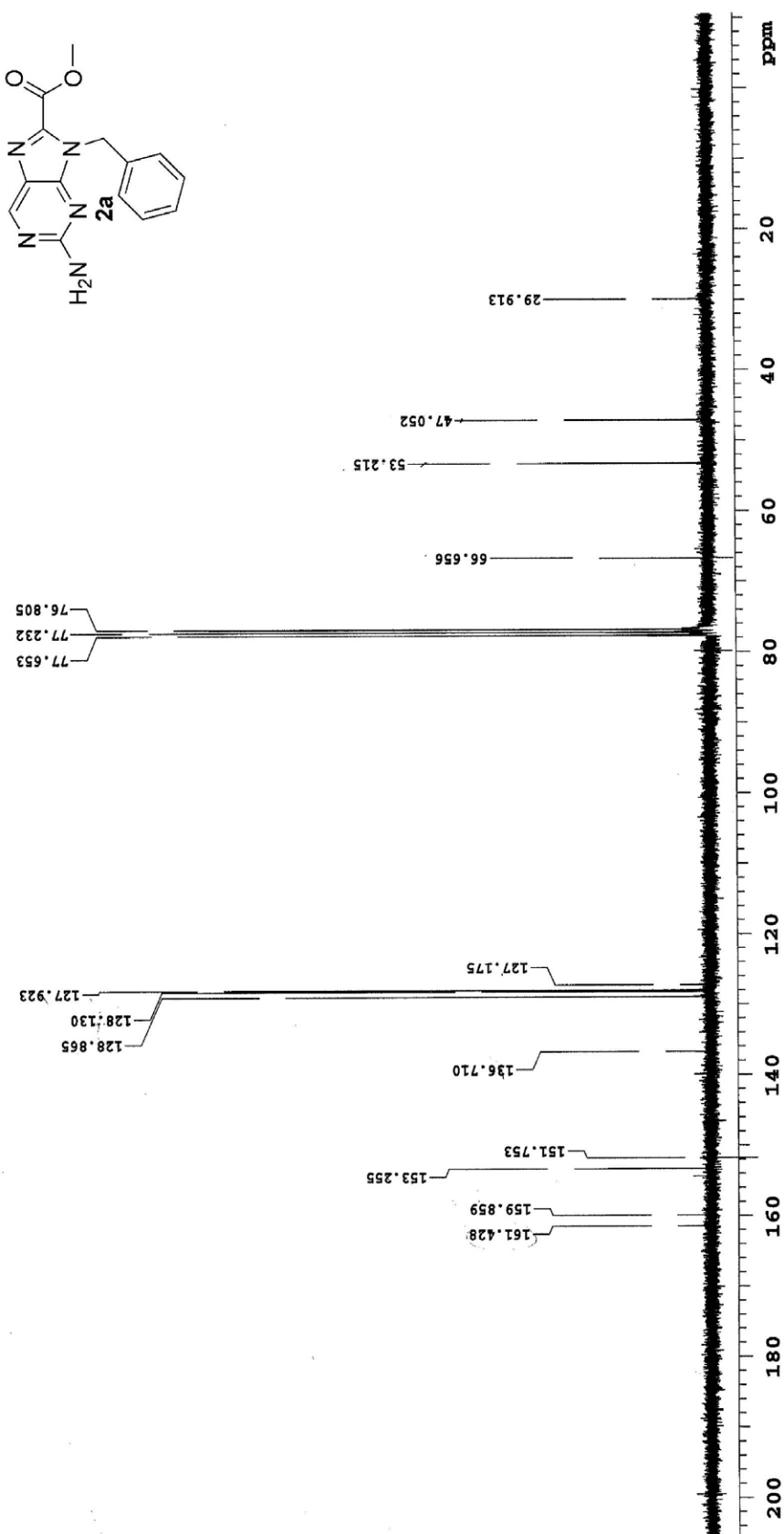
**$^{13}\text{C}$  NMR Spectra**



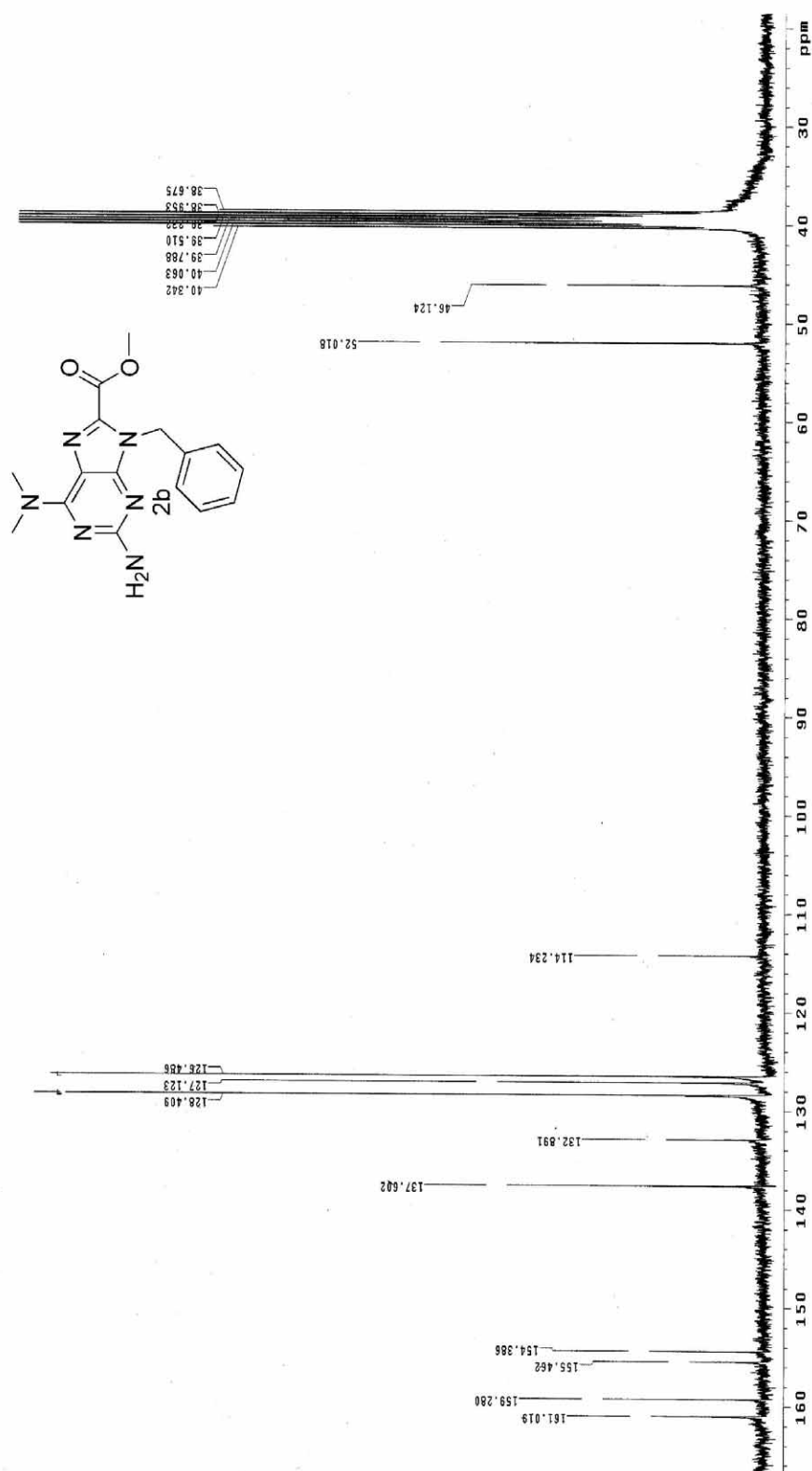


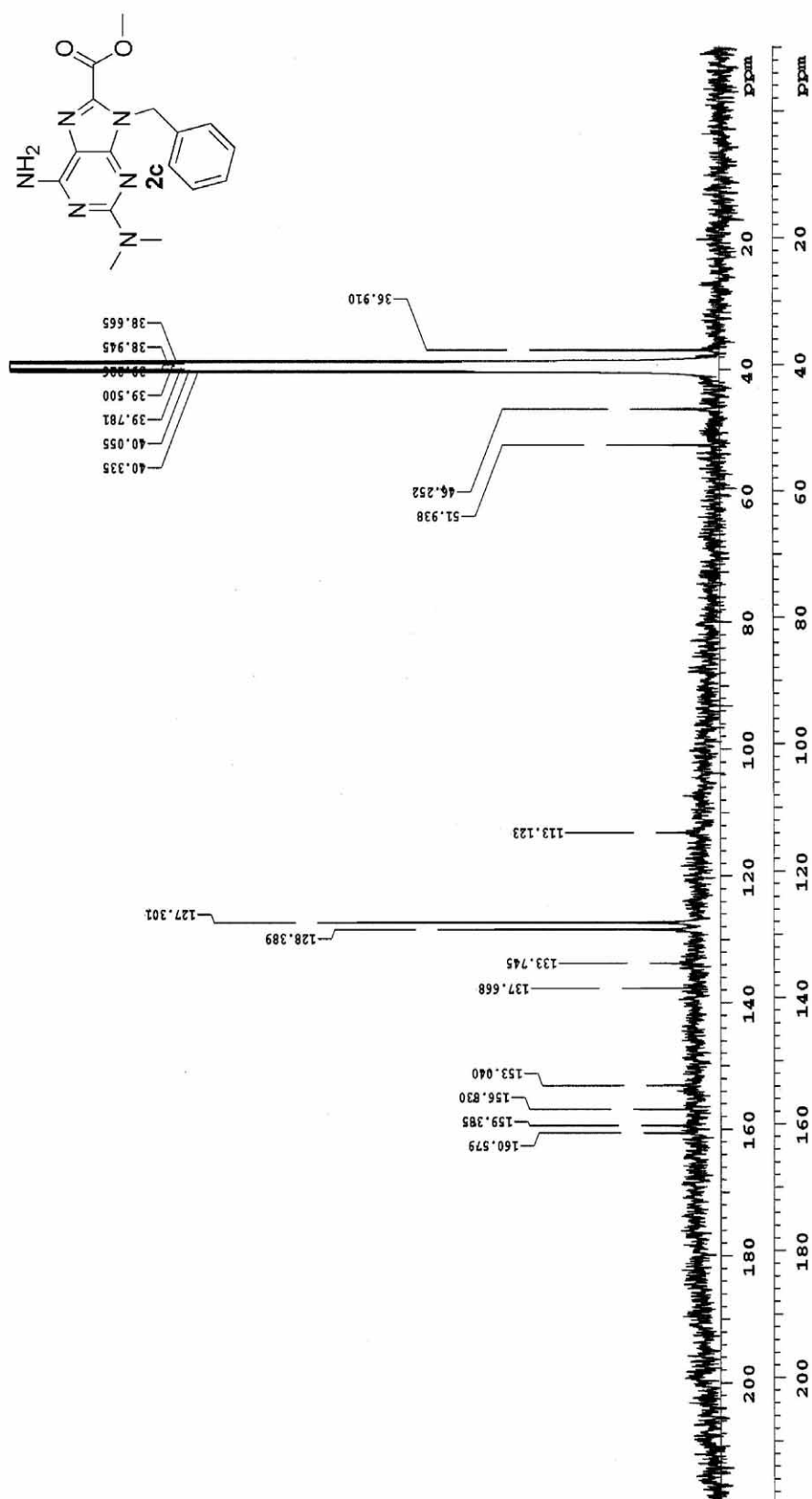


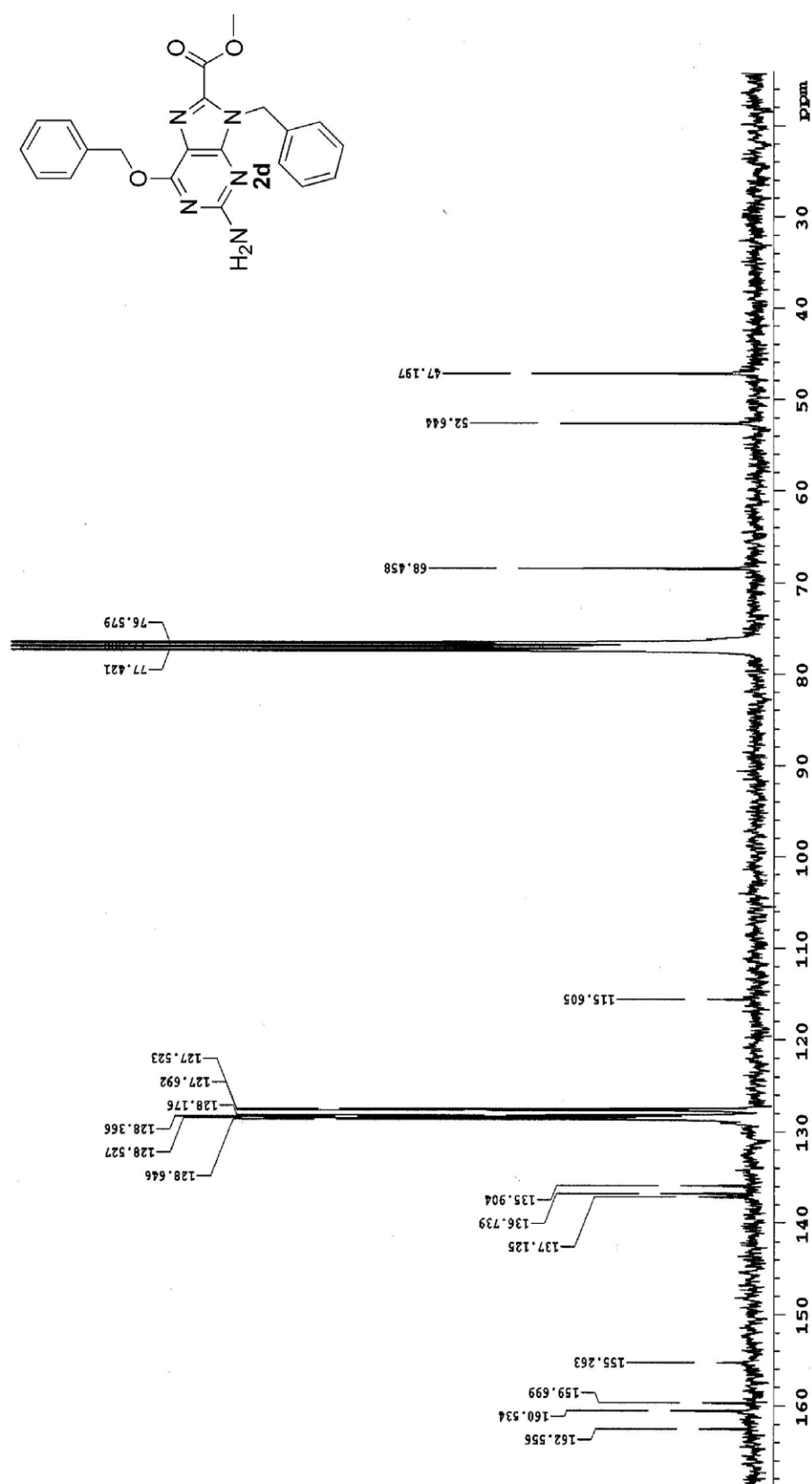


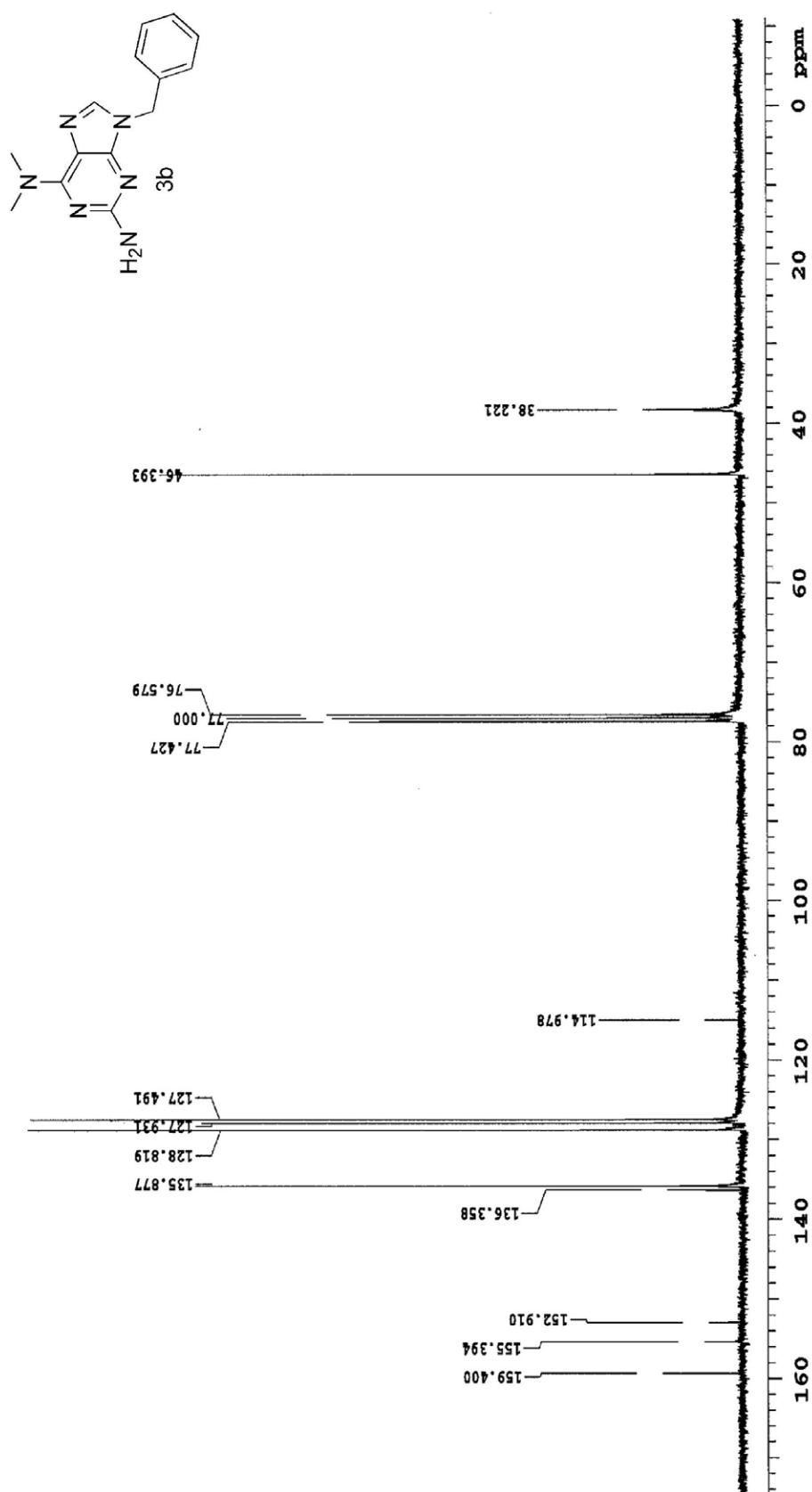


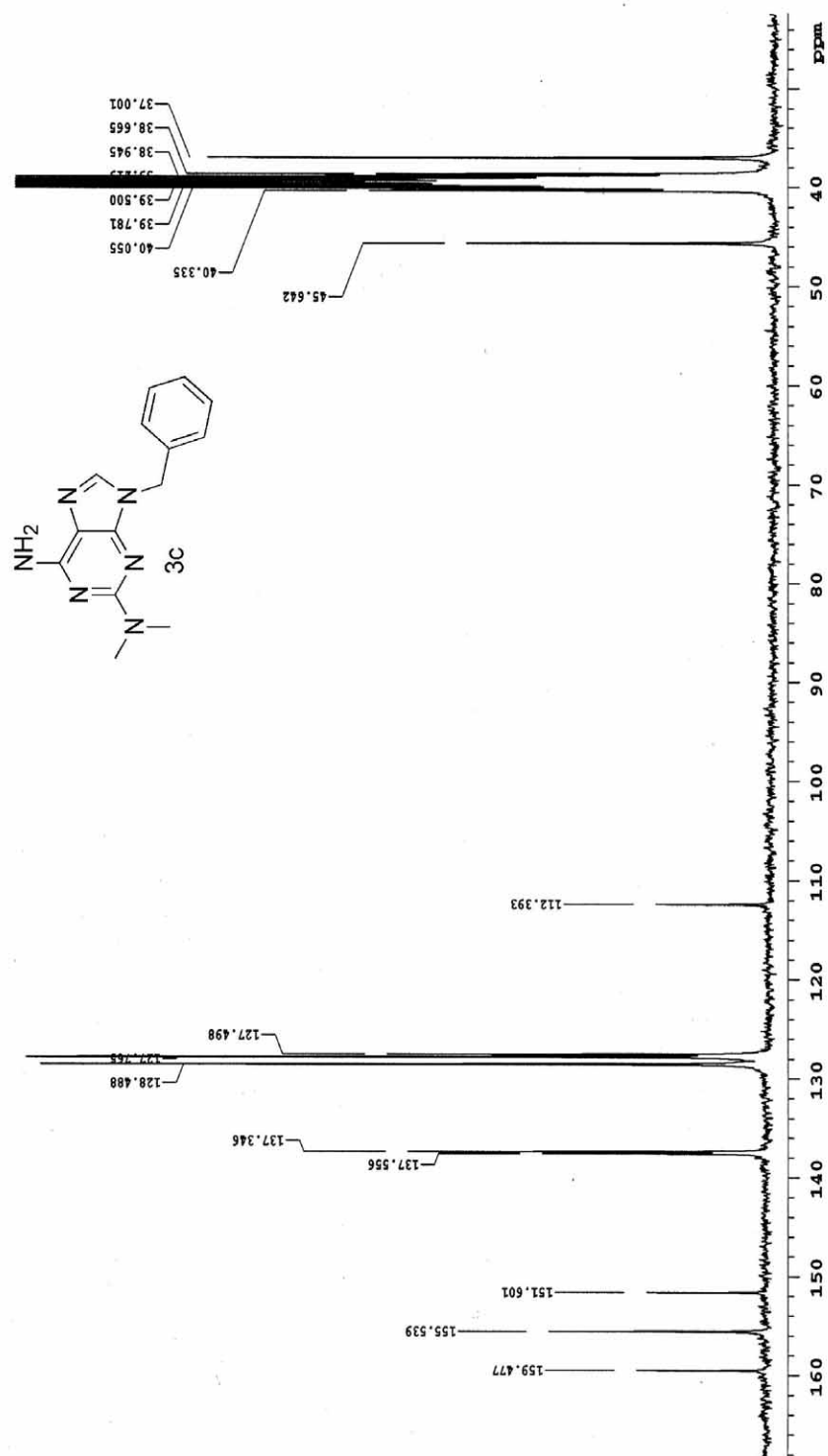


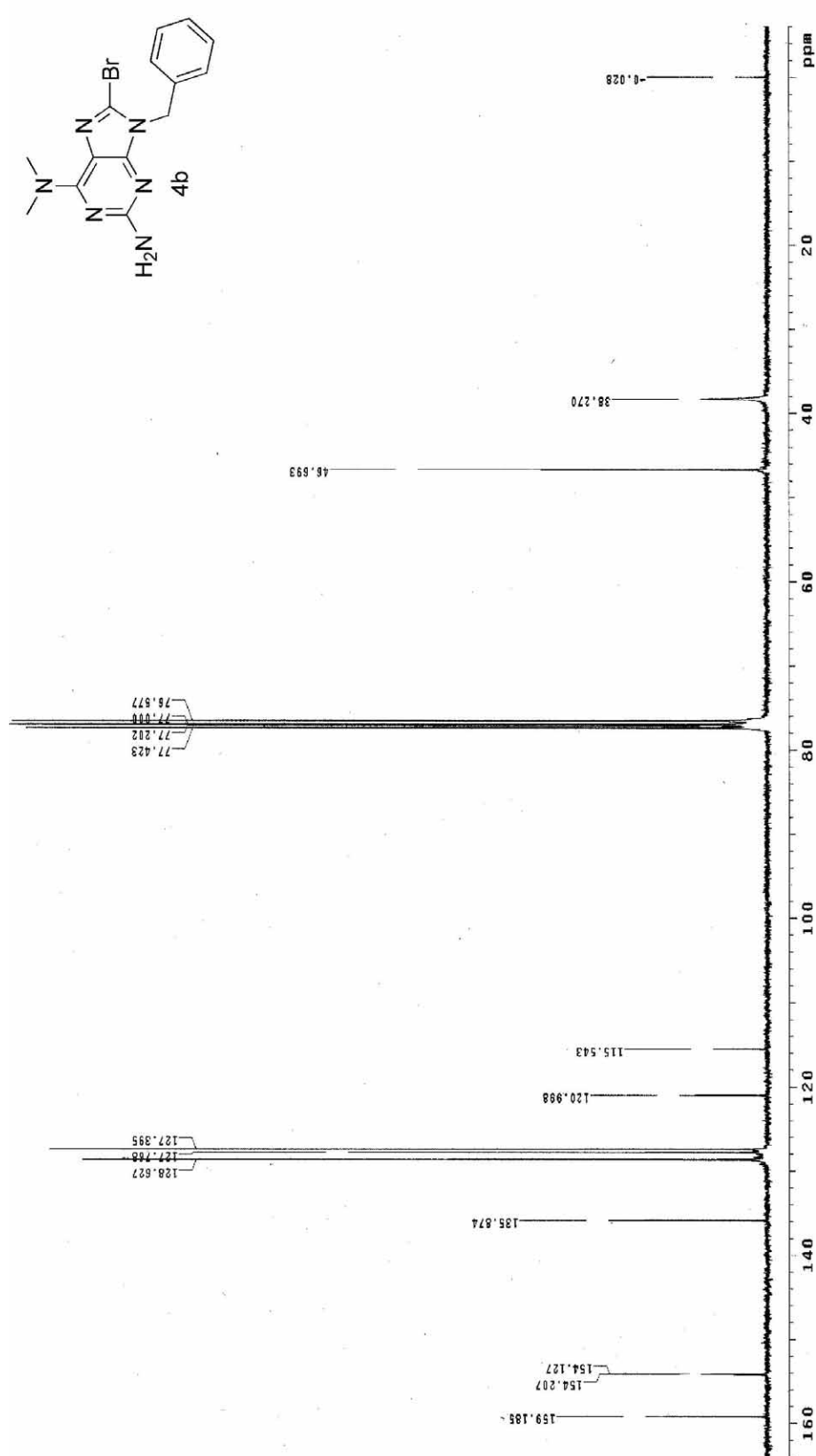


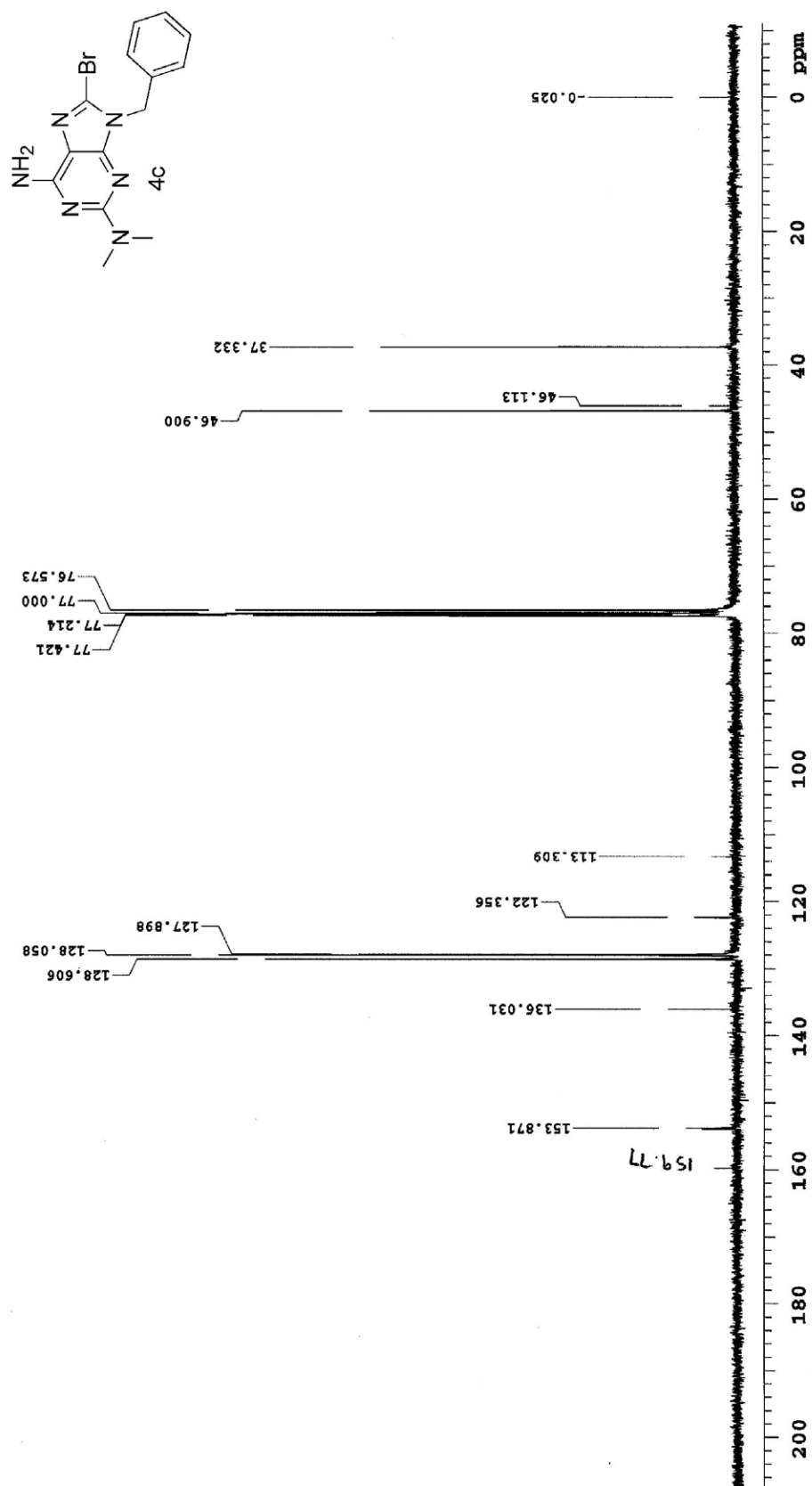


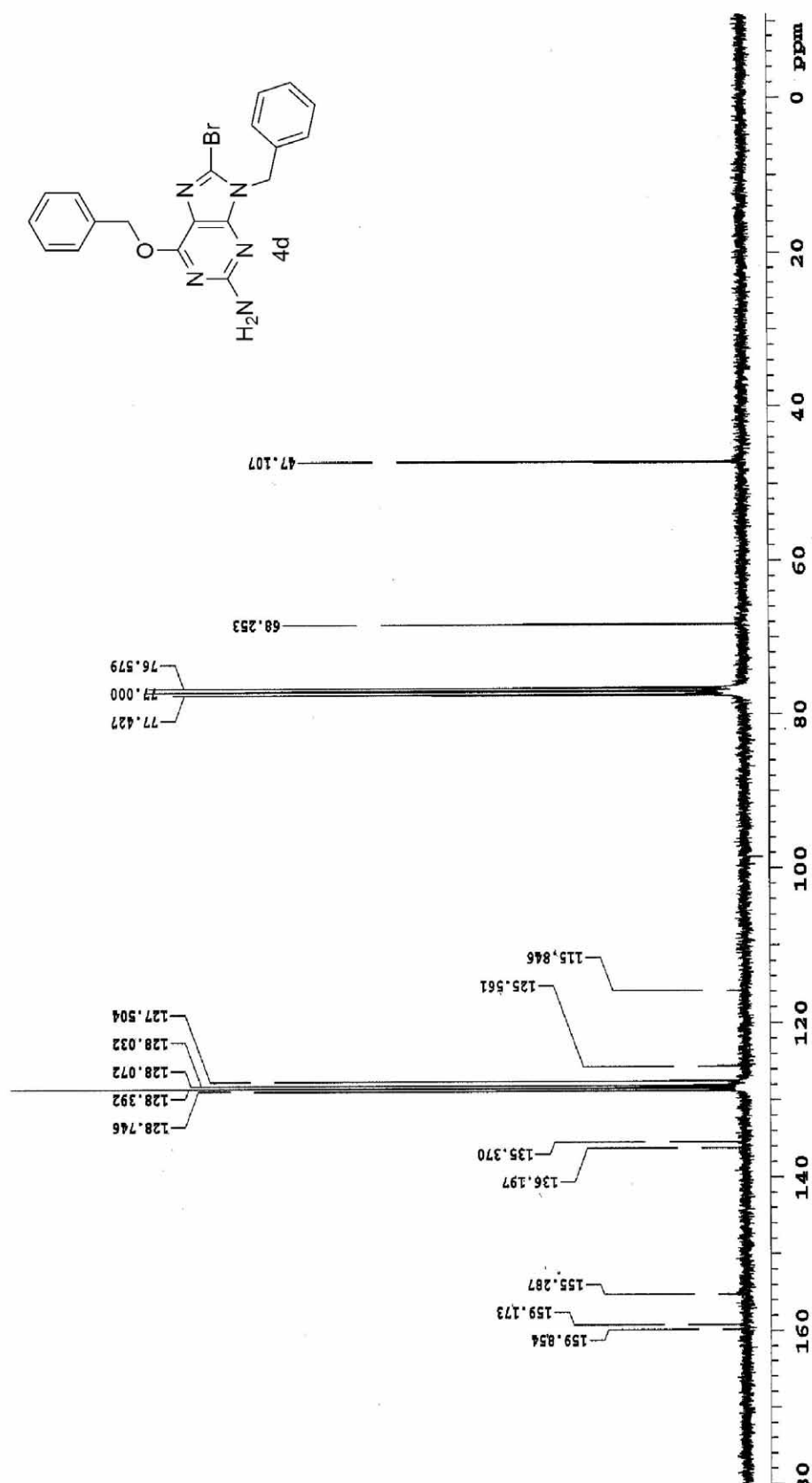














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