

## Supporting Information

### **Ethynyl-substituted benzosiloxaboroles: the role of C( $\pi$ )...B interactions in their crystal packing and use in Huisgen 1,3-dipolar cycloaddition**

P. Pacholak,<sup>a,b</sup> K. Durka\*,<sup>a</sup> K. Woźniak<sup>b</sup> J. Krajewska,<sup>c</sup> A. E. Laudy\*,<sup>c</sup> S. Luliński\*<sup>a</sup>

<sup>[a]</sup> *Faculty of Chemistry, Warsaw University of Technology, Noakowskiego 3, 00-664 Warsaw, Poland*

<sup>[b]</sup> *University of Warsaw, Faculty of Chemistry, Warsaw, Poland, Pasteura 1, 02-093*

<sup>[c]</sup> *Department of Pharmaceutical Microbiology and Bioanalysis, Medical University of Warsaw, Banacha 1b, 02-097 Warsaw, Poland*

Email address: sergiusz.lulinski@pw.edu.pl (S. Luliński).

Email address: alaudy@wp.pl (A. E. Laudy).

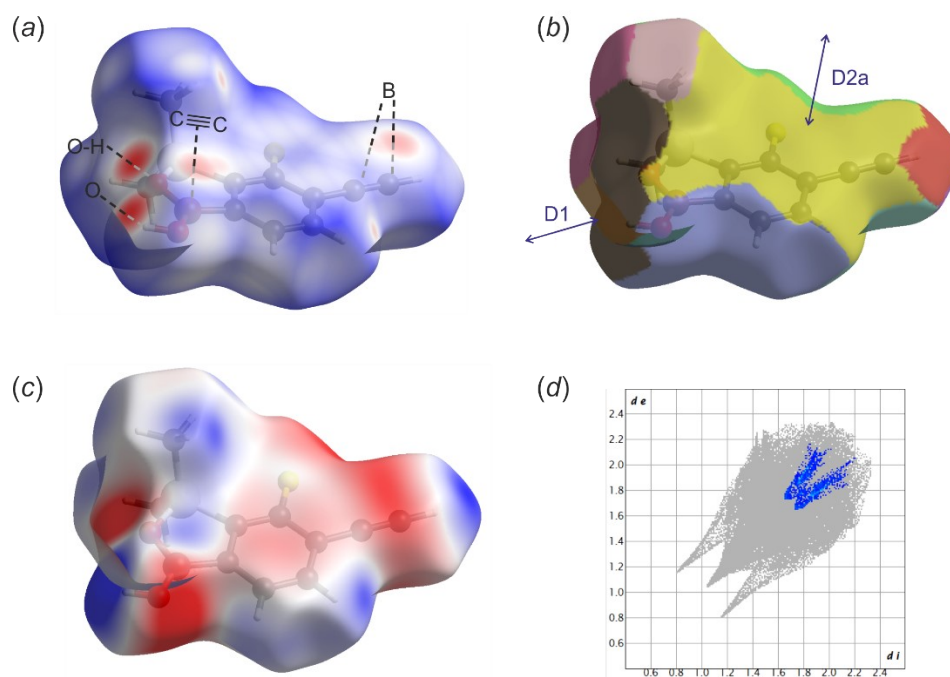
## 1. Structural and computational studies

**Table S1.1.** Selected crystal data, data collection and refinement parameters for **1c**, **1c·MeCN** and **2c**.

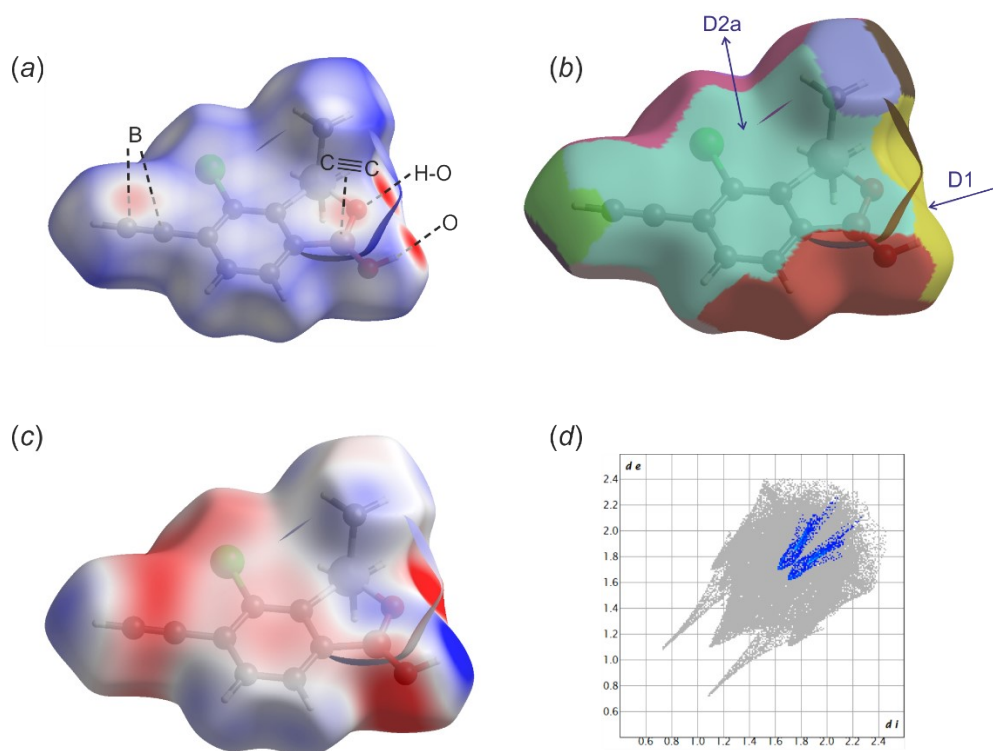
	<b>1c</b>	<b>1c·MeCN</b>	<b>2c</b>
Formula	C <sub>10</sub> H <sub>10</sub> BO <sub>2</sub> SiF	C <sub>12</sub> H <sub>13</sub> BO <sub>2</sub> SiNF	C <sub>10</sub> H <sub>10</sub> O <sub>2</sub> ClSiB
Weight	220.08	261.13	236.53
<i>T</i> / K	100(2)	100(2)	100(2)
Crystal system	triclinic	monoclinic	triclinic
Space group	<i>P</i> -1	<i>P</i> 2 <sub>1</sub> / <i>m</i>	<i>P</i> -1
<i>a</i> / Å	6.9862(2)	8.6127(2)	6.9876(3)
<i>b</i> / Å	7.4451(2)	7.0177(2)	7.7137(4)
<i>c</i> / Å	10.5854(2)	11.1617(3)	10.6833(5)
$\alpha$ / °	92.553(2)	90	85.701(4)
$\beta$ / °	97.855(2)	97.153(2)	81.858(4)
$\gamma$ / °	91.766(2)	90	89.401(4)
<i>V</i> / Å <sup>3</sup>	544.49(2)	669.38(3)	568.42(5)
<i>Z</i>	2	2	2
$\rho_{\text{calc}}$ / g·cm <sup>-3</sup>	1.342	1.296	1.382
$\mu$ / mm <sup>-1</sup>	0.204	0.179	3.788
F(000)	228.0	272.0	244.0
Crystal size / mm <sup>3</sup>	0.342 × 0.129 × 0.089	0.212 × 0.174 × 0.074	0.12 × 0.11 × 0.10
Radiation	MoK $\alpha$ ( $\lambda$ = 0.71073)	MoK $\alpha$ ( $\lambda$ = 0.71073)	Cu K $\alpha$ ( $\lambda$ = 1.54184)
2 $\Theta$ range for data collection / °	3.888 to 57.656	4.766 to 67.666	8.384 to 154.988
Index ranges	-9 ≤ <i>h</i> ≤ 9, -10 ≤ <i>k</i> ≤ 9, -14 ≤ <i>l</i> ≤ 14	-13 ≤ <i>h</i> ≤ 12, -10 ≤ <i>k</i> ≤ 10, -16 ≤ <i>l</i> ≤ 17	-8 ≤ <i>h</i> ≤ 8, -9 ≤ <i>k</i> ≤ 9, -13 ≤ <i>l</i> ≤ 13
Reflections collected	24998	14332	5118
Independent reflections	2683 [ <i>R</i> <sub>int</sub> = 0.0255, <i>R</i> <sub>sigma</sub> = 0.0128]	2733 [ <i>R</i> <sub>int</sub> = 0.0292, <i>R</i> <sub>sigma</sub> = 0.0228]	2404 [ <i>R</i> <sub>int</sub> = 0.0154, <i>R</i> <sub>sigma</sub> = 0.0148]
Data/restraints/parameters	2683/0/139	2733/1/114	2404/1/141
Goodness-of-fit	1.046	1.085	1.084
Final <i>R</i> indexes [I >= 2 $\sigma$ (I)]	<i>R</i> <sub>1</sub> = 0.0284, <i>wR</i> <sub>2</sub> = 0.0768	<i>R</i> <sub>1</sub> = 0.0335, <i>wR</i> <sub>2</sub> = 0.0916	<i>R</i> <sub>1</sub> = 0.0300, <i>wR</i> <sub>2</sub> = 0.0773
Final <i>R</i> indexes [all data]	<i>R</i> <sub>1</sub> = 0.0304, <i>wR</i> <sub>2</sub> = 0.0783	<i>R</i> <sub>1</sub> = 0.0402, <i>wR</i> <sub>2</sub> = 0.0971	<i>R</i> <sub>1</sub> = 0.0302, <i>wR</i> <sub>2</sub> = 0.0775
Largest diff. peak/hole / e·Å <sup>-3</sup>	0.44/-0.23	0.47/-0.34	0.38/-0.37

**Table S1.2.** Selected crystal data, data collection and refinement parameters for **3b**, **4d** and **5b**.

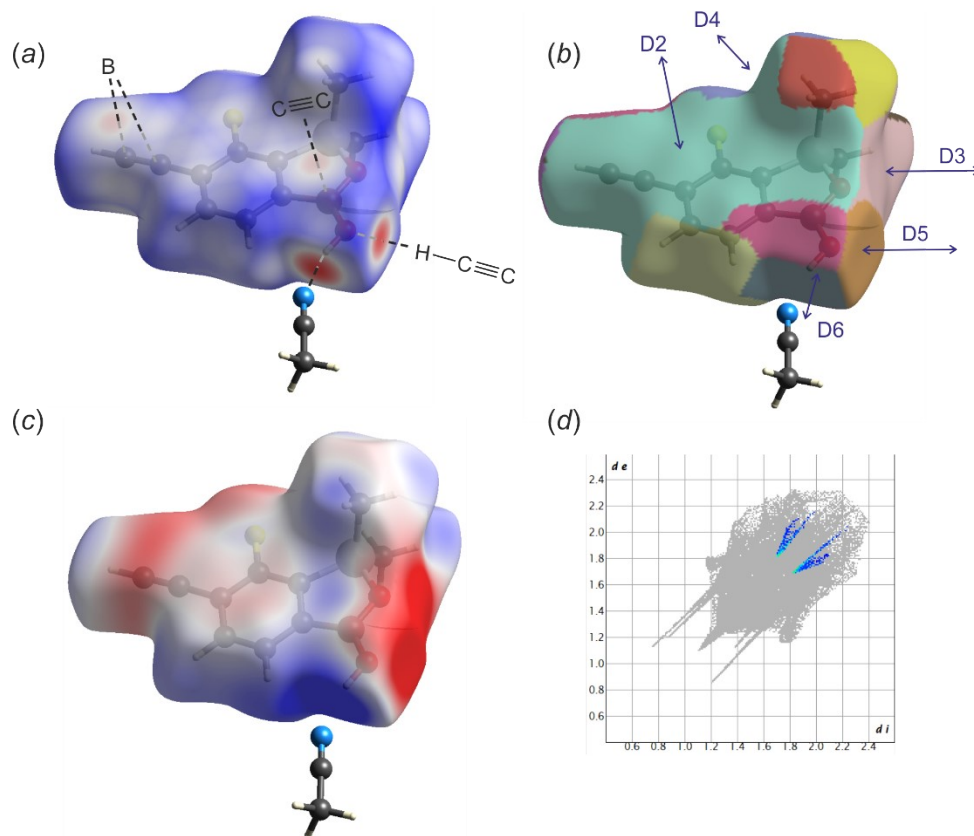
	<b>3b</b>	<b>4d</b>	<b>5b</b>
Formula	C <sub>16</sub> H <sub>16</sub> BN <sub>3</sub> O <sub>5</sub> SiFClS	C <sub>17</sub> H <sub>15</sub> N <sub>3</sub> O <sub>4</sub> SiSBF <sub>2</sub> Cl <sub>3</sub>	C <sub>36</sub> H <sub>40</sub> N <sub>8</sub> O <sub>5</sub> F <sub>2</sub> Si <sub>2</sub> S <sub>2</sub>
Weight	455.73	540.63	823.06
<i>T</i> / K	100(2)	100(2)	100(2)
Crystal system	monoclinic	triclinic	monoclinic
Space group	<i>P2/c</i>	<i>P</i> -1	<i>P2<sub>1</sub>/c</i>
<i>a</i> / Å	15.7378(4)	8.8484(4)	12.2547(2)
<i>b</i> / Å	5.96710(10)	11.1678(4)	21.3124(4)
<i>c</i> / Å	21.2618(5)	12.0146(5)	14.7890(2)
$\alpha$ / °	90	89.089(3)	90
$\beta$ / °	97.318(2)	77.254(3)	91.9810(10)
$\gamma$ / °	90	80.690(3)	90
<i>V</i> / Å <sup>3</sup>	1980.41(8)	1142.50(8)	3860.24(11)
<i>Z</i>	4	2	4
$\rho_{\text{calc}}$ / g·cm <sup>-3</sup>	1.528	1.572	1.416
$\mu$ / mm <sup>-1</sup>	3.677	0.592	2.388
F(000)	936.0	548.0	1720.0
Crystal size / mm <sup>3</sup>	0.15 × 0.11 × 0.10	0.21 × 0.107 × 0.104	0.231 × 0.146 × 0.087
Radiation	CuK $\alpha$ ( $\lambda$ = 1.54178)	MoK $\alpha$ ( $\lambda$ = 0.71073)	CuK $\alpha$ ( $\lambda$ = 1.54178)
2 $\Theta$ range for data collection / °	5.662 to 145.38	4.784 to 60.008	7.218 to 153.4
Index ranges	-19 ≤ <i>h</i> ≤ 19, -7 ≤ <i>k</i> ≤ 7, -21 ≤ <i>l</i> ≤ 26	-11 ≤ <i>h</i> ≤ 12, -15 ≤ <i>k</i> ≤ 15, -16 ≤ <i>l</i> ≤ 16	-15 ≤ <i>h</i> ≤ 15, -26 ≤ <i>k</i> ≤ 26, -18 ≤ <i>l</i> ≤ 12
Reflections collected	15611	19660	22446
Independent reflections	3890 [ <i>R</i> <sub>int</sub> = 0.0270, <i>R</i> <sub>sigma</sub> = 0.0222]	6007 [ <i>R</i> <sub>int</sub> = 0.0317, <i>R</i> <sub>sigma</sub> = 0.0475]	8052 [ <i>R</i> <sub>int</sub> = 0.0357, <i>R</i> <sub>sigma</sub> = 0.0380]
Data/restraints/parameters	3890/1/265	6007/1/313	8052/0/496
Goodness-of-fit	1.040	1.035	1.017
Final <i>R</i> indexes [ <i>I</i> ≥ 2 $\sigma$ ( <i>I</i> )]	<i>R</i> <sub>1</sub> = 0.0298, <i>wR</i> <sub>2</sub> = 0.0764	<i>R</i> <sub>1</sub> = 0.0489, <i>wR</i> <sub>2</sub> = 0.1085	<i>R</i> <sub>1</sub> = 0.0445, <i>wR</i> <sub>2</sub> = 0.1170
Final <i>R</i> indexes [all data]	<i>R</i> <sub>1</sub> = 0.0360, <i>wR</i> <sub>2</sub> = 0.0810	<i>R</i> <sub>1</sub> = 0.0670, <i>wR</i> <sub>2</sub> = 0.1181	<i>R</i> <sub>1</sub> = 0.0497, <i>wR</i> <sub>2</sub> = 0.1226
Largest diff. peak/hole / e·Å <sup>-3</sup>	0.35/-0.37	0.74/-0.68	0.56/-0.36



**Figure S1.1** Hirshfeld surfaces generated for **1c** with mapped (a)  $d_{\text{norm}}$  property value over the range  $-0.50$  to  $1.30$ , (b) fragment patch, (c) electrostatic potential over the range  $-0.06$  to  $0.08$  a.u. (d) Fingerprint plots with marked C...B contacts.



**Figure S1.2.** Hirshfeld surfaces generated for **2c** with mapped (a)  $d_{\text{norm}}$  property value over the range  $-0.50$  to  $1.30$ , (b) fragment patch, (c) electrostatic potential over the range  $-0.06$  to  $0.08$  a.u. (d) Fingerprint plots with marked C...B contacts.



**Figure S1.3.** Hirshfeld surfaces generated for **1c**·MeCN with mapped (a)  $d_{\text{norm}}$  property value over the range  $-0.50$  to  $1.30$ , (b) fragment patch, (c) electrostatic potential over the range  $-0.06$  to  $0.08$  a.u. (d) Fingerprint plots with marked C...B contacts.

**Table S1.3.** The intermolecular donor-acceptor orbital  $C(\pi)\dots B$  interaction energies ( $\text{kJ}\cdot\text{mol}^{-1}$ ) estimated by 2<sup>nd</sup>-order perturbation theory within NBO analysis.

Structure	<b>1c</b>		<b>2c</b>		<b>1c</b> ·MeCN
Motif	D2a	D2b	D2a	D2b	D2
$E(\pi_{\text{CC}(\text{ethynyl})} \rightarrow \text{p}_B)$	5.1	2.6	5.4	2.3	2.7
$E(\pi_{\text{CC}(\text{aromatic})} \rightarrow \pi^*_{\text{CC}(\text{ethynyl})})$	0.6	0.8	0.6	1.0	0.5

**Table S1.4.** Electron density ( $\rho$ ,  $\text{e}\cdot\text{\AA}^{-3}$ ) and negative Laplacian ( $\nabla^2\rho$ ,  $\text{e}\cdot\text{\AA}^{-5}$ ) at BCPs of  $C(\pi)\dots B$  interactions.

Structure	<b>1c</b>		<b>2c</b>		<b>1c</b> ·MeCN
Motif	D2a	D2b	D2a	D2b	D2
$\rho$	0.033	0.035	0.035	0.036	0.036
$\nabla^2\rho$	0.33	0.36	0.35	0.36	0.35

## 2. Antimicrobial activity.

**Table S2.1.** The antibacterial activity of tested agents against standard Gram-positive strains

Agent tested	MIC in mg·L <sup>-1</sup> [MBC in mg·L <sup>-1</sup> ] <sup>a</sup> (Diameter of inhibition zone in mm)					
	<i>S. aureus</i> ATCC 6538P	<i>S. aureus</i> ATCC 43300 MRSA	<i>S. epidermidis</i> ATCC 12228	<i>E. faecalis</i> ATCC 29212	<i>E. faecium</i> ATCC 6057	<i>B. subtilis</i> ATCC 6633 <sup>b</sup>
<b>1c</b>	25 [50/400] <sup>c</sup> (25)	25 [50] (20)	<b>12.5</b> [25/>400] <sup>c</sup> (29)	100 (15)	100 (16)	NT (26)
<b>2c</b>	<b>12.5</b> [25/200] <sup>c</sup> (21)	<b>12.5</b> [25] (23)	<b>12.5</b> [400] (26)	50 (20)	50 (21)	NT (27)
<b>3a</b>	25 (22)	50 (20)	50 (24)	200 (-)	200 (11)	NT (21)
<b>3b</b>	50 (20)	50 (22)	50 (23)	200 (-)	200 (-)	NT (21)
<b>3c</b>	<b>12.5</b> [25/>400] <sup>c</sup> (22)	<b>12.5</b> [25] (23)	<b>12.5</b> [25/>400] <sup>c</sup> (29)	100 (16)	50 (16)	NT (27)
<b>4a</b>	25 (20)	50 (20)	50 (25)	200 (-)	200 (-)	NT (23)
<b>4b</b>	25 (25)	50 (21)	50 (20)	200 (-)	200 (-)	NT (18)
<b>4c</b>	25 (20)	50 (22)	50 (24)	200 (-)	200 (-)	NT (20)
<b>5a</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	NT (-)
<b>5b</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	NT (-)
<b>LIN<sup>d</sup></b>	1 [>128] (25)	2 [>128] (25)	1 [>128] (26)	2 [>128] (15)	2 [>128] (14)	NT (30)

The highest activity against Gram-positive bacteria indicated by the low MIC values ( $\leq 12.5$  mg·L<sup>-1</sup>) is shown in boldface.

(-): The inhibition zone was not observed in the disc-diffusion method. The diameter of the paper discs was 9 mm.

<sup>a</sup> Only the MBC values  $\leq 400$  mg·L<sup>-1</sup> are presented.

<sup>b</sup> The growth type of *B. subtilis* in the MHB medium prevented reading the MIC values of tested substances.

<sup>c</sup> The Eagle effect [1,2] was observed during the determination of the MBC value of the same tested agents against *Staphylococcus* spp. strains. The Eagle effect is shown in the italic face.

<sup>d</sup> LIN, linezolid was used as a reference agent active against Gram-positive bacteria. The diameter of a commercial disc containing 0.03 mg of linezolid was 6 mm; the MIC of linezolid was determined according to the CLSI recommendations [3].

**Table S2.2.** The antibacterial activity of tested agents against standard Gram-negative strains.

Agent tested	MIC in mg·L <sup>-1</sup> [MBC in mg·L <sup>-1</sup> ] <sup>a</sup> / x-fold reduction of MIC in the presence of PAβN <sup>b</sup> (Diameter of inhibition zone in mm)										
	<i>E. coli</i> ATCC 25922	<i>K. pneumoniae</i> ATCC 13883	<i>P. mirabilis</i> ATCC 12453	<i>E. cloacae</i> DSM 6234	<i>S. marcescens</i> ATCC 13880	<i>A. baumannii</i> ATCC 19606	<i>P. aeruginosa</i> ATCC 27853	<i>S. maltophilia</i> ATCC 13637	<i>S. maltophilia</i> ATCC 12714	<i>B. cepacia</i> ATCC 25416 <sup>c</sup>	<i>B. bronchiseptica</i> ATCC 4617 <sup>c</sup>
<b>1c</b>	400 [400]/4 (13)	400 [400]/2 (-)	400/2 (-)	400/2 (-)	>400/8 (-)	>400 (-)	>400 (-)	200 [400]/2 (11)	100 [400] (12)	>400 (-)	400 (-)
<b>2c</b>	>400/8 (-)	>400 (-)	>400/2 (-)	>400/2 (-)	>400/4 (-)	>400/4 (-)	>400 (-)	400 [400]/8 (13)	400/4 (-)	>400 (-)	400 (-)
<b>3a</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>3b</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400/2 (-)	>400 (-)	>400 (-)	400 (-)
<b>3c</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	400/4 (15)	>400/4 (-)	>400 (-)	>400 (-)
<b>4a</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>4b</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400/2 (-)	>400/2 (-)	>400 (-)	400 (-)
<b>4c</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400/2 (-)	>400 (-)	>400 (-)	>400 (-)
<b>5a</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>5b</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>Nf<sup>d</sup></b>	8 [8] (24)	32 [32] (23)	128 [ <b>&gt;128</b> ] (9)	32 [32] (17)	128 [ <b>&gt;128</b> ] (12)	64 [128] (9)	>128 [ <b>&gt;128</b> ] (-)	128 [ <b>&gt;128</b> ] (-)	128 [ <b>&gt;128</b> ] (-)	32 [32] (12)	64 [128] (-)

PAβN: efflux pump inhibitor. The significant decreases (at least a 4-fold) in the MIC values of tested compounds after the addition of PAβN are shown in boldface. The test was performed in the MHB medium supplemented with 1 mM MgSO<sub>4</sub>.

(-): The inhibition zone was not observed in the disc-diffusion method. The diameter of the paper discs was 9 mm.

<sup>a</sup> Only the MBC values ≤400 mg·L<sup>-1</sup> are presented.

<sup>b</sup> In the table, only at least 2-fold decreases in the MIC values of tested compounds after the addition of PAβN are presented.

<sup>c</sup> The growth of *B. cepacia* ATCC 25416 and *B. bronchiseptica* ATCC 4617 strains was inhibited in the MHB medium supplemented with 1 mM MgSO<sub>4</sub> and 20 mg·L<sup>-1</sup> PAβN.

<sup>d</sup> Nf, nitrofurantoin was used as a reference agent active against Gram-negative bacteria. The diameter of a commercial disc containing 0.3 mg of nitrofurantoin was 6 mm; the MIC of nitrofurantoin was determined according to the CLSI recommendations [3].

**Table S2.3.** The antifungal activity of tested agents against standard yeasts strains.

Agent tested	MIC in mg·L <sup>-1</sup> [MFC in mg·L <sup>-1</sup> ] <sup>a</sup> (Diameter of inhibition zone in mm)						
	<i>C. albicans</i> ATCC 90028	<i>C. parapsilosis</i> ATCC 22019	<i>C. tropicalis</i> ATCC 750	<i>C. tropicalis</i> IBA 171	<i>C. guilliermondii</i> IBA 155	<i>C. krusei</i> ATCC 6258	<i>S. cerevisiae</i> ATCC 9763
<b>1c</b>	200 (20)	400 (15)	100 (20)	<b>25</b> [400] (24)	400 (23)	400 (23)	200 (14)
<b>2c</b>	100 [400] (22)	200 (21)	50 [400] (20)	<b>12.5</b> [200] (25)	100 (37)	100 (15)	<b>12.5</b> [200] (28)
<b>3a</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	400 (-)	>400 (-)	>400 (-)
<b>3b</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>3c</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>4a</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>4b</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>4c</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>5a</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>5b</b>	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)	>400 (-)
<b>Fl<sup>b</sup></b>	1 (43)	2 (32)	0.38 (40)	0.38 (39)	0.75 (40)	64 <sup>c</sup> (16)	16 <sup>d</sup> (12)

The highest activity against yeasts indicated by the low MIC values (25 mg·L<sup>-1</sup>) is shown in boldface.

(-): The inhibition zone was not observed in the disc-diffusion method. The diameter of the paper discs was 9 mm.

<sup>a</sup> Only the MFC values ≤400 mg·L<sup>-1</sup> are presented.

<sup>b</sup> FL, fluconazole was used as a reference antifungal agent; the diameter of a commercial disc containing 0.025 mg of fluconazole was 6 mm; the MIC value of fluconazole was determined by the Etest method [4].

<sup>c</sup> The ellipse was visible pointing the MIC value 64 mg·L<sup>-1</sup>. However, with macro-colonies up to a concentration ≥256 mg·L<sup>-1</sup>. In accordance with the recommendations for the Etest method, the MIC value of fluconazole against *C. krusei* can also be interpreted as ≥256 mg·L<sup>-1</sup> [4,5]. *C. krusei* is intrinsically resistant to fluconazole.

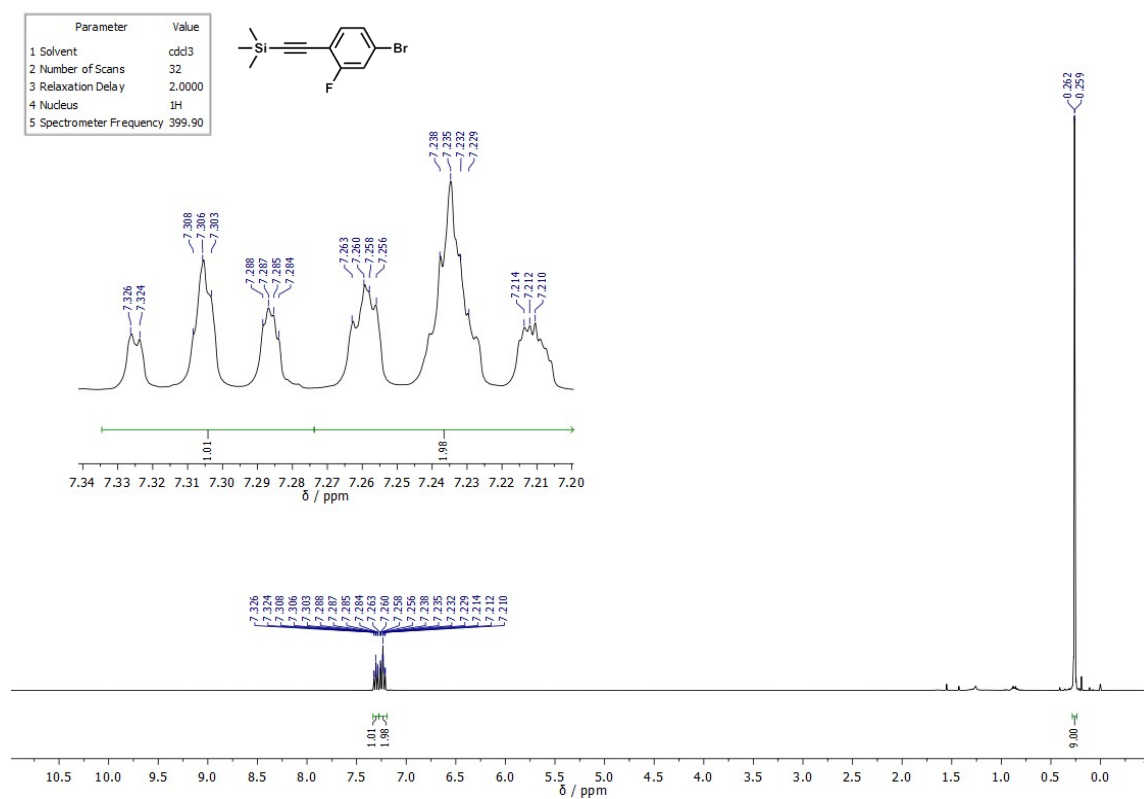
<sup>d</sup> The ellipse was visible pointing the MIC value 16 mg·L<sup>-1</sup>, with colonies up to concentration ≥256 mg·L<sup>-1</sup>. There are no recommendations for the Etest method interpretation of the MIC value of fluconazole against *S. cerevisiae*. The obtained MIC 16 mg·L<sup>-1</sup> is in line with the published results [6].



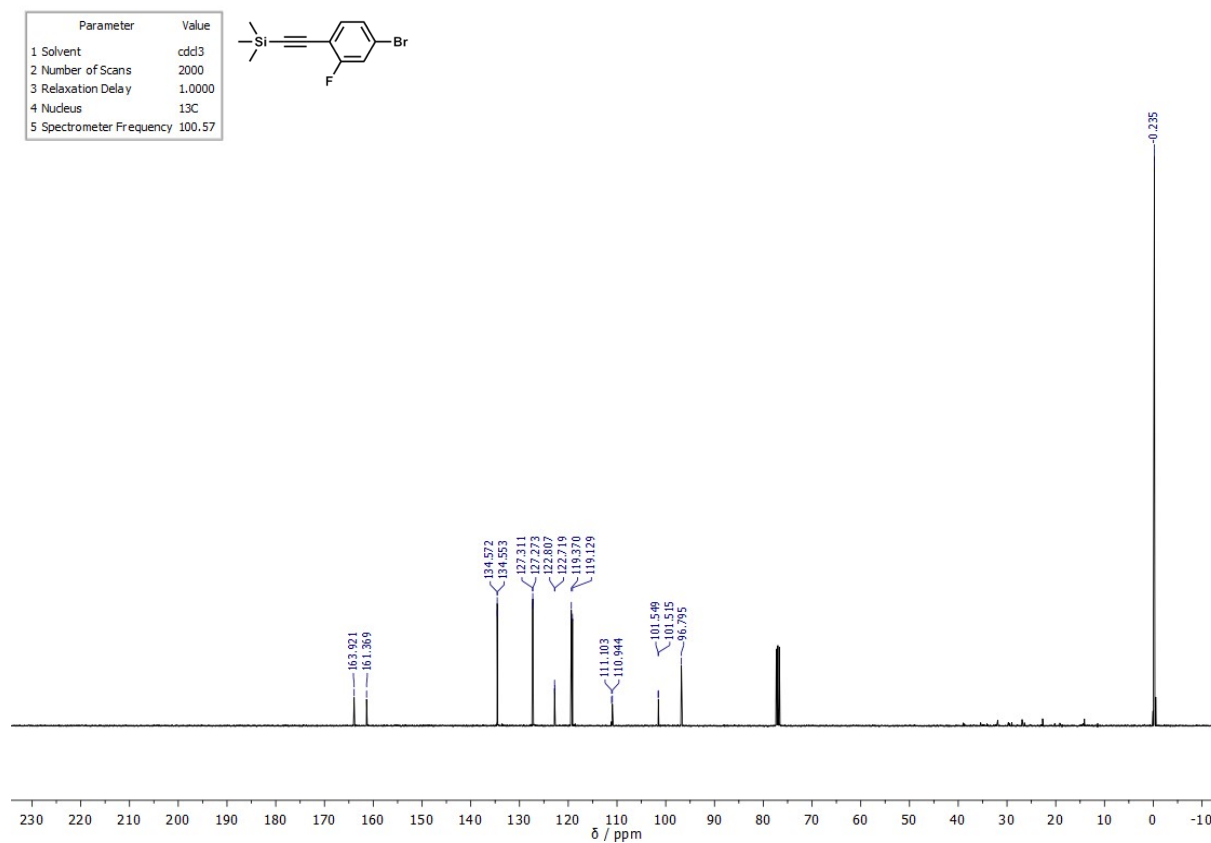
### 3. References

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6. M. A. Pfaller, M. Bale, B. Buschelman, M. Lancaster, A. Espinel-Ingroff, J. H. Rex and M. G. Rinaldi. Selection of candidate quality control isolates and tentative quality control ranges for *in vitro* susceptibility testing of yeast isolates by National Committee for Clinical Laboratory Standards proposed standard methods, *J. Clin. Microbiol.*, 1994, **32**, 1650–1653.

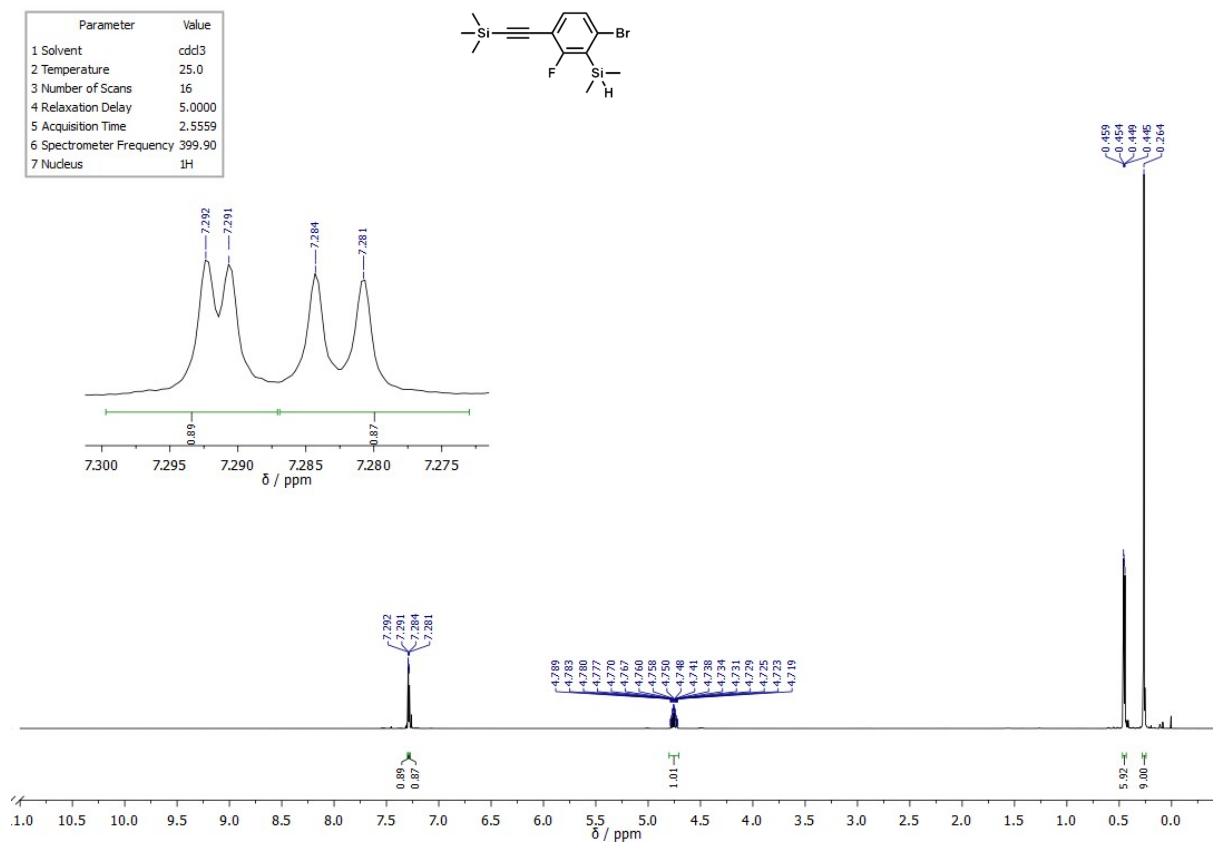
## 4. NMR spectra of new compounds



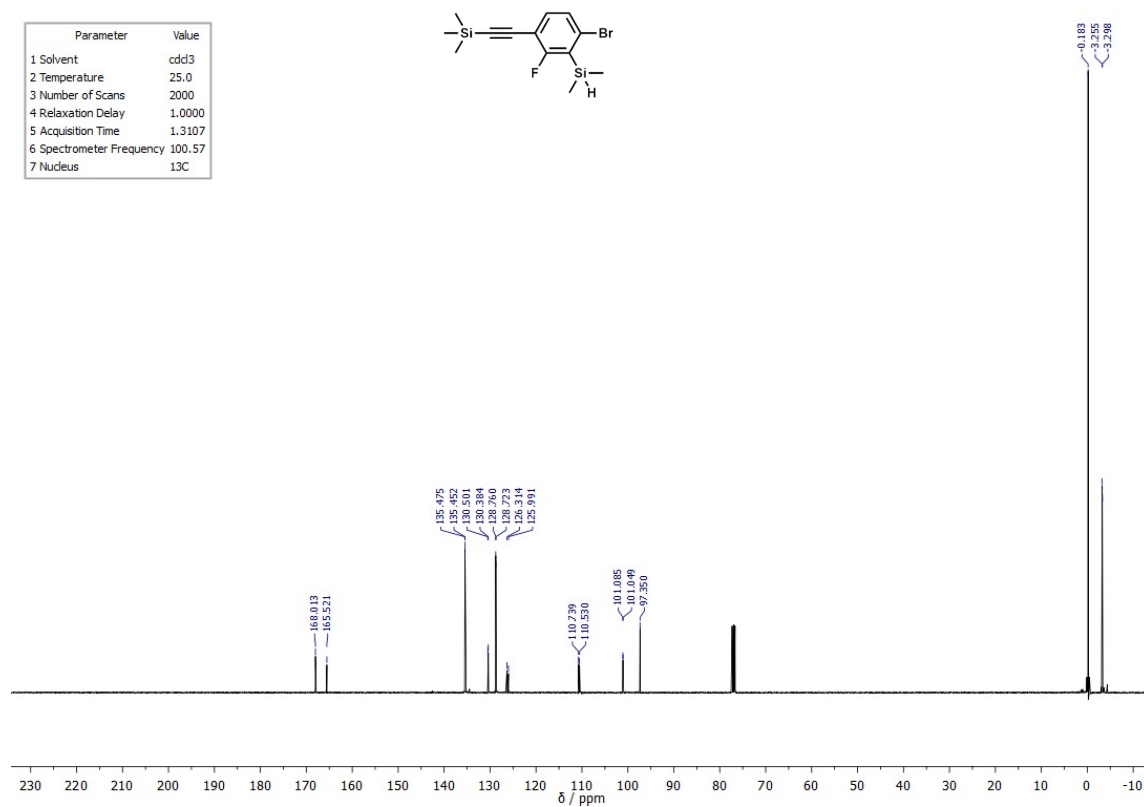
**Figure S4.1.** <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) spectrum of **1a**.



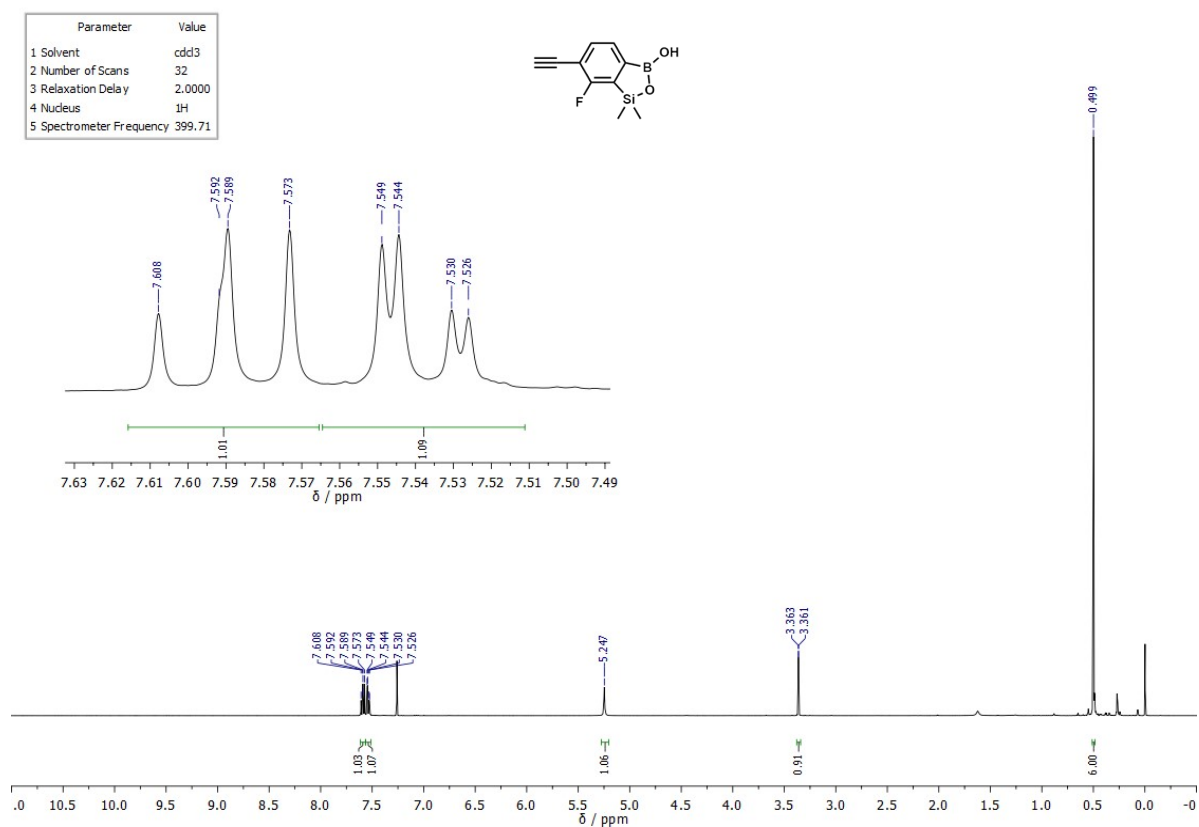
**Figure S4.2.** <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>) spectrum of **1a**.



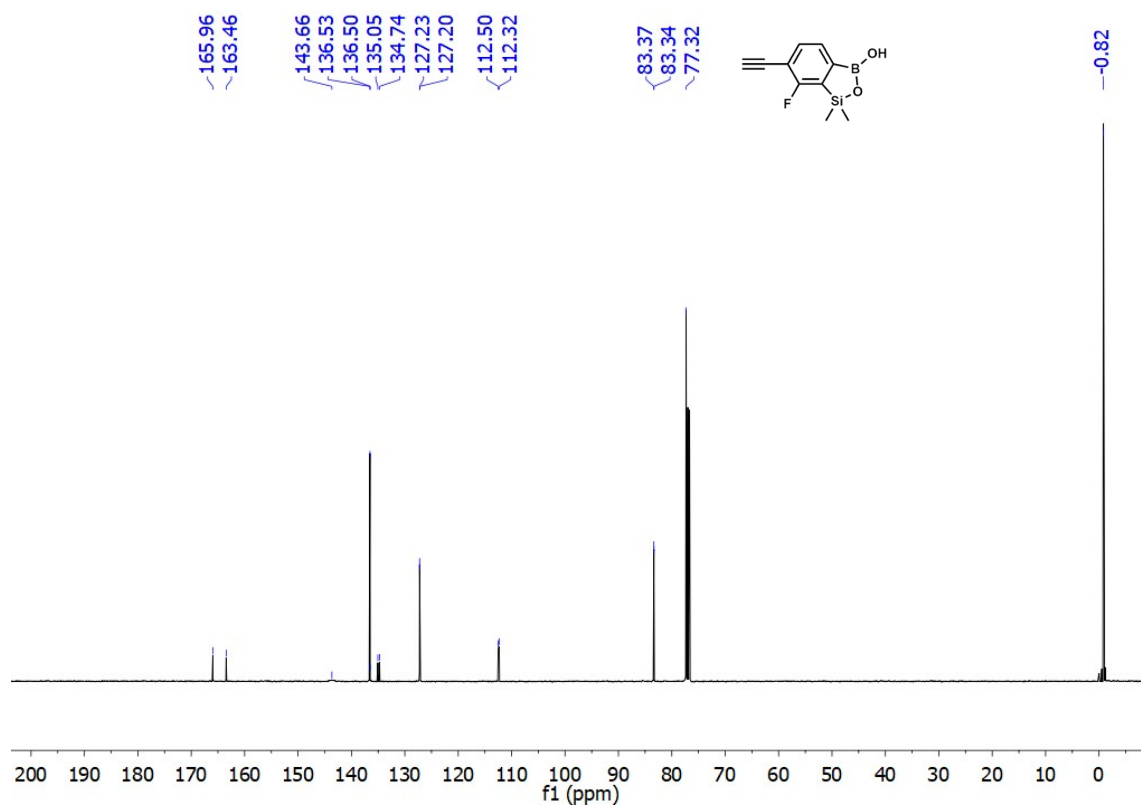
**Figure S4.3.** <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) spectrum of **1b**.



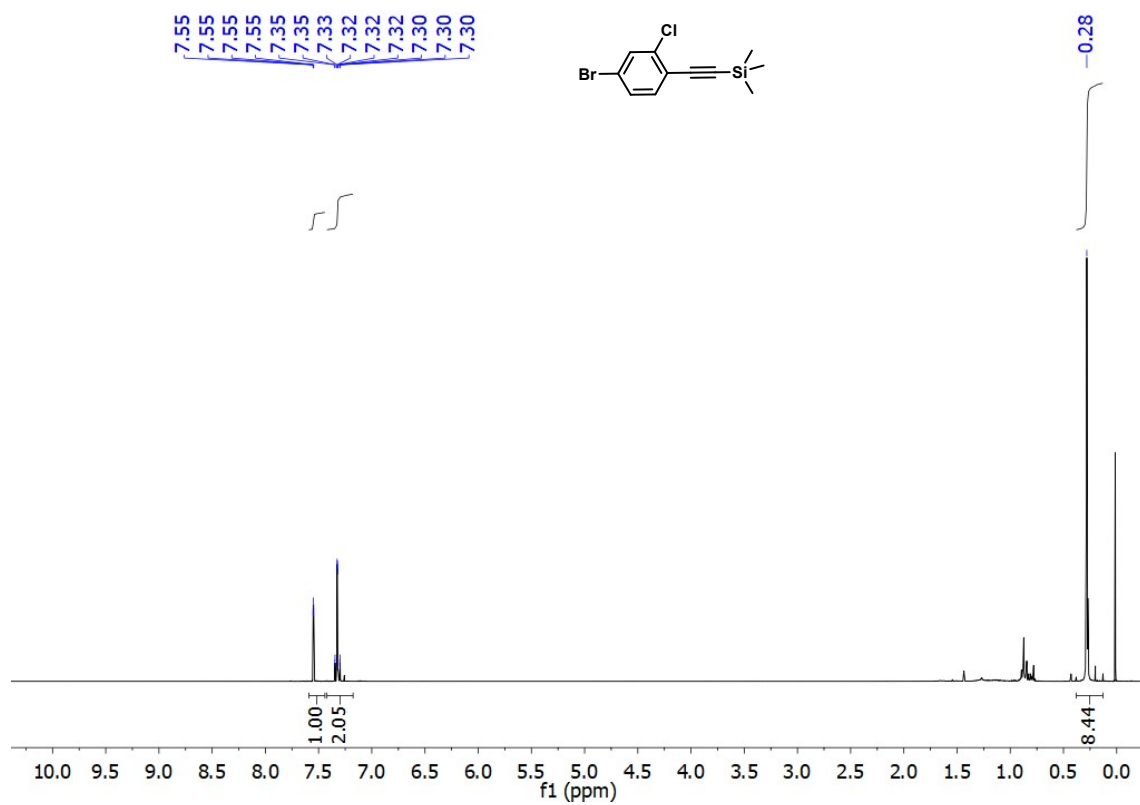
**Figure S4.4.** <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>) spectrum of **1b**.



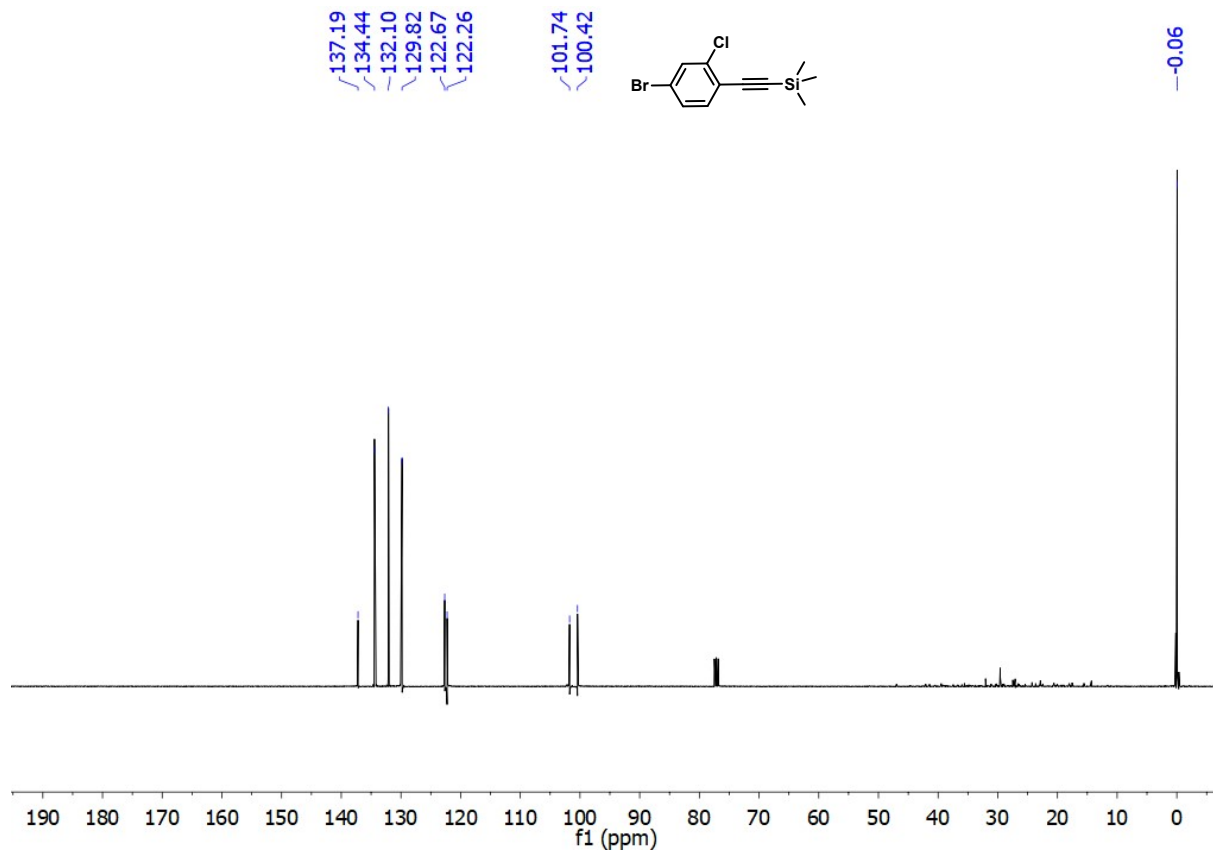
**Figure S4.5.** <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) spectrum of **1c**.



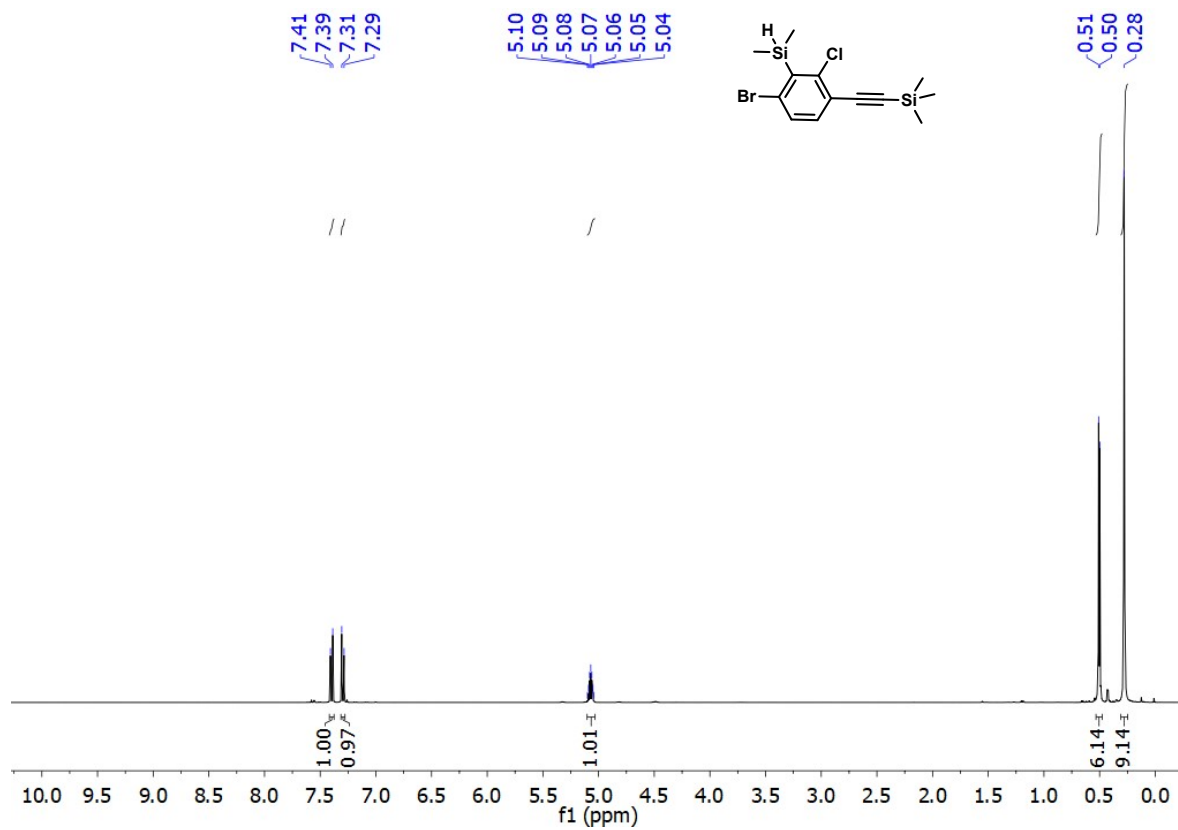
**Figure S4.6.** <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>) spectrum of **1c**.



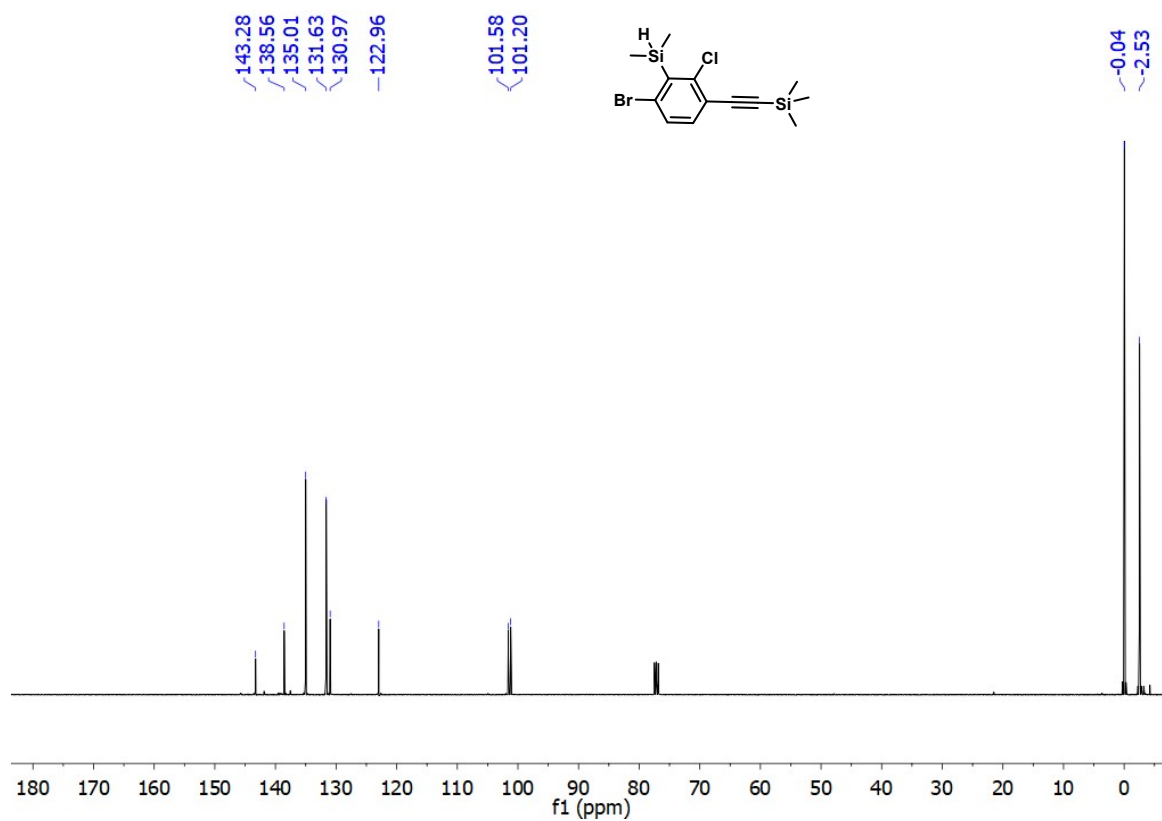
**Figure S4.7.**  $^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ ) spectrum of **2a**.



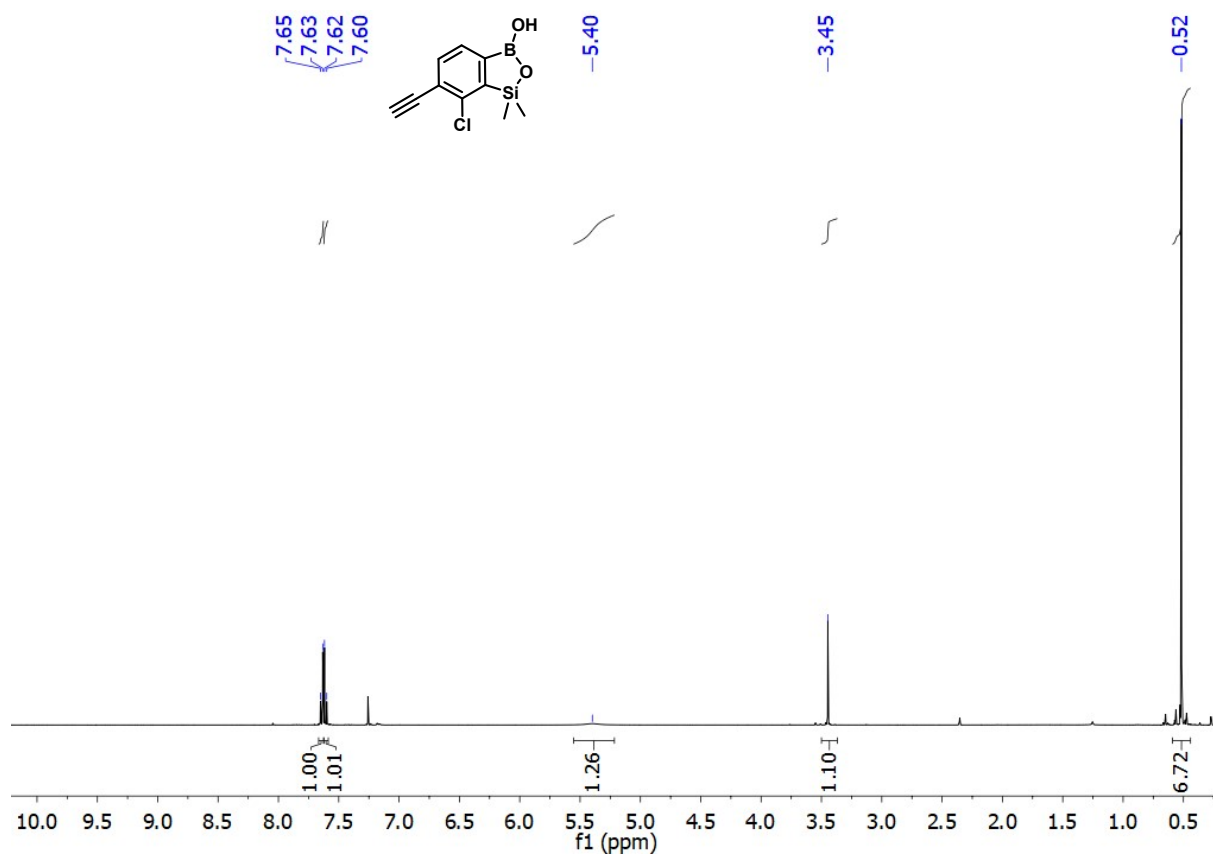
**Figure S4.8.**  $^{13}\text{C}$  NMR (101 MHz,  $\text{CDCl}_3$ ) spectrum of **2a**.



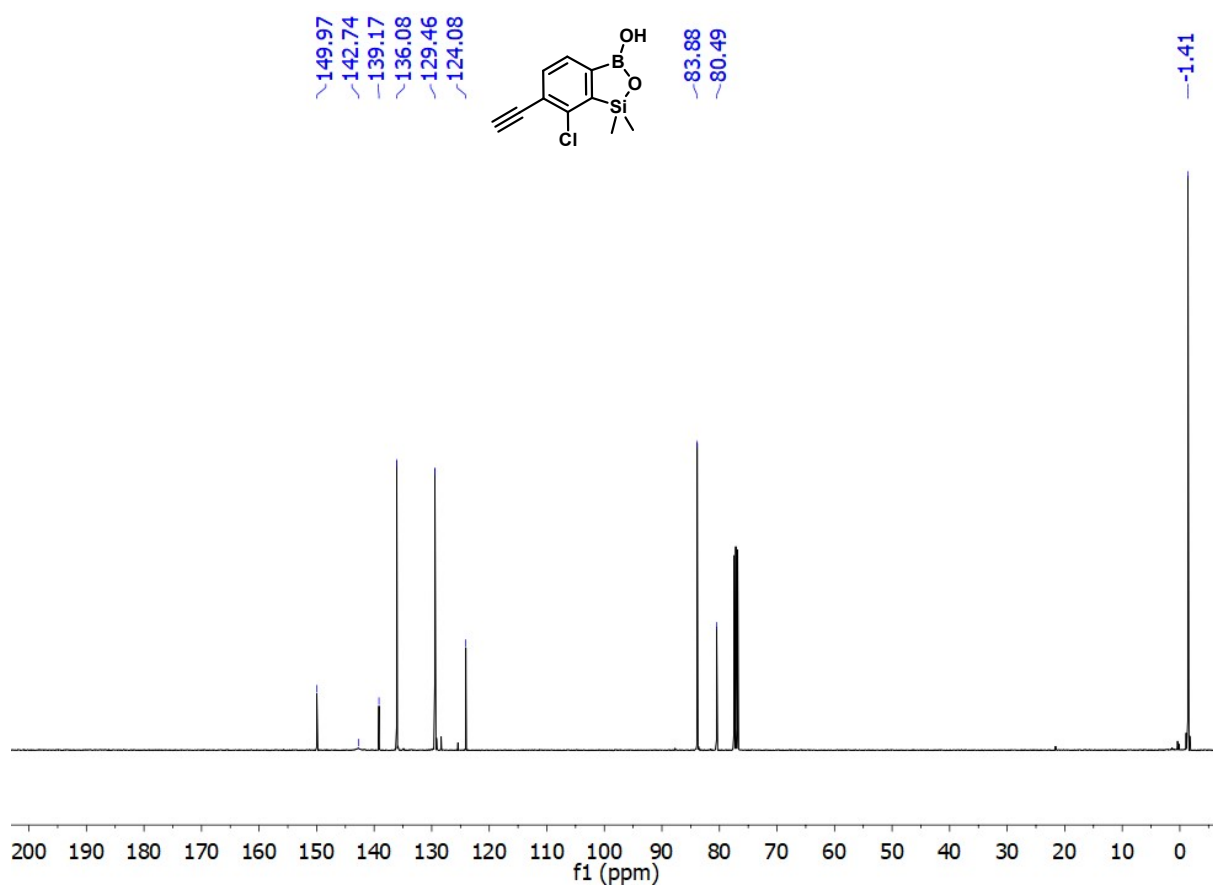
**Figure S4.9.** <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) spectrum of **2b**.



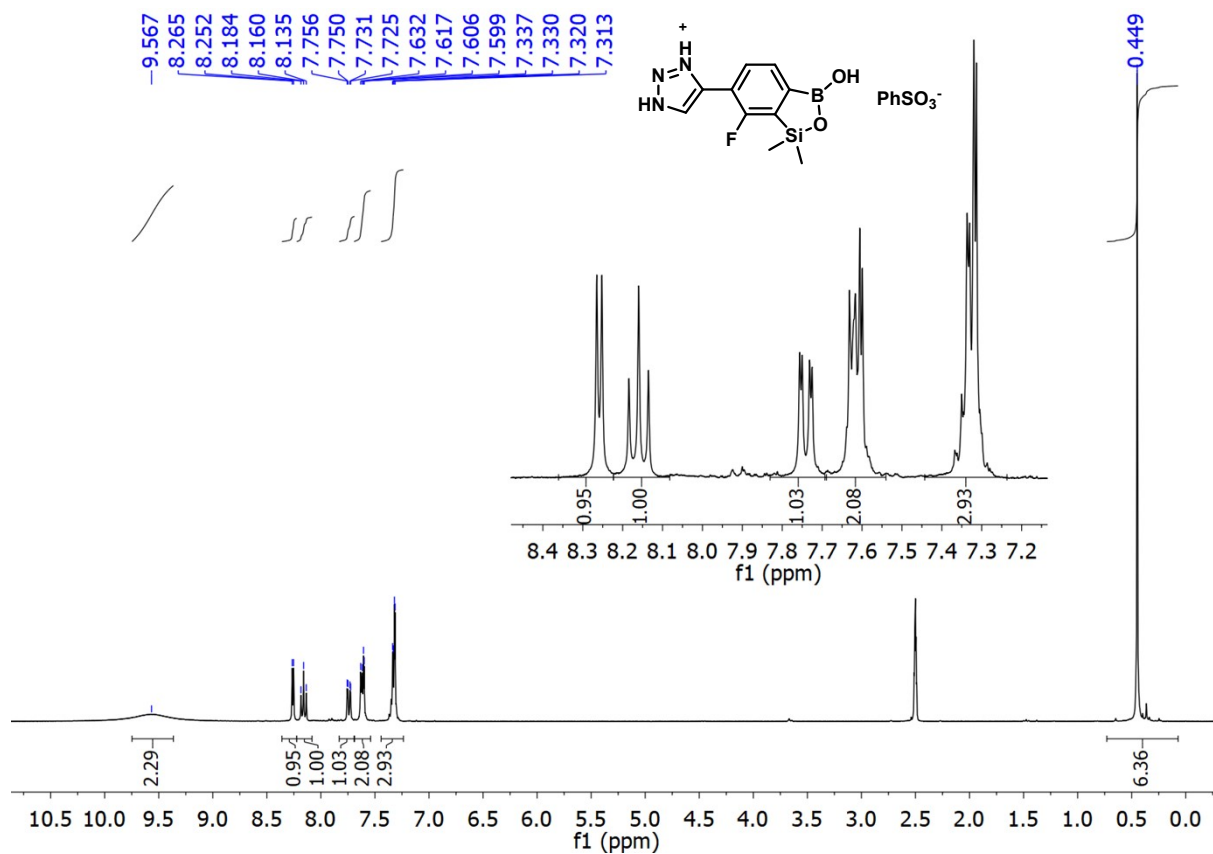
**Figure S4.10.** <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>) spectrum of **2b**.



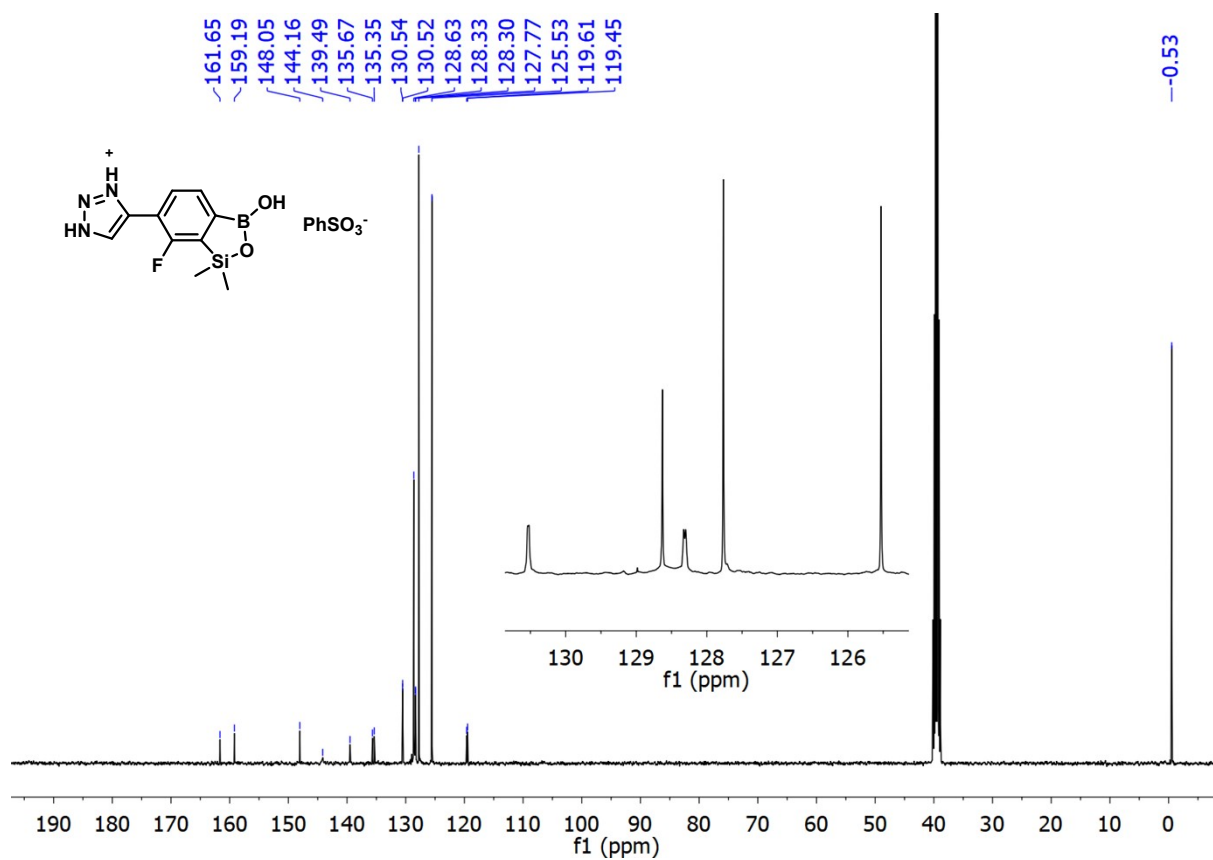
**Figure S4.11.** <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) spectrum of **2c**.



**Figure S4.12.** <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>) spectrum of **2c**.

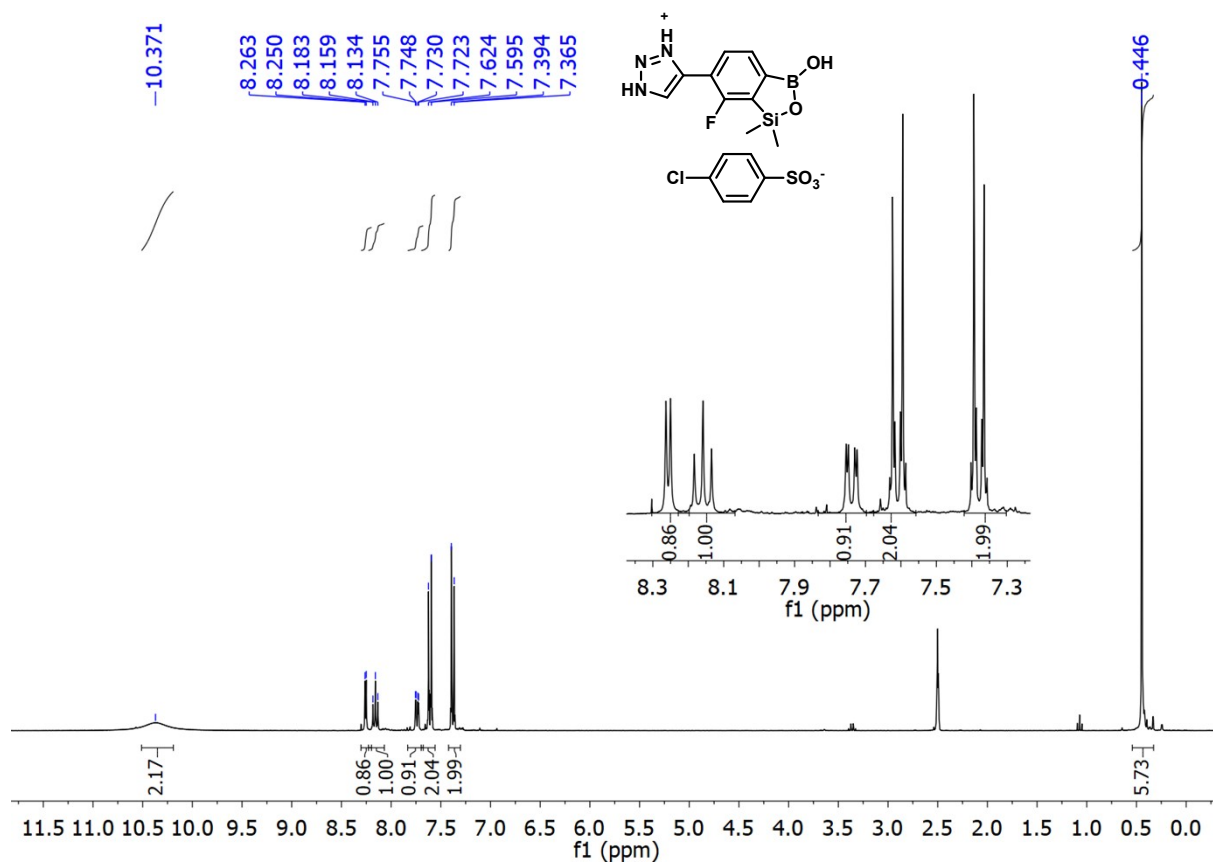


**Figure S4.13.**  $^1\text{H NMR}$  (400 MHz,  $\text{DMSO-}d_6$ ) spectrum of **3a**.

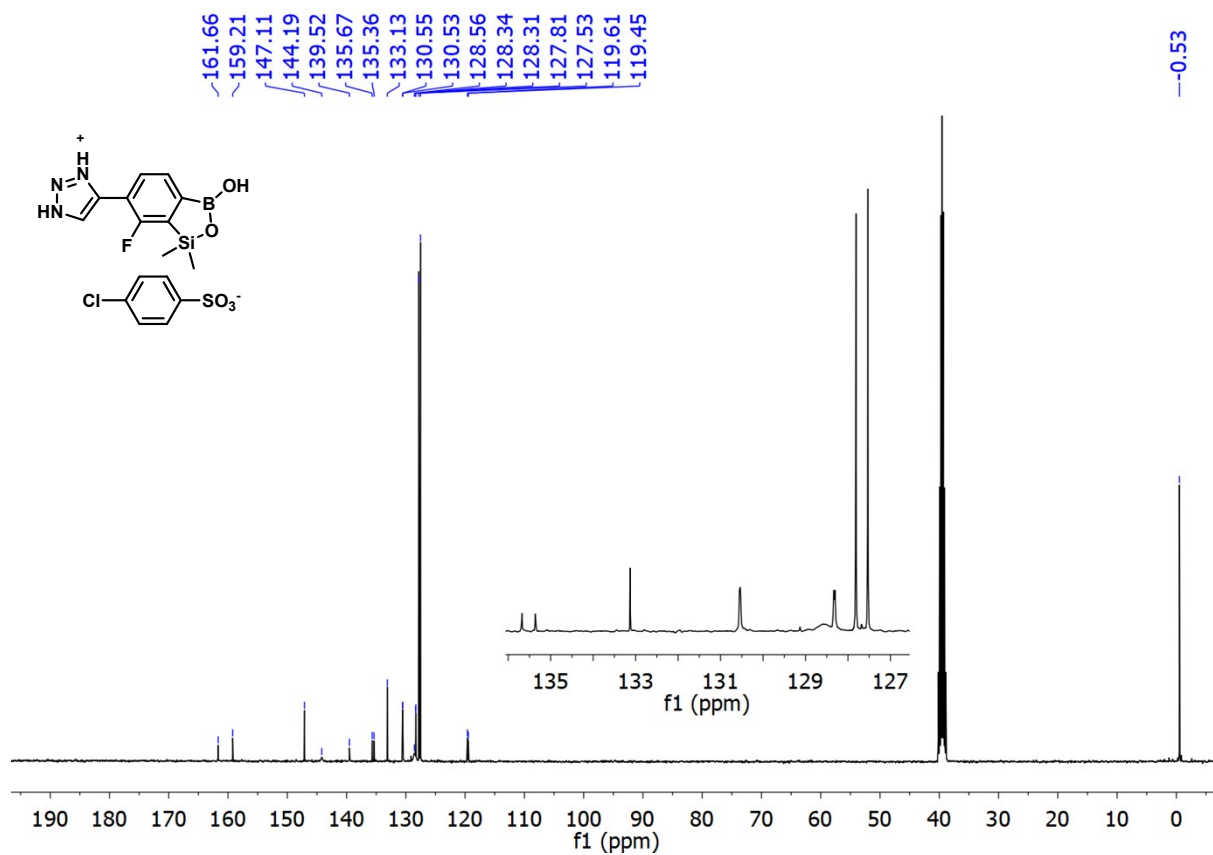


**Figure S4.14.**  $^{13}\text{C NMR}$  (101 MHz,  $\text{DMSO-}d_6$ ) spectrum of **3a**.





**Figure S4.15.**  $^1\text{H NMR}$  (400 MHz,  $\text{DMSO-}d_6$ ) spectrum of **3b**.



**Figure S4.16.**  $^{13}\text{C NMR}$  (101 MHz,  $\text{DMSO-}d_6$ ) spectrum of **3b**.

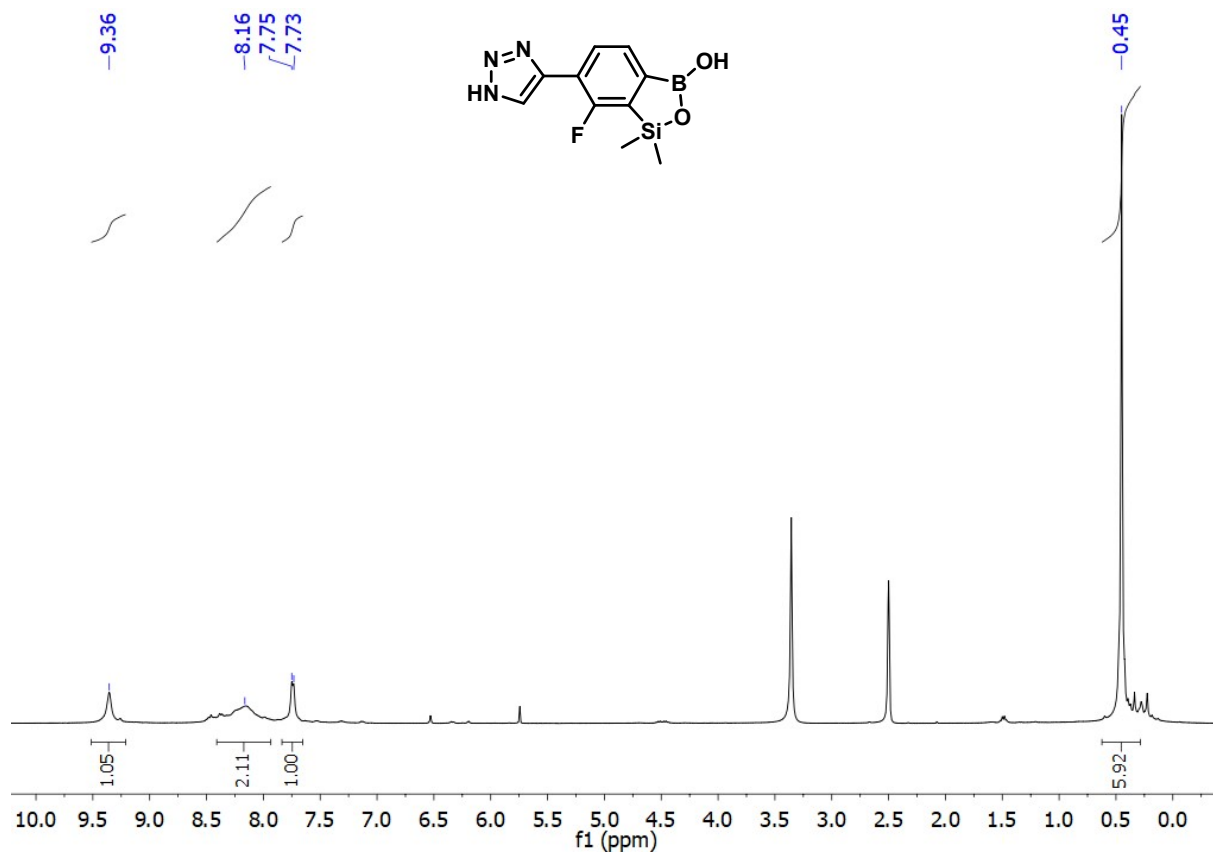


Figure S4.17.  $^1\text{H}$  NMR (400 MHz,  $\text{DMSO-}d_6$ ) spectrum of **3c**.

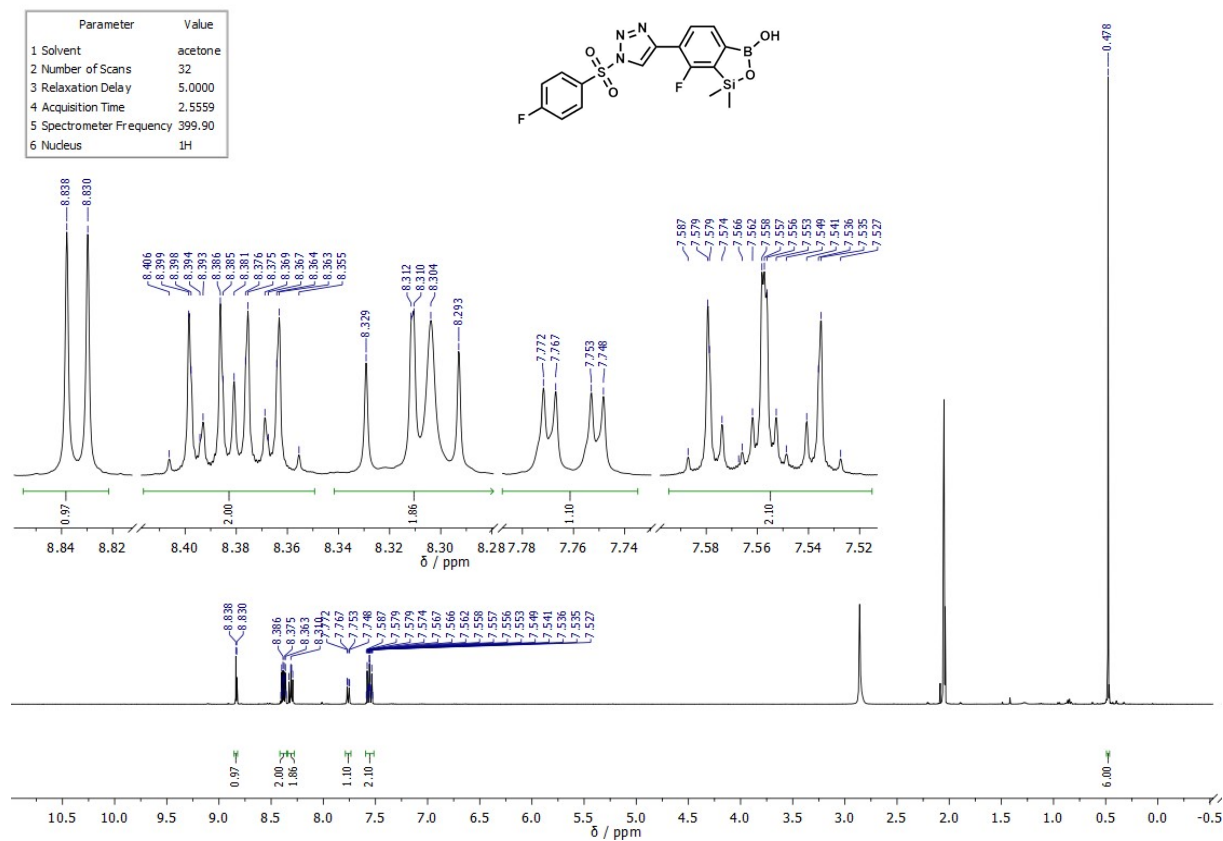


Figure S4.18.  $^1\text{H}$  NMR (400 MHz,  $\text{acetone-}d_6$ ) spectrum of **4a**.

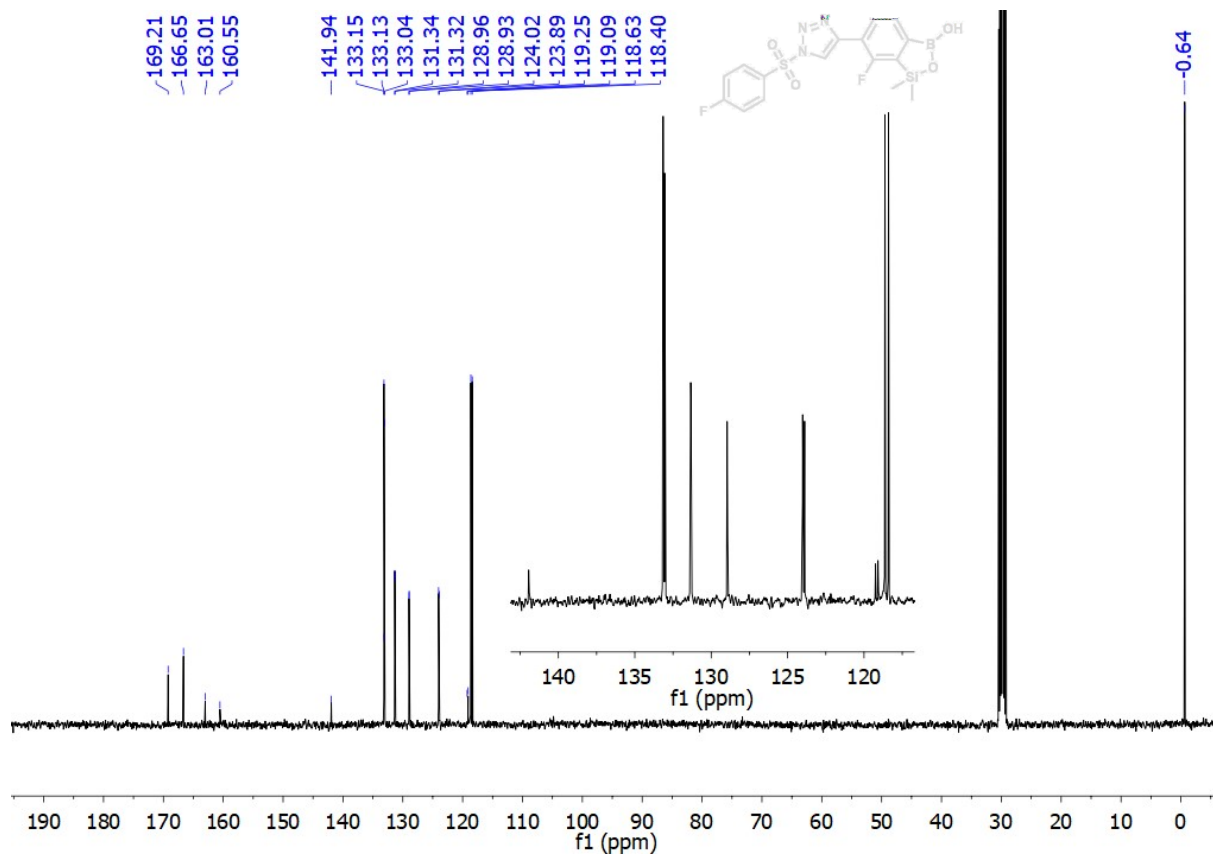


Figure S4.19. <sup>13</sup>C NMR (400 MHz, acetone-*d*<sub>6</sub>) spectrum of 4a.

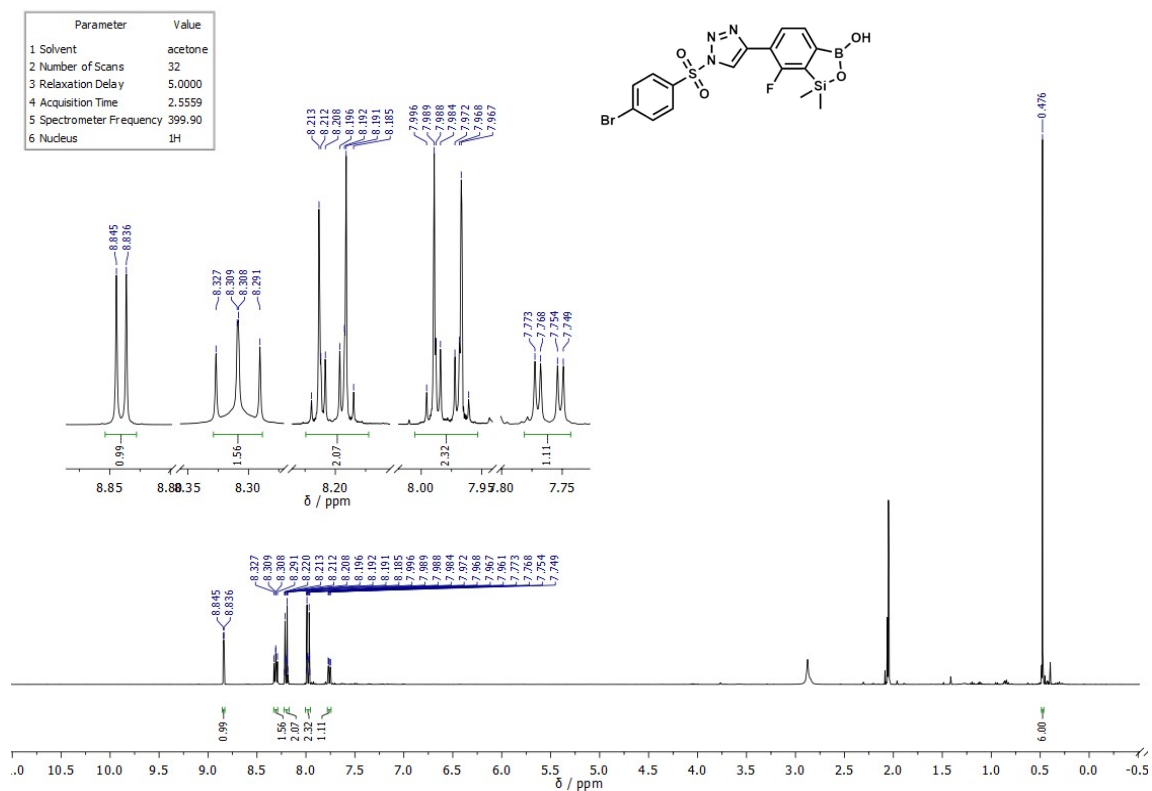


Figure S4.20. <sup>1</sup>H NMR (400 MHz, acetone-*d*<sub>6</sub>) spectrum of 4b.

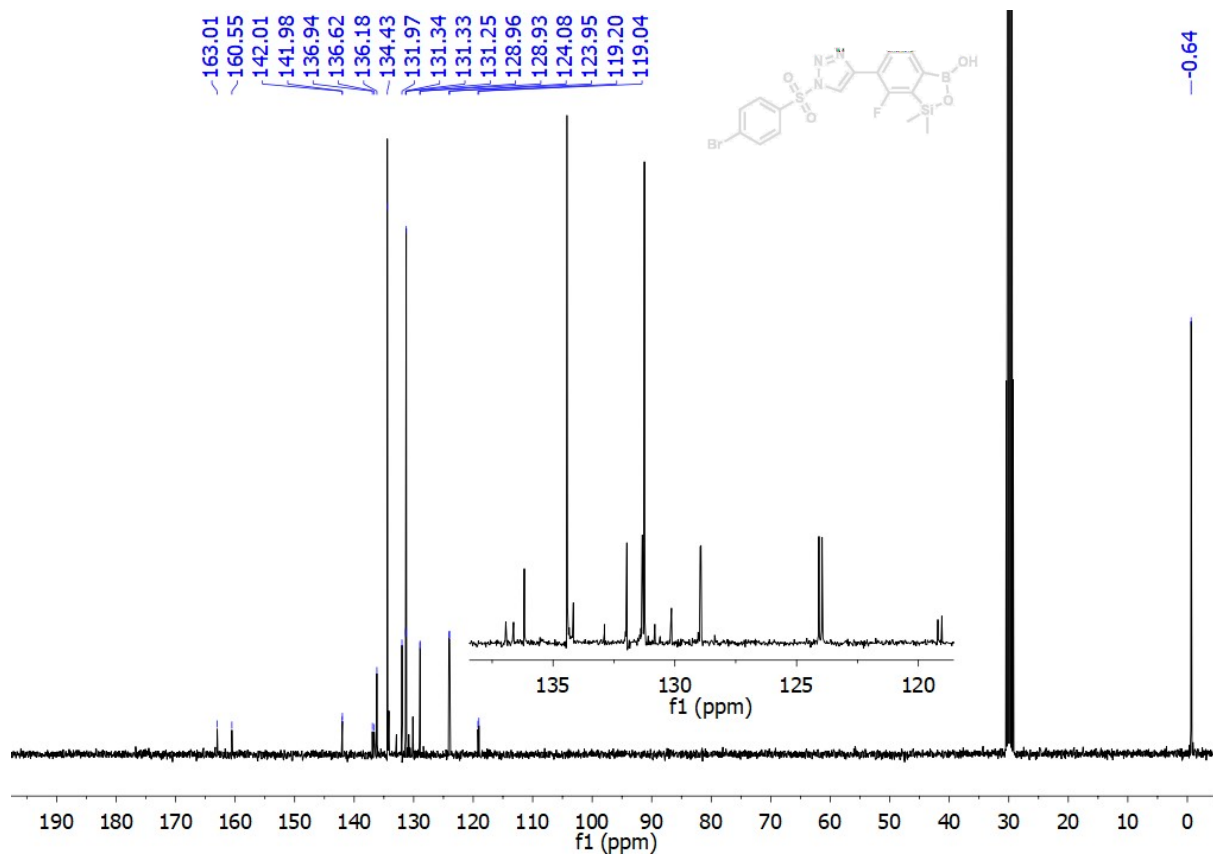


Figure S4.21. <sup>13</sup>C NMR (400 MHz, acetone-*d*<sub>6</sub>) spectrum of **4b**.

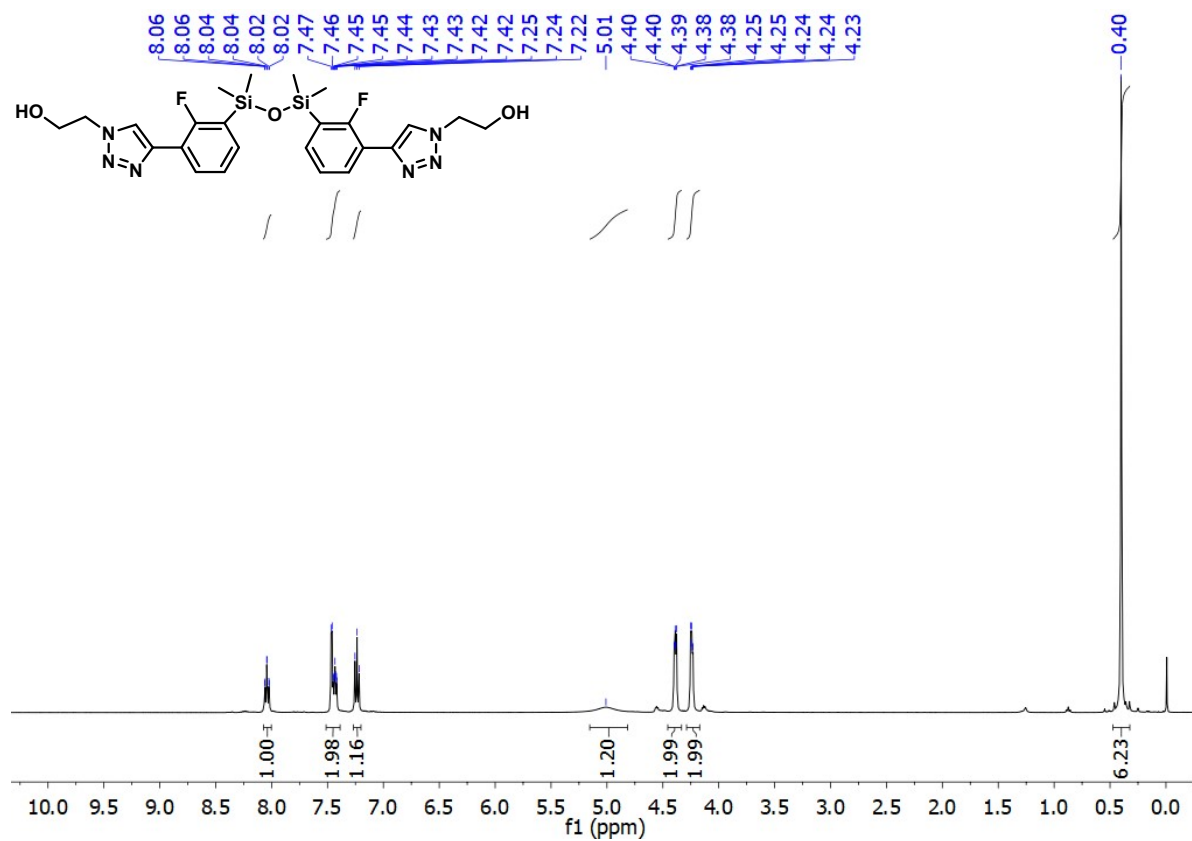
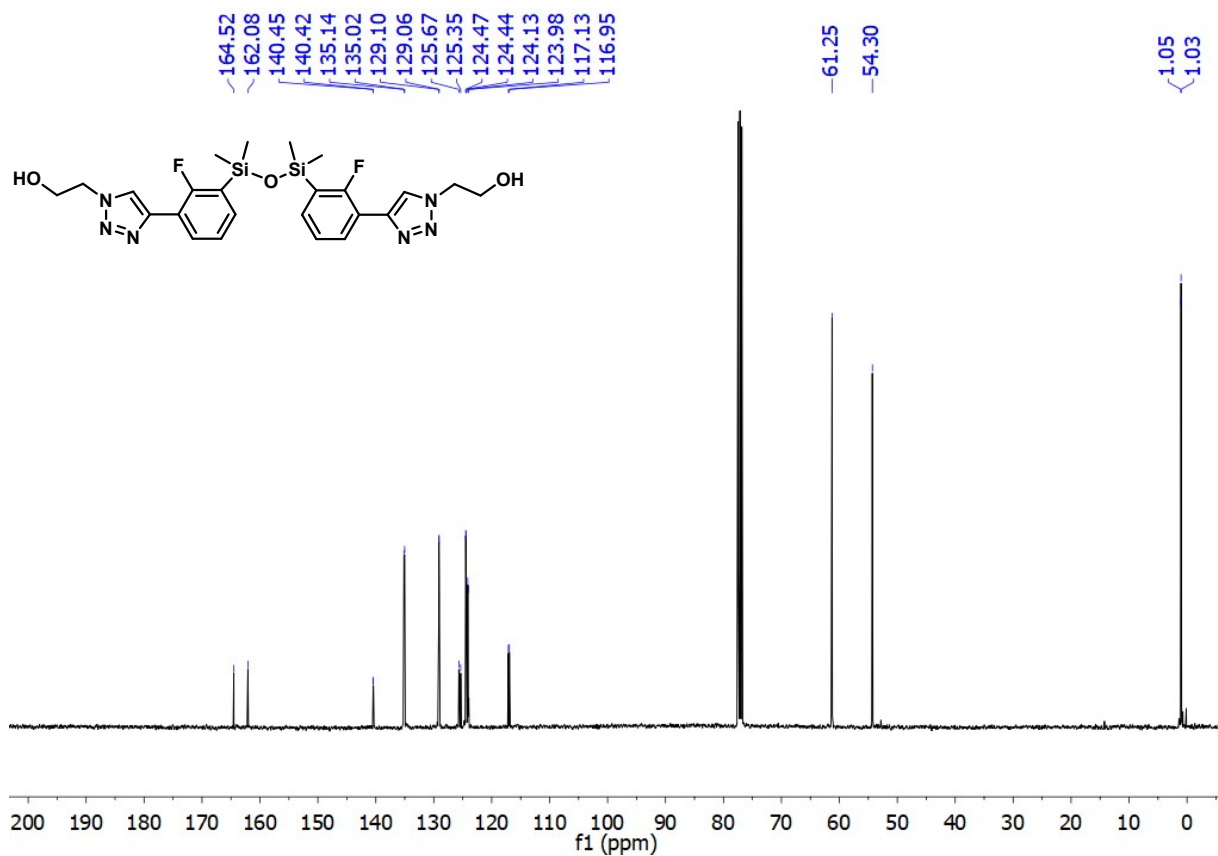
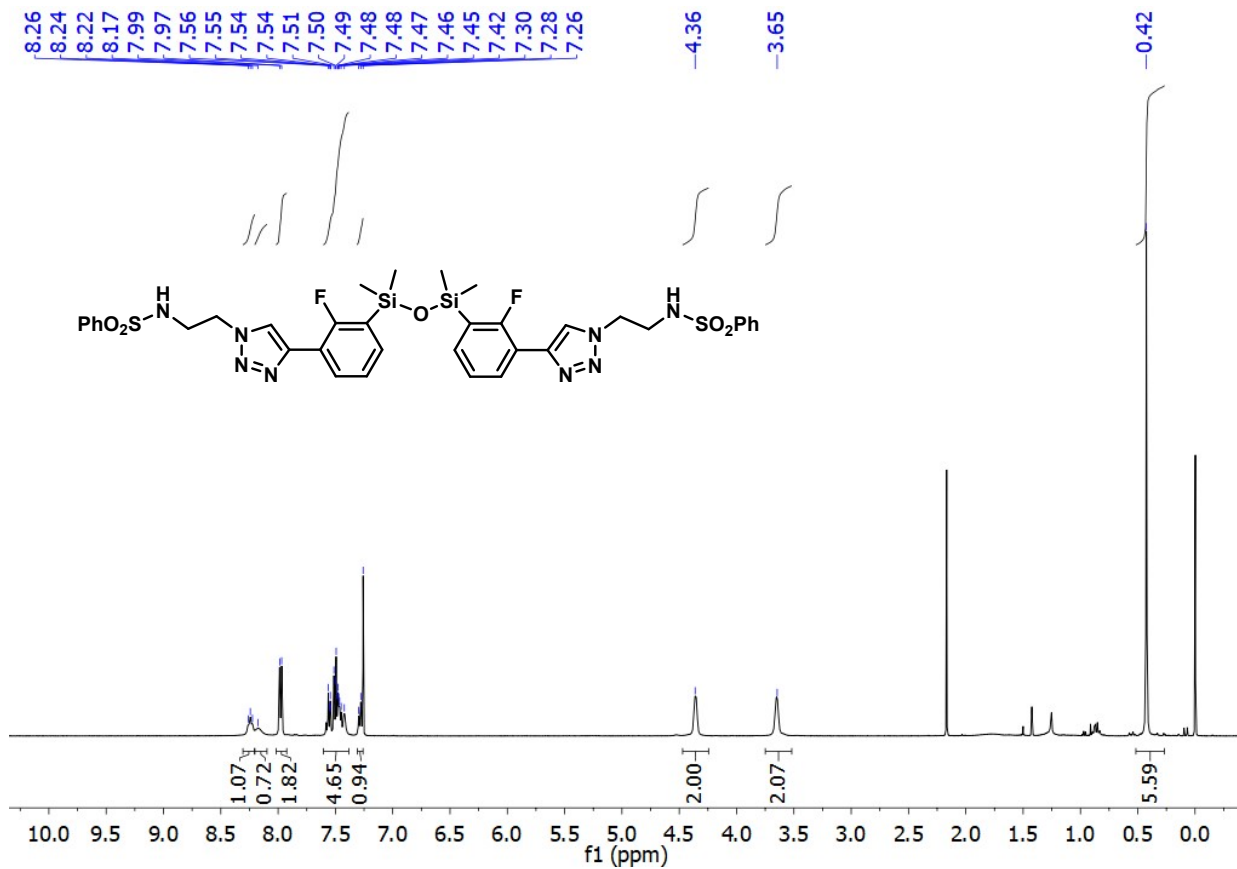


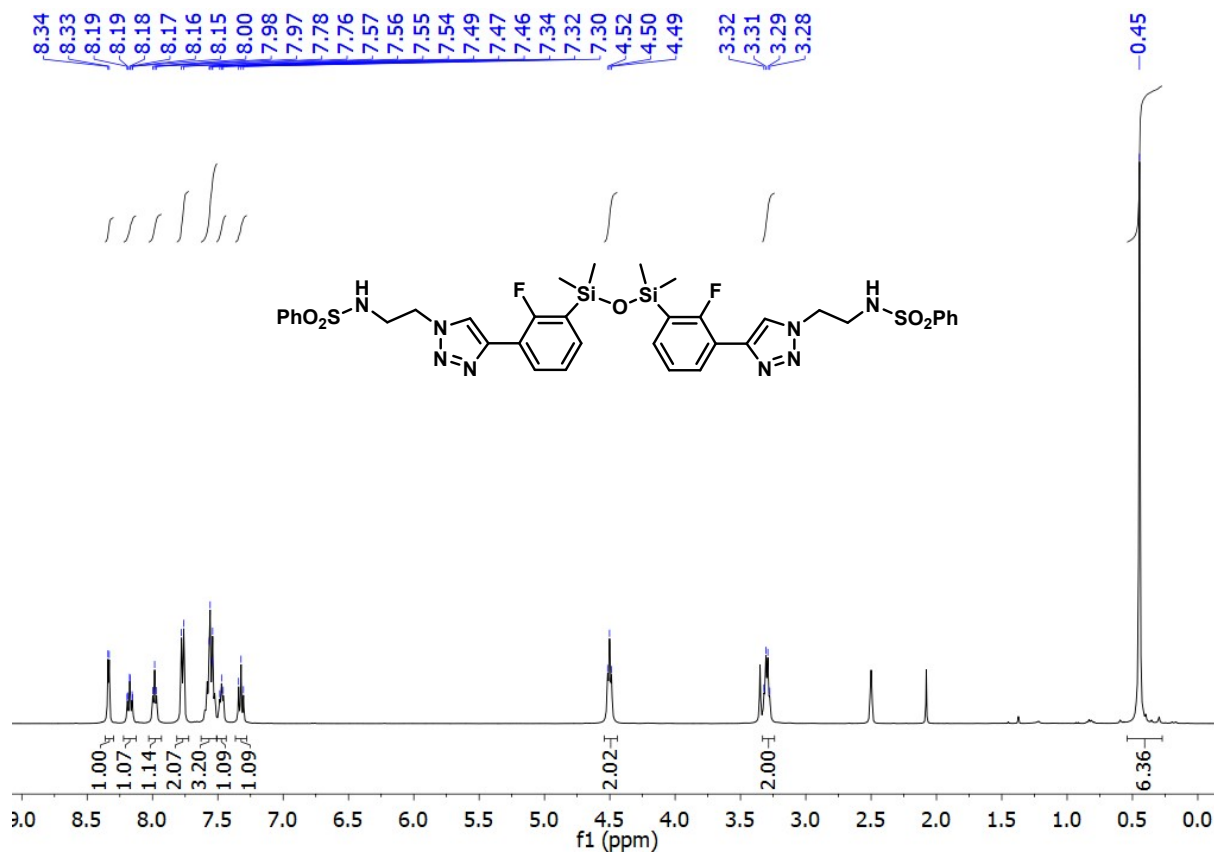
Figure S4.22. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) spectrum of **5a**.



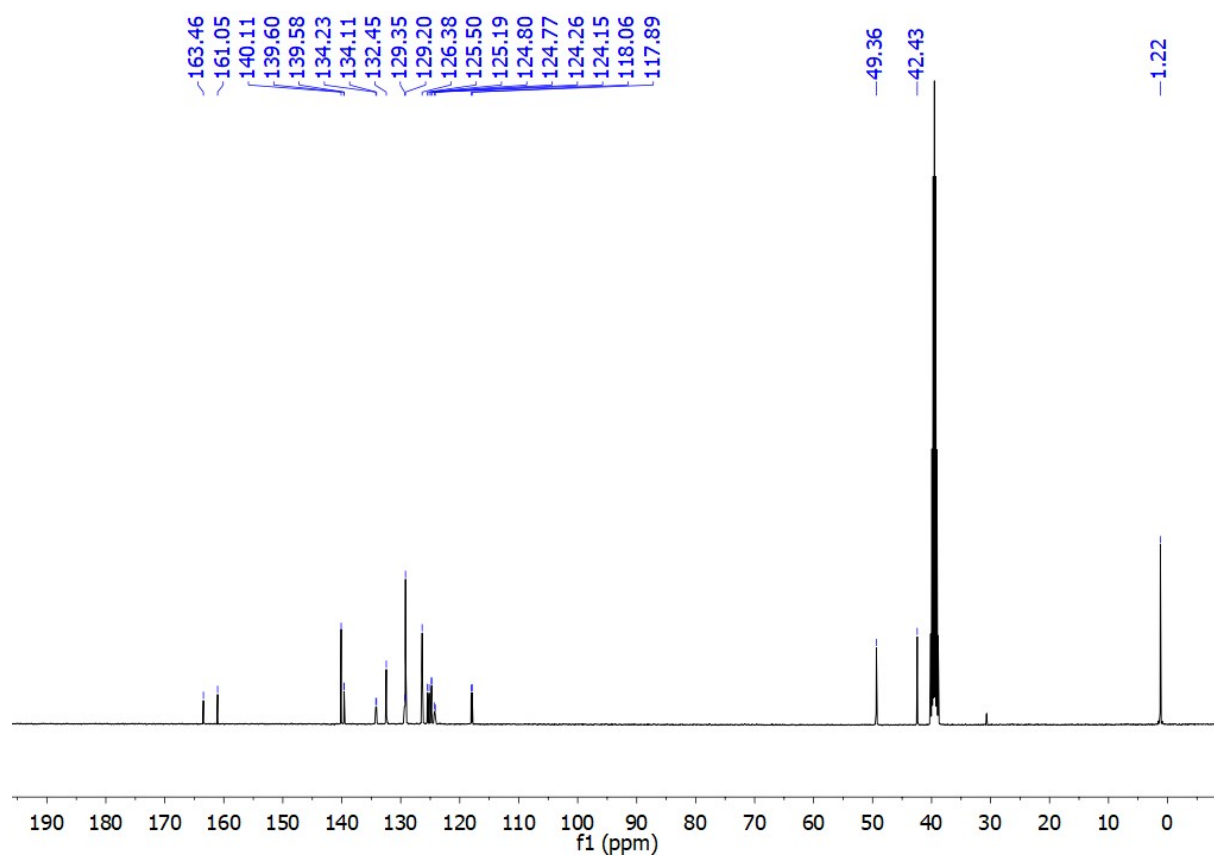
**Figure S4.23.** <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>) spectrum of **5a**.



**Figure S4.24.** <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) spectrum of **5b**.



**Figure S4.25.**  $^1\text{H NMR}$  (400 MHz,  $\text{DMSO-}d_6$ ) spectrum of **5b**.



**Figure S4.26.**  $^{13}\text{C NMR}$  (101 MHz,  $\text{DMSO-}d_6$ ) spectrum of **5b**.