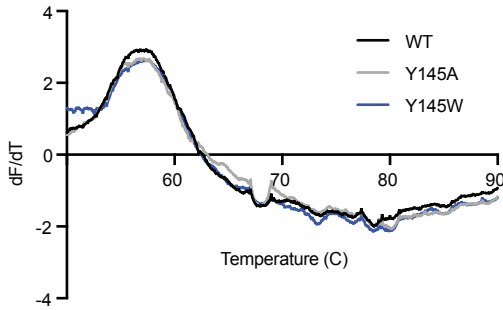


Supplemental Figure 2. Y148W, but not A223W or D225W, partially mirrors the motions in the NBD that accompany the ADP-bound or compound-bound states. Two replicates are shown and the position of loop 222 (see Fig 1) is highlighted.

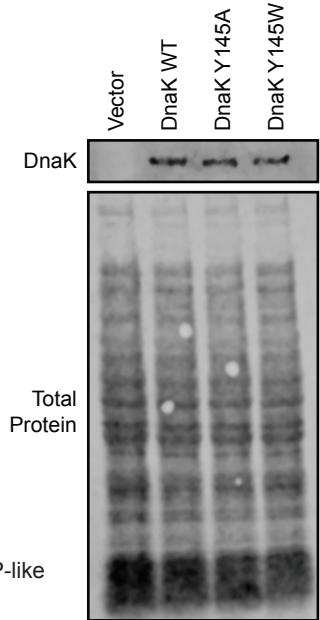
(A) DnaK mutants are properly folded, based on DSF



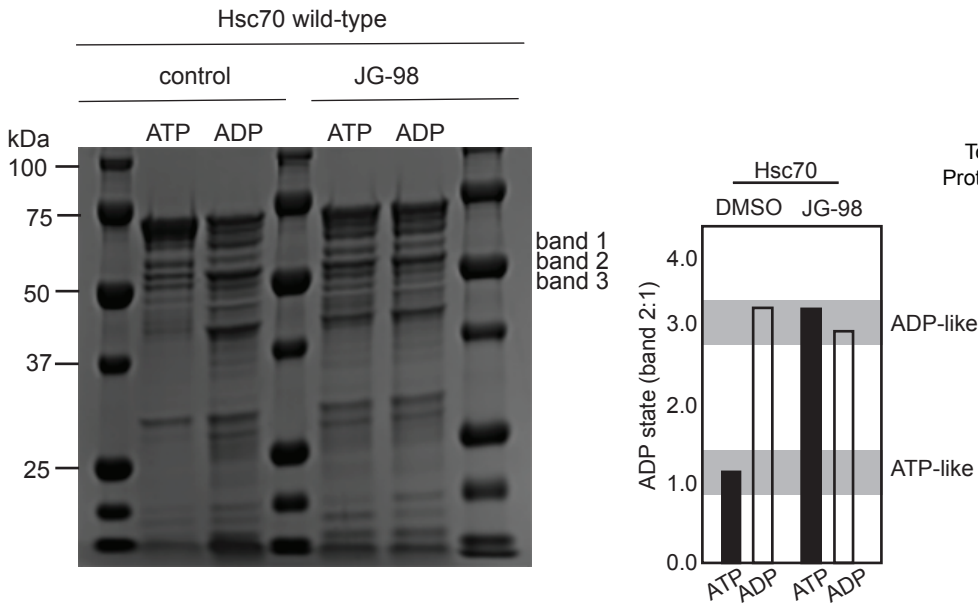
(B) ATPase activity of DnaK Y145W is relatively resistant to JG-48

DnaK	JG-48 inhibition of V_{maxJ} (%)
WT	35%
Y145W	8%

(D) DnaK mutants are expressed at similar levels

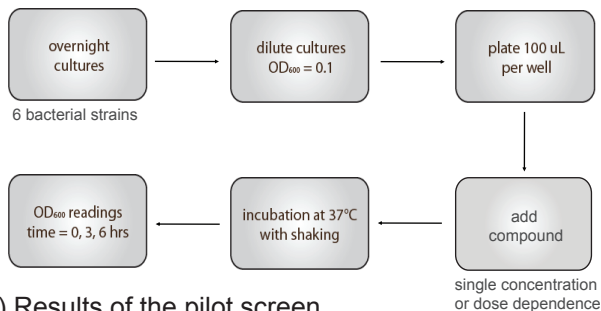


(C) JG-98 stabilizes the ADP-like conformer of Hsc70

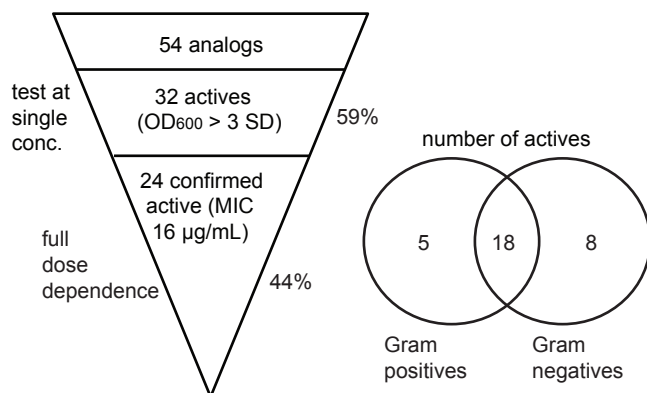


Supplemental Figure 3. Controls for partial proteolysis and bacterial complementation experiments. (A) Differential scanning fluorimetry results on DnaK and its mutants show that the proteins are properly folded (T_m values $\sim 58^\circ\text{C}$). Results are the average of experiments performed in triplicate. (B) ATPase activity of WT DnaK and Y145W DnaK was measured in the presence of DnaJ and JG-48. The Trp mutant was relatively resistant to the inhibitor, as expected. (C) Partial proteolysis experiments showing that JG-98 ($10\ \mu\text{M}$) is able to stabilize the ADP-like state, even if ATP ($1\ \text{mM}$) is added. The quantification is the result of duplicate experiments and is the ratio of band intensity from band 2:1. Band intensities were determined using Image J. (D) Arabinose-mediated induction of DnaK and its mutants in the $\Delta dnaK$ *E. coli* strain, showing similar expression levels.

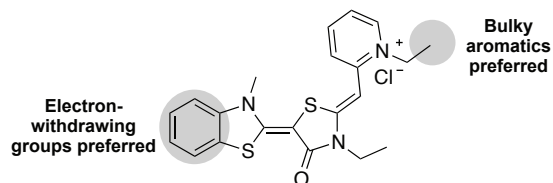
(A) Workplan for MIC measurements



(B) Results of the pilot screen



(D) Structure-activity relationships from the pilot screen



(C) Confirmed actives (MIC values < 16 µg/mL)

analog	<i>B. subtilis</i>	<i>B. anthracis</i>	<i>B. cereus</i>	<i>S. aureus</i>	<i>E. coli</i>	<i>H. influenza</i>
MKT077	---	---	>16	>16	---	16
YM1	---	>16	---	>16	---	16
YM9	---	---	---	---	---	>16
YM21	---	---	---	---	---	>16
JG12	>16	>16	---	>16	---	16
JG18	---	>16	---	---	---	>16
JG31	---	>16	---	---	---	16
JG33	>16	16	16	16	---	8
JG35	---	>16	---	16	---	16
JG37	---	---	---	---	---	>16
JG38	---	>16	>16	---	---	16
JG41	---	16	>16	>16	---	8
JG43	16	8	8	8	---	8
JG58	---	---	---	---	---	---
JG60	---	>16	>16	>16	---	>16
JG61	---	---	---	---	---	16
JG66	---	>16	---	>16	---	---
JG70	8	8	8	8	---	16
JG71	>16	---	---	4	>16	---
JG73	16	8	16	16	>16	8
JG74	---	8	8	8	>16	4
JG78	---	4	8	4	---	8
JG81	>16	>16	>16	>16	---	16
JG83	8	4	16	8	---	4
JG86	---	16	---	>16	---	8
JG91	>16	16	>16	16	---	4
JG96	>16	8	8	8	---	16
JG97	16	4	8	8	---	16
JG98	2	---	---	---	---	---
JG111	---	16	---	>16	---	---
JG121	>16	>16	---	---	---	16
JG123	---	---	---	---	---	>16
tetracycline	2	0.25	0.5	1	4	1
DMSO	>16	>16	>16	>16	>16	>16

Supplemental Figure 4. Pilot screen of JG-98 analogs against a panel of bacteria. (A) Schematic of the method for determination of anti-bacterial activity. In the pilot screen, 54 compounds were tested at a single concentration (16 µg/mL). If the compound reduced growth by 3SD of the negative control (1% DMSO) for any of the 6 bacterial strains, then it was selected for MIC determination (12 concentrations). Confirmed actives were defined as those with MIC values of 16 µg/mL or better. Compounds that were not active in the initial screen were not retested for dose dependence (---). All pilot screen experiments performed at 37 °C. Compounds are described in Li *et al.* 2013 *ACS Med. Chem. Lett.* 4:1042.

01152014_1
STANDARD 1H OBSERVE

8.25
8.25
8.24
8.24

7.89
7.88
7.84
7.84
7.63
7.61
7.43
7.42
7.42
7.41
7.36
7.34
6.52

5.75

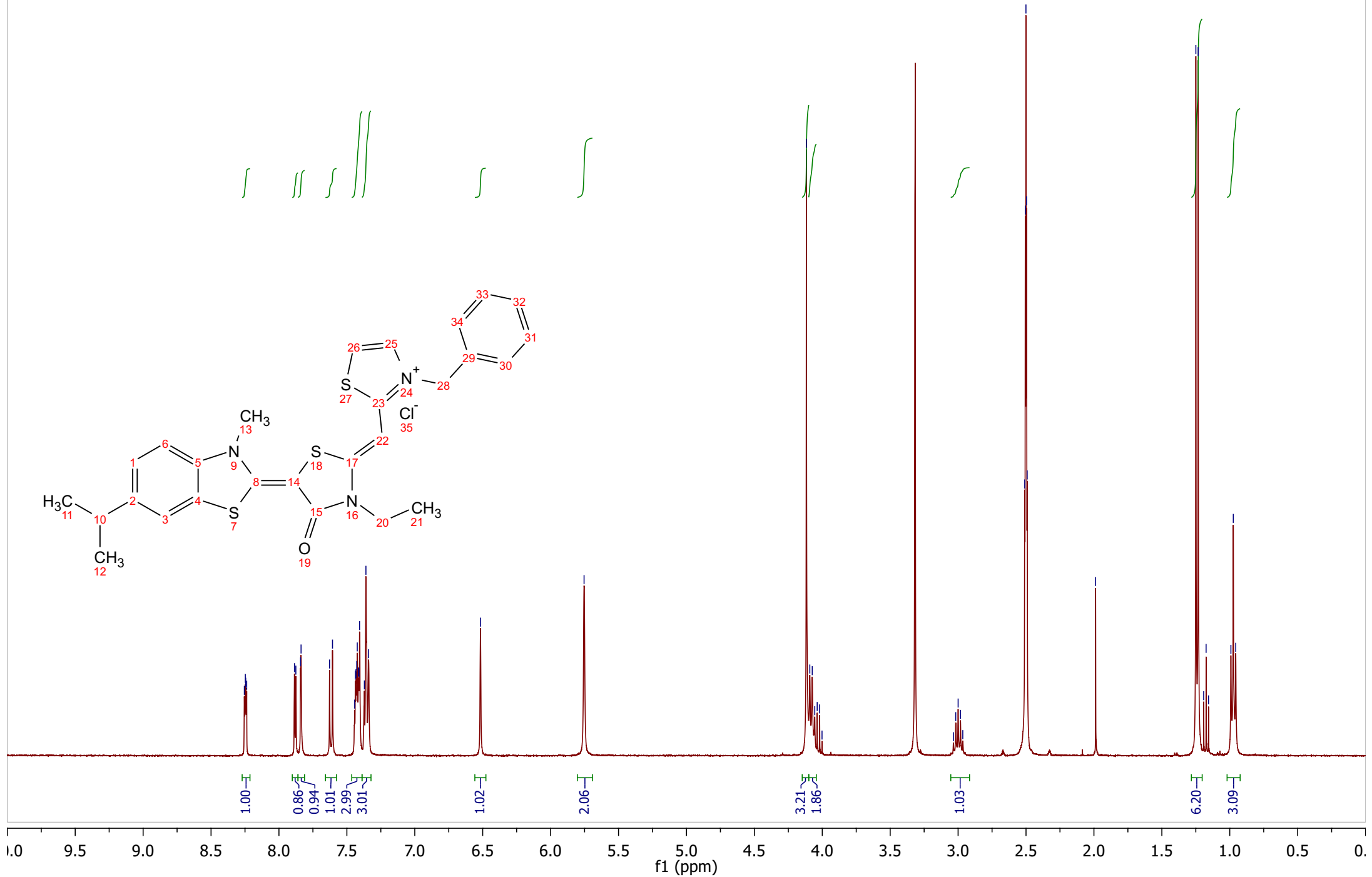
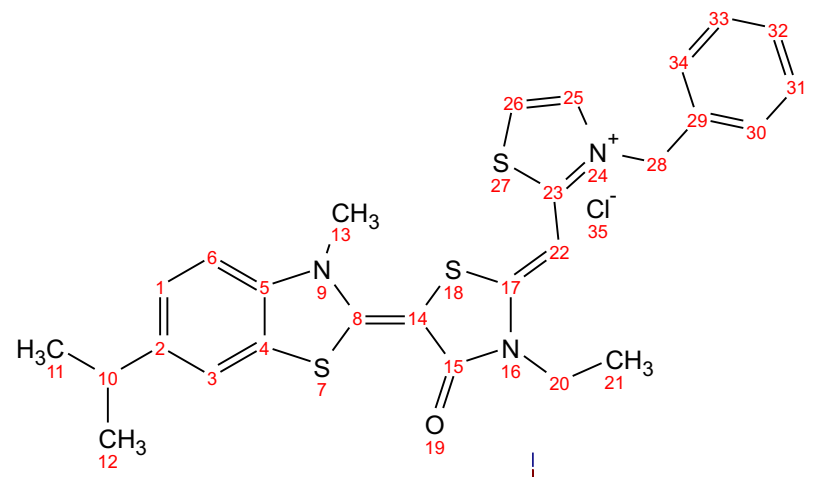
4.12
4.09
4.07
4.06
4.04
4.02
4.00

3.03
3.02
3.00
2.98
2.97
2.51
2.50
2.50
2.50
2.49

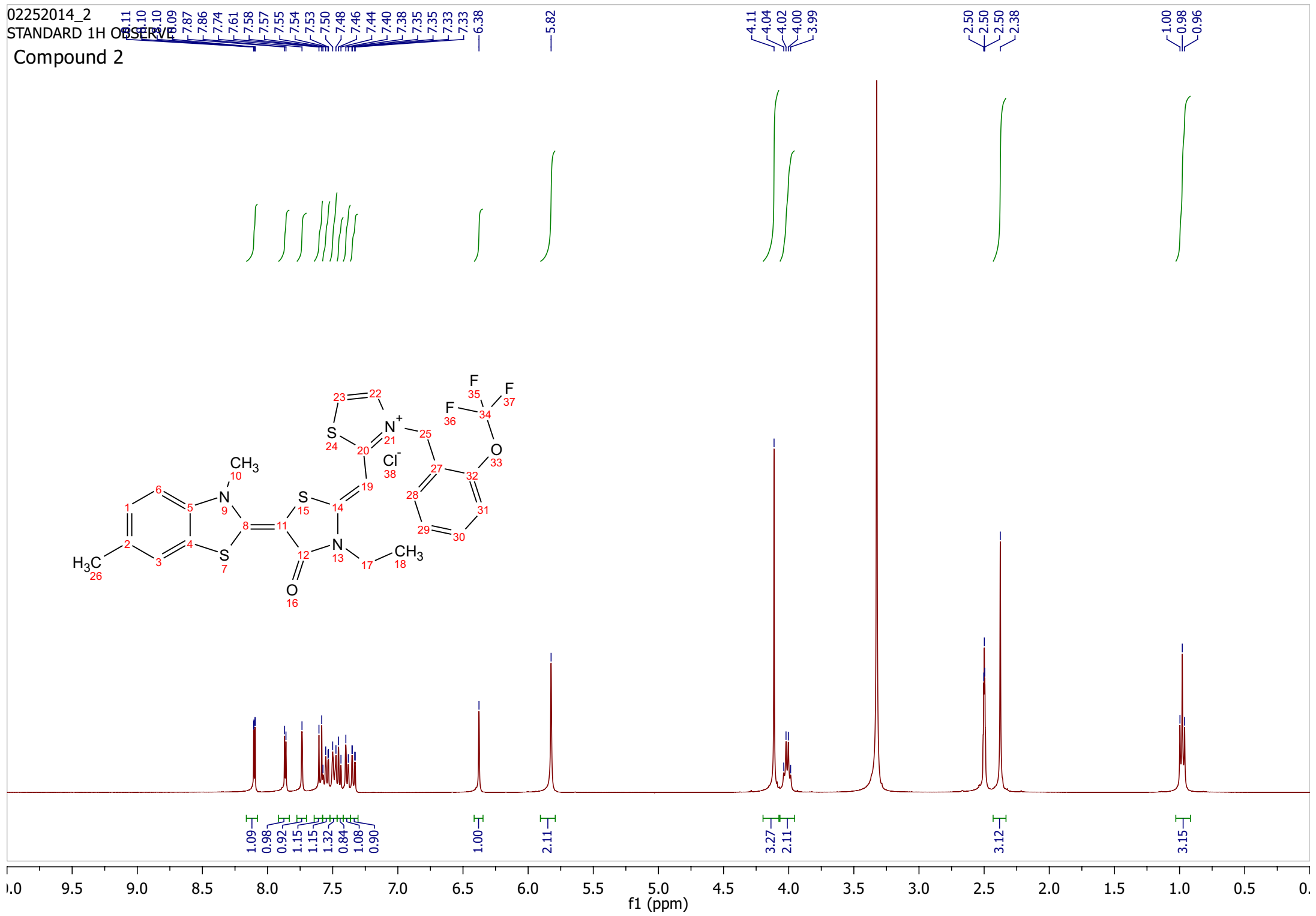
1.99

1.25
1.23
1.19
1.17
1.16
0.99
0.97
0.96

Compound 1

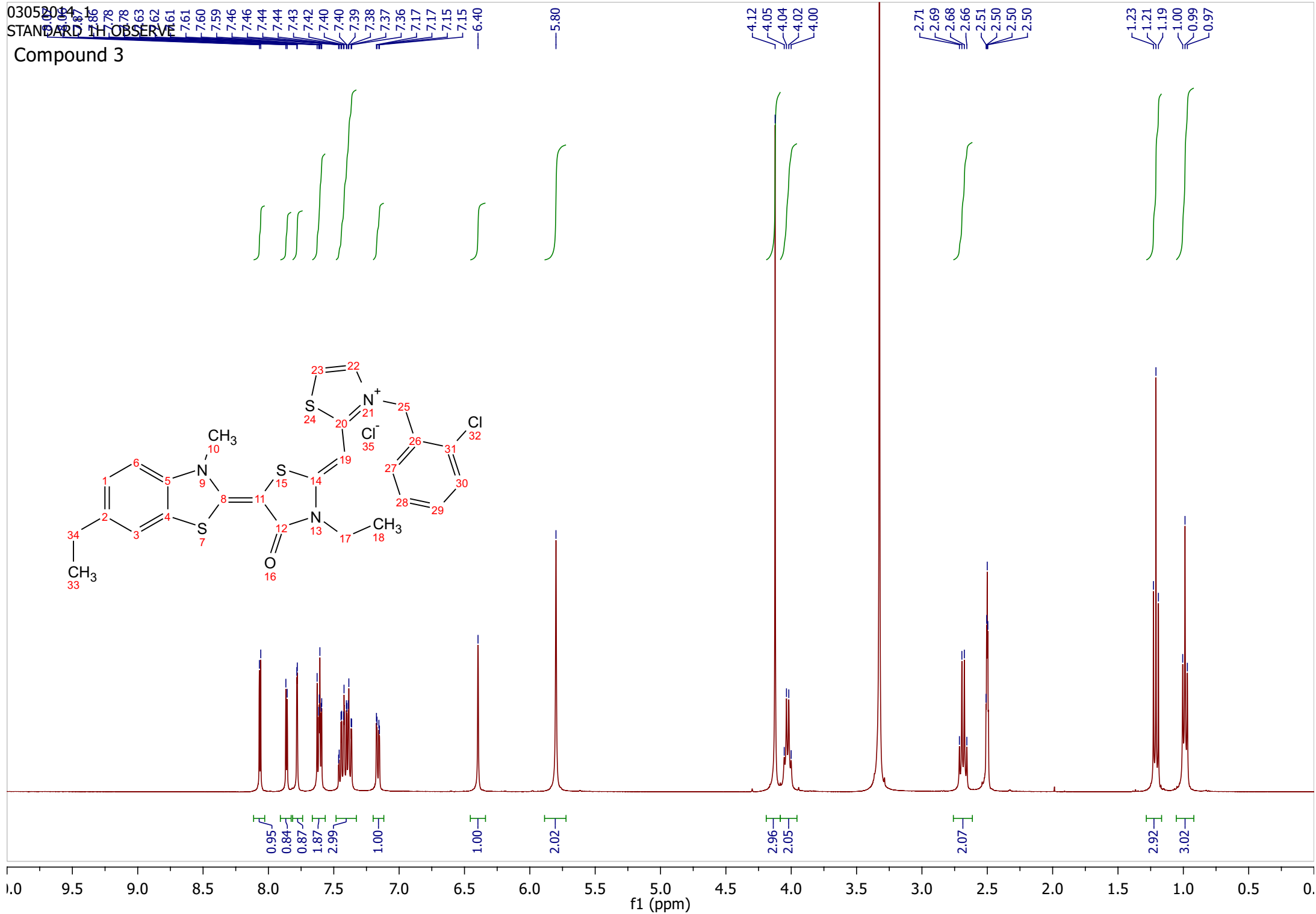


Compound 2



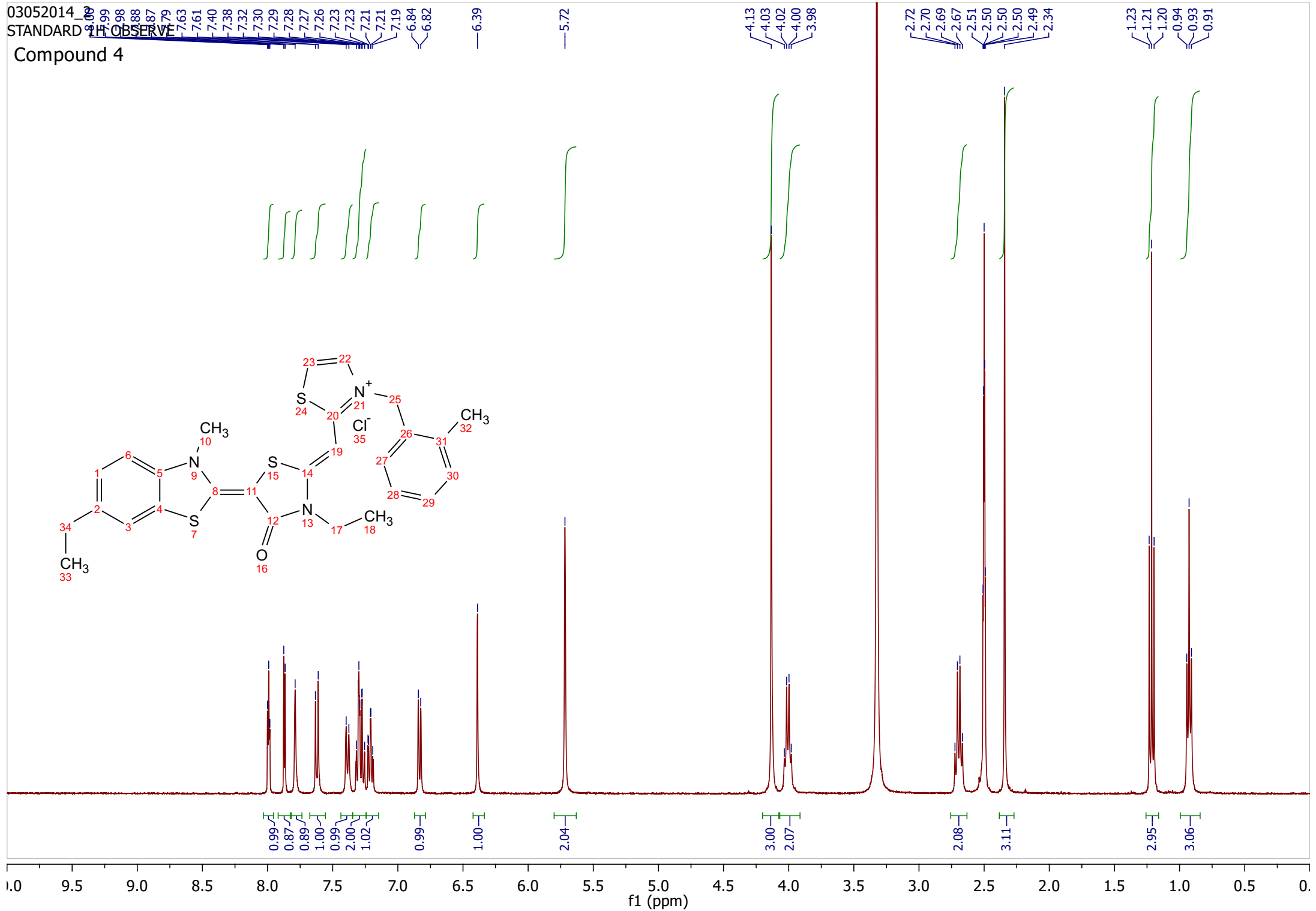
03052011
STANDARD 1H-OBSERVE

Compound 3

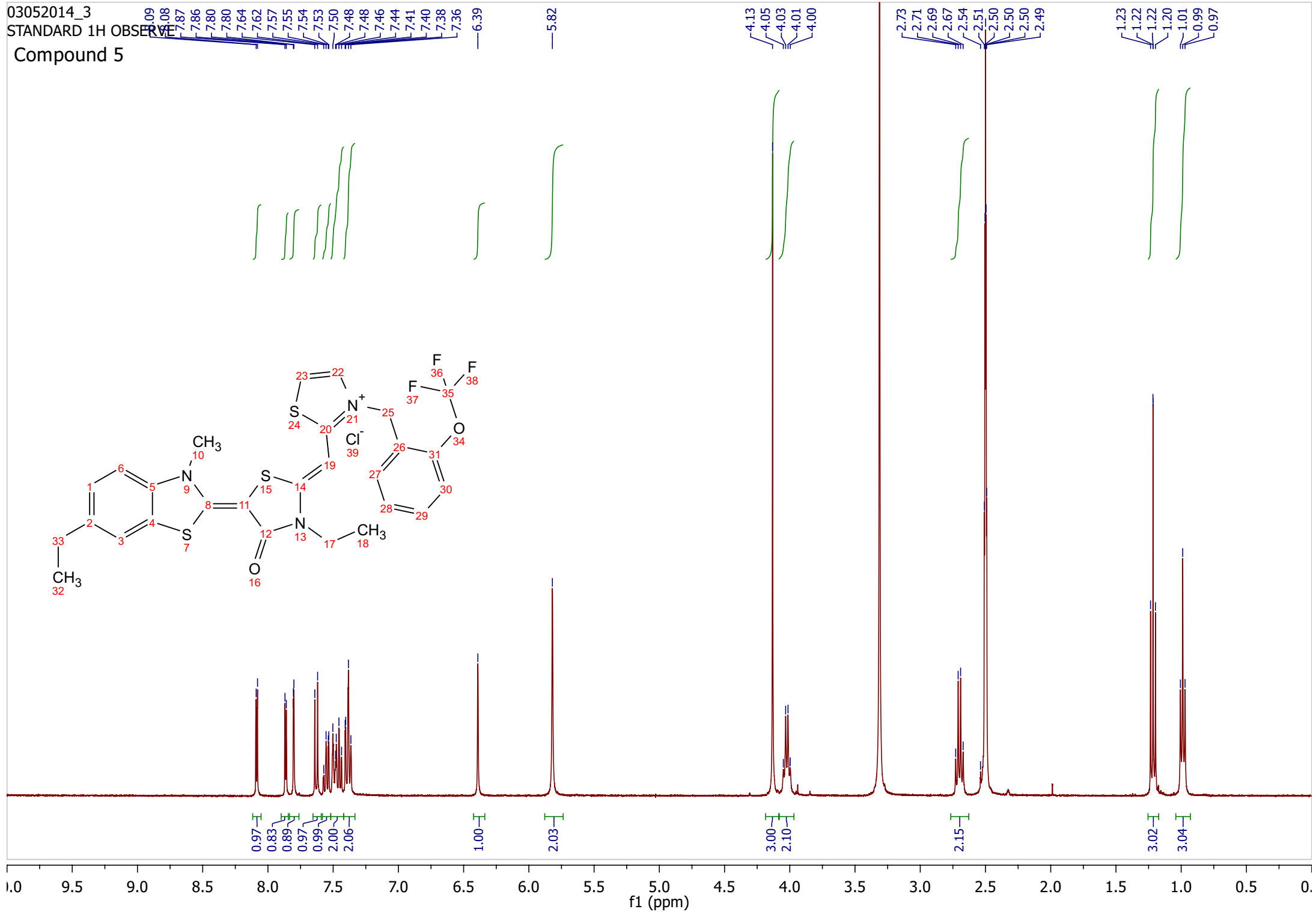


03052014_00
STANDARD 1H OBSERVE

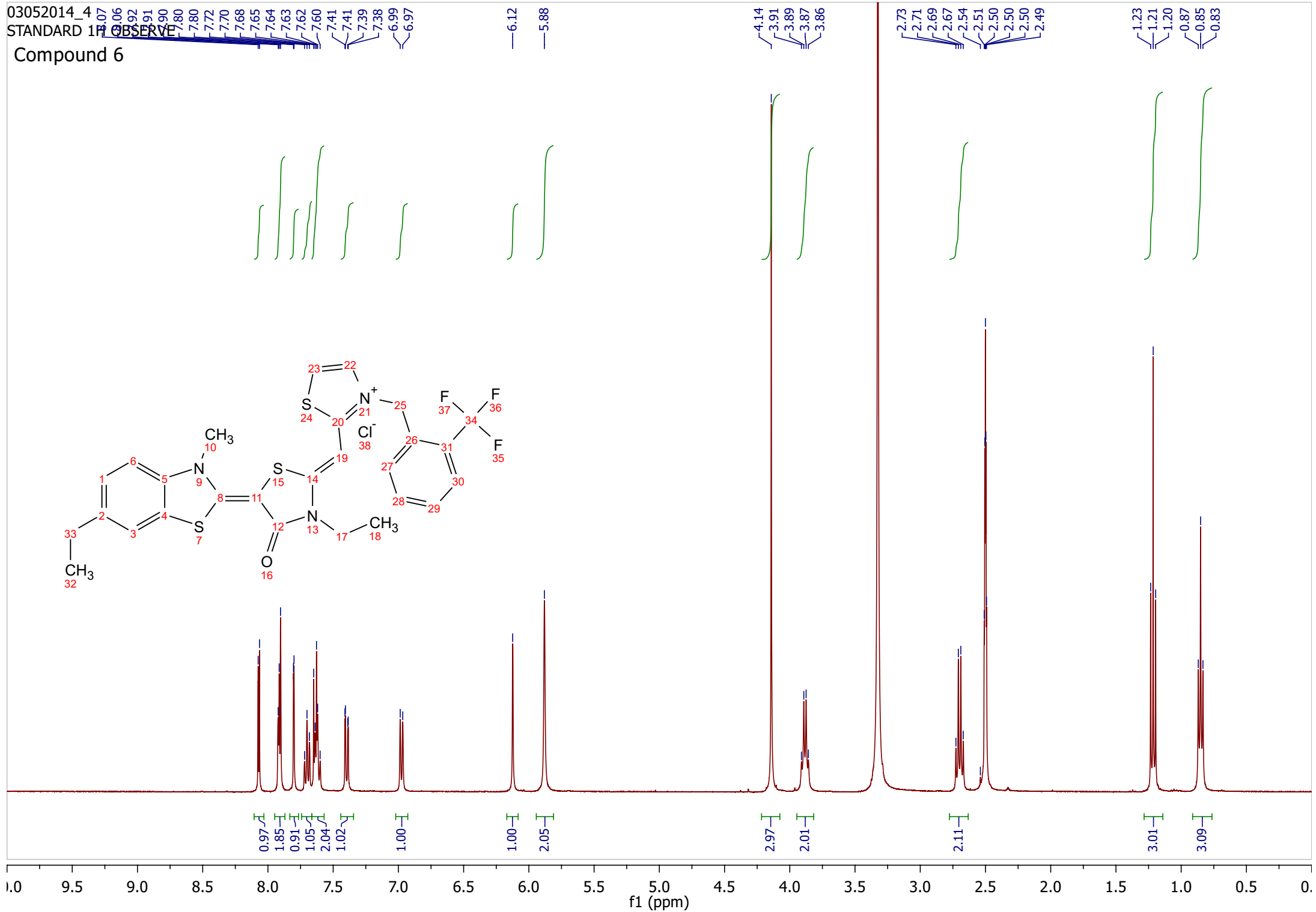
Compound 4



Compound 5

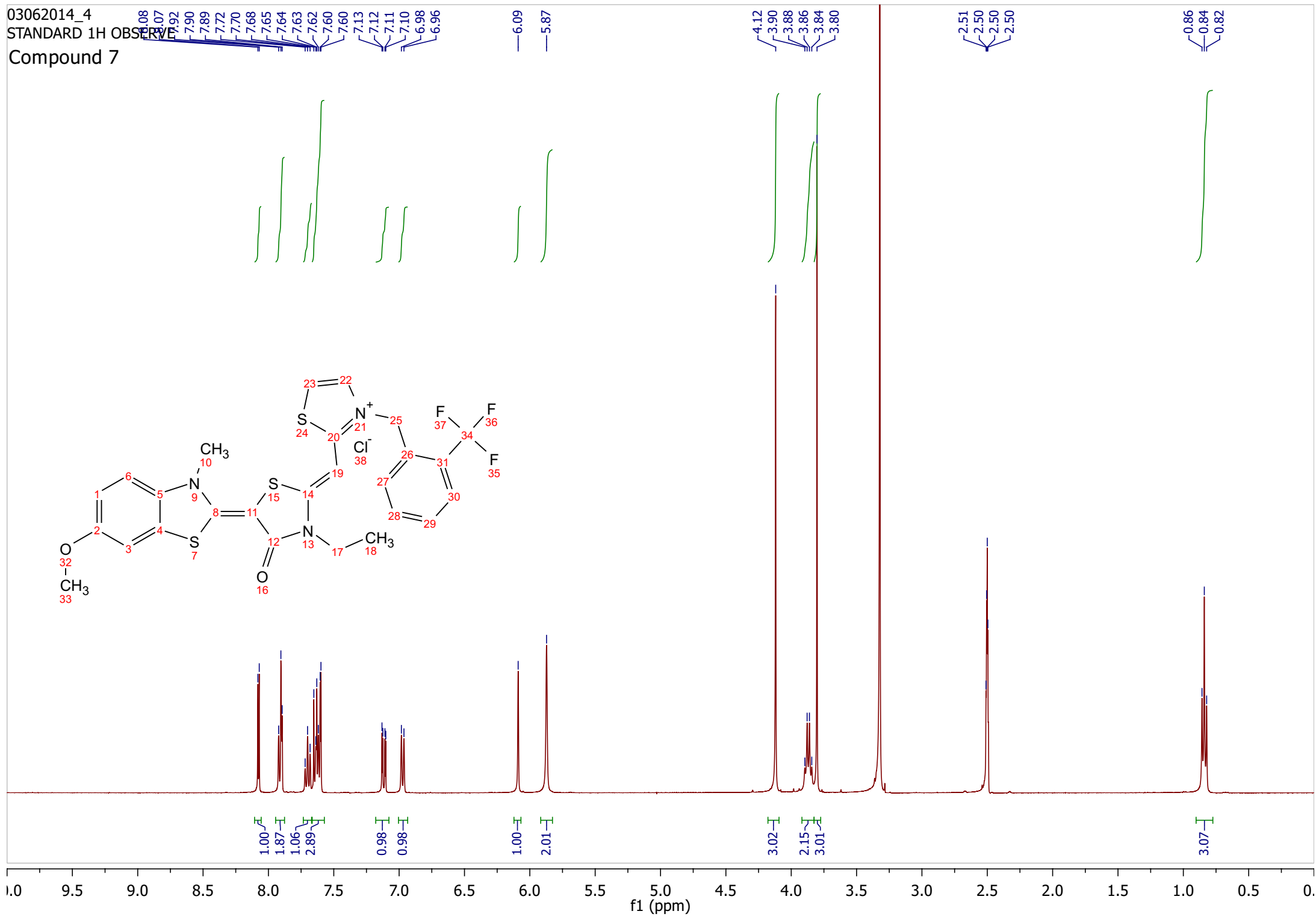


Compound 6



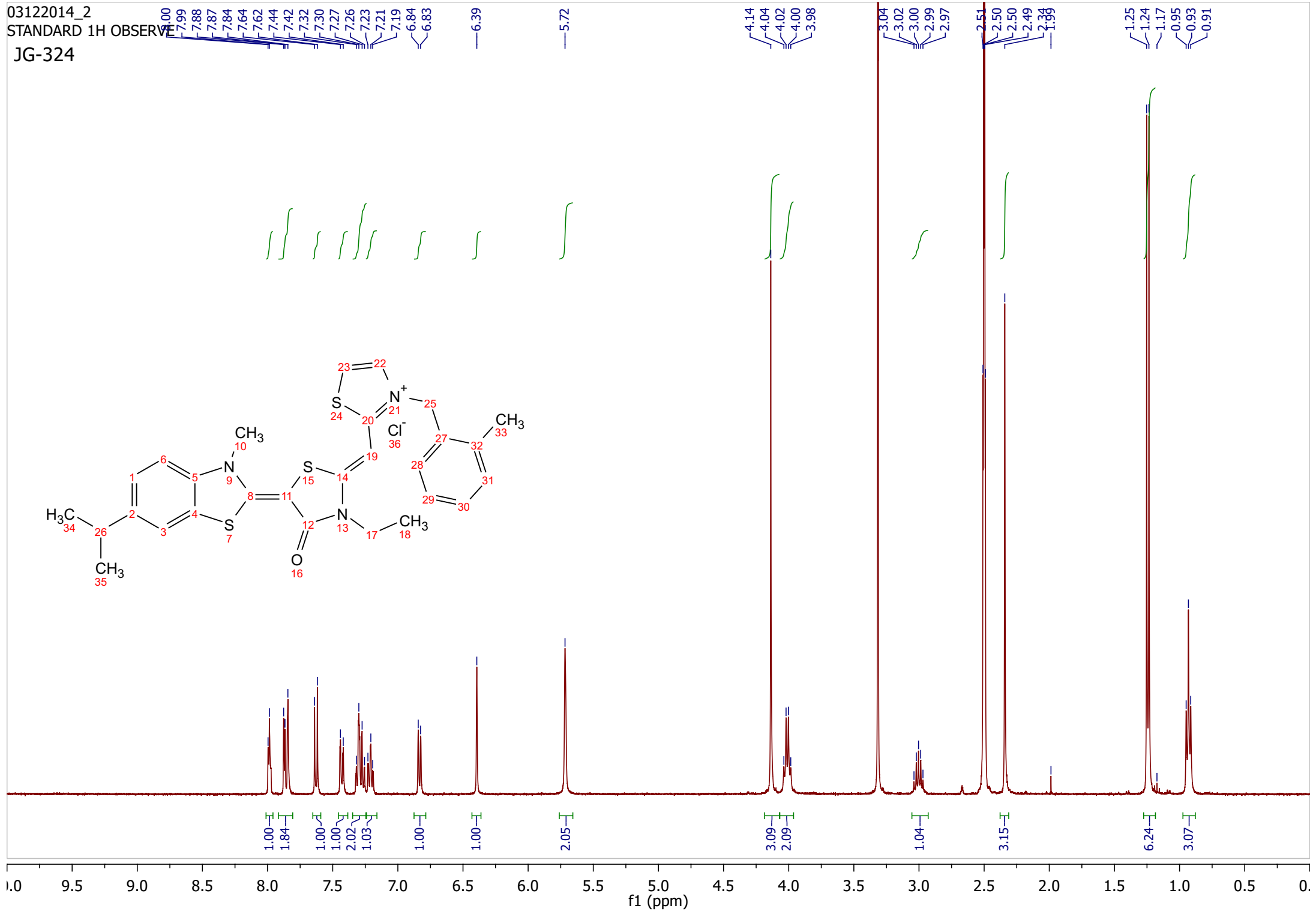
03062014_4
STANDARD 1H OBSERVE

Compound 7



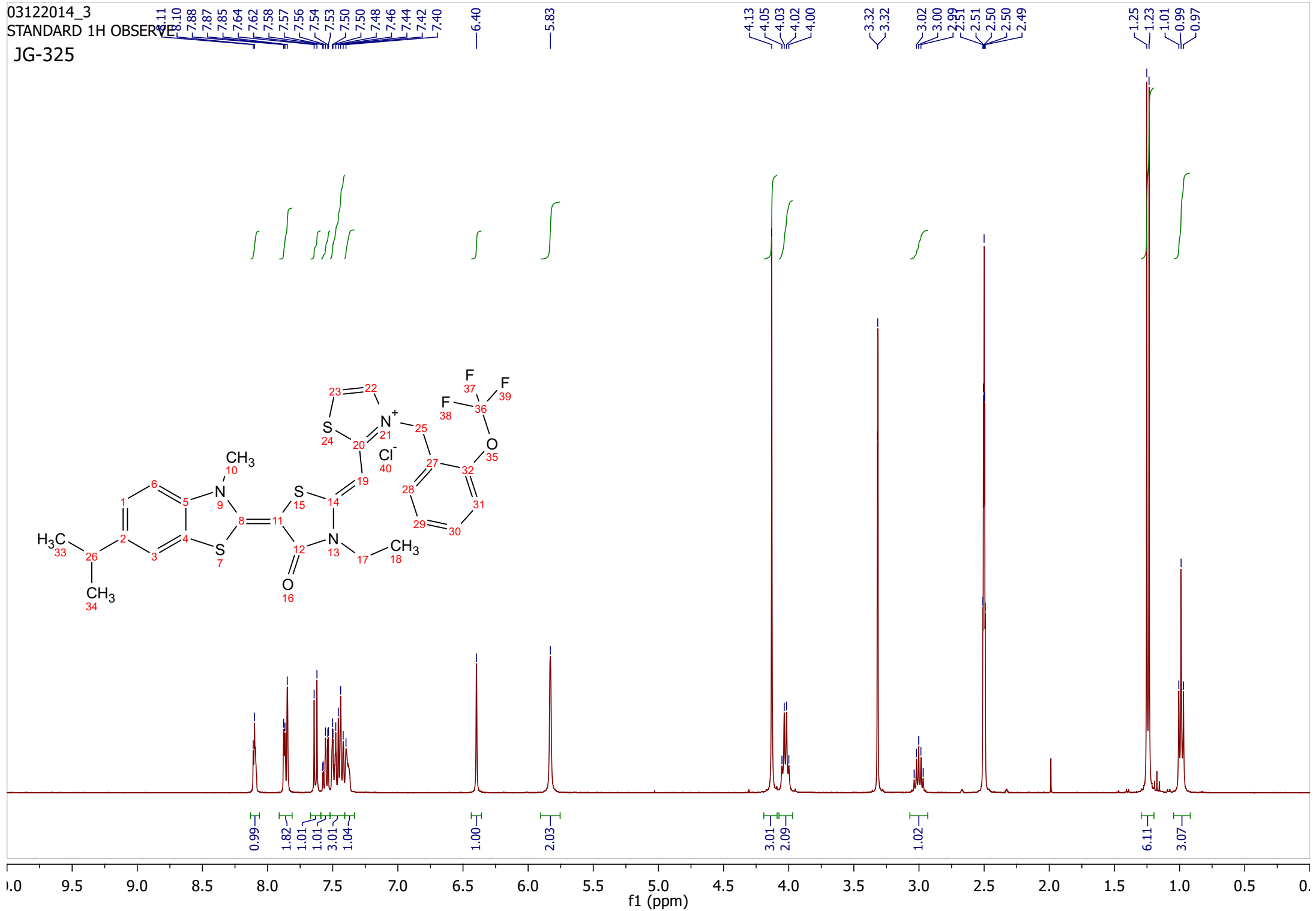
03122014_2
STANDARD 1H OBSERVE

JG-324

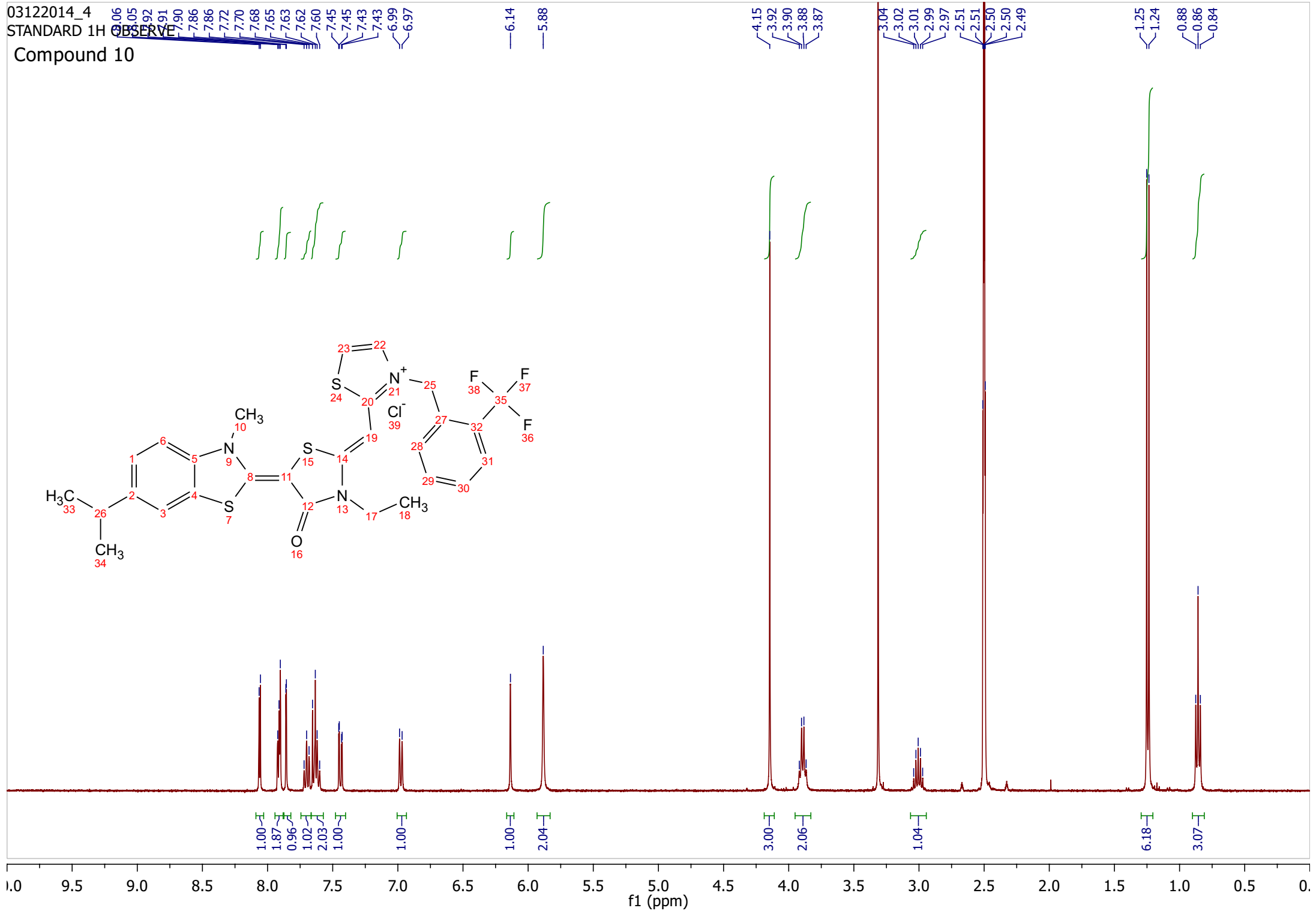


03122014_3
STANDARD 1H OBSERVE

JG-325

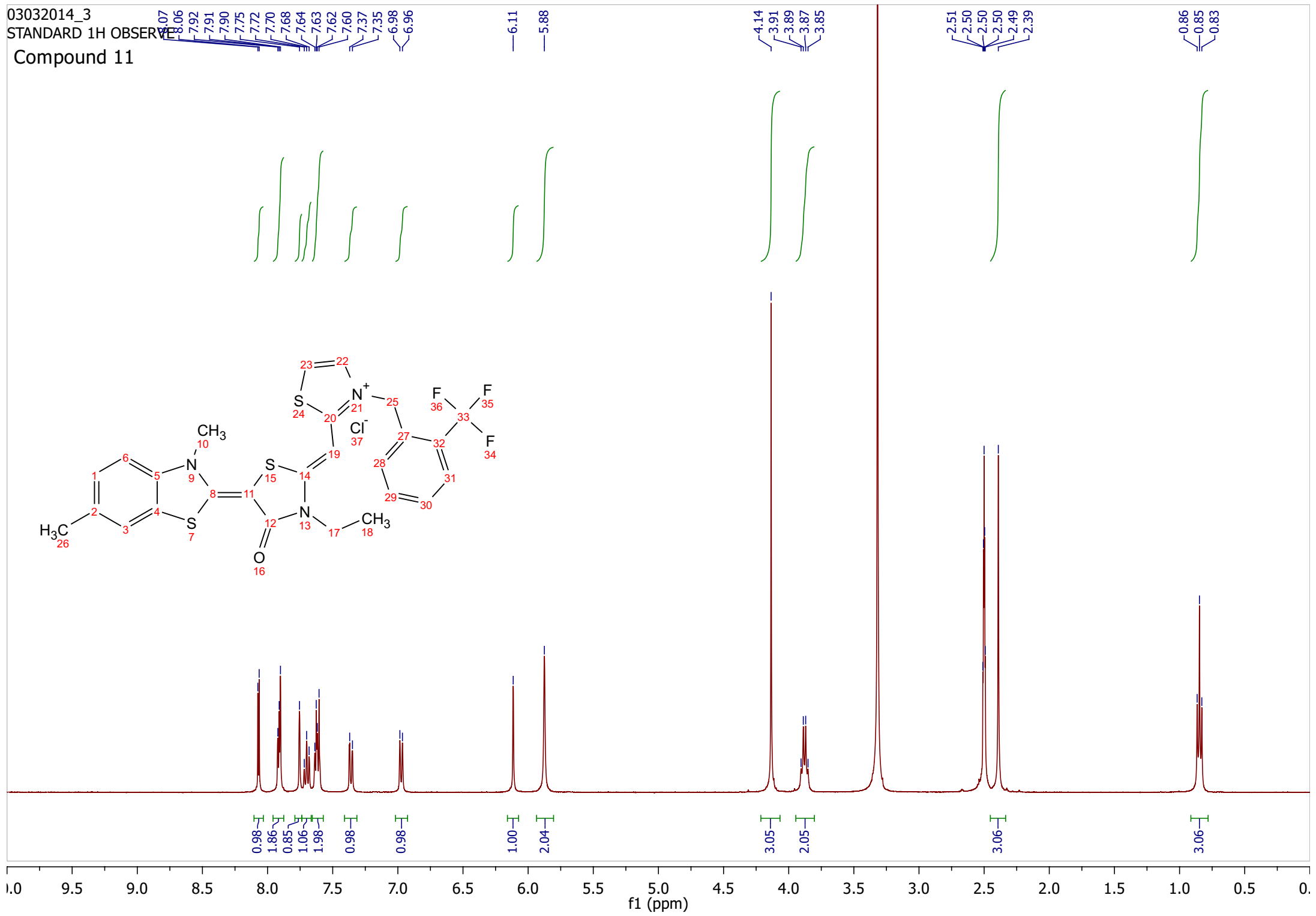


Compound 10



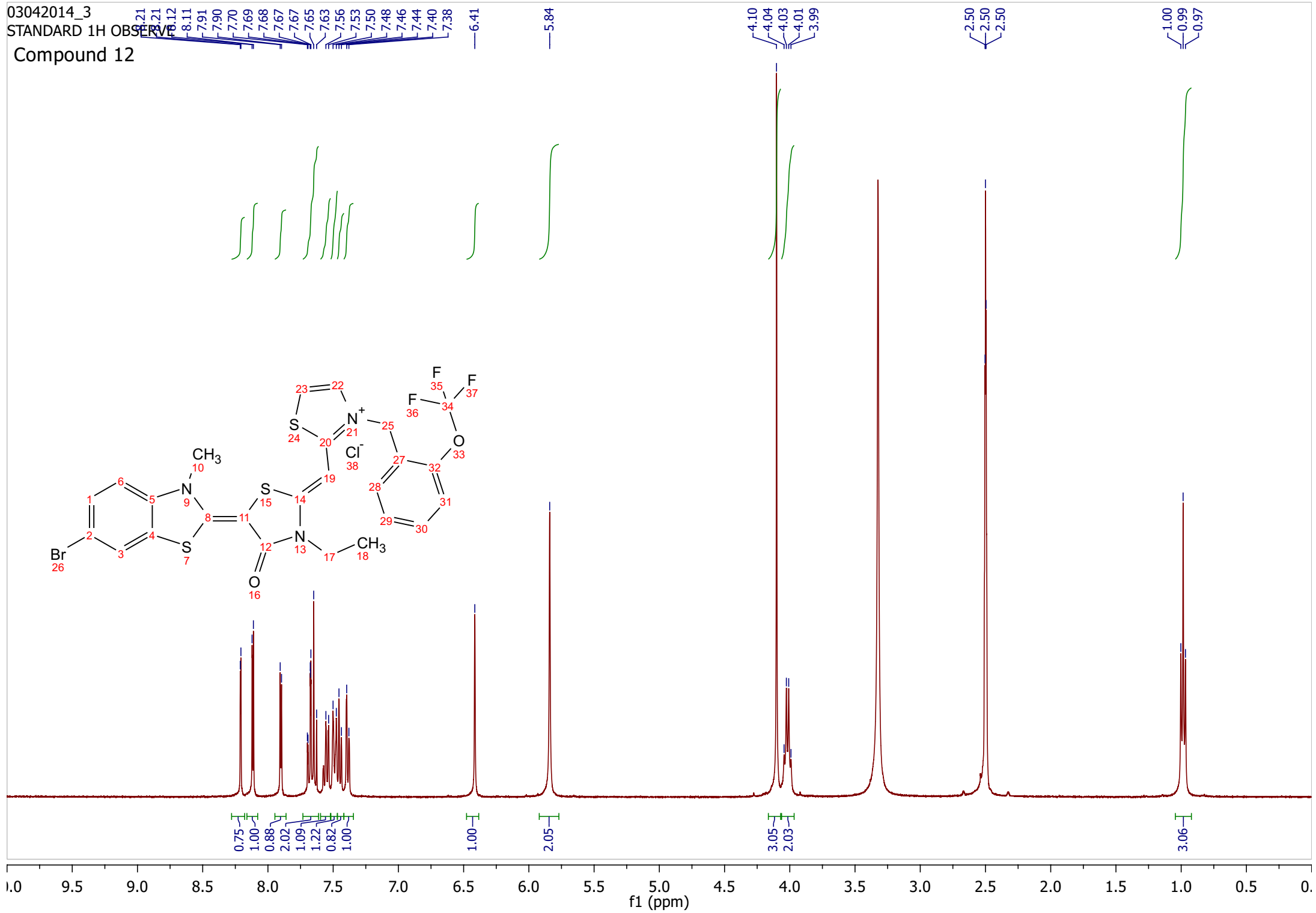
03032014_3
STANDARD 1H OBSERVE

Compound 11



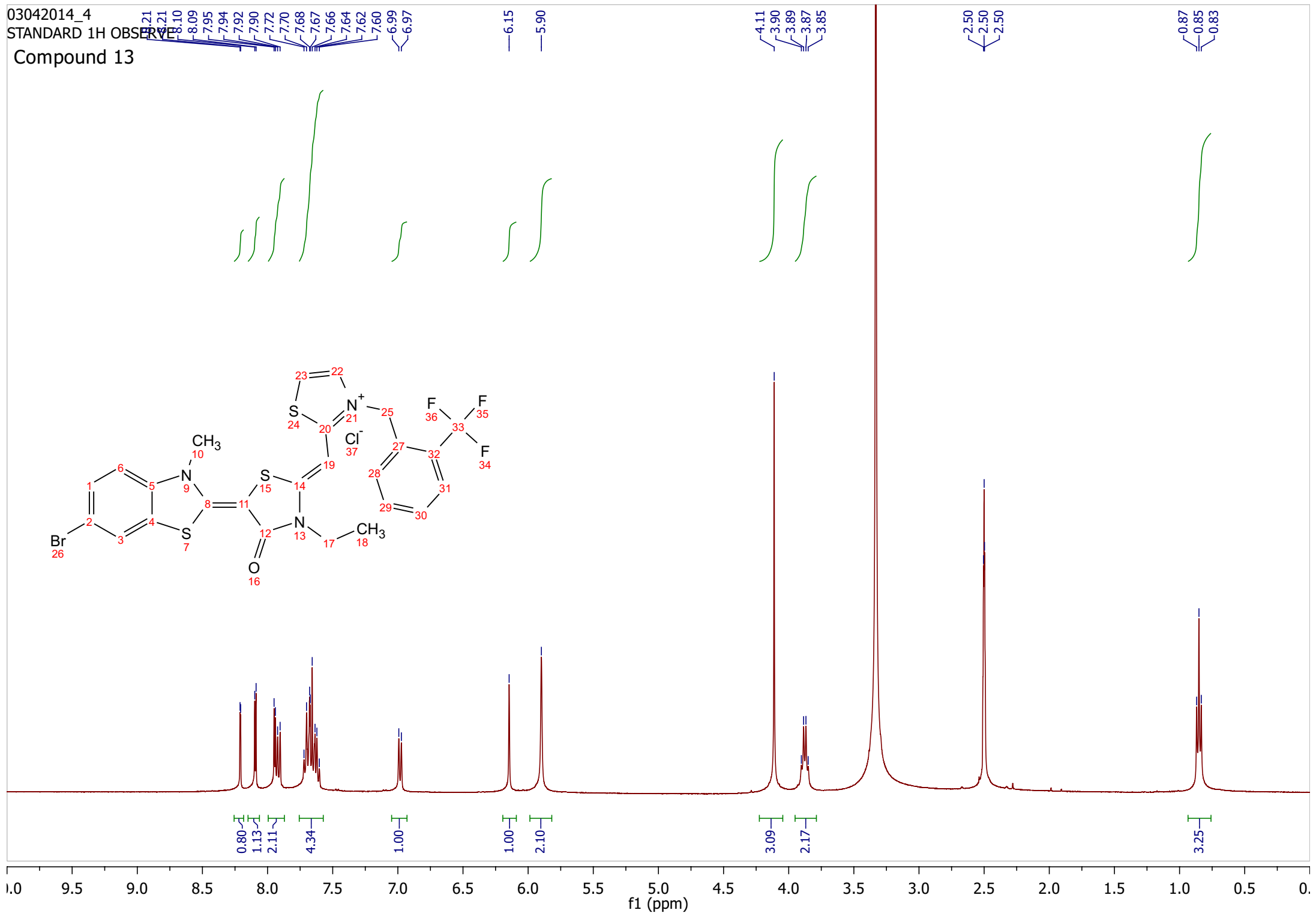
03042014_3
STANDARD 1H OBSERVE

Compound 12



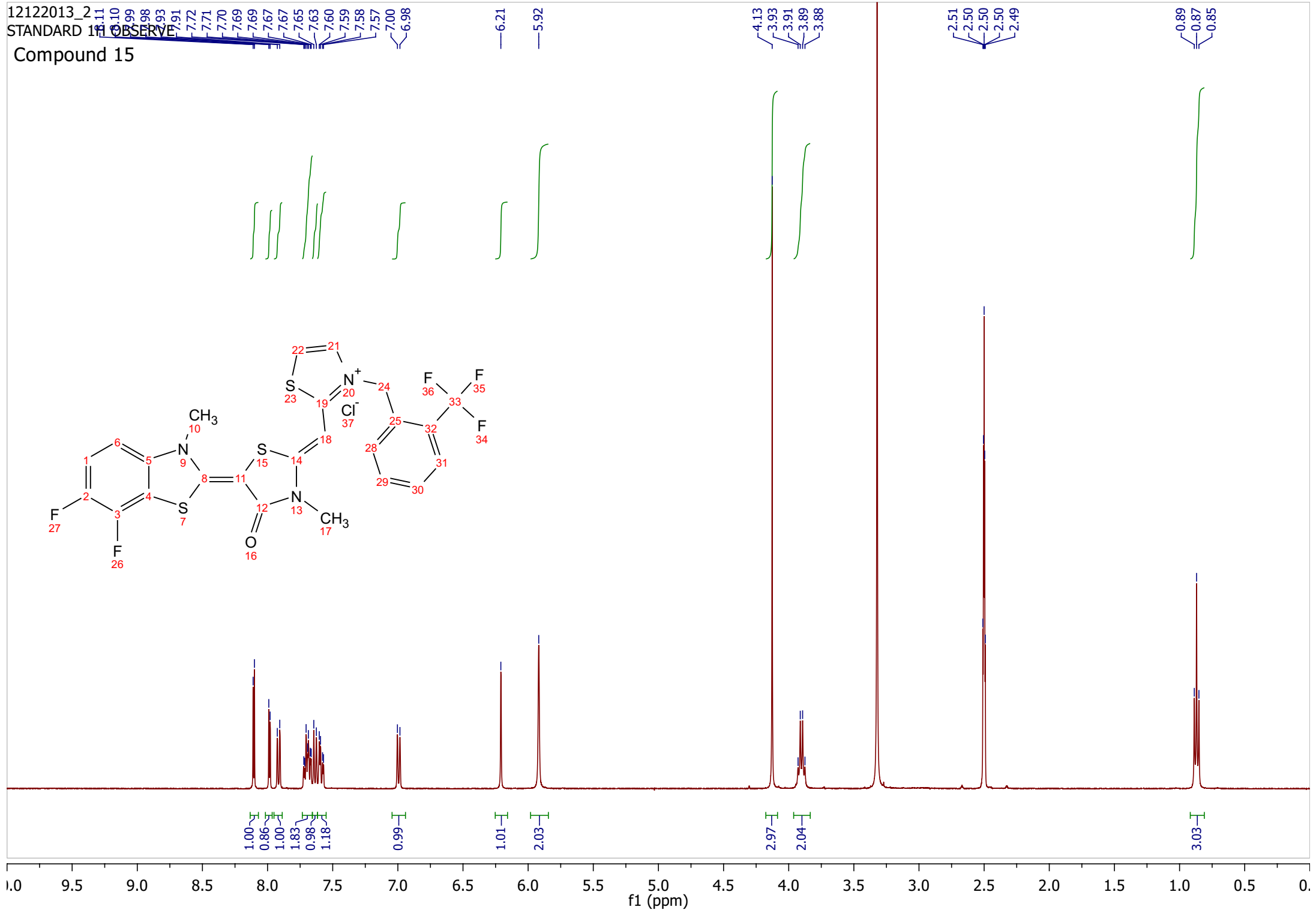
03042014_4
STANDARD 1H OBSERVE

Compound 13



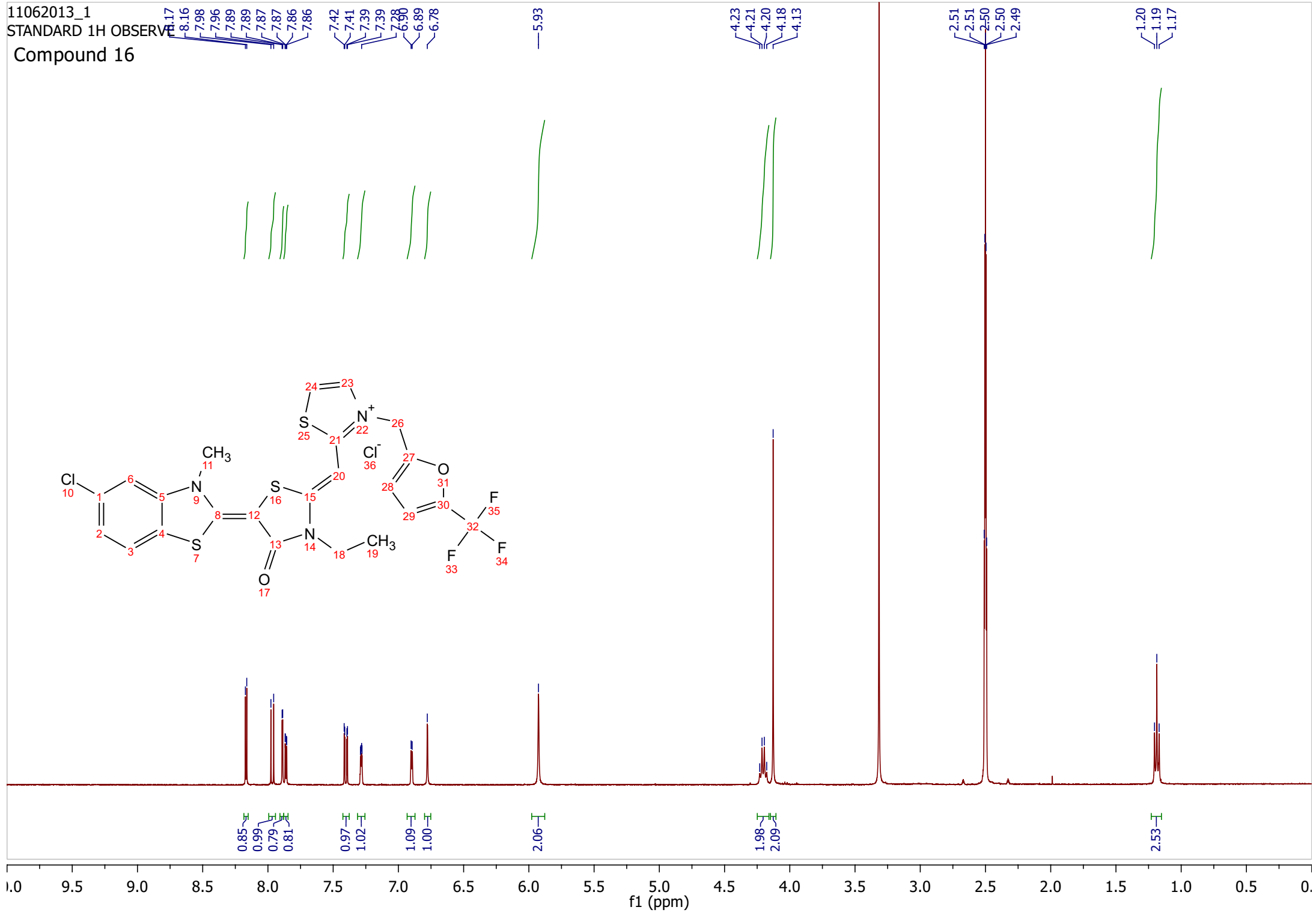
12122013_2
STANDARD 1H OBSERVE

Compound 15



11062013_1
STANDARD 1H OBSERVE

Compound 16



12132013_1
STANDARD 1H OBSERVE
Compound 17

8.23
8.22

7.92
7.91

7.65
7.58

7.29
7.28

6.93
6.83

5.98

4.24
4.22

4.21
4.19

3.32

2.50
2.50

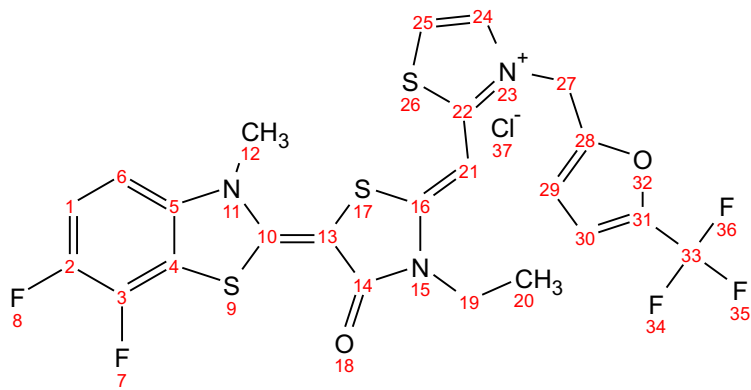
2.50
2.50

1.99

1.21
1.21

1.19
1.19

1.17
1.17



1.00

0.87

1.01

0.98

0.86

0.95

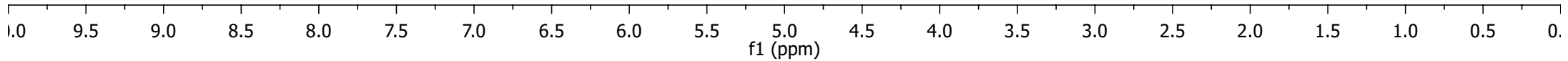
1.00

2.05

2.10

3.03

3.18



03122014_6
STANDARD 1H OBSERVE
Compound 18

8.17
8.17
8.16
8.16

7.85
7.85

7.84
7.83

7.64
7.62
7.44

6.92
6.91
6.77

5.93

4.24
4.22

4.20
4.18
4.13

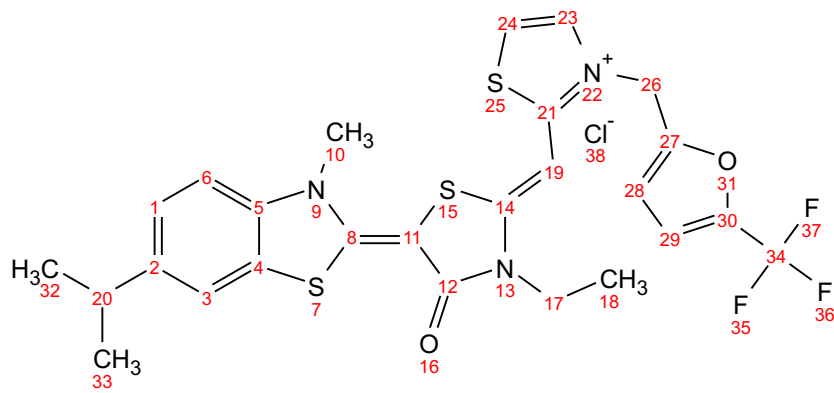
3.04
3.02

3.00
2.99

2.97
2.51
2.50
2.50
2.50
2.49

1.25
1.24
1.20

1.19
1.17



0.99

1.79

0.98

0.99

0.87

0.94

1.00

2.01

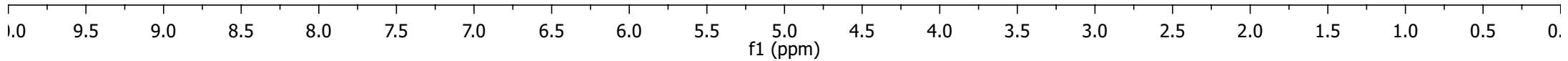
2.07

2.99

1.03

6.14

3.12



03052014_6
STANDARD 1H OBSERVE
Compound 19

8.15
8.14
8.14

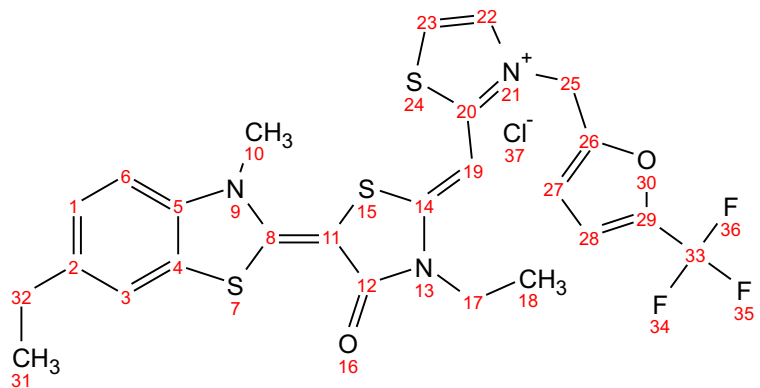
7.83
7.82
7.81
7.64
7.62
7.29
7.28
6.90
6.75

5.91

4.23
4.23
4.21
4.21
4.20
4.20
4.18
4.18
4.14
4.14

3.31

2.69
2.69
2.51
2.51
2.50
2.50
2.50
2.50
2.49
2.49
2.49
1.24
1.24
1.22
1.22
1.21
1.21
1.20
1.20
1.19
1.19
1.17
1.17



0.99

0.87

0.89

0.99

0.98

0.88

0.95

1.00

2.05

2.14

3.03

2.20

3.01

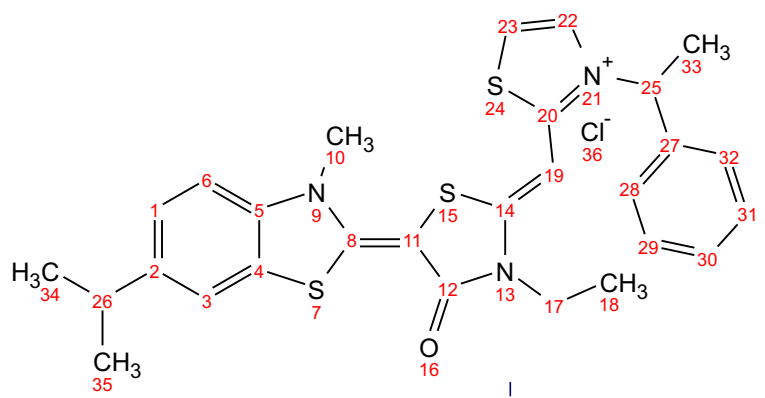
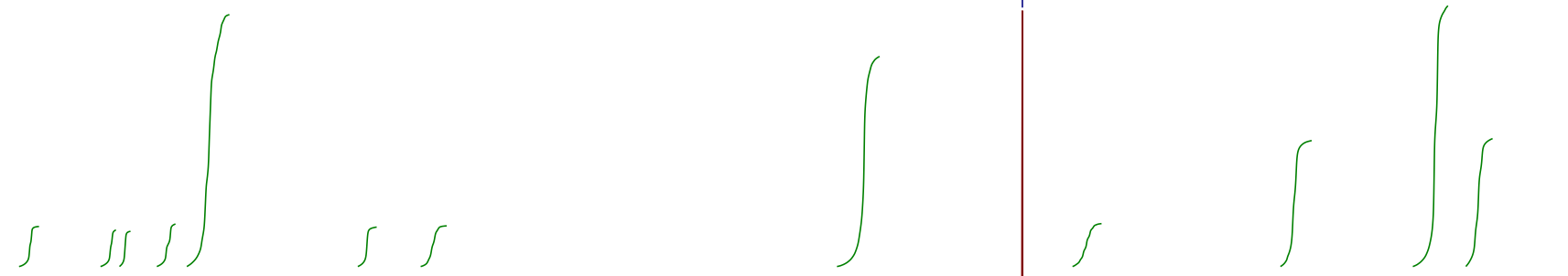
3.05

1.0 9.5 9.0 8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5 0.0
f1 (ppm)

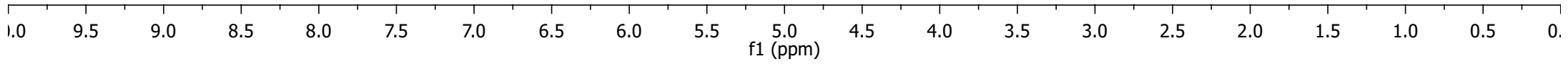
03102014_3
STANDARD 1H OBSERVE
JG-319

8.32
8.30
7.91
7.90
7.84
7.63
7.61
7.45
7.43
7.41
7.41
7.39
7.37
7.35
6.29
6.28
6.26

4.12
4.04
4.02
3.50
3.32
3.03
3.01
3.00
2.98
2.96
2.50
2.50
1.99
1.96
1.94
1.25
1.23
1.19
1.17
1.15
1.04
1.02
1.01

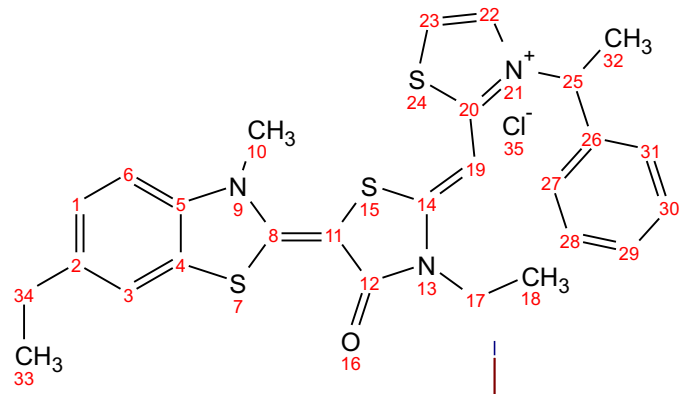
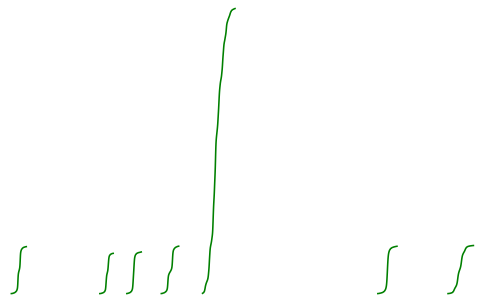


1.02
0.93
0.89
1.08
6.38
1.00
1.03
5.32
1.09
3.19
6.60
3.24



03102014_4
STANDARD 1H OBSERVE
JG-320

8.31 8.30 7.91 7.78 7.78 7.62 7.60 7.43 7.41 7.41 7.40 7.39 7.39 6.61 6.30 6.29 6.27 6.25 4.11 4.10 4.08 2.72 2.70 2.68 2.66 2.51 2.51 2.50 2.50 2.49 1.96 1.94 1.23 1.21 1.19 1.04 1.02 1.01



0.99

0.85

0.88

1.00

6.01

1.00

1.02

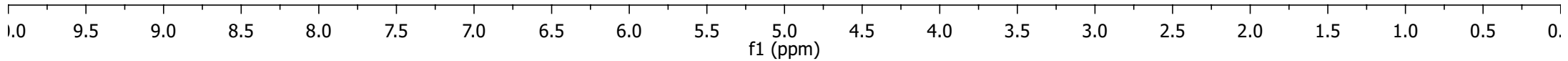
5.05

2.11

3.03

3.09

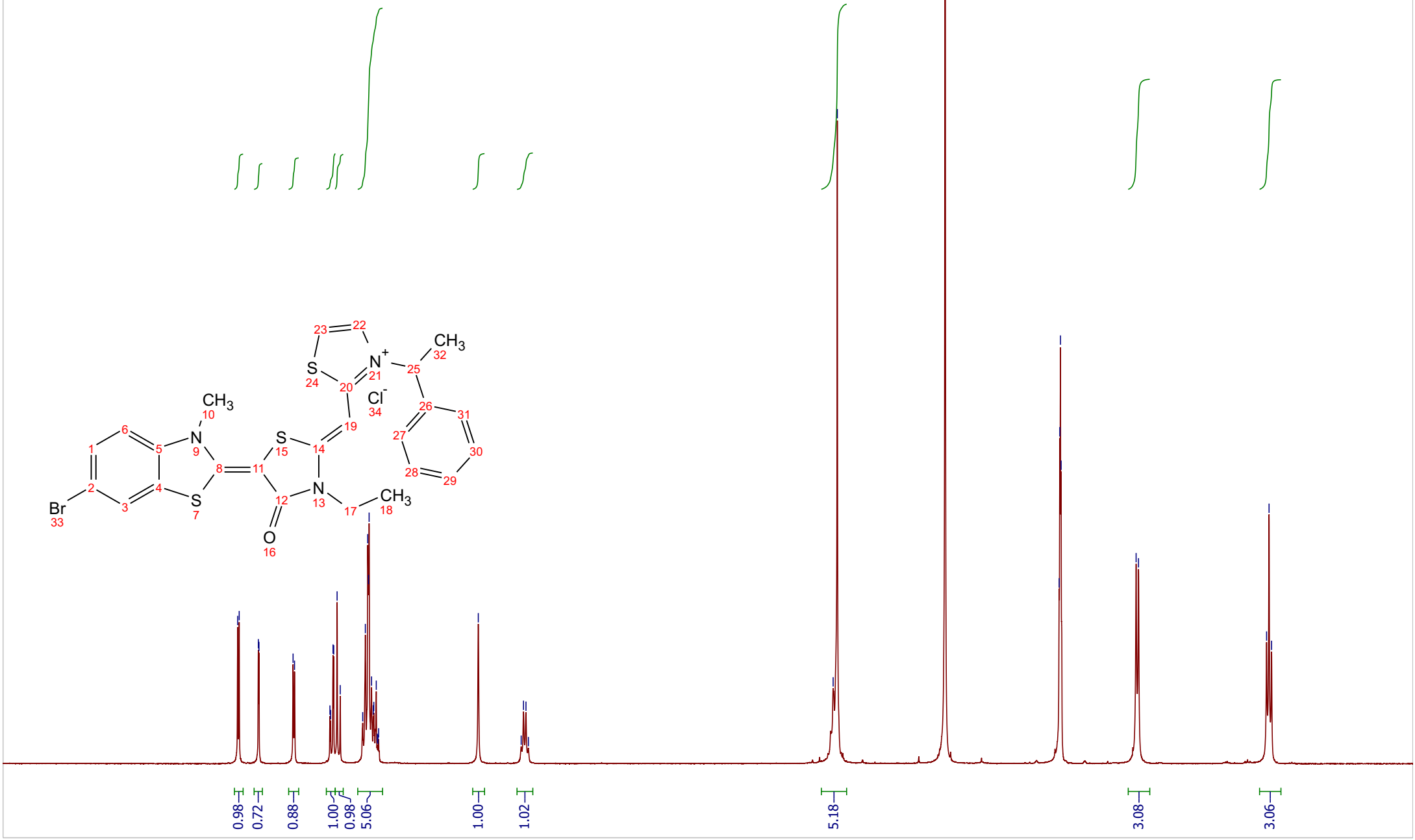
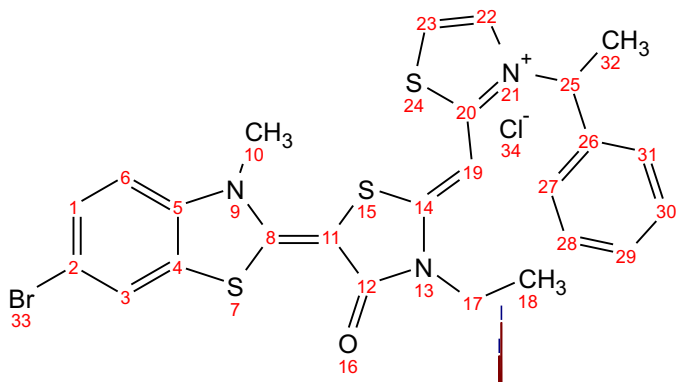
3.04



03112014_1
STANDARD 1H OBSERVE
Compound 22

8.34
8.32
8.19
8.18
7.94
7.93
7.66
7.65
7.63
7.43
7.41
7.41
7.40
7.39
7.25
6.32
6.31
6.29
6.27

4.11
4.08
2.51
2.50
2.50
2.50
1.96
1.95
1.04
1.02
1.00



1.0 9.5 9.0 8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5 0.0
f1 (ppm)

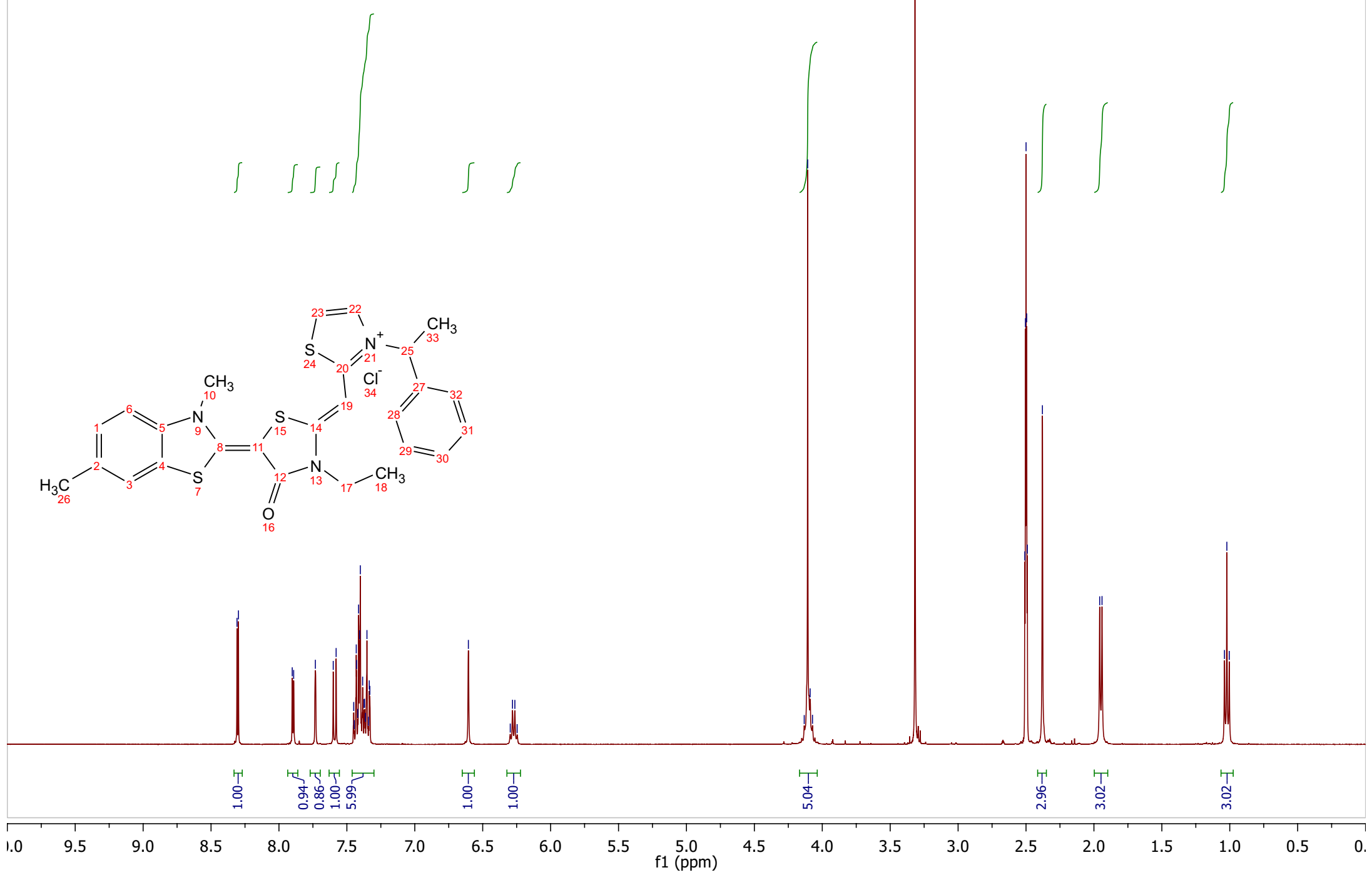
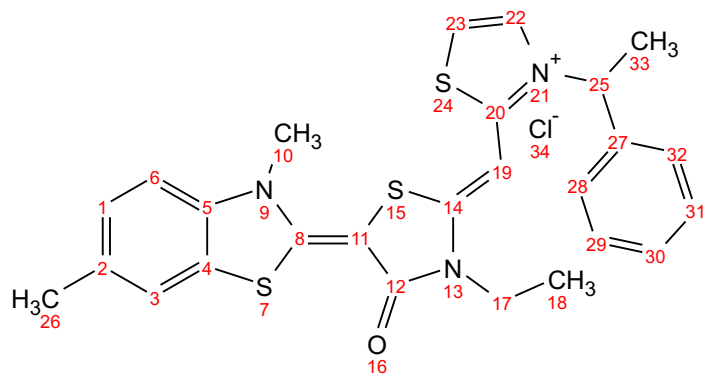
03102014_2
STANDARD 1H OBSERVE
Compound 23

8.31
8.30
7.90
7.89
7.73
7.60
7.58
7.43
7.43
7.41
7.41
7.40
7.38
7.35
6.61
6.30
6.28
6.26
6.25

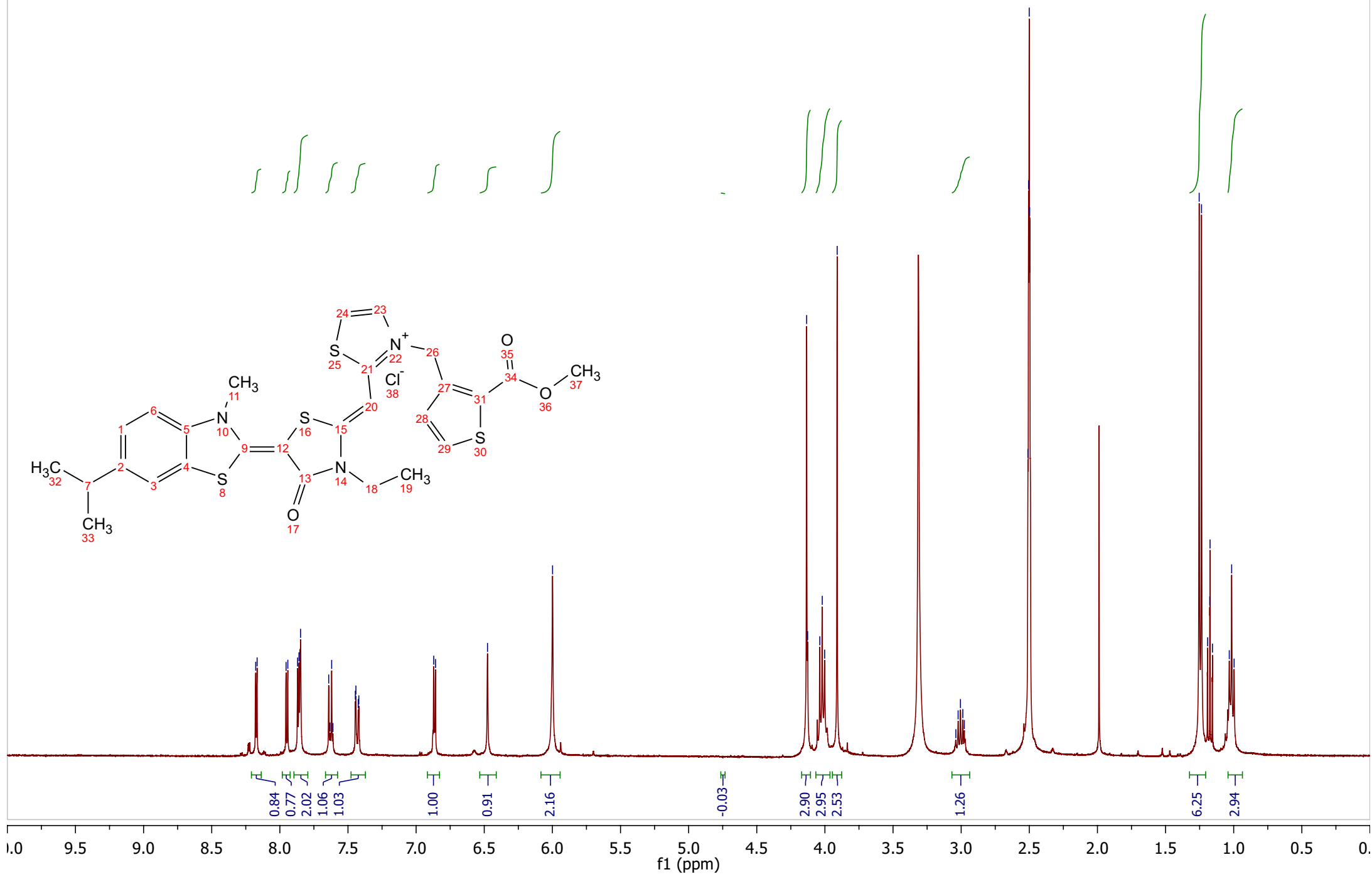
4.13
4.11
4.09
4.07

2.51
2.50
2.50
2.49
2.38
1.96
1.94

1.04
1.02
1.00

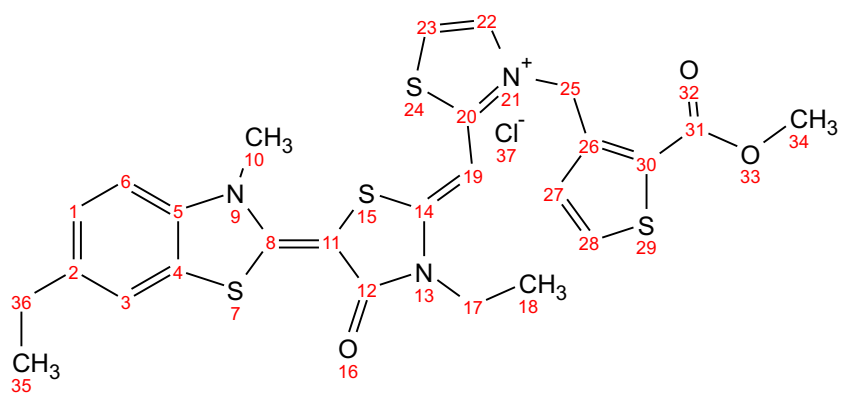
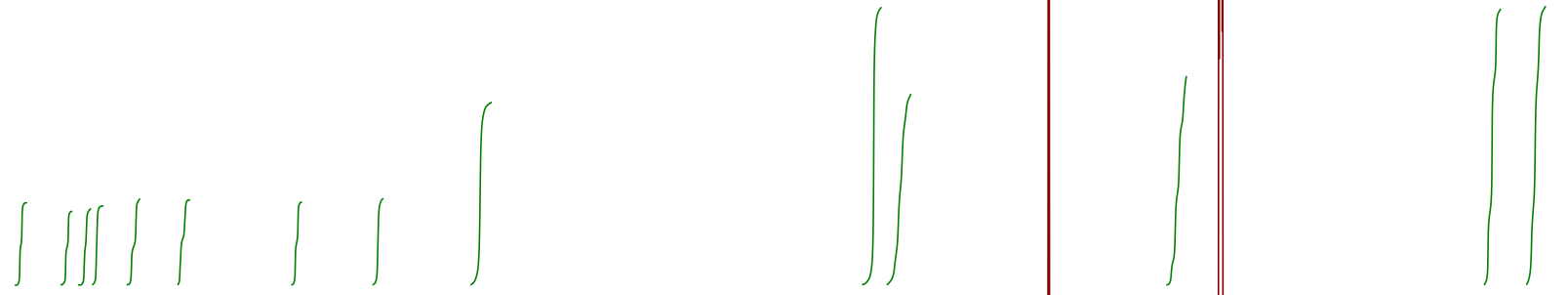


Compound 24

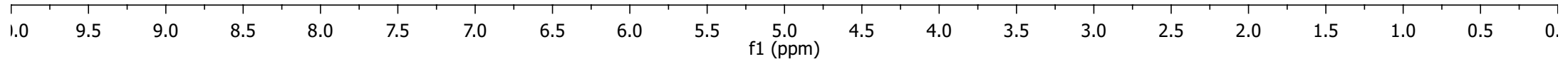


03052014_5
STANDARD 1H OBSERVE
Compound 25

8.17 8.16 7.95 7.94 7.87 7.85 7.80 7.64 7.62 7.41 7.39 6.87 6.85 6.48 6.00 4.14 4.02 4.00 3.91 3.83 2.73 2.71 2.69 2.67 2.54 2.51 2.50 2.50 2.49 2.49 2.33 1.24 1.22 1.20 1.03 1.01 1.00



0.96 0.85 0.89 0.92 1.00 0.99 0.96 1.00 2.12 3.23 2.22 2.43 3.21 3.24



12122013_1
STANDARD 1H OBSERVE
Compound 26

8.29
8.28

7.94
7.93

7.68
7.63

7.62
7.61

7.61
7.10

6.68

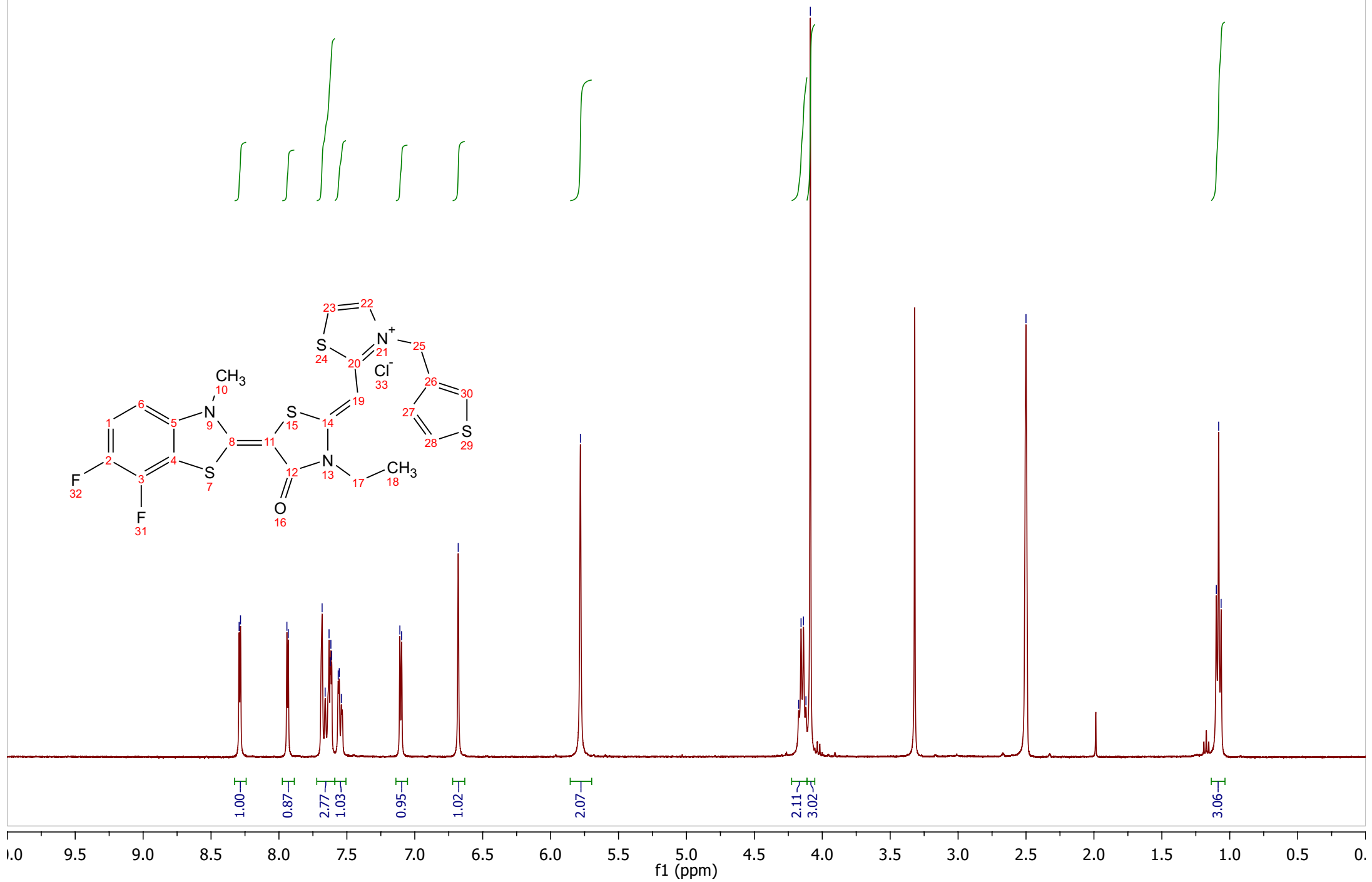
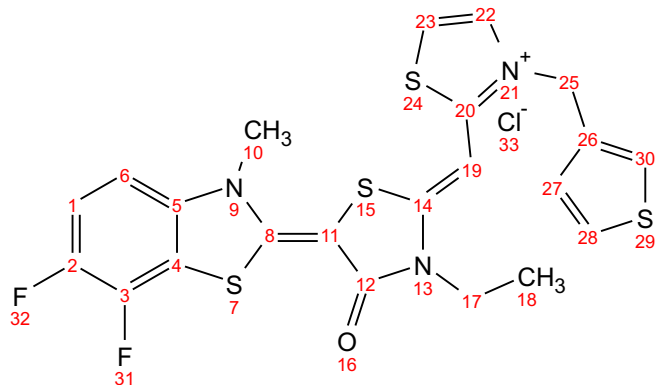
5.78

4.17
4.16

4.14
4.12
4.09

2.50

1.10
1.08
1.06



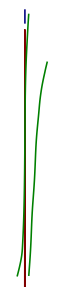
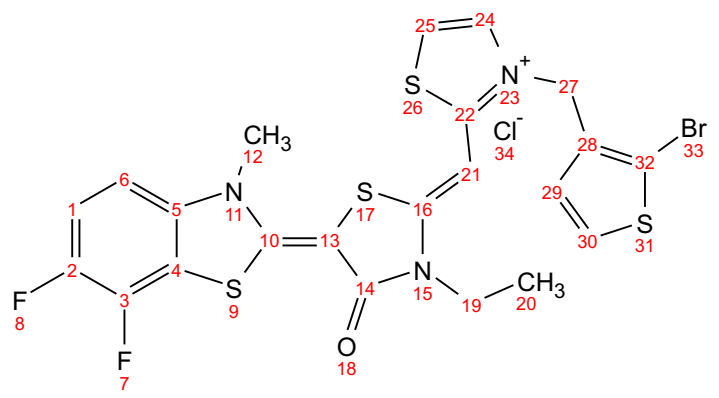
12132013_2
STANDARD 1H OBSERVE
Compound 27

8.16
8.15
7.94
7.93
7.72
7.70
7.68
7.66
7.64
7.59
7.58
7.57
7.56
6.94
6.93
6.45

4.11
4.08
4.06
4.05

2.50
2.50
2.50

1.13
1.11
1.09



1.00

0.95

0.76

1.03

1.01

0.89

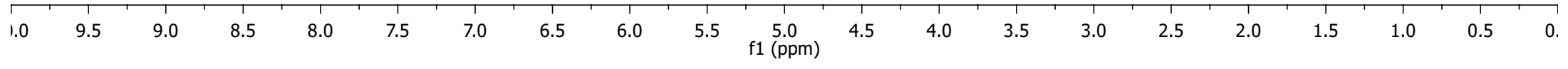
1.02

2.17

2.13

1.74

2.31



12062013_6
STANDARD 1H OBSERVE
Compound 28

8.22
8.21
7.92
7.91
7.69
7.66
7.64
7.58
7.56
7.55

6.68

6.35

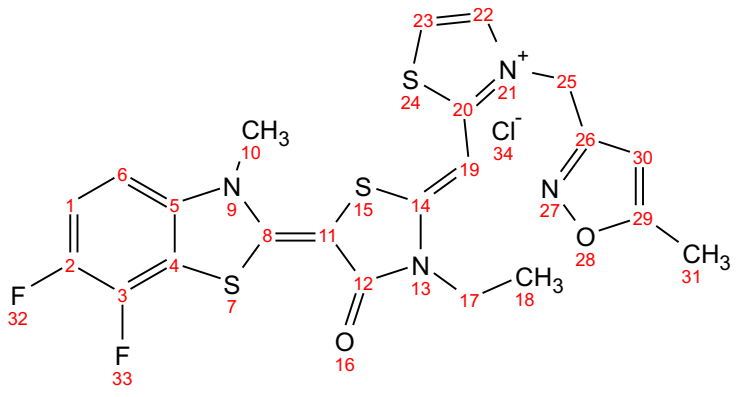
5.88

4.13
4.11

3.32

2.51
2.50
2.50
2.50
2.49
2.40
2.40

1.16
1.14
1.12



1.00

0.85

0.98

1.00

1.02

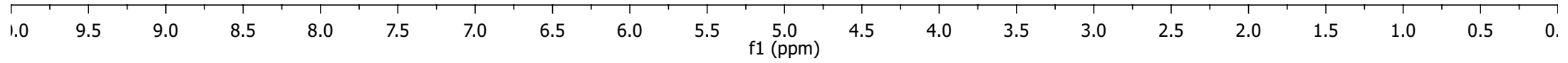
0.85

2.02

5.10

2.96

3.11



02252014_5
STANDARD 1H OBSERVE
JG-273

8.48
8.47
7.84
7.83
7.61
7.60
7.59
7.58
7.57
7.56
7.18
6.92
6.90
6.89

6.08

3.81
3.80
3.67
3.65
3.64
3.33
3.32
3.10
3.08
3.06

2.50

1.16
1.14
1.12

