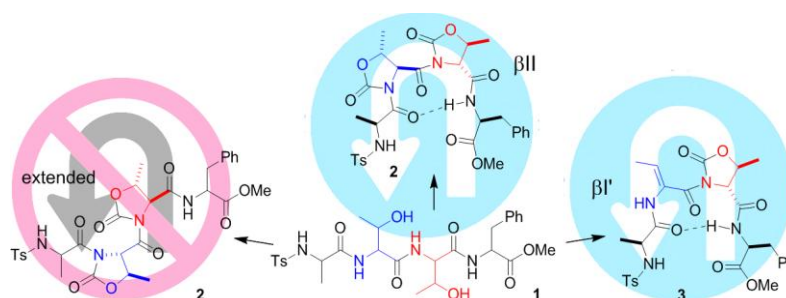


In-Peptide Synthesis of Di-Oxazolidinone and Dehydroamino Acid-Oxazolidinone Motives as β -Turns Inducers *Supporting Information*

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Constraining a peptide in a snap: linear peptides **1** give rise in a single step to sequences Oxd²-Oxd³ (**2**) or Δ Abu²-Oxd³ (**3**); these scaffolds adopt well defined extended or folded conformations, in particular normal or inverse β -turns of type I or II.

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¹H-NMR characterization of the model compounds **2a,c,d**, and **3b,c** at 400 MHz in 8:2 DMSO-d₆/H₂O.

Ts-Ala-(5'-Me-Oxd²)-(5'-Me-Oxd³)-PheOMe (**2a**). ¹H-NMR δ: 1.18 (d, J=6.7 Hz, 3H, AlaMe), 1.44-1.47 (m, 6H, Oxd³Me+Oxd²Me), 2.37 (s, 3H, TsMe), 2.94 (dd, J=8.4, 13.9 Hz, 1H, PheHβ), 3.01 (dd, J=5.7, 13.9 Hz, 1H, PheHβ), 3.56 (s, 3H, OMe), 4.48-4.52 (m, 2H, PheHα+Oxd³H₅), 4.57 (d, J=1.9 Hz, 1H, Oxd³H₄), 4.66 (dq, J=2.0, 7.0 Hz, 1H, Oxd²H₅), 5.02 (dq, J=6.7, 8.5 Hz, 1H, AlaHα), 5.17 (d, J=2.0 Hz, 1H, Oxd²H₄), 7.11 (d, J=7.6 Hz, 2H, PheArH), 7.14-7.26 (m, 3H, PheArH) 7.37 (d, J=8.0 Hz, 2H, TsArH), 7.63 (d, J=8.0 Hz, 2H, TsArH), 8.36 (d, J=8.5 Hz, 1H, AlaNH), 8.98 (d, J=7.0 Hz, 1H, PheNH).

Ts-Ala-ΔAbu-(5'-Me-Oxd)-PheOMe (**3b**). ¹H-NMR δ: 1.03 (d, J=7.2 Hz, 3H, AlaMe), 1.36 (d, J=6.2 Hz, 3H, OxdMe), 1.66 (d, J=7.2 Hz, 3H, ΔAbuMe), 2.40 (s, 3H, TsMe), 2.98 (dd, J=7.2, 14.0 Hz, 1H, PheHβ), 3.04 (dd, J=6.0, 14.0 Hz, 1H, PheHβ), 3.59 (s, 3H, OMe), 3.90 (quint, J=6.4 Hz, 1H, AlaHα), 4.33 (dq, J=2.8, 6.2 Hz, 1H, OxdH₅), 4.42 (d, J=2.8 Hz, 1H, OxdH₄), 4.54 (q, J=7.4 Hz, 1H, PheHα), 5.71 (q, J=7.2 Hz, 1H, ΔAbuHβ), 7.18-7.25 (m, 3H, PheArH), 7.31 (t, J=7.4 Hz, 2H, PheArH), 7.38 (d, J=8.1 Hz, 2H, TsArH), 7.68 (d, J=8.1 Hz, 2H, TsArH), 7.94 (d, J=7.6 Hz, 1H, AlaNH), 8.78 (d, J=7.2 Hz, 1H, PheNH), 9.53 (s, 1H, ΔAbuNH).

Ts-Ala-(5'-Me-Oxd²)-D-(5'-Me-Oxd³)-Phe-OMe (**2c**). ¹H-NMR δ: 1.17 (d, J=6.8 Hz, 3H, AlaMe), 1.35 (d, J=6.2 Hz, 3H, Oxd²Me), 1.39 (d, J=6.4 Hz, 3H, Oxd³Me), 2.36 (s, 3H, TsMe), 2.83 (dd, J=8.1, 13.8 Hz, 1H, PheHβ), 3.11 (dd, J=8.1, 13.8 Hz, 1H, PheHβ), 3.63 (s, 3H, OMe), 3.84 (quint, J=5.6 Hz, 1H, Oxd³H₅), 4.36 (d, J=4.8 Hz, 1H, Oxd³H₄), 4.51 (dq, J=8.1, 9.6 Hz, 1H, PheHα), 4.95-5.03 (m, 2H, Oxd²H₄+AlaHα), 5.06 (q, 1H, J=6.2 Hz, Oxd²H₅), 7.17 (d, J=6.8 Hz, 2H, PheArH), 7.19-7.30 (m, 3H, PheArH), 7.37 (d, J=8.2 Hz, 2H, TsArH), 7.62 (d, J=8.2 Hz, 2H, TsArH), 8.42 (d, J=9.2 Hz, 1H, AlaNH), 8.94 (d, J=8.1 Hz, 1H, PheNH).

Ts-Ala-ΔAbu-D-(5'-MeOxd)-PheOMe (**3c**). ¹H-NMR δ: 1.03 (d, J=7.2 Hz, 3H, AlaMe), 1.20 (d, J=6.0 Hz, 3H, OxdMe), 1.67 (d, J=7.2 Hz, 3H, ΔAbuMe), 2.36 (s, 3H, TsMe), 2.85 (dd, J=10.4, 13.5 Hz, 1H, PheHβ), 3.12 (dd, J=4.8, 13.5 Hz, 1H, PheHβ), 3.58-3.69 (m, 4H, OxdH₅+OMe), 3.99 (quint, J=7.1 Hz, 1H, AlaHα), 4.25 (d, J=2.0 Hz, 1H, OxdH₄), 4.58 (m, 1H, PheHα), 5.65 (q, J=6.8 Hz, 1H, ΔAbuHβ), 7.13-7.24 (m, 2H, PheArH), 7.28 (t, J=7.0 Hz, 2H, PheArH), 7.35 (d, J=6.6 Hz, 2H, TsArH), 7.65 (d, J=6.6 Hz, 2H, TsArH), 7.93 (d, J=8.4 Hz, 1H, AlaNH), 8.85 (d, J=8.4 Hz, 1H, PheNH), 9.73 (s, 1H, ΔAbuNH).

Ts-Ala-(5'-Me-Oxd²)-D-(5'-Me-Oxd³)-D-PheOMe (**2d**). ¹H-NMR δ: 1.15 (d, J=6.6 Hz, 3H, AlaMe), 1.40 (d, J=6.2 Hz, 3H, Oxd²Me), 1.46 (d, J=6.0 Hz, 3H, Oxd³Me), 2.37 (s, 3H, TsMe), 2.97 (d, J=6.8 Hz, 2H, PheHβ), 3.58 (s, 3H, OMe), 4.46-4.52 (m, 3H, PheHα+Oxd³H_{4,5}), 5.02 (m, 1H, AlaHα), 5.03 (d, J=1.6 Hz, 1H, Oxd²H₄), 5.10 (dq, J=1.6, 6.2 Hz, 1H, Oxd²H₅), 7.18 (d, J=6.7 Hz, 2H, PheArH), 7.19-7.23 (m, 3H, PheArH), 7.37 (d, J=8.2 Hz, 2H, TsArH), 7.62 (d, J=8.2 Hz, 2H, TsArH), 8.41 (d, J=9.6 Hz, 1H, AlaNH), 8.89 (d, J=7.6 Hz, 1H, PheNH).

TableS1. Selected ¹H-NMR chemical shifts (δ) of the model compds **2a**, **2c**, **2d**, and **3b**, **3c**, solvents: S1 = CDCl₃; S2 = 8:2 DMSO-d₆/H₂O; amino acid stereochemistry has been omitted.

	2a		2c		2d		3b		3c	
	S1	S2	S1	S2	S1	S2	S1	S2	S1	S2
Ala ¹ NH	5.39	8.36	5.44	8.42	5.61	8.41	5.92	7.94	5.66	7.93
Ala ¹ H α	5.22	5.02	5.11	5.0	5.16	5.02	3.85	3.90	3.87	3.99
Oxd ² H4	5.26	5.17	5.24	5.0	5.42	5.03	-	-	-	-
Oxd ² H5	4.58	4.66	4.53	5.06	4.58	5.10	-	-	-	-
Δ AbuNH	-	-	-	-	-	-	8.66	9.53	8.28	9.73
Δ AbuH β	-	-	-	-	-	-	6.04	5.71	6.27	5.65
Oxd ³ H4	4.26	4.57	4.29	4.36	4.3	4.5	4.38	4.42	4.4	4.25
Oxd ³ H5	4.75	4.5	4.64	3.84	4.3	4.5	4.4	4.33	4.6	3.6
Phe ⁴ NH	6.28	8.98	6.55	8.94	6.63	8.89	7.64	8.78	7.98	8.85
Phe ⁴ H α	4.85	4.5	4.82	4.51	4.88	4.5	4.84	4.54	4.79	4.58

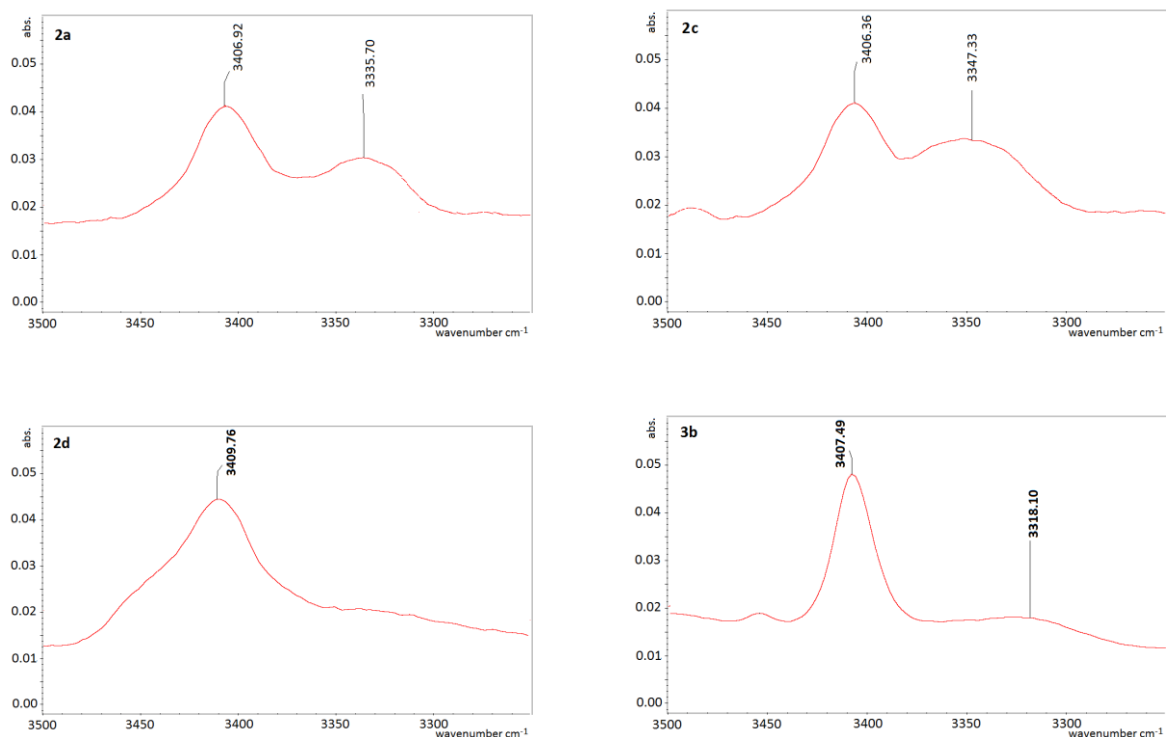


Figure S1. Amide NH stretching regions of the IR absorption spectra for samples of tetrapeptides **2a**, **2d** and **3b** (3 mM in DCM) at room temperature.

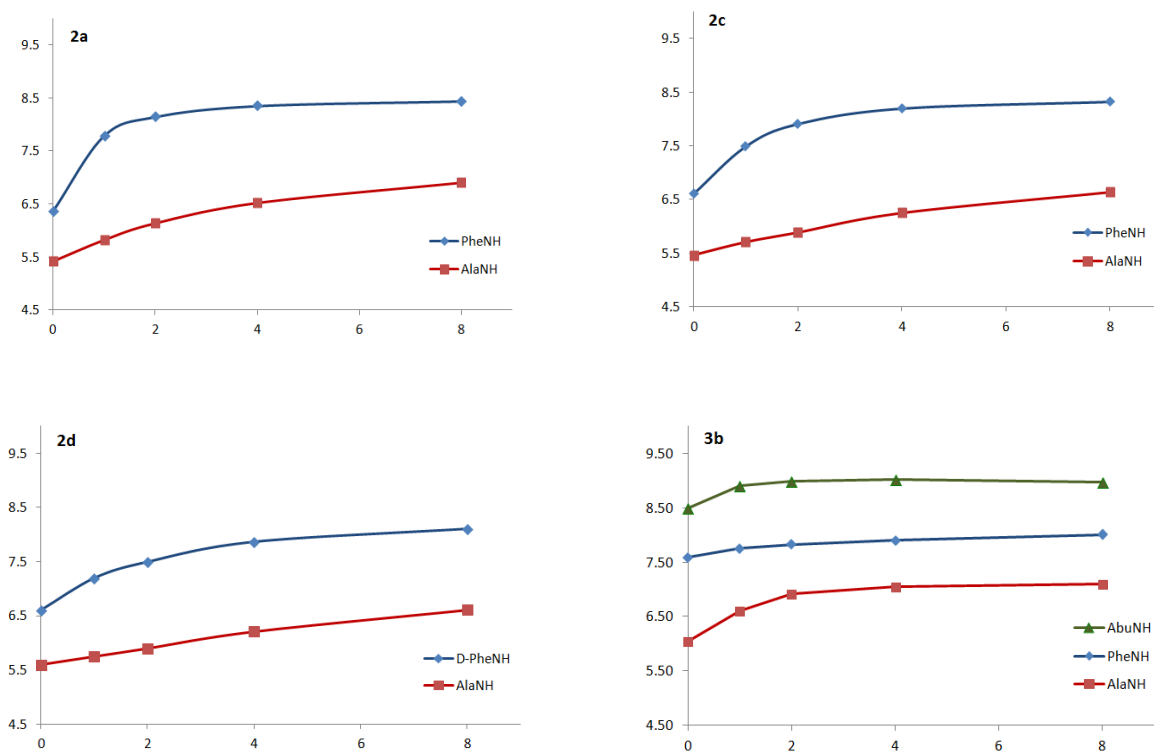


Figure S2. Variation of NH proton chemical shift (p.p.m.) of **2a**, **2c**, **2d** and **3b** as a function of increasing percentages of DMSO-d₆ to the CDCl₃ solution (v/v).

Table S2. Non-obvious ROESY cross-peaks observed for **2a**.^a

Cross-peak	intensity	Cross-peak	intensity
PheNH-Oxd ³ Me	w	PheNH-PheHβ(up)	s
PheNH-PheHβ(dw)	m	PheNH-PheHα	m
PheNH-Oxd ³ H4	s	PheNH-Oxd ³ H5	m
PheNH-Oxd ² H5	w	PheNH-PheArH	m
AlaNH-AlaMe	vs	AlaNH-AlaHα	s
AlaNH-TsArH2,6	s	TsArH2,6-AlaMe	w
TsArH2,6-Oxd ² Me	w	TsArH2,6-AlaHα	m
TsArH3,5-Oxd ² Me	s	TsArH3,5-Oxd ² H4	w
PheArH-PheHα	s	Oxd ² H4-Oxd ³ H4	w
Oxd ² H4-AlaHα	w	AlaHα-Oxd ² H5	w
Oxd ² H5-Oxd ³ H5	w	COOMe-PheHα	w

^a Stereochemistry has been omitted; ^b up = upfield, dw = downfield; ^c vs = very strong, s = strong, m = medium, w = weak.

Table S3. Non-obvious ROESY cross-peaks observed for **2c**.^a

Cross-peak	intensity	Cross-peak	intensity
PheNH-PheArH	w	PheNH-PheHα	m
PheNH-Oxd ³ H4	vs	PheNH-Oxd ³ H5	w
PheNH-PheHβ(dw)	w	PheNH-PheHβ(up)	s
PheNH-Oxd ³ Me	w	PheNH-COOMe	w
AlaNH-TsArH2,6	m	AlaNH-AlaHα	m
AlaNH-AlaMe	s	TsArH2,6-AlaMe	w
TsArH2,6-Oxd ² Me	w	TsArH2,6-AlaHα	m
TsArH3,5-Oxd ² Me	w	TsArH3,5-AlaHα	w
PheArH2,4,6-Oxd ³ Me	w	PheArH2,4,6-PheHβ(up)	s
PheArH2,4,6-PheHβ(dw)	s	PheArH2,4,6-AlaMe	w
PheArH2,4,6-PheHα	m	PheArH2,4,6-COOMe	w
PheArH2,4,6-Oxd ³ H5	w	Oxd ² H4-Oxd ³ H5	w
Oxd ² H4-Oxd ³ H4	w	Oxd ² H5-Oxd ³ H4	w
Oxd ² H5-PheHα	w	AlaHα-PheHα	w
PheHα-PheHβ(dw)	s	PheHα-PheHβ(up)	m
PheHα-COOMe	w	PheHα-Oxd ³ H5	w
PheHα-Oxd ³ H4	w	Oxd ³ H5-PheHβ(dw)	w
COOMe-AlaMe	m	COOMe-TsMe	w
COOMe-PheHβ(dw)	m	COOMe-PheHβ(up)	m
TsMe-Oxd ² Me	w		

^a Stereochemistry has been omitted; ^b up = upfield, dw = downfield; ^c vs = very strong, s = strong, m = medium, w = weak.

Table S4. Non-obvious ROESY cross-peaks observed for **2d**.^a

Cross-peak	intensity	Cross-peak	intensity
PheNH-AlaMe	w	PheNH-Oxd ³ Me	w
PheNH-PheH β	vs	PheNH-COOMe	w
PheNH-PheH α	m	PheNH-Oxd ³ H4	m
PheNH-Oxd ³ H5	w	PheNH-PheArH	s
AlaNH-AlaMe	s	AlaNH-COOMe	w
AlaNH-AlaH α	s	AlaNH-TsArH2,6	s
AlaNH-TsArH3,5	w	TsArH2,6-AlaMe	m
TsArH2,6-Oxd ² Me	w	TsArH2,6-AlaH α	s
TsArH2,6-OxdH4	w	TsArH3,5-Oxd ² Me	s
TsArH3,5-AlaH α	m	PheArH3,5-AlaMe	m
PheArH3,5-COOMe	w	PheArH3,5-PheH α	m
PheArH-Oxd ³ Me	w	PheArH-COOMe	s
Oxd ² H4-TsMe	m	PheH α -AlaMe	w
COOMe-AlaMe	w	COOMe-PheH β	s
PheH β -AlaMe	m	TsMe-AlaMe	w
Oxd ² Me-AlaMe	w		

^a Stereochemistry has been omitted; ^b up = upfield, dw = downfield; ^c vs = very strong, s = strong, m = medium, w = weak.

Table S5. Non-obvious ROESY cross-peaks observed for **3b**.^a

Cross-peak	intensity	Cross-peak	intensity
Δ AbuNH-AlaMe	m	Δ AbuNH- Δ AbuMe	s
Δ AbuNH-AlaH α	vs	Δ AbuNH-AlaNH	s
Δ AbuNH-PheNH	w	Δ AbuNH-TsArH2,6	w
PheNH-Oxd ³ H5	w	PheNH-PheH β	vs
PheNH-Oxd ³ H4	vs	PheNH-PheH α	s
PheNH-PheArH	m	PheNH-AlaNH	w
AlaNH-AlaMe	vs	AlaNH-AlaH α	s
AlaNH-TsArH2,6	w	TsArH2,6-AlaMe	w
TsArH2,6-AlaH α	m	PheArH-COOMe	w
PheArH-PheH β	vs	PheArH-PheH α	s
PheArH-Oxd ³ H5	w	Δ AbuHb-Oxd ³ H4	w
COOMe-PheH β	m	COOMe-PheH α	w
PheH β -Oxd ³ H4	w	COOMe-Oxd ³ Me	w
Δ AbuMe-TsMe	w	Δ AbuMe-AlaMe	s
Δ AbuMe-Oxd ³ Me	w		

^a Stereochemistry has been omitted; ^b up = upfield, dw = downfield; ^c vs = very strong, s = strong, m = medium, w = weak.

Table S6. Non-obvious ROESY cross-peaks observed for **3c**.^a

Cross-peak	intensity	Cross-peak	intensity
ΔAbuNH-AlaMe	m	ΔAbuNH-ΔAbuMe	s
ΔAbuNH-Oxd ³ Me	w	ΔAbuNH-AlaNH	vs
ΔAbuNH-ΔAbuHβ	w	ΔAbuNH-TsArH2,6	w
ΔAbuNH-PheNH	w	PheNH-PheArH	m
PheNH-PheHβ(up)	s	PheNH-PheHβ(dw)	w
PheNH-Oxd ³ H5	m	PheNH-Oxd ³ H4	s
PheNH-PheHα	s	PheNH-PheArH	m
AlaNH-AlaMe	m	AlaNH-AlaHα	m
TsArH2,6-AlaMe	m	TsArH2,6-AlaHα	s
TsArH2,6-AlaNH	m	TsArH3,5-AlaNH	w
PheArH3,5-Oxd ³ Me	w	PheArH3,5-PheHβ(up)	w
PheArH3,5-PheHα	w	PheArH2,6-Oxd ³ Me	w
PheArH2,6-PheHβ(up)	vs	PheArH2,6-PheHβ(dw)	vs
PheArH2,6-Oxd ³ H5	w	PheArH2,6-PheHα	s
PheHα-PheHβ(dw)	vs	PheHα-PheHβ(up)	s
Oxd ³ H5-PheHβ(up)	w	COOMe-PheNH	w
COOMe-PheHα	m	COOMe-PheHβ(dw)	w
COOMe-PheHβ(up)	w		

^a Stereochemistry has been omitted; ^b up = upfield, dw = downfield; ^c vs = very strong, s = strong, m = medium, w = weak.

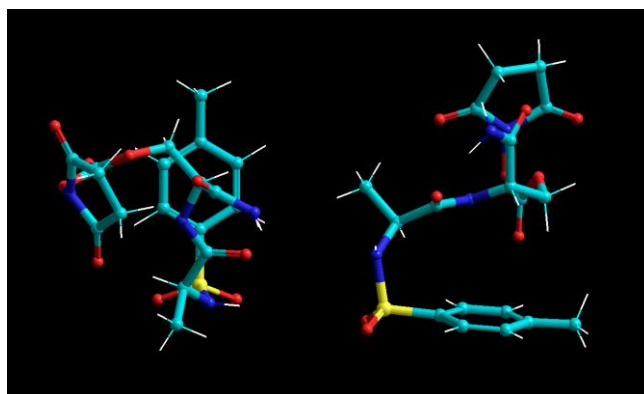


Figure S3. Top (left) and side (right) view of the intermediate anion **A** of Scheme 1, calculated for the cyclization of the model peptide Ts-Ala-Ser-NH₂ with DSC and DIPEA, employing ab initio molecular orbital (MO) theory. A systematic conformational analysis for the structures was done at the HF/6-31G* level. The conformers were re-optimized at the HF/6-31G** level. Backbones are rendered in balls-and-cylinders, hydrogen atoms in sticks. Optimization was performed by conjugate gradient algorithm, convergence at 0.001; energies are expressed in Kcal mol⁻¹. DIPEAH⁺ was included in the computations, but is not visualized for clarity.

Table S7. Angles ϕ and ψ of the residues $i+1$, $i+2$ observed for the compounds analyzed and ideal values.

compd	ϕ_{i+1}	ψ_{i+1}	χ_{i+1}	ϕ_{i+2}	ψ_{i+2}	χ_{i+2}
2a (A)	-73	158	-114	-81	76	-115
2a (B)	-104	-179	-90	-40	-50	-140
2c (C)	-45	129	-122	69	-86	118
2c (D)	-51	144	-117	77	-27	110
2d	-55	139	-117	77	-29	113
3b	-38	-59	-2	-96	29	-114
3c (E)	61	55	2	93	-2	117
3c (F)	37	63	0	81	-36	113
I	-60	-30	-	-90	0	-
II	-60	120	-	80	0	-

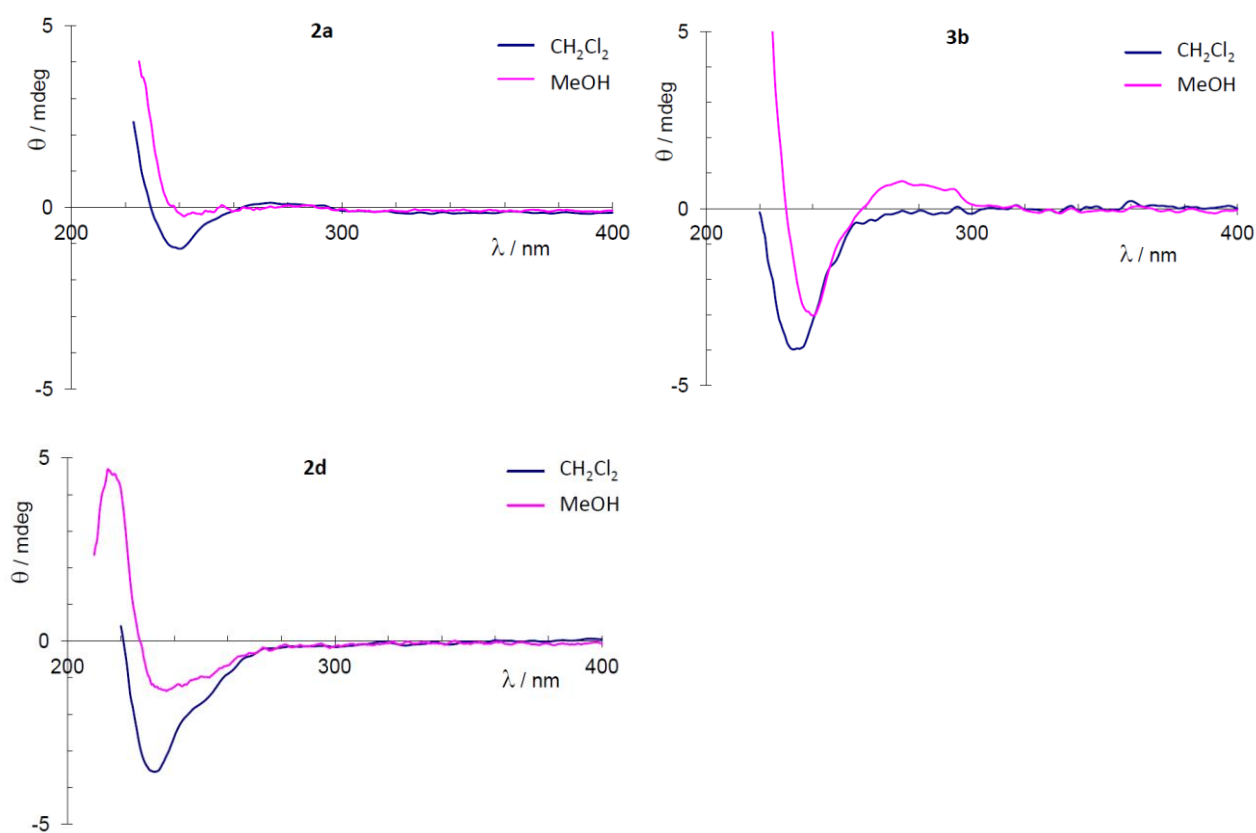


Figure S4. ECD spectra of **2a**, **3b**, and **2d**; 1 mM in DCM or MeOH, path length 0.1 cm.

Figure S5.

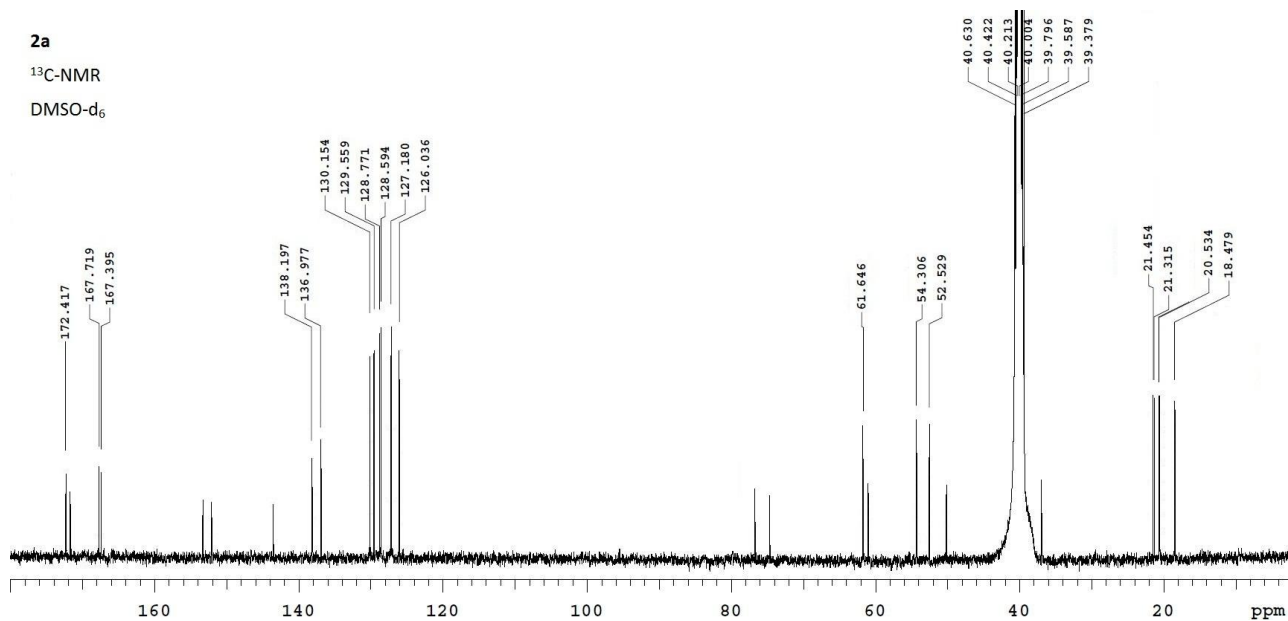
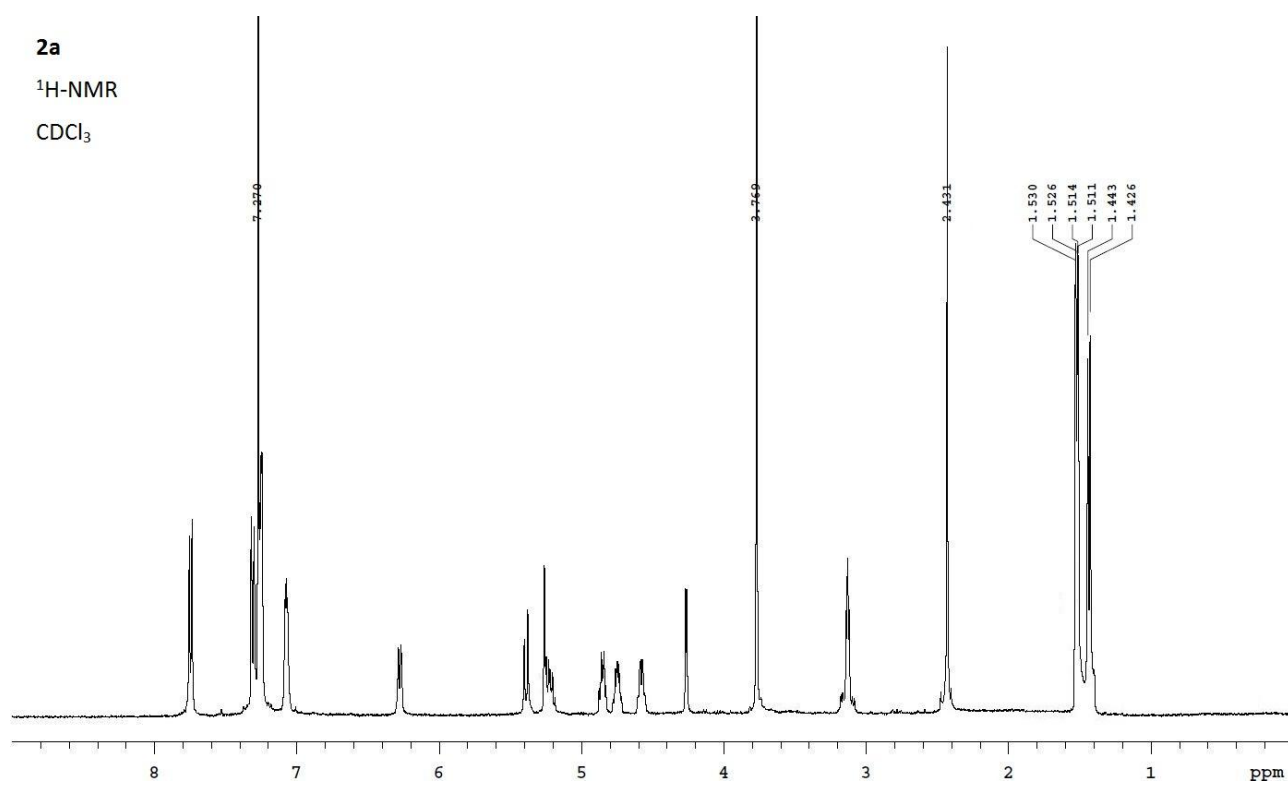


Figure S6.

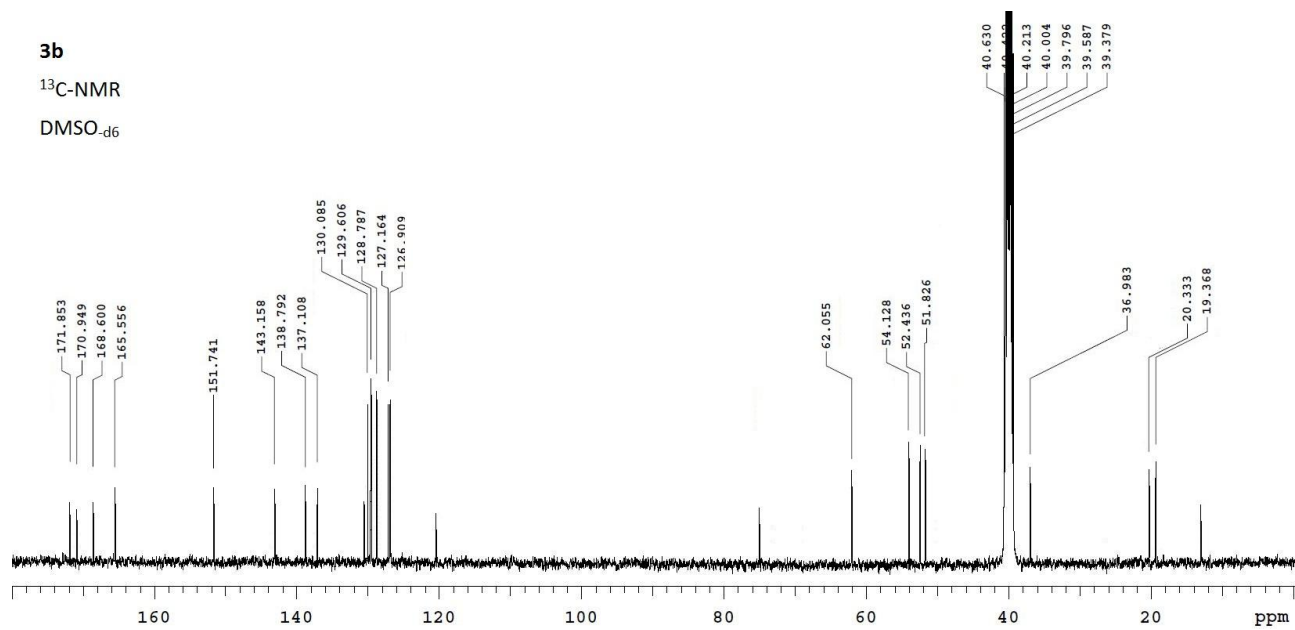
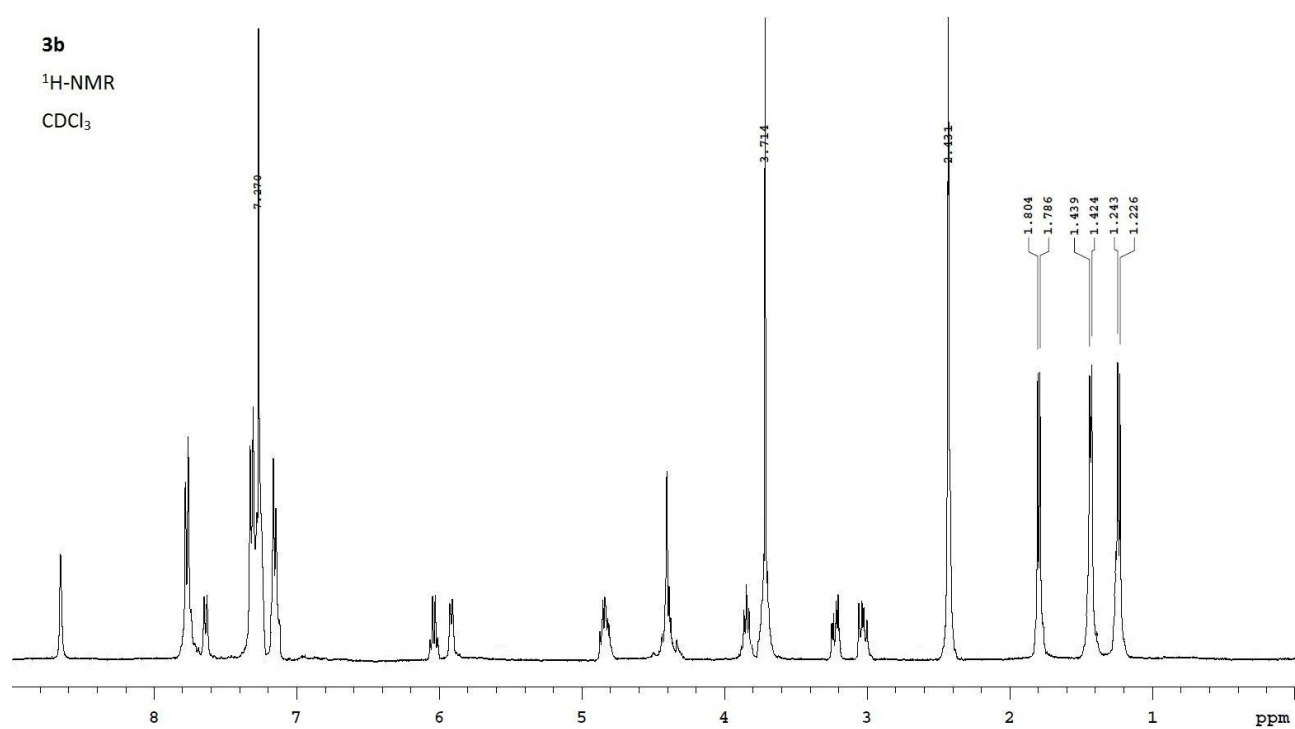


Figure S7.

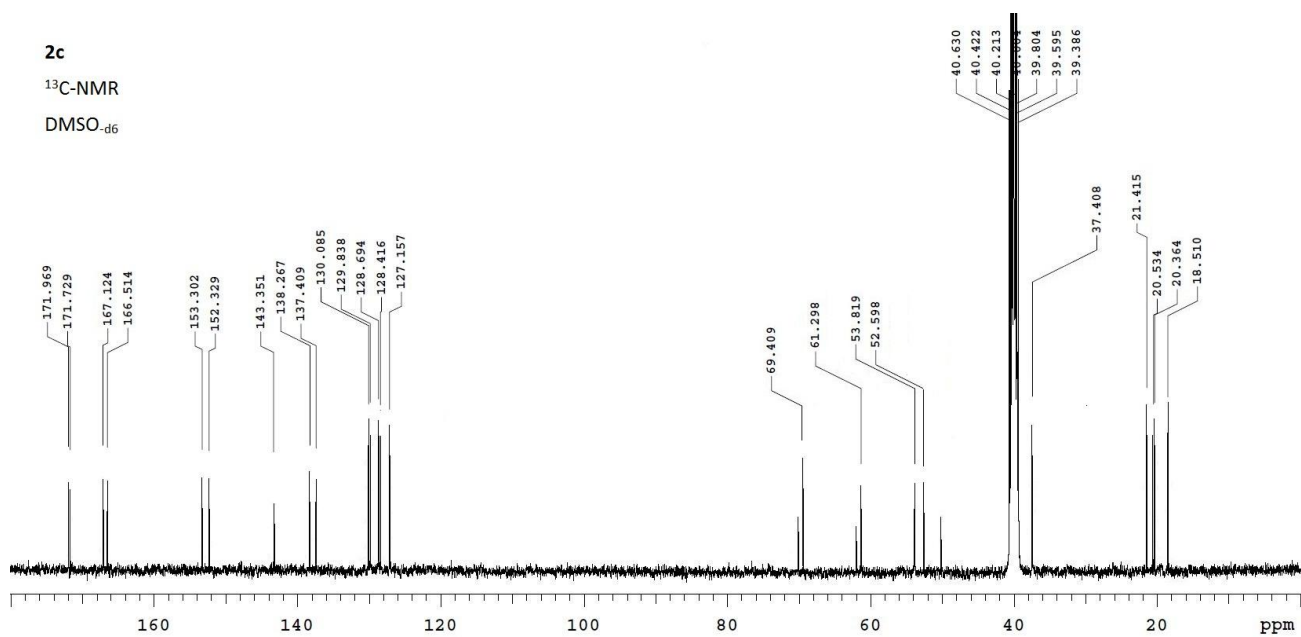
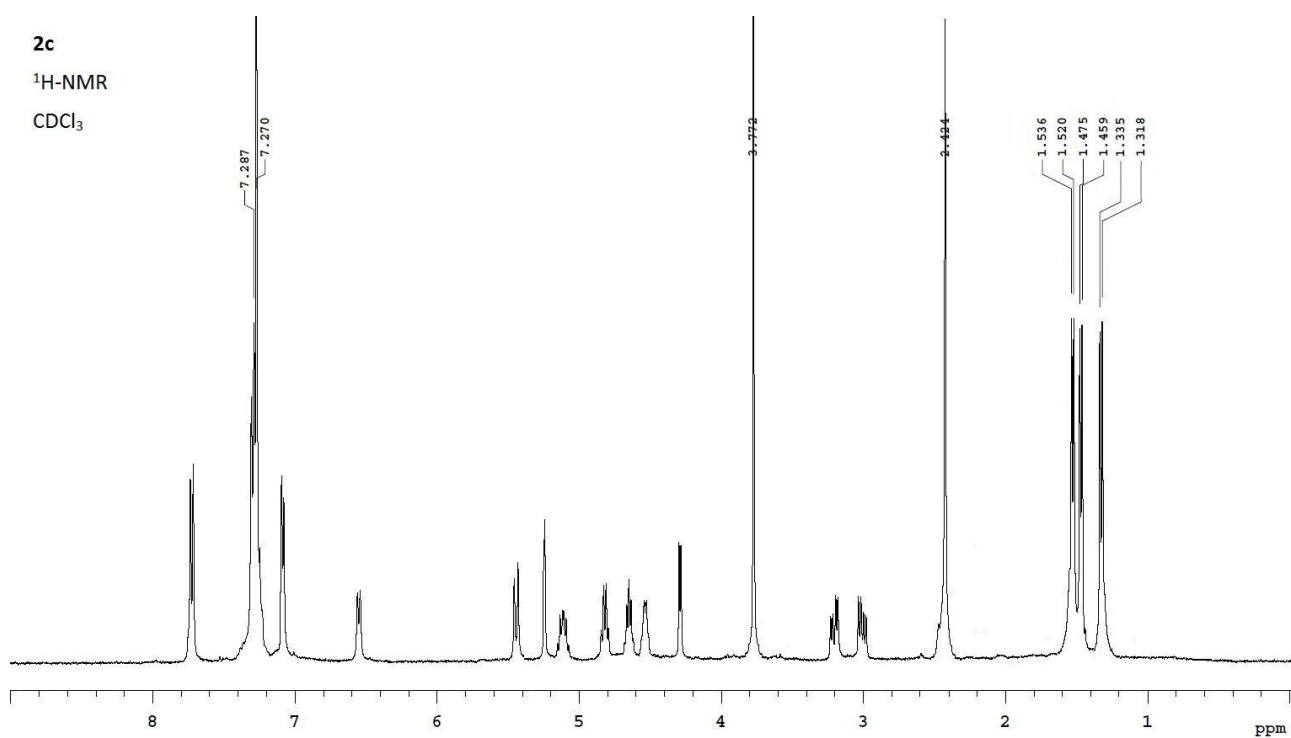


Figure S8.

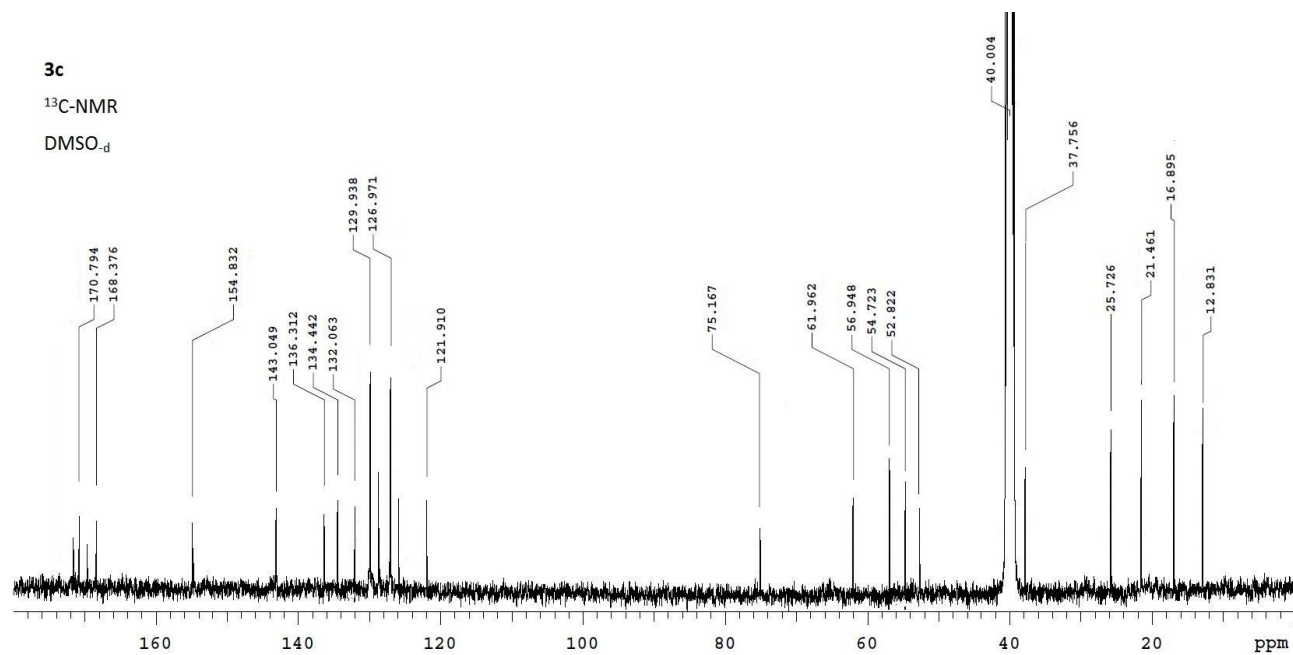
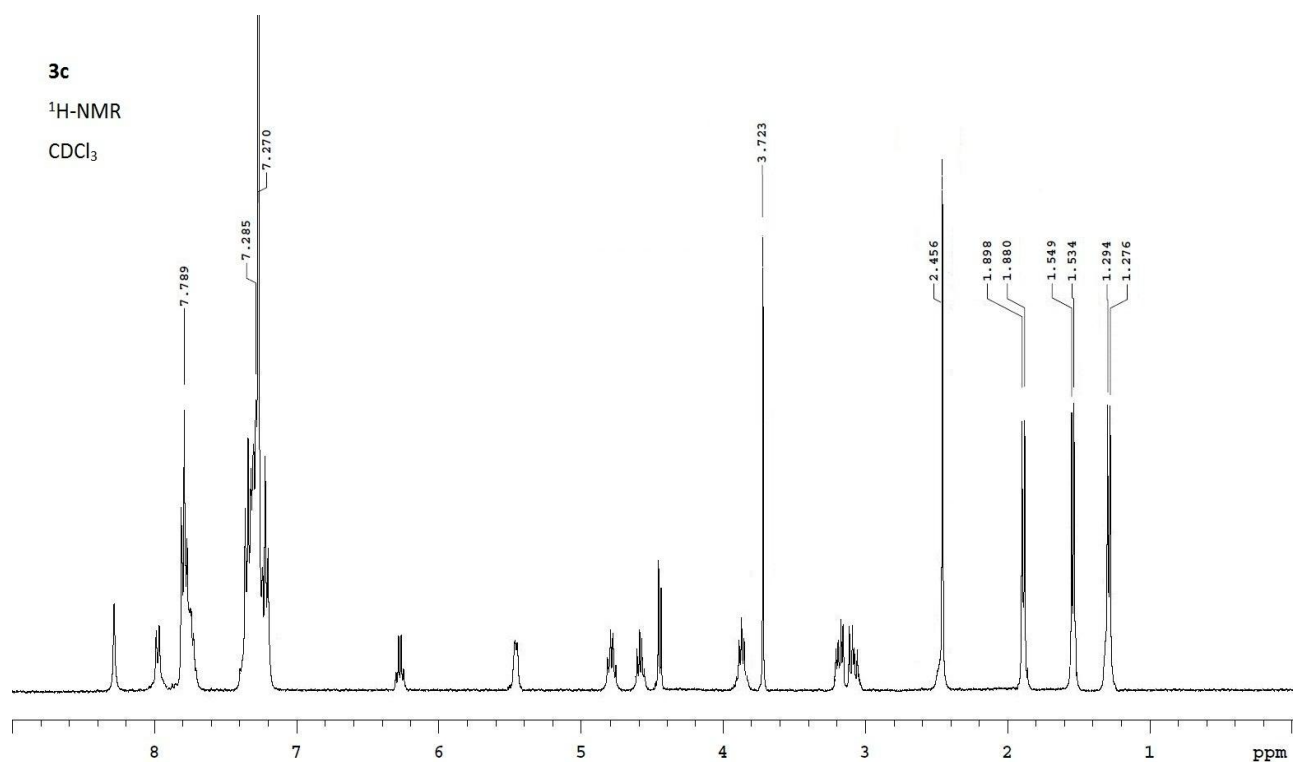


Figure S9.

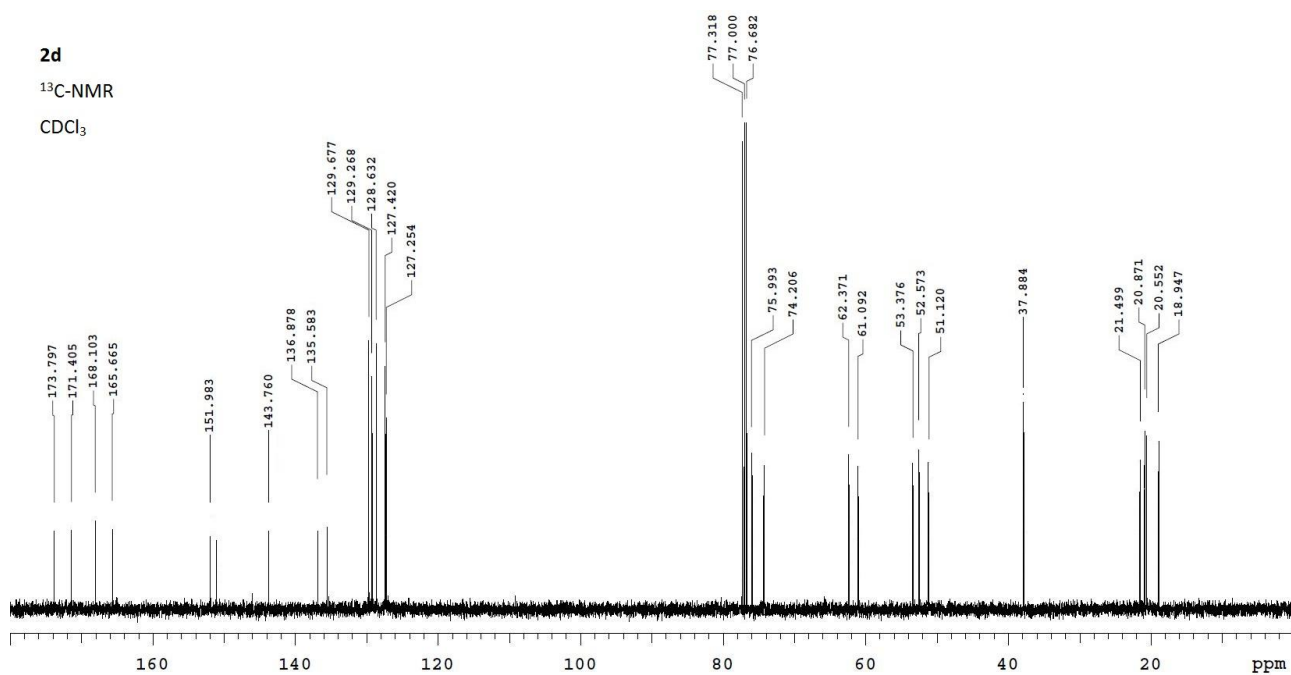
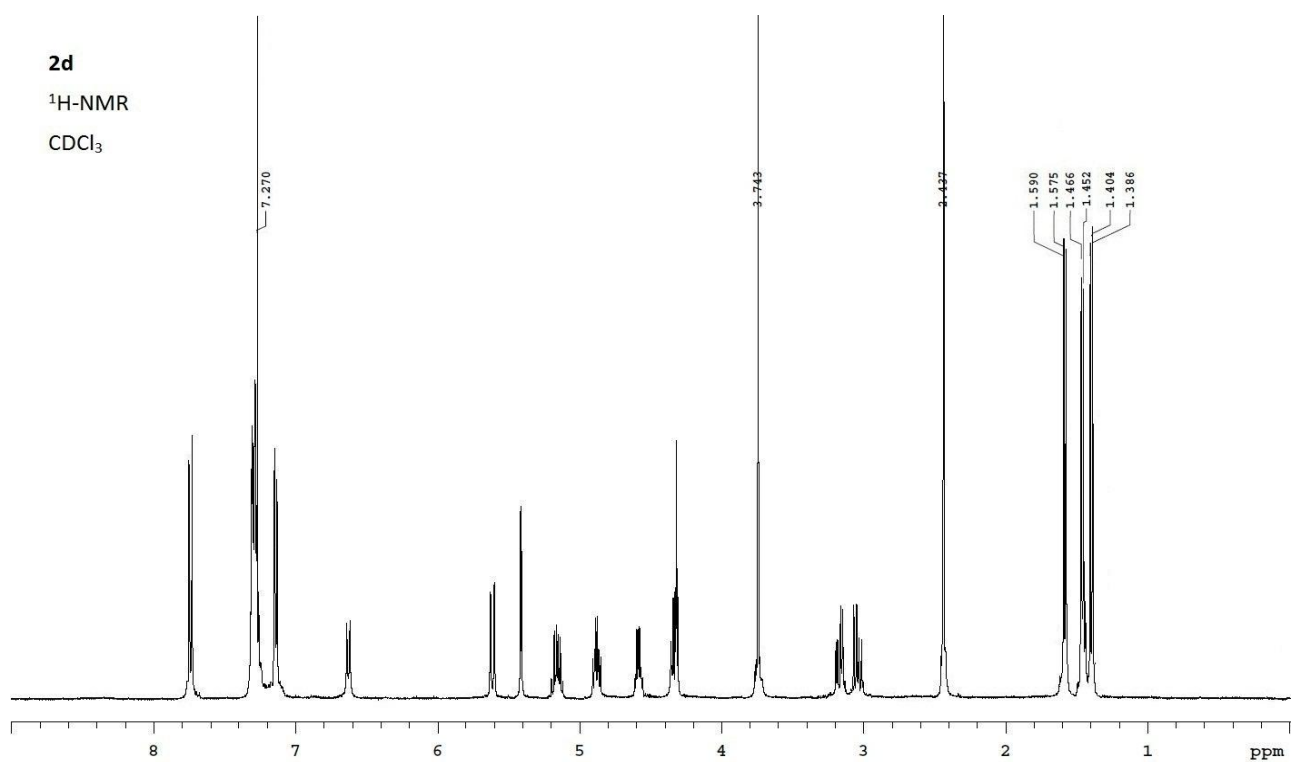


Figure S10.

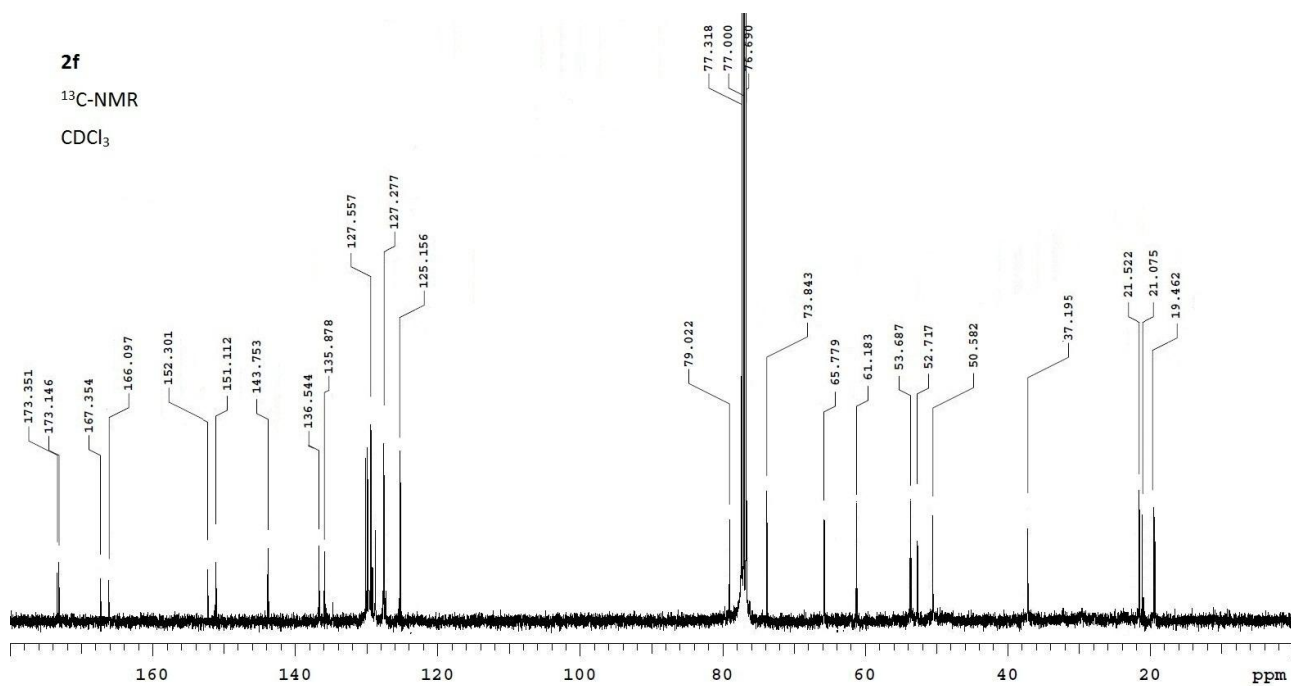
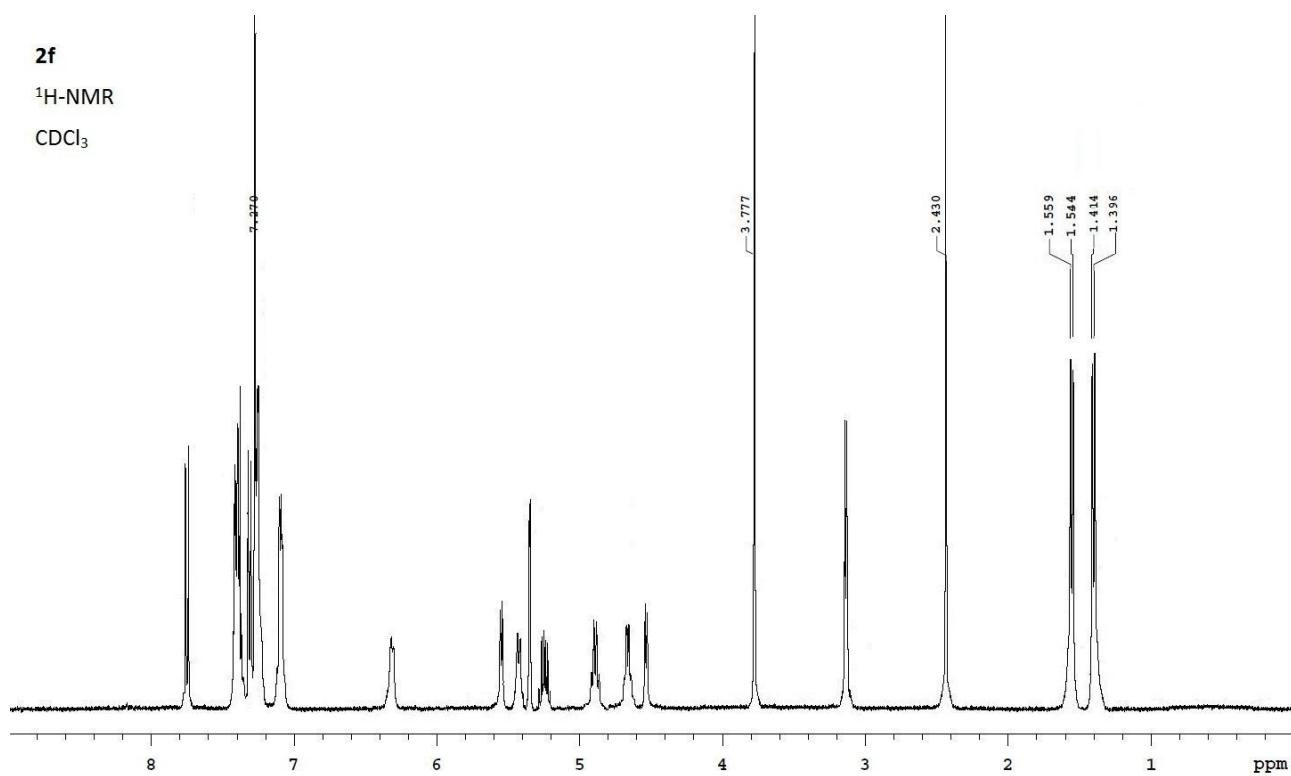


Figure S11.

