

Electronic Supporting Information

Novel Foldamer Structural Architecture from Cofacial Aromatic Building Blocks

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General Methods

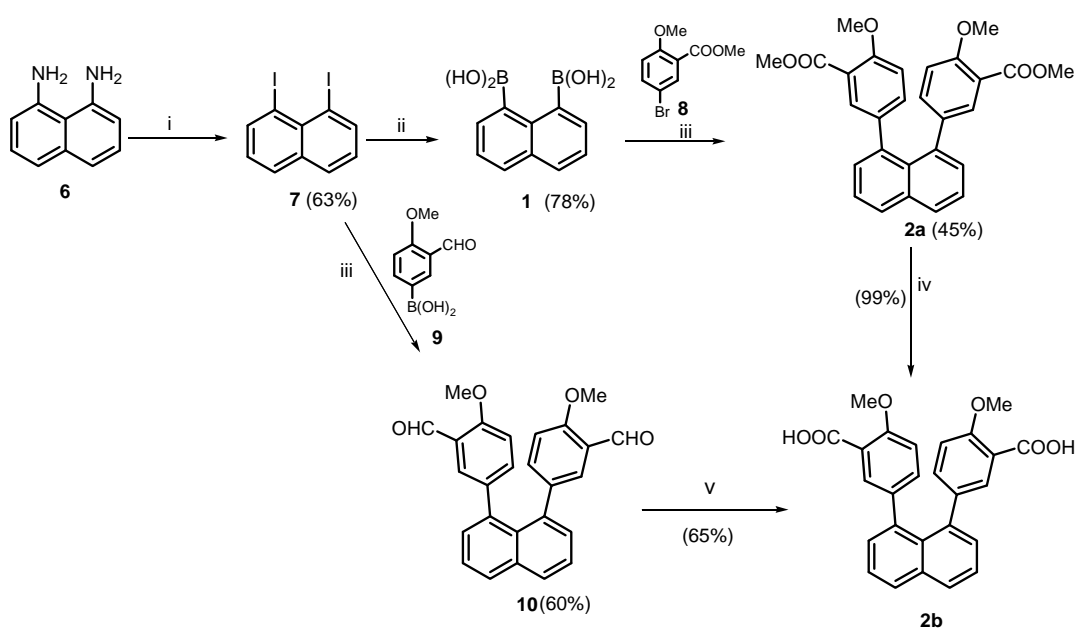
Unless otherwise stated, all the chemicals and reagents were obtained commercially. Dichloromethane was dried by distilling over P_2O_5 and keeping over 4 Å mol sieves. Analytical Thin Layer Chromatography was done on precoated silica gel plates (Kieselgel 60F₂₅₄, Merck). Column chromatographic purifications were done with 100-200 Mesh Silica gel. NMR spectra were recorded in $CDCl_3$ on AV 200 MHz or AV 400 MHz or AV 500 MHz Bruker NMR spectrometers. All chemical shifts are reported in δ ppm downfield to TMS and peak multiplicities as singlet (s), doublet (d), quartet (q), broad (b), broad singlet (bs), and multiplet (m). Elemental analyses were performed on an Elementar-Vario (Heraeus Company Ltd., Germany). IR spectra were recorded in nujol or $CHCl_3$ using Shimadzu FTIR-8400 spectrophotometer. Melting points were determined on a Buchi Melting Point B 540. MALDI-TOF spectrometric measurements were done with Voyager-DE STR Biospectrometer workstation using α -cyano-4-hydroxycinnamic acid (CHCA) as the matrix. Single crystal X-ray data were collected on a Bruker SMART APEX CCD Area diffractometer with graphite monochromatized ($Mo\ K\alpha = 0.71073\text{\AA}$) radiation at room temperature. Since the quality of the crystals of **3** was poor and the diffraction at high angles was not observed, off limit is changed in the refinement. In the asymmetric unit, there is mainly one molecule along with 0.5 $CHCl_2$ and 0.25 water molecules. There are voids in the structure when the solvent molecules are not there. Merohedral twinning was checked in RLATT and the crystals are not twinned but the quality of crystals was poor and the high angle diffraction was not good which resulted in high R. All the data were corrected for Lorentzian, polarization, and

absorption effects using Bruker's SAINT and SADABS programs. SHELX-97¹ was used for structure solution and full matrix least squares refinement on F2. Hydrogen atoms were included in the refinement as per the riding model.

Experimental Methods

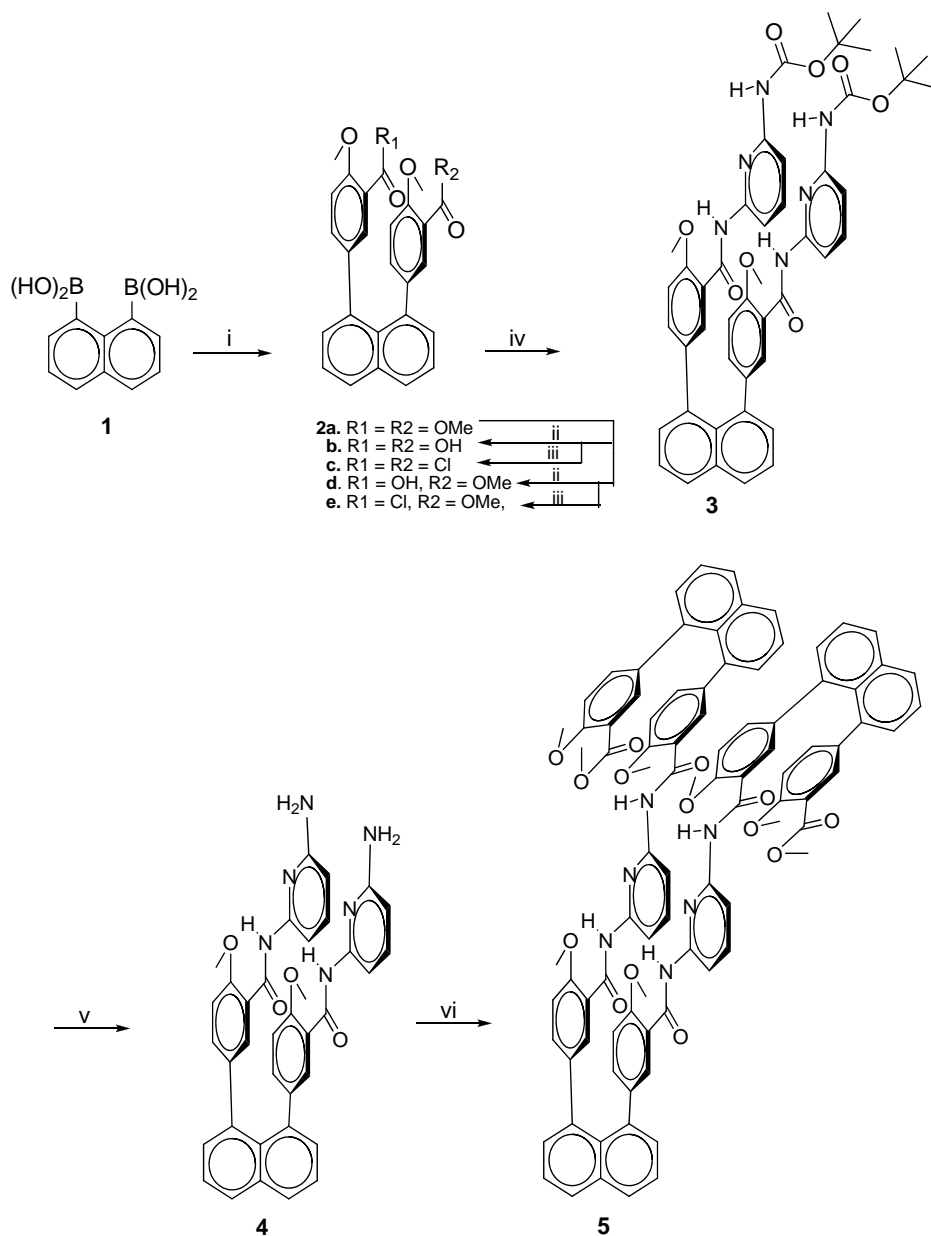
1,8-diaminonaphthalene **6** and 3-formyl-4-methoxyphenyl boronic acid **9** were obtained from Aldrich. 1,8-diiodonaphthalene **7**² and the naphthalene diboronic acid **1**³ were prepared according to the literature procedures. Suzuki coupling of **1** with **8**, yielding **2a**, followed a recent procedure.⁴ Alternatively, the acid **2b** could also be obtained from the reported bis-formyl compound⁵ **10** (obtained by the Suzuki coupling of **7** with **9**) by Jones oxidation.

Scheme 1: Synthesis of the diacid **2b**^a



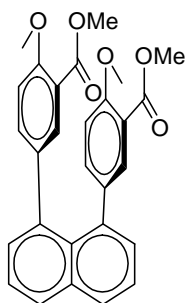
^a**Reagents and conditions:** (i) (a) NaNO₂, H₂SO₄, H₂O, -15 to -20 °C; (b) KI in H₂O; (ii) BuLi, Trimethyl borate, -78 °C; (iii) Pd(OAc)₂ (12 mol%), K₂CO₃ (8 equiv.), PEG-400, DABCO (24 mol%), TBAB (0.1 equiv.), 110 °C, 15 h.; (iv) LiOH (2N), Dioxane/ H₂O; (v) CrO₃, H₂SO₄, acetone, rt, 3 h (Jones oxidation).

Scheme 2: Syntheses of **3** and **5^b**



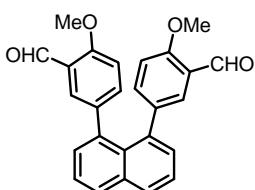
^b**Reagents and conditions:** i) methyl-5-bromo-2-methoxy benzoic acid methyl ester, DABCO, Pd (OAc)₂, K₂CO₃, TBAB, PEG-400, 110 °C 15 h; ii) LiOH, dioxane, H₂O; iii) (COCl)₂, DMF (cat.), DCM, rt, 3h; iv) mono-boc-2,6-diamino pyridine, Et₃N, rt, 12 h; v) TFA, DCM, rt, 30 min; vi) **2e**, Et₃N, DCM, rt, 12 h.

1,8-Bis-(4-methoxy-3-methylcarboxyphenyl)-naphthalene **2a**:



The Suzuki coupling of **1** with **8** yielding **2a** followed a recently reported procedure⁴ as follows: A reaction mixture in PEG-400 (6 mL) containing naphthalene diboronic acid **1** (0.2 g, 0.9 mmol, 1 equiv.), methyl-5-bromo-2-methoxy-benzoic methyl ester **8** (0.68 g, 2.7 mmol, 3 equiv.), Pd(OAc)₂ (12 mol%), DABCO (24 mol%), TBAB (0.1 equiv.), and K₂CO₃ (8 equiv.) was sealed and heated at 110 °C for 15 h. After cooling to room temperature, the product was extracted into DCM (100 mL), and the crude compound obtained was purified by column chromatography (eluent: AcOEt / pet.ether: 25/75; R_f: 0.3) to yield **2a** (0.19 g, 45%). mp: 158-160 °C; Found: C, 73.82; H, 5.42. Calc. for C₂₈H₂₄O₆: C, 73.67; H, 5.30%; ν_{\max} (CHCl₃) / (cm⁻¹): 3020, 2951, 2841, 1724, 1717, 1611, 1501, 1437, 1308, 1277, 1238, 1217, 1182, 1086, 1026; ¹H NMR (200 MHz, CDCl₃) δ : 7.98-7.93 (dd, *J* = 1.37 Hz, 2H), 7.59-7.51 (m, 2H), 7.46-7.39 (m, 4H), 7.08-7.03 (dd, *J* = 2.28 Hz, 1H), 6.98-6.93 (dd, *J* = 2.43 Hz, 1H), 3.85-3.81 (m, 12 H); ¹³C NMR (50 MHz, CDCl₃) δ : 166.5, 166.0, 157.6, 157.4, 138.7, 135.4, 135.2, 135.1, 134.9, 134.4, 133.6, 132.6, 131.0, 130.7, 128.9, 125.2, 118.6, 111.4, 111.3, 56.3, 56.1, 51.9, 51.8; ESI-MS: 457.35 (M+H)⁺, 479.28 (M+Na)⁺, 496.26 (M+K)⁺.

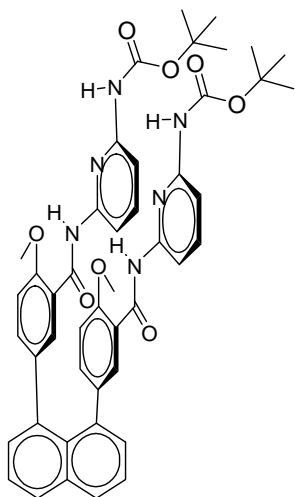
1,8-Bis-(3-formyl-4-methoxy-phenyl)-naphthalene **10**⁵



Following the same procedure for the preparation of **2a**, the Suzuki coupling of **7** (0.4 g, 1.5 mmol, 1 equiv.) with **9** (0.57 g, 3.1 mmol, 3 equiv.) yielded **10** (60%). ¹³C (50 MHz,

CDCl₃) δ : 189.3, 159.7, 138.1, 137.3, 135.5, 135.2, 130.6, 129.4, 128.9, 125.1, 123.2, 110.7, 55.7; ESI-MS: 397.31 (M+H)⁺, 419.30 (M+Na)⁺, 435.28 (M+K)⁺;

Synthesis of the short oligomer 3: To a stirred solution of the diacid **2b** (1g, 2.3

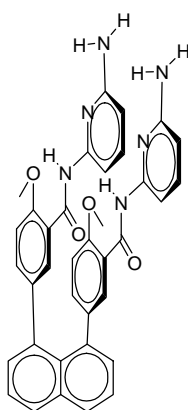


mmol, 1 equiv.) in dry DCM (15 ml), 2 drops of DMF were added followed by the drop-wise addition of oxalyl chloride (1.2 ml, 9.2, 4 equiv.). After 3 h at room temperature, solvent and excess oxalyl chloride were removed from the reaction mixture under reduced pressure to get the crude acid chloride **2c** which was directly reacted with a reaction mixture containing mono-Boc-2,6-diaminopyridine⁶ (9.2 mmol, 4 equiv.) and TEA (9.2 mmol, 4 equiv.) in DCM (10

ml). The reaction mixture was stirred overnight at rt. Work-up followed by column chromatographic purification afforded **3** (eluent: 80% dichloromethane / pet.ether, R_f: 0.4) as a white solid (1.41 g, 75%), which could be crystallized from a solution of dichloromethane and MeOH. mp: 172 °C (decom.); Found: C, 68.26; H, 5.84; N, 10.44. Calc. for C₄₆H₄₆N₆O₈: C, 68.13; H, 5.72; N, 10.36%; ν (CHCl₃)/ (cm⁻¹): 3421, 3356, 3018, 1730, 1587, 1506, 1497, 1456, 1369, 1305, 1217, 1153, 1053, 1028; ¹H NMR (CDCl₃, 400 MHz) δ : 10.12, (s, 2H), 8.48 (bs, 2H), 8.08-8.06 (m, 2H), 7.98-7.96, *J* = 8.28 Hz, 2H), 7.77-7.73 (m, 6H), 7.58-7.54 (t, *J* = 7.54 Hz, 2H), 7.40-7.38 (d, *J* = 7.04 Hz, 2H), 7.18-7.16 (dd, *J* = 2.26 Hz, *J* = 6.29 Hz, 2H), 6.69-6.67 (d, *J* = 8.80 Hz, 2H), 3.73 (s, 6H), 1.38 (s, 18H); ¹³C NMR (CDCl₃, 100 MHz) δ : 163.1, 155.4, 152.5, 150.8, 150.1, 140.9, 138.7, 136.7, 135.6, 135.3, 133.2, 130.6, 130.0,

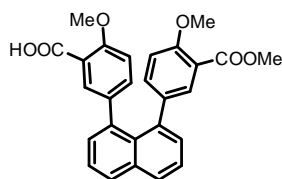
128.8, 125.2, 119.4, 110.6, 108.9, 107.7, 81.1, 56.1, 28.1; ESI MS: 810.96 (M+H)⁺, 832.94 (M+Na)⁺; 848.95 (M+K)⁺.

Synthesis of bis-amine 4:



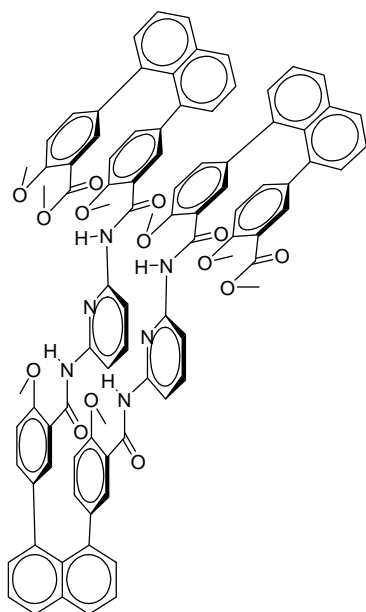
The compound **3** (0.5 g, 0.6 mmol) was subjected to the Boc-deprotection using TFA/DCM. After complete consumption of **3**, the solvent was stripped off from the reaction mixture. The residue was neutralized with sat. sodium bicarbonate solution and the white precipitate obtained was filtered, washed with water, and the residue was dried in KOH desiccator and taken for the further reaction without any purification.

Synthesis of monoacid 2d:



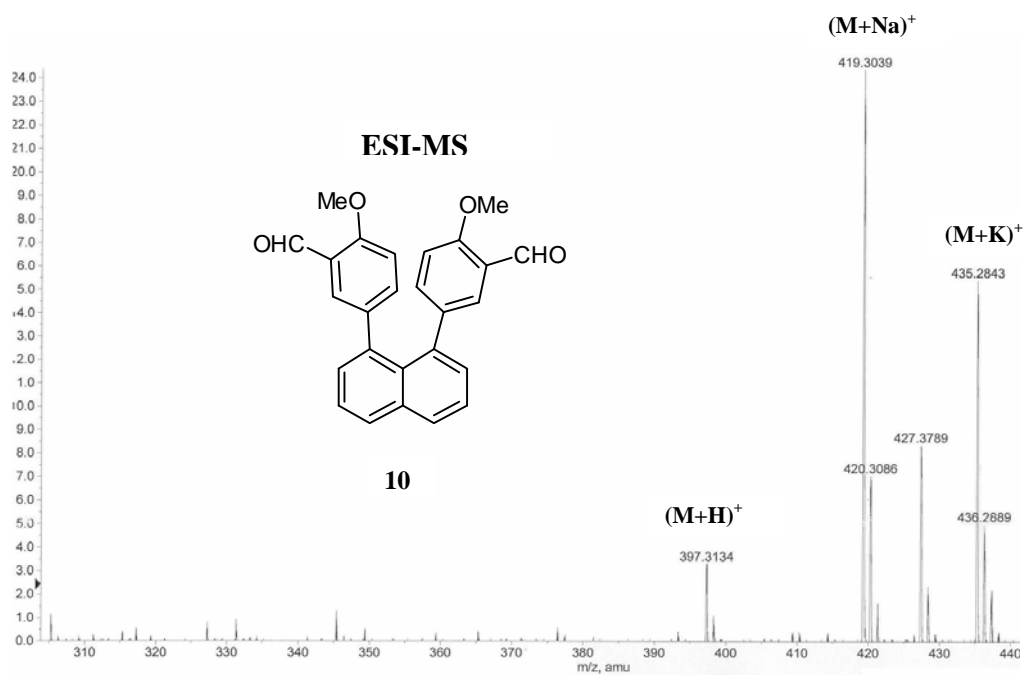
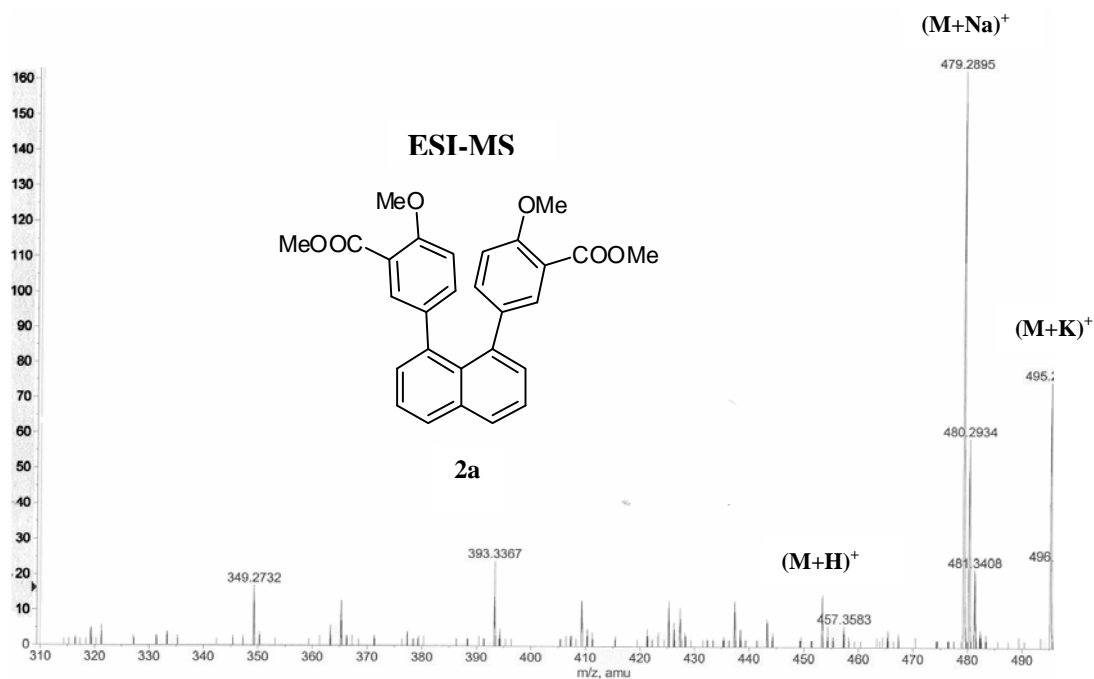
The bis-ester **2a** (0.6 g, 1.3 mmol, 1 equiv.) was subjected to careful saponification using LiOH·H₂O (56 mg, 1.3 mmol, 1 equiv.) dissolved in MeOH/H₂O (9:1 mL). After stirring for 6 h, the reaction mixture was stripped off the solvent under reduced pressure. The residue was suspended in water and washed with DCM to remove any unreacted bis-ester **6**. The aqueous layer was neutralized with dil. HCl to get an off-white precipitate, which was taken in DCM and filtered to remove the undissolved material, which is presumably the bis-acid **2b**. The filtrate was evaporated to get the crude acid which was used for the next reaction without further purification.

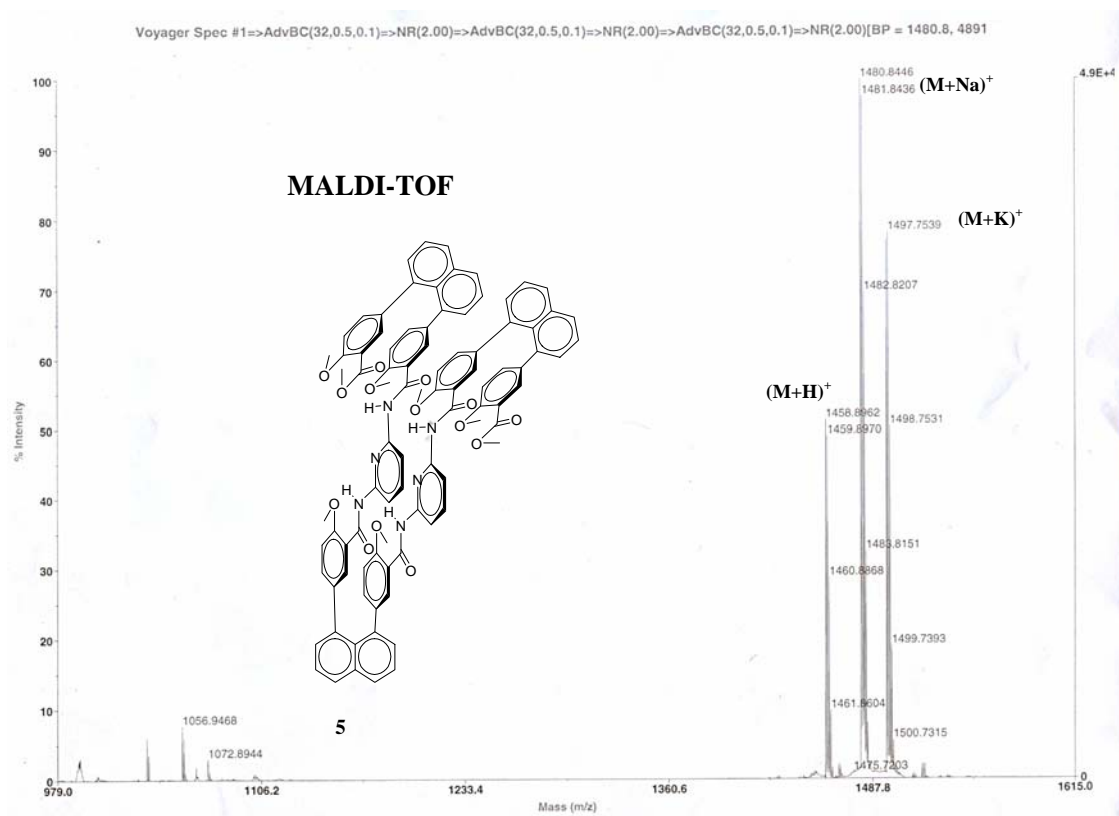
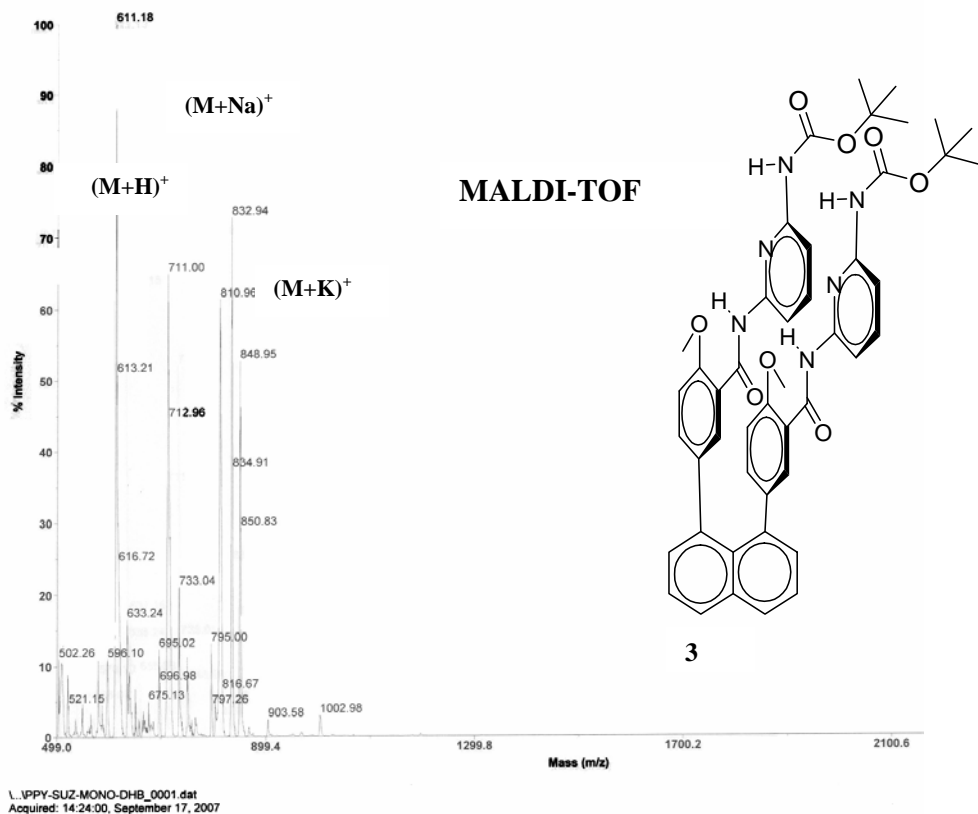
Synthesis of 5: The monoacid **2d** (0.36 g, 0.8 mmol, 3 equiv.) was converted to the corresponding acid chloride **2e** using oxalyl chloride (4 equiv.) and cat. amount of

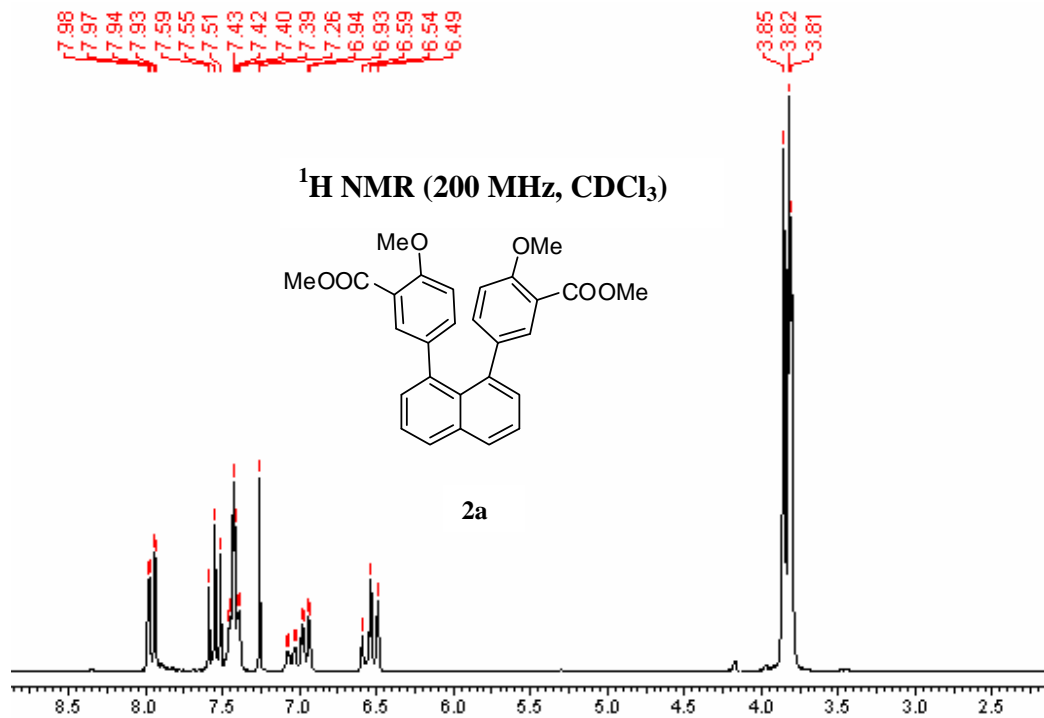
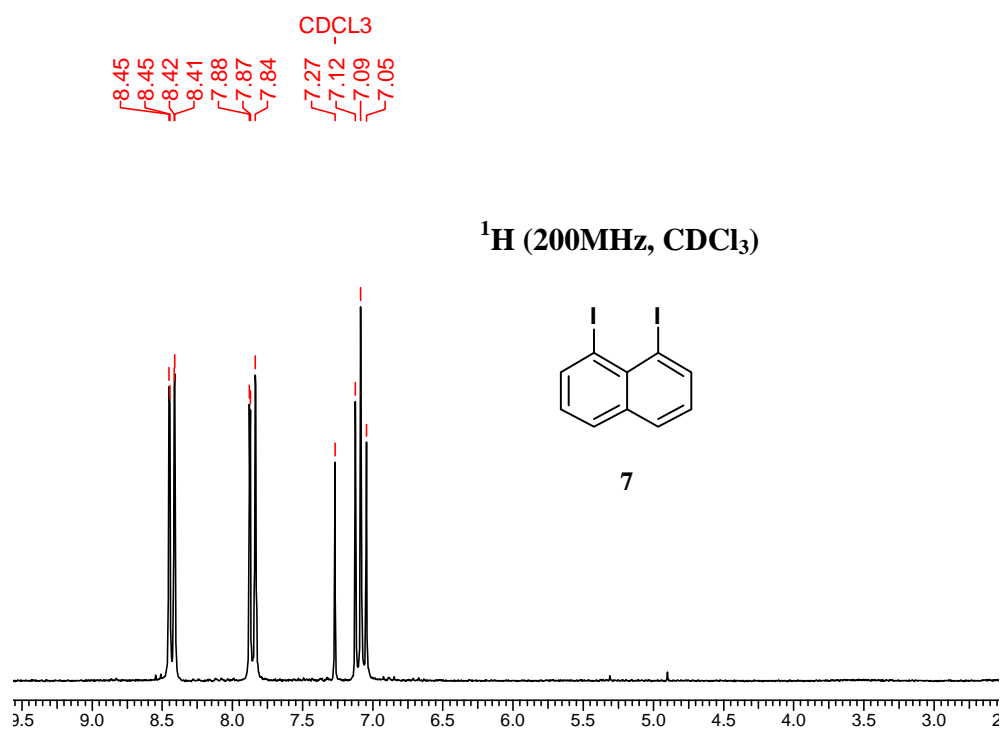


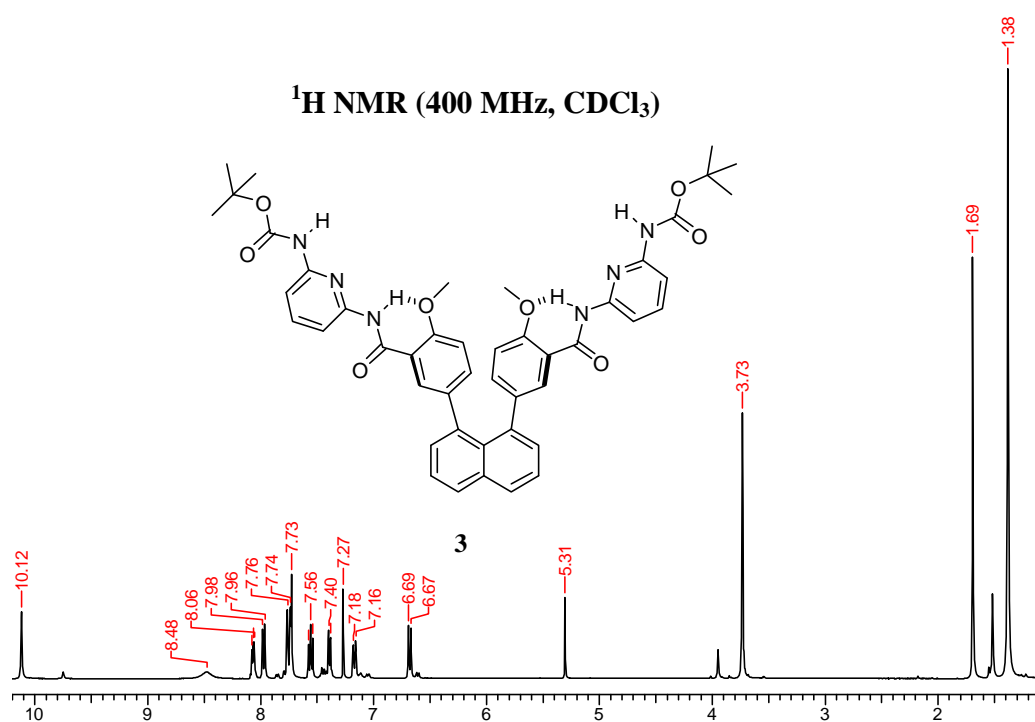
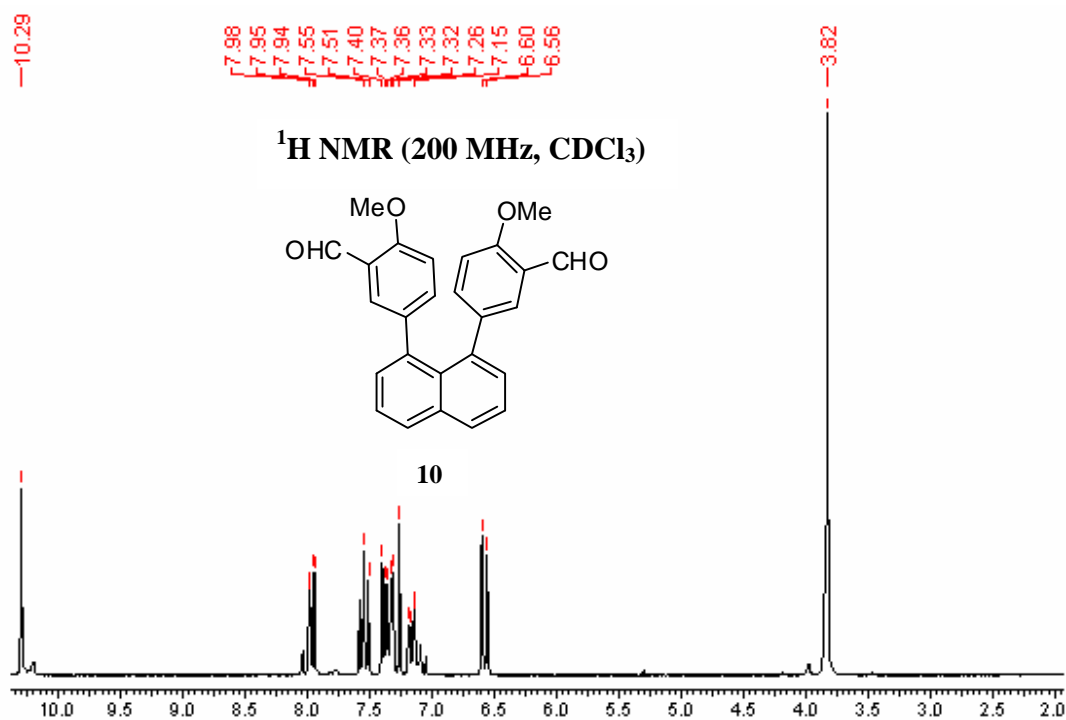
DMF in dichloromethane (15 mL) as the solvent. After stirring the reaction mixture for 3h at room temperature, the solvent and excess oxalyl chloride were removed under reduced pressure to get the crude acid chloride **2e** which was taken in a solution containing the bis-amine **4** (0.166 g, 0.27 mmol, 1 equiv.) and TEA (4 equiv.). The reaction mixture was allowed to stir for 12 h at rt. The insoluble product was filtered off and washed with DCM to get an off-white residue (62%) which could not be further purified

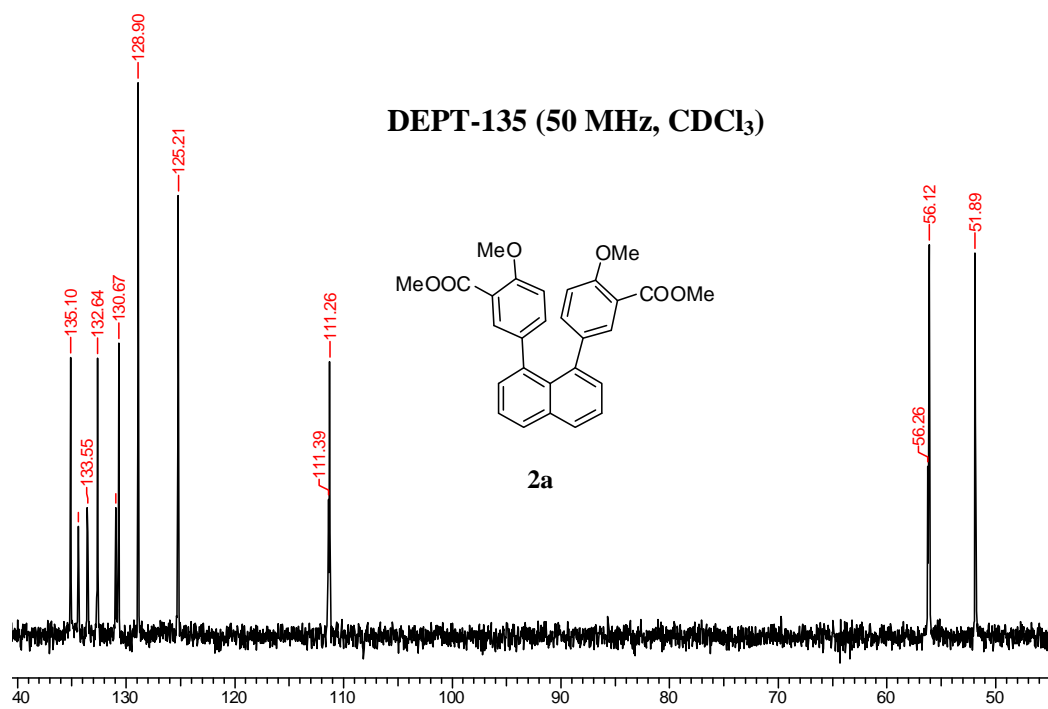
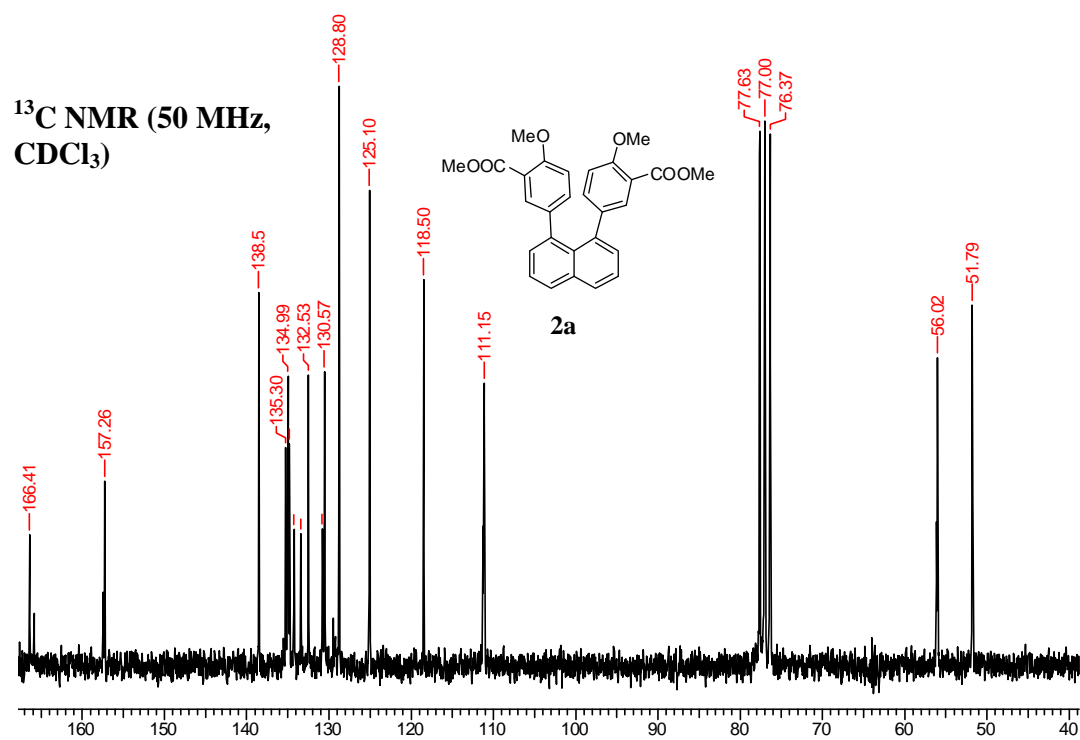
owing to its inadequate solubility in various organic solvents. mp: > 295 °C, Found: 1459.89 (M+H)⁺; 1481.84 (M+Na)⁺; 1497.75 (M+K)⁺. MALDI-TOF calc. for C₉₀H₇₀N₆O₁₄, 1459.50 (M+H)⁺; ν (nujol) / (cm⁻¹): 3321, 3159, 2954, 2922, 2852, 2725, 1716, 1703, 1670, 1458, 1377, 1305, 1242, 1169, 1153, 1078, 1016.

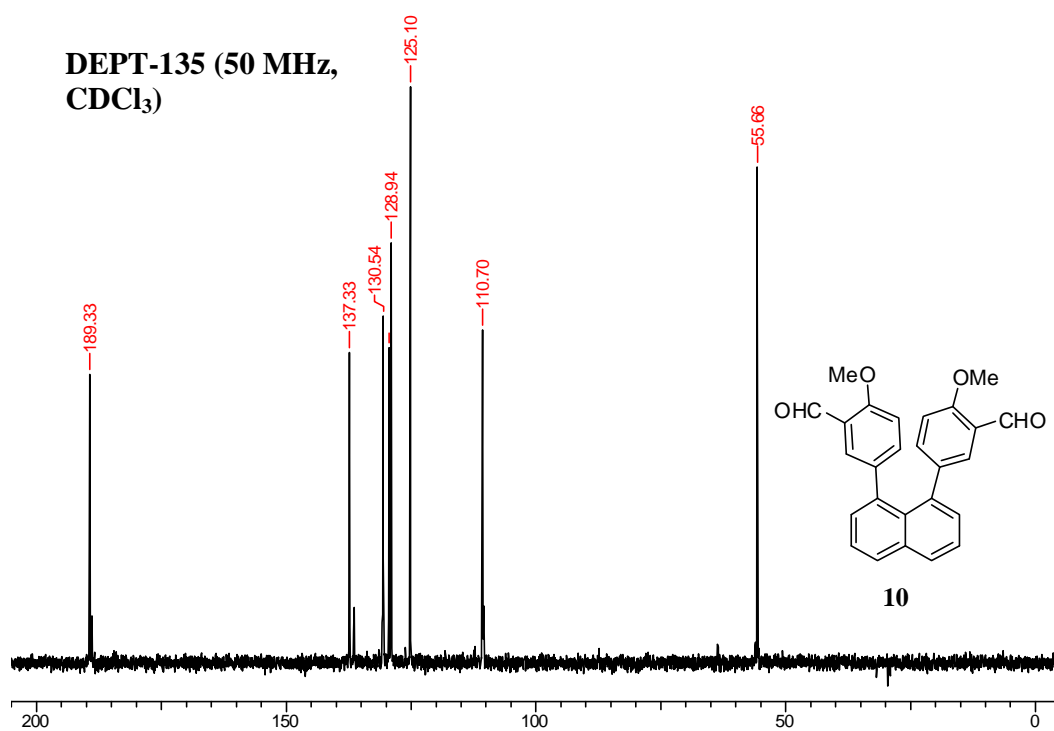
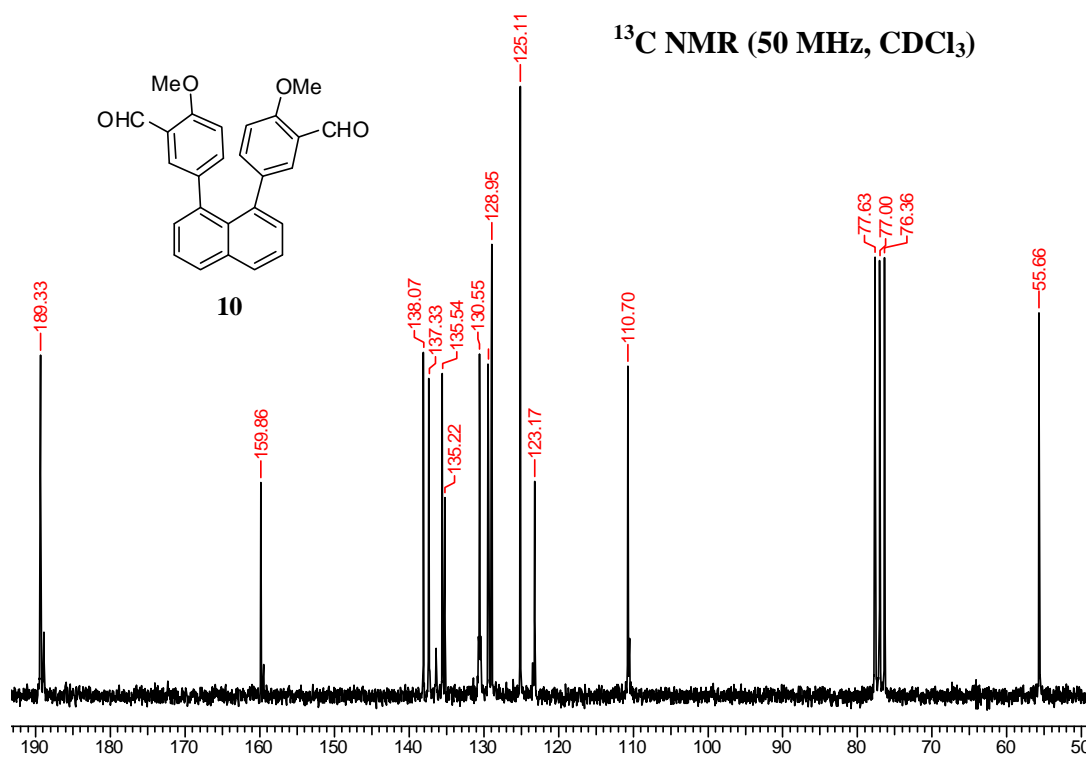


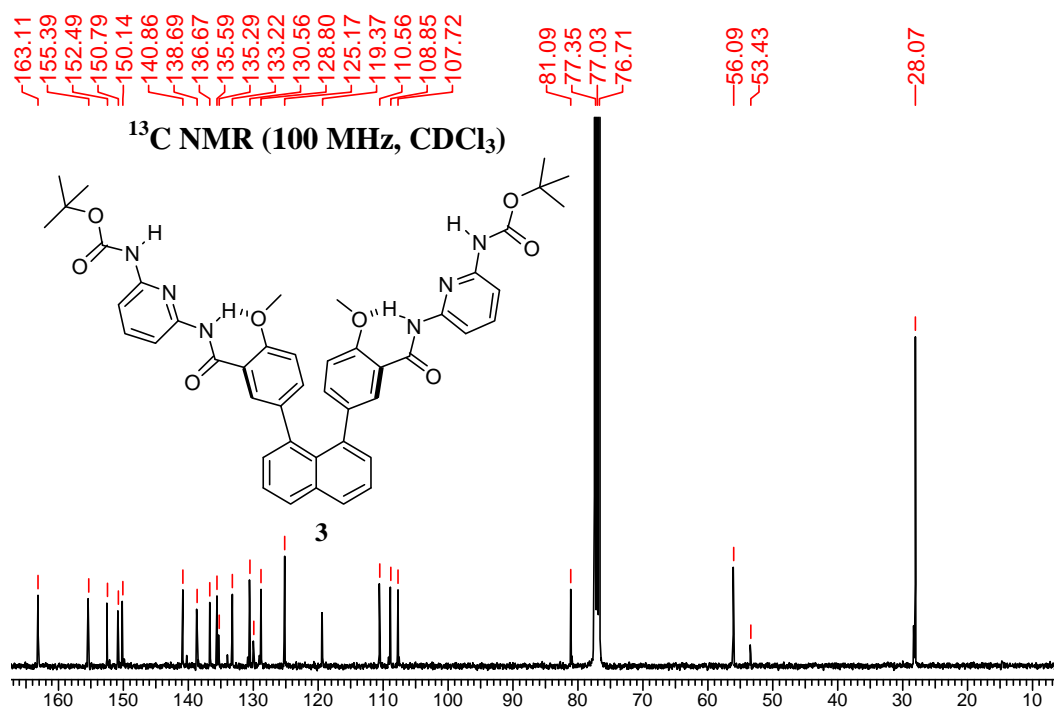












the peak at 53.43 ppm corresponds to DCM in the crystal lattice

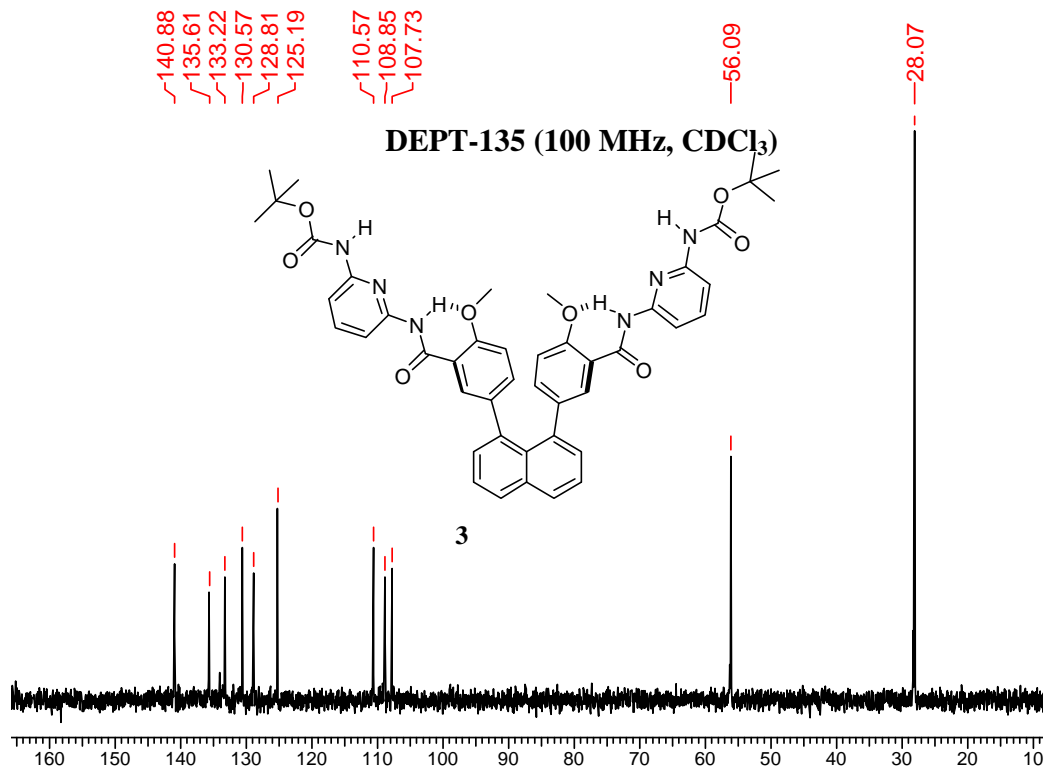
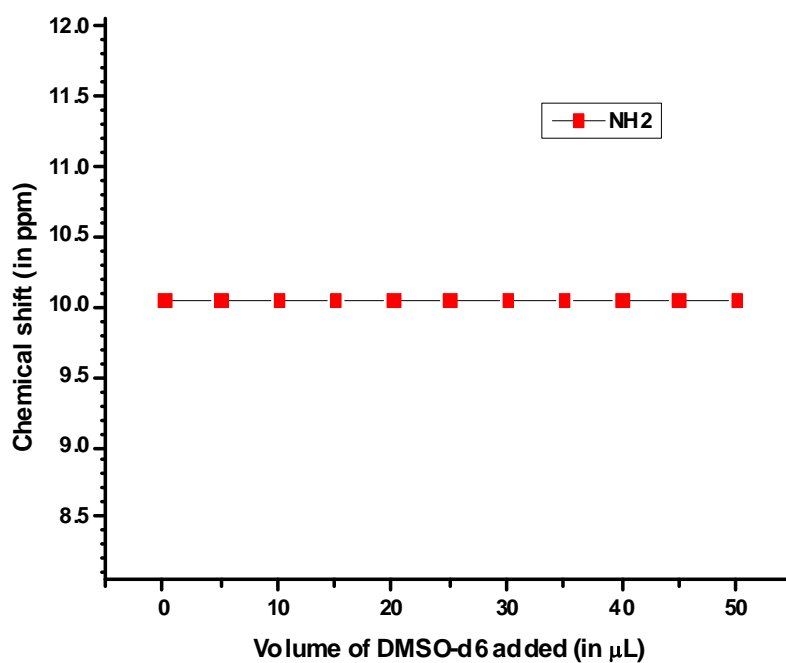
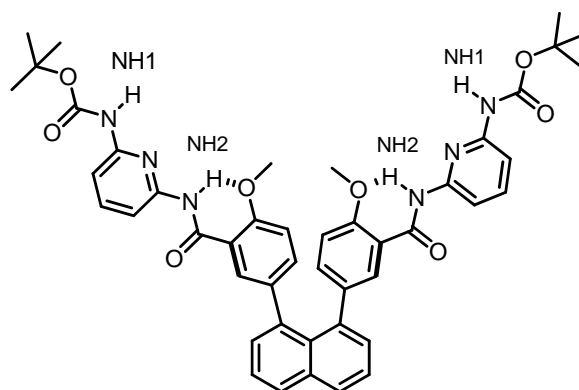


Table 1: Titration study of **3** in CDCl₃ (V_{DMSO-d6} added at each step = 5 μl)

| No | V _{DMSO-d6} (in μl) | δ _{NH2} |
|----|---------------------------------|------------------|
| 1 | 0 | 10.05 |
| 2 | 5 | 10.05 |
| 3 | 10 | 10.05 |
| 4 | 15 | 10.05 |
| 5 | 20 | 10.05 |
| 6 | 25 | 10.05 |
| 7 | 30 | 10.05 |
| 8 | 35 | 10.05 |
| 9 | 40 | 10.05 |
| 10 | 45 | 10.05 |
| 11 | 50 | 10.05 |



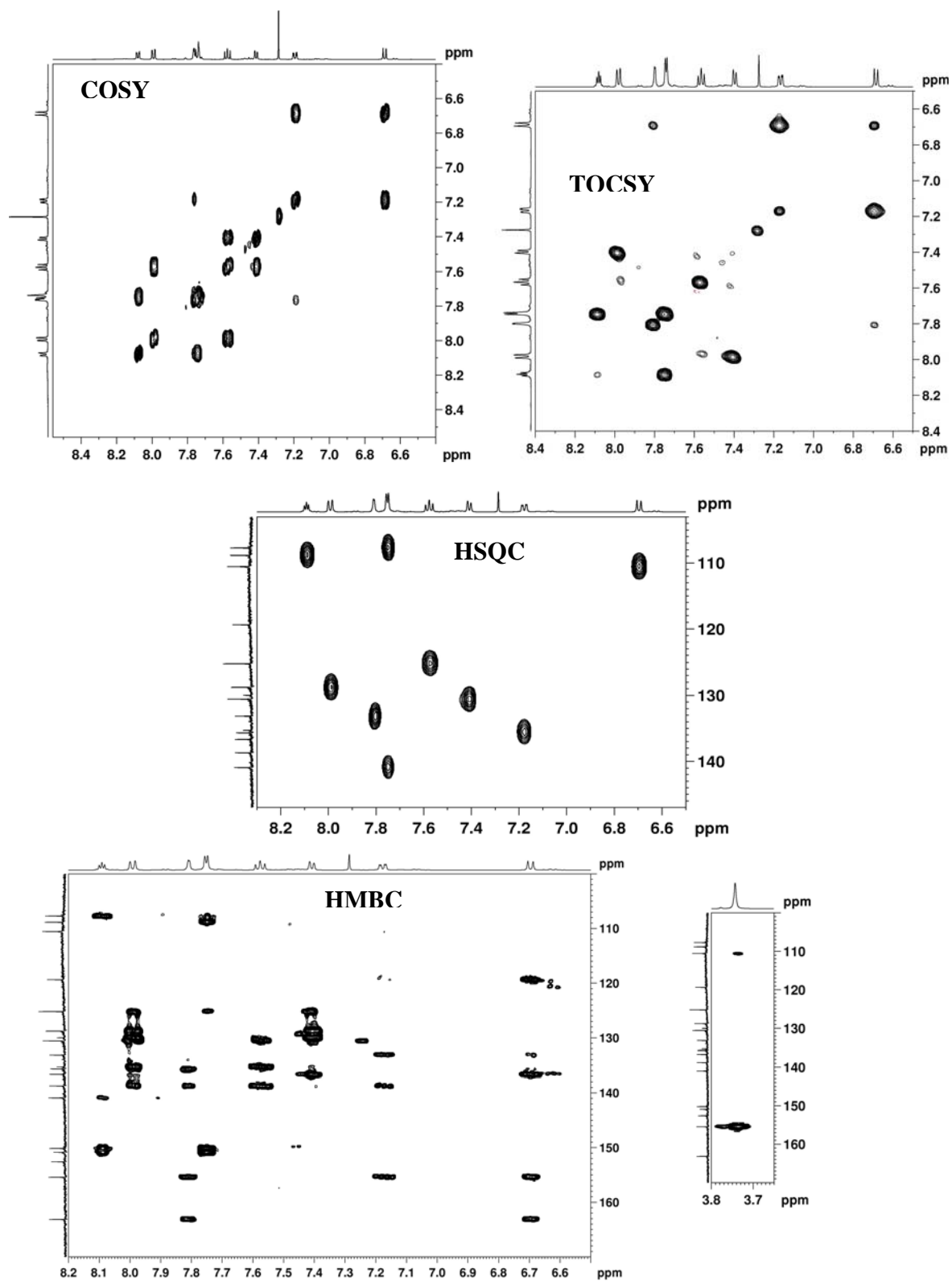


Figure S1. Partial COSY, TOCSY, HSQC, and HMBC spectra of **3**

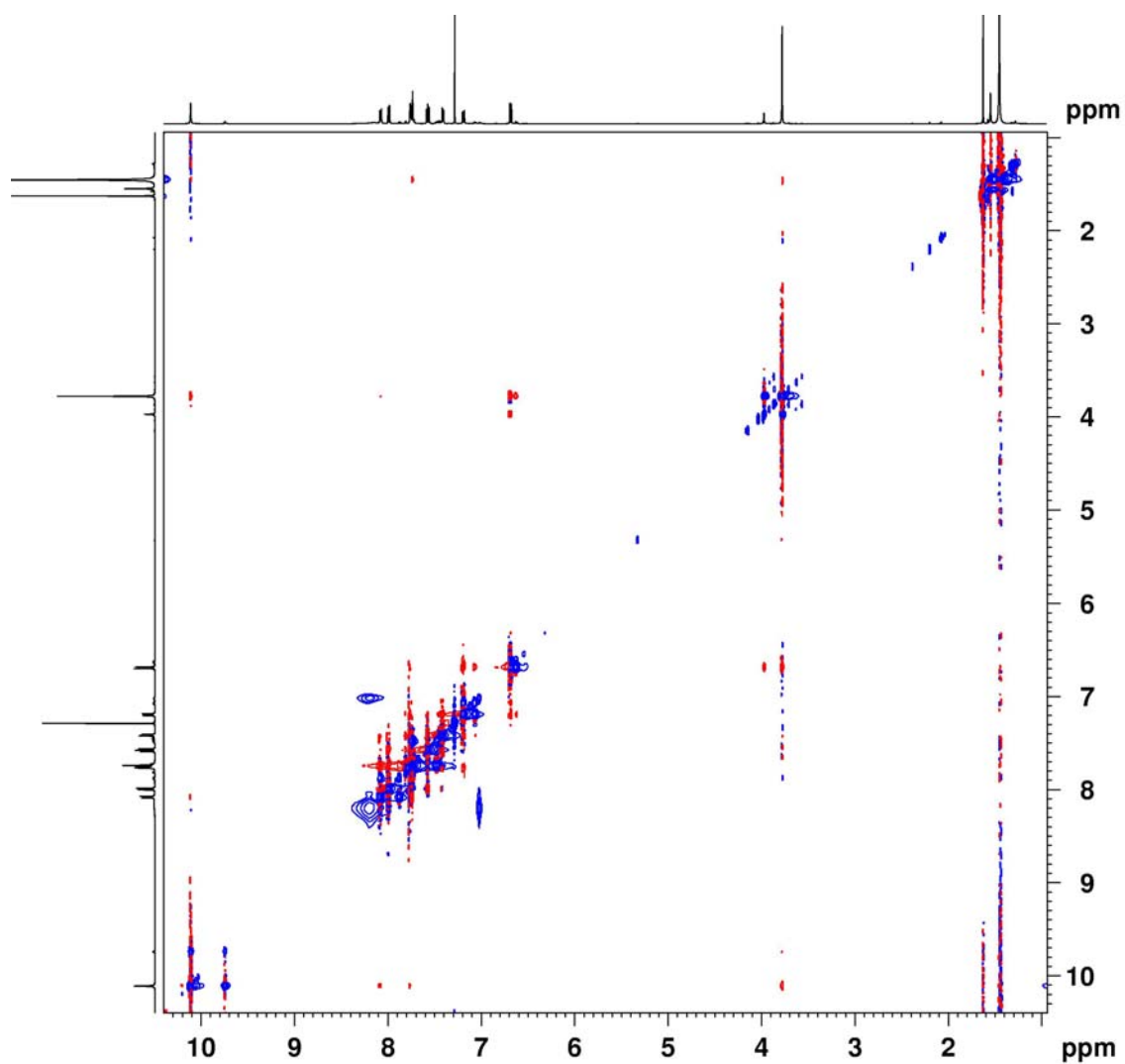


Figure S3. NOESY spectrum of **3** (500 MHz, CDCl₃)

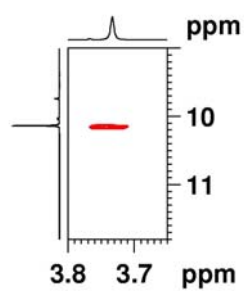


Figure S4. 2D NOESY extract of **3** showing nOe between OMe and NH (500 MHz, CDCl₃)

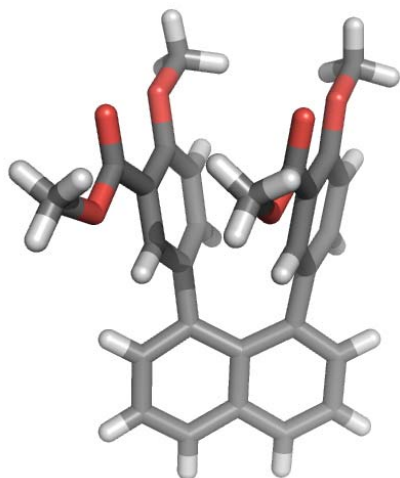


Figure S5. Crystal structure of **2a** with cofacial aryl rings containing the ester groups
| in *syn* orientation.

Details of the *ab initio* calculations

At first, a systematic conformational analysis for compound **3** was performed at the HF/6-31G* level of *ab initio* MO theory to test the reliability of the quantum chemical methods. Several conformational alternatives were considered as starting conformations of geometry optimization resulting from different torsion angles of the single bonds in the molecule. According to this analysis the anti-conformer with the side chains on the phenyl rings in the 1,8-positions pointing into opposite directions referred to the naphthalene ring is slightly favored over the syn-conformer, which corresponds to the X-ray structure having these groups in parallel orientation ($\Delta E = 10.0$ kJ/mol in favour to anti). Considering correlation energy at the MP2//HF/6-31G* level, the stability order is reversed. The experimentally found syn-structure is clearly favoured by 9.0 kJ/mol over the anti-form now. This can be expected because of the influence of dispersion interactions between the parallel aromatic side chains in the syn-form, which is not considered at the Hartree-Fock level. Therefore, this structure was taken as the basis for the optimization of the much larger oligomer **5**. The geometry of compounds **3** (syn and anti) and **5** is given as pdb-files. All calculations were performed employing the Gaussian03 software package.

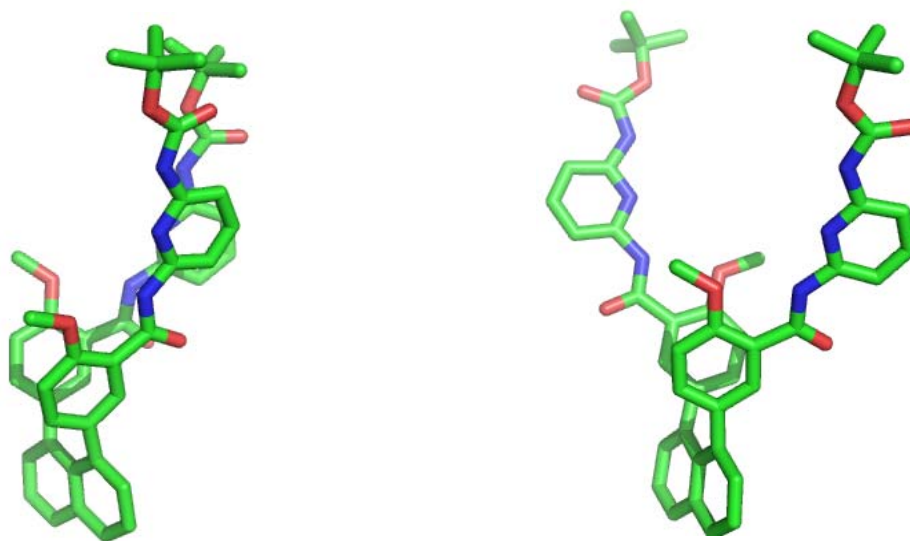


Figure S6. Possible isomers (*ab initio* models) of compound **3**: *Syn* (left) and *Anti* (right). *Note: The overlap of side arms (Pyridyl rings) is noticeable in the Syn isomer.*

Pdb-files for the most stable conformers of **3** and **5**

Compound 3 *Syn* ($E_T(\text{HF}/6\text{-}31\text{G}^*)$) = -2694.483943 a.u.;
 $E_T(\text{MP2}/\text{HF}/6\text{-}31\text{G}^*)$ = -2702.710143 a.u.)

```
HEADER
REMARK exported
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HETATM 2 N UNK 0001 -4.393 2.488 -0.366
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HETATM 4 C UNK 0001 -3.496 1.685 -1.076
HETATM 5 N UNK 0001 -2.240 1.969 -0.790
HETATM 6 O UNK 0001 -6.191 3.422 0.437
HETATM 7 C UNK 0001 -3.867 0.705 -1.985
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HETATM 9 O UNK 0001 1.351 0.531 -2.312
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HETATM 12 O UNK 0001 1.291 3.393 0.701
HETATM 13 C UNK 0001 -1.509 0.297 -2.321
HETATM 14 N UNK 0001 -1.037 -2.846 0.200
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HETATM 16 C UNK 0001 -1.269 1.301 -1.391
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HETATM 18 O UNK 0001 -5.393 -2.724 0.697
HETATM 19 C UNK 0001 1.211 1.316 -1.412
HETATM 20 O UNK 0001 3.044 -3.080 -0.609
HETATM 21 C UNK 0001 2.431 1.880 -0.719
HETATM 22 O UNK 0001 1.766 -1.504 2.957
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| | | | | | | | |
|--------|----|---|-----|------|--------|--------|--------|
| HETATM | 23 | C | UNK | 0001 | 2.460 | 2.851 | 0.293 |
| HETATM | 24 | H | UNK | 0001 | -8.402 | -2.108 | -0.696 |
| HETATM | 25 | C | UNK | 0001 | 3.676 | 3.231 | 0.844 |
| HETATM | 26 | C | UNK | 0001 | 4.854 | 2.657 | 0.404 |
| HETATM | 27 | C | UNK | 0001 | 4.861 | 1.699 | -0.597 |
| HETATM | 28 | C | UNK | 0001 | 3.640 | 1.339 | -1.145 |
| HETATM | 29 | C | UNK | 0001 | 6.161 | 1.190 | -1.144 |
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| HETATM | 31 | C | UNK | 0001 | 7.757 | 1.394 | -2.970 |
| HETATM | 32 | C | UNK | 0001 | 8.715 | 0.787 | -2.231 |
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| HETATM | 34 | C | UNK | 0001 | 9.466 | -0.301 | -0.173 |
| HETATM | 35 | C | UNK | 0001 | 9.221 | -0.804 | 1.059 |
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| HETATM | 37 | C | UNK | 0001 | 6.863 | -0.236 | 0.869 |
| HETATM | 38 | C | UNK | 0001 | 7.121 | 0.423 | -0.388 |
| HETATM | 39 | C | UNK | 0001 | 5.493 | -0.491 | 1.420 |
| HETATM | 40 | C | UNK | 0001 | 5.114 | -0.062 | 2.679 |
| HETATM | 41 | C | UNK | 0001 | 3.877 | -0.393 | 3.211 |
| HETATM | 42 | C | UNK | 0001 | 2.984 | -1.164 | 2.485 |
| HETATM | 43 | C | UNK | 0001 | 3.347 | -1.628 | 1.211 |
| HETATM | 44 | C | UNK | 0001 | 4.598 | -1.288 | 0.718 |
| HETATM | 45 | C | UNK | 0001 | 2.511 | -2.478 | 0.283 |
| HETATM | 46 | C | UNK | 0001 | 0.171 | -3.071 | -0.288 |
| HETATM | 47 | C | UNK | 0001 | 0.394 | -3.815 | -1.440 |
| HETATM | 48 | C | UNK | 0001 | -0.727 | -4.319 | -2.072 |
| HETATM | 49 | C | UNK | 0001 | -2.002 | -4.092 | -1.586 |
| HETATM | 50 | C | UNK | 0001 | -2.091 | -3.332 | -0.428 |
| HETATM | 51 | C | UNK | 0001 | -4.564 | -3.265 | -0.179 |
| HETATM | 52 | C | UNK | 0001 | 1.267 | 4.423 | 1.654 |
| HETATM | 53 | C | UNK | 0001 | 1.383 | -1.168 | 4.264 |
| HETATM | 54 | C | UNK | 0001 | -7.599 | 3.698 | 0.642 |
| HETATM | 55 | C | UNK | 0001 | -7.569 | 4.867 | 1.622 |
| HETATM | 56 | C | UNK | 0001 | -8.291 | 2.491 | 1.272 |
| HETATM | 57 | C | UNK | 0001 | -8.253 | 4.119 | -0.672 |
| HETATM | 58 | C | UNK | 0001 | -6.836 | -2.821 | 0.584 |
| HETATM | 59 | C | UNK | 0001 | -7.274 | -4.281 | 0.670 |
| HETATM | 60 | C | UNK | 0001 | -7.315 | -2.045 | 1.807 |
| HETATM | 61 | C | UNK | 0001 | -7.317 | -2.144 | -0.697 |
| HETATM | 62 | H | UNK | 0001 | -6.939 | -2.499 | 2.718 |
| HETATM | 63 | H | UNK | 0001 | -8.399 | -2.041 | 1.847 |
| HETATM | 64 | H | UNK | 0001 | -6.994 | -2.685 | -1.575 |
| HETATM | 65 | H | UNK | 0001 | -3.957 | 3.169 | 0.214 |
| HETATM | 66 | H | UNK | 0001 | -0.038 | 2.404 | -0.248 |
| HETATM | 67 | H | UNK | 0001 | -3.174 | -2.432 | 1.018 |
| HETATM | 68 | H | UNK | 0001 | -0.707 | -0.230 | -2.786 |
| HETATM | 69 | H | UNK | 0001 | 0.816 | -1.987 | 1.276 |
| HETATM | 70 | H | UNK | 0001 | 3.717 | 3.975 | 1.614 |
| HETATM | 71 | H | UNK | 0001 | 5.783 | 2.974 | 0.845 |
| HETATM | 72 | H | UNK | 0001 | 3.599 | 0.603 | -1.923 |
| HETATM | 73 | H | UNK | 0001 | 5.766 | 2.192 | -2.953 |
| HETATM | 74 | H | UNK | 0001 | 7.966 | 1.748 | -3.963 |
| HETATM | 75 | H | UNK | 0001 | 9.709 | 0.653 | -2.620 |
| HETATM | 76 | H | UNK | 0001 | 10.451 | -0.350 | -0.602 |
| HETATM | 77 | H | UNK | 0001 | 10.006 | -1.258 | 1.638 |

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|--------|-----|----|-----|------|--------|--------|--------|
| HETATM | 78 | H | UNK | 0001 | 7.697 | -1.299 | 2.485 |
| HETATM | 79 | H | UNK | 0001 | 5.783 | 0.546 | 3.260 |
| HETATM | 80 | H | UNK | 0001 | 3.622 | -0.038 | 4.190 |
| HETATM | 81 | H | UNK | 0001 | 4.866 | -1.670 | -0.247 |
| HETATM | 82 | H | UNK | 0001 | 1.381 | -3.980 | -1.810 |
| HETATM | 83 | H | UNK | 0001 | -0.604 | -4.901 | -2.968 |
| HETATM | 84 | H | UNK | 0001 | -2.874 | -4.476 | -2.068 |
| HETATM | 85 | H | UNK | 0001 | -4.894 | 0.500 | -2.197 |
| HETATM | 86 | H | UNK | 0001 | -3.068 | -0.759 | -3.303 |
| HETATM | 87 | H | UNK | 0001 | -6.946 | -1.127 | -0.752 |
| HETATM | 88 | H | UNK | 0001 | -6.968 | -1.020 | 1.766 |
| HETATM | 89 | H | UNK | 0001 | 1.657 | 4.086 | 2.607 |
| HETATM | 90 | H | UNK | 0001 | 0.229 | 4.695 | 1.769 |
| HETATM | 91 | H | UNK | 0001 | 1.828 | 5.285 | 1.314 |
| HETATM | 92 | H | UNK | 0001 | 2.066 | -1.591 | 4.991 |
| HETATM | 93 | H | UNK | 0001 | 0.402 | -1.594 | 4.404 |
| HETATM | 94 | H | UNK | 0001 | 1.330 | -0.094 | 4.395 |
| HETATM | 95 | H | UNK | 0001 | -7.068 | 4.581 | 2.540 |
| HETATM | 96 | H | UNK | 0001 | -8.578 | 5.180 | 1.864 |
| HETATM | 97 | H | UNK | 0001 | -7.040 | 5.710 | 1.193 |
| HETATM | 98 | H | UNK | 0001 | -8.338 | 1.658 | 0.586 |
| HETATM | 99 | H | UNK | 0001 | -9.302 | 2.763 | 1.557 |
| HETATM | 100 | H | UNK | 0001 | -7.761 | 2.180 | 2.167 |
| HETATM | 101 | H | UNK | 0001 | -7.703 | 4.941 | -1.118 |
| HETATM | 102 | H | UNK | 0001 | -9.266 | 4.457 | -0.478 |
| HETATM | 103 | H | UNK | 0001 | -8.291 | 3.301 | -1.376 |
| HETATM | 104 | H | UNK | 0001 | -6.951 | -4.844 | -0.193 |
| HETATM | 105 | H | UNK | 0001 | -8.356 | -4.329 | 0.730 |
| HETATM | 106 | H | UNK | 0001 | -6.866 | -4.744 | 1.563 |
| CONECT | 1 | 2 | 3 | 6 | | | |
| CONECT | 2 | 1 | 4 | 65 | | | |
| CONECT | 3 | 1 | | | | | |
| CONECT | 4 | 2 | 5 | 7 | | | |
| CONECT | 5 | 4 | 16 | | | | |
| CONECT | 6 | 1 | 54 | | | | |
| CONECT | 7 | 4 | 10 | 85 | | | |
| CONECT | 8 | 16 | 19 | 66 | | | |
| CONECT | 9 | 19 | | | | | |
| CONECT | 10 | 7 | 13 | 86 | | | |
| CONECT | 11 | 50 | 51 | 67 | | | |
| CONECT | 12 | 23 | 52 | | | | |
| CONECT | 13 | 10 | 16 | 68 | | | |
| CONECT | 14 | 46 | 50 | | | | |
| CONECT | 15 | 51 | | | | | |
| CONECT | 16 | 5 | 8 | 13 | | | |
| CONECT | 17 | 45 | 46 | 69 | | | |
| CONECT | 18 | 51 | 58 | | | | |
| CONECT | 19 | 8 | 9 | 21 | | | |
| CONECT | 20 | 45 | | | | | |
| CONECT | 21 | 19 | 23 | 28 | | | |
| CONECT | 22 | 42 | 53 | | | | |
| CONECT | 23 | 12 | 21 | 25 | | | |
| CONECT | 24 | 61 | | | | | |
| CONECT | 25 | 23 | 26 | 70 | | | |
| CONECT | 26 | 25 | 27 | 71 | | | |

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CONNECT 102 57
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CONNECT 104 59
CONNECT 105 59
CONNECT 106 59
END

Compound 3 (*Anti*) ($E_T(\text{HF}/6\text{-}31\text{G}^*)$) = -2694.487753 a.u.;

$E_T(\text{MP2//}6\text{-}31\text{G}^*)$ = -2702.706700 a.u.)

HEADER
REMARK exported
HETATM 1 C UNK 0001 -4.696 6.200 -0.028
HETATM 2 N UNK 0001 -3.530 5.520 0.148
HETATM 3 O UNK 0001 -4.960 6.895 -0.963
HETATM 4 C UNK 0001 -2.408 5.473 -0.679
HETATM 5 N UNK 0001 -1.457 4.700 -0.190
HETATM 6 O UNK 0001 -5.484 5.963 1.007
HETATM 7 C UNK 0001 -2.289 6.156 -1.882
HETATM 8 N UNK 0001 0.568 3.712 -0.201
HETATM 9 O UNK 0001 2.298 3.650 -1.640
HETATM 10 C UNK 0001 -1.097 5.989 -2.563
HETATM 11 N UNK 0001 -3.538 -5.515 -0.148
HETATM 12 O UNK 0001 0.770 1.894 1.785
HETATM 13 C UNK 0001 -0.083 5.186 -2.073
HETATM 14 N UNK 0001 -1.465 -4.698 0.190
HETATM 15 O UNK 0001 -4.970 -6.888 0.963
HETATM 16 C UNK 0001 -0.325 4.555 -0.859
HETATM 17 N UNK 0001 0.562 -3.711 0.201
HETATM 18 O UNK 0001 -5.492 -5.957 -1.008

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|--------|----|---|-----|------|--------|--------|--------|
| HETATM | 19 | C | UNK | 0001 | 1.800 | 3.301 | -0.601 |
| HETATM | 20 | O | UNK | 0001 | 2.291 | -3.650 | 1.641 |
| HETATM | 21 | C | UNK | 0001 | 2.563 | 2.385 | 0.324 |
| HETATM | 22 | O | UNK | 0001 | 0.766 | -1.893 | -1.784 |
| HETATM | 23 | C | UNK | 0001 | 2.065 | 1.712 | 1.448 |
| HETATM | 24 | H | UNK | 0001 | -7.704 | -8.461 | -1.454 |
| HETATM | 25 | C | UNK | 0001 | 2.905 | 0.882 | 2.178 |
| HETATM | 26 | C | UNK | 0001 | 4.222 | 0.709 | 1.798 |
| HETATM | 27 | C | UNK | 0001 | 4.743 | 1.352 | 0.685 |
| HETATM | 28 | C | UNK | 0001 | 3.897 | 2.187 | -0.024 |
| HETATM | 29 | C | UNK | 0001 | 6.198 | 1.232 | 0.345 |
| HETATM | 30 | C | UNK | 0001 | 6.943 | 2.367 | 0.502 |
| HETATM | 31 | C | UNK | 0001 | 8.349 | 2.370 | 0.394 |
| HETATM | 32 | C | UNK | 0001 | 8.998 | 1.201 | 0.185 |
| HETATM | 33 | C | UNK | 0001 | 8.276 | -0.007 | -0.001 |
| HETATM | 34 | C | UNK | 0001 | 8.996 | -1.215 | -0.187 |
| HETATM | 35 | C | UNK | 0001 | 8.344 | -2.383 | -0.395 |
| HETATM | 36 | C | UNK | 0001 | 6.939 | -2.377 | -0.503 |
| HETATM | 37 | C | UNK | 0001 | 6.196 | -1.241 | -0.346 |
| HETATM | 38 | C | UNK | 0001 | 6.856 | -0.005 | -0.001 |
| HETATM | 39 | C | UNK | 0001 | 4.740 | -1.358 | -0.685 |
| HETATM | 40 | C | UNK | 0001 | 4.220 | -0.715 | -1.798 |
| HETATM | 41 | C | UNK | 0001 | 2.902 | -0.885 | -2.177 |
| HETATM | 42 | C | UNK | 0001 | 2.061 | -1.714 | -1.448 |
| HETATM | 43 | C | UNK | 0001 | 2.559 | -2.387 | -0.323 |
| HETATM | 44 | C | UNK | 0001 | 3.893 | -2.192 | 0.024 |
| HETATM | 45 | C | UNK | 0001 | 1.793 | -3.302 | 0.602 |
| HETATM | 46 | C | UNK | 0001 | -0.333 | -4.553 | 0.860 |
| HETATM | 47 | C | UNK | 0001 | -0.092 | -5.184 | 2.075 |
| HETATM | 48 | C | UNK | 0001 | -1.107 | -5.985 | 2.564 |
| HETATM | 49 | C | UNK | 0001 | -2.298 | -6.152 | 1.883 |
| HETATM | 50 | C | UNK | 0001 | -2.416 | -5.469 | 0.680 |
| HETATM | 51 | C | UNK | 0001 | -4.705 | -6.195 | 0.028 |
| HETATM | 52 | C | UNK | 0001 | 0.185 | 1.151 | 2.825 |
| HETATM | 53 | C | UNK | 0001 | 0.181 | -1.149 | -2.824 |
| HETATM | 54 | C | UNK | 0001 | -6.809 | 6.539 | 1.141 |
| HETATM | 55 | C | UNK | 0001 | -7.286 | 5.969 | 2.473 |
| HETATM | 56 | C | UNK | 0001 | -7.709 | 6.064 | 0.002 |
| HETATM | 57 | C | UNK | 0001 | -6.718 | 8.061 | 1.211 |
| HETATM | 58 | C | UNK | 0001 | -6.817 | -6.532 | -1.142 |
| HETATM | 59 | C | UNK | 0001 | -7.717 | -6.055 | -0.004 |
| HETATM | 60 | C | UNK | 0001 | -7.293 | -5.963 | -2.475 |
| HETATM | 61 | C | UNK | 0001 | -6.728 | -8.054 | -1.211 |
| HETATM | 62 | H | UNK | 0001 | -6.035 | -8.355 | -1.990 |
| HETATM | 63 | H | UNK | 0001 | -8.291 | -6.321 | -2.700 |
| HETATM | 64 | H | UNK | 0001 | -7.315 | -4.879 | -2.440 |
| HETATM | 65 | H | UNK | 0001 | -3.449 | 4.979 | 0.979 |
| HETATM | 66 | H | UNK | 0001 | 0.214 | 3.349 | 0.656 |
| HETATM | 67 | H | UNK | 0001 | -3.456 | -4.975 | -0.979 |
| HETATM | 68 | H | UNK | 0001 | 0.839 | 5.053 | -2.592 |
| HETATM | 69 | H | UNK | 0001 | 0.208 | -3.349 | -0.655 |
| HETATM | 70 | H | UNK | 0001 | 2.542 | 0.360 | 3.040 |
| HETATM | 71 | H | UNK | 0001 | 4.848 | 0.061 | 2.383 |
| HETATM | 72 | H | UNK | 0001 | 4.259 | 2.702 | -0.892 |
| HETATM | 73 | H | UNK | 0001 | 6.443 | 3.280 | 0.767 |

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|--------|-----|----|-----|------|--------|--------|--------|
| HETATM | 74 | H | UNK | 0001 | 8.892 | 3.288 | 0.524 |
| HETATM | 75 | H | UNK | 0001 | 10.073 | 1.166 | 0.152 |
| HETATM | 76 | H | UNK | 0001 | 10.071 | -1.182 | -0.155 |
| HETATM | 77 | H | UNK | 0001 | 8.885 | -3.303 | -0.526 |
| HETATM | 78 | H | UNK | 0001 | 6.436 | -3.290 | -0.768 |
| HETATM | 79 | H | UNK | 0001 | 4.847 | -0.068 | -2.384 |
| HETATM | 80 | H | UNK | 0001 | 2.540 | -0.362 | -3.039 |
| HETATM | 81 | H | UNK | 0001 | 4.254 | -2.707 | 0.892 |
| HETATM | 82 | H | UNK | 0001 | 0.830 | -5.052 | 2.593 |
| HETATM | 83 | H | UNK | 0001 | -0.965 | -6.495 | 3.501 |
| HETATM | 84 | H | UNK | 0001 | -3.088 | -6.769 | 2.252 |
| HETATM | 85 | H | UNK | 0001 | -3.077 | 6.775 | -2.251 |
| HETATM | 86 | H | UNK | 0001 | -0.954 | 6.499 | -3.499 |
| HETATM | 87 | H | UNK | 0001 | -6.630 | -6.265 | -3.277 |
| HETATM | 88 | H | UNK | 0001 | -6.403 | -8.477 | -0.272 |
| HETATM | 89 | H | UNK | 0001 | 0.266 | 0.088 | 2.637 |
| HETATM | 90 | H | UNK | 0001 | -0.856 | 1.435 | 2.841 |
| HETATM | 91 | H | UNK | 0001 | 0.637 | 1.391 | 3.780 |
| HETATM | 92 | H | UNK | 0001 | 0.265 | -0.086 | -2.636 |
| HETATM | 93 | H | UNK | 0001 | -0.860 | -1.431 | -2.839 |
| HETATM | 94 | H | UNK | 0001 | 0.632 | -1.390 | -3.779 |
| HETATM | 95 | H | UNK | 0001 | -7.309 | 4.886 | 2.438 |
| HETATM | 96 | H | UNK | 0001 | -8.284 | 6.328 | 2.698 |
| HETATM | 97 | H | UNK | 0001 | -6.622 | 6.271 | 3.276 |
| HETATM | 98 | H | UNK | 0001 | -7.392 | 6.466 | -0.949 |
| HETATM | 99 | H | UNK | 0001 | -8.728 | 6.384 | 0.192 |
| HETATM | 100 | H | UNK | 0001 | -7.701 | 4.980 | -0.055 |
| HETATM | 101 | H | UNK | 0001 | -6.025 | 8.361 | 1.990 |
| HETATM | 102 | H | UNK | 0001 | -7.694 | 8.469 | 1.454 |
| HETATM | 103 | H | UNK | 0001 | -6.392 | 8.484 | 0.272 |
| HETATM | 104 | H | UNK | 0001 | -7.402 | -6.457 | 0.947 |
| HETATM | 105 | H | UNK | 0001 | -8.737 | -6.374 | -0.194 |
| HETATM | 106 | H | UNK | 0001 | -7.709 | -4.972 | 0.052 |
| CONECT | 1 | 2 | 3 | 6 | | | |
| CONECT | 2 | 1 | 4 | 65 | | | |
| CONECT | 3 | 1 | | | | | |
| CONECT | 4 | 2 | 5 | 7 | | | |
| CONECT | 5 | 4 | 16 | | | | |
| CONECT | 6 | 1 | 54 | | | | |
| CONECT | 7 | 4 | 10 | 85 | | | |
| CONECT | 8 | 16 | 19 | 66 | | | |
| CONECT | 9 | 19 | | | | | |
| CONECT | 10 | 7 | 13 | 86 | | | |
| CONECT | 11 | 50 | 51 | 67 | | | |
| CONECT | 12 | 23 | 52 | | | | |
| CONECT | 13 | 10 | 16 | 68 | | | |
| CONECT | 14 | 46 | 50 | | | | |
| CONECT | 15 | 51 | | | | | |
| CONECT | 16 | 5 | 8 | 13 | | | |
| CONECT | 17 | 45 | 46 | 69 | | | |
| CONECT | 18 | 51 | 58 | | | | |
| CONECT | 19 | 8 | 9 | 21 | | | |
| CONECT | 20 | 45 | | | | | |
| CONECT | 21 | 19 | 23 | 28 | | | |
| CONECT | 22 | 42 | 53 | | | | |

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END

Compound 5 ($E_T(\text{HF}/6\text{-}31\text{G}^*) = -4823.810231$ a.u.)

HEADER

REMARK exported

| | | | | | | | |
|--------|----|---|-----|------|--------|--------|--------|
| HETATM | 1 | N | UNK | 0001 | 5.098 | 2.233 | 0.066 |
| HETATM | 2 | N | UNK | 0001 | 0.709 | 2.895 | -0.660 |
| HETATM | 3 | N | UNK | 0001 | 4.056 | -1.990 | -0.906 |
| HETATM | 4 | N | UNK | 0001 | 1.808 | -2.031 | -0.754 |
| HETATM | 5 | N | UNK | 0001 | -0.382 | -2.255 | -0.269 |
| HETATM | 6 | N | UNK | 0001 | 2.944 | 2.653 | -0.457 |
| HETATM | 7 | C | UNK | 0001 | 12.434 | -0.698 | -0.148 |
| HETATM | 8 | C | UNK | 0001 | 11.069 | -0.758 | 0.232 |
| HETATM | 9 | C | UNK | 0001 | 12.873 | -1.306 | -1.353 |
| HETATM | 10 | C | UNK | 0001 | 13.380 | -0.027 | 0.671 |
| HETATM | 11 | C | UNK | 0001 | 10.199 | -1.593 | -0.560 |
| HETATM | 12 | C | UNK | 0001 | 10.666 | 0.023 | 1.376 |
| HETATM | 13 | C | UNK | 0001 | 9.243 | 0.350 | 1.714 |
| HETATM | 14 | C | UNK | 0001 | 8.838 | -2.050 | -0.127 |
| HETATM | 15 | C | UNK | 0001 | 10.670 | -2.142 | -1.719 |
| HETATM | 16 | C | UNK | 0001 | 11.620 | 0.639 | 2.135 |
| HETATM | 17 | C | UNK | 0001 | 12.005 | -1.976 | -2.145 |

| | | | | | | | |
|--------|----|---|-----|------|--------|--------|--------|
| HETATM | 18 | C | UNK | 0001 | 12.991 | 0.593 | 1.809 |
| HETATM | 19 | C | UNK | 0001 | 7.704 | -1.776 | -0.870 |
| HETATM | 20 | C | UNK | 0001 | 6.450 | -2.286 | -0.532 |
| HETATM | 21 | C | UNK | 0001 | 8.710 | -2.880 | 0.983 |
| HETATM | 22 | C | UNK | 0001 | 6.358 | -3.126 | 0.577 |
| HETATM | 23 | C | UNK | 0001 | 5.307 | -1.833 | -1.411 |
| HETATM | 24 | C | UNK | 0001 | 7.486 | -3.414 | 1.328 |
| HETATM | 25 | C | UNK | 0001 | 8.466 | 1.063 | 0.816 |
| HETATM | 26 | C | UNK | 0001 | 7.171 | 1.475 | 1.113 |
| HETATM | 27 | C | UNK | 0001 | 8.703 | 0.065 | 2.962 |
| HETATM | 28 | C | UNK | 0001 | 6.656 | 1.175 | 2.373 |
| HETATM | 29 | C | UNK | 0001 | 6.454 | 2.200 | -0.002 |
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| HETATM | 107 | O | UNK | 0001 | -3.389 | -4.384 | -2.784 |
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| HETATM | 173 | H | UNK | 0001 | 3.966 | -2.448 | -0.026 |
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| CONECT | 2 | 35 | 39 | 132 | | | |

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CONNECT 5 66 70 133
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END

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