

Resolution of a Configurationally Stable Hetero[4]helicene

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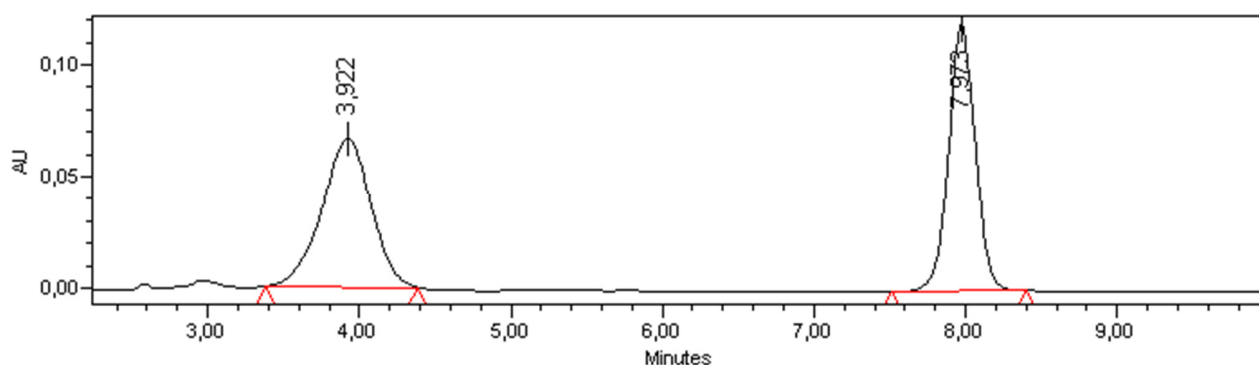
HPLC Analysis

The HPLC resolution of products was performed on a HPLC Waters Alliance 2695 equipped with a 200 μ L loop injector and a spectrophotometer UV Waters PDA 2996.

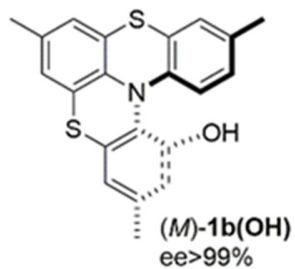
CHIRALPAK® IA (250 x 4.6 mm/ 5 μ m) purchased from Chiral Technologies Europe.

The mobile phase, delivered at a flow rate 1.2 mL/min, was hexane/CH₂Cl₂ 70/30 v/v + 1% MeOH.

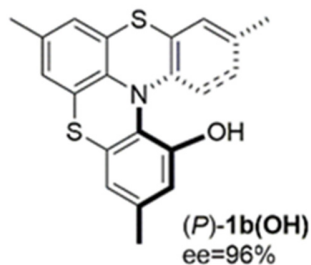
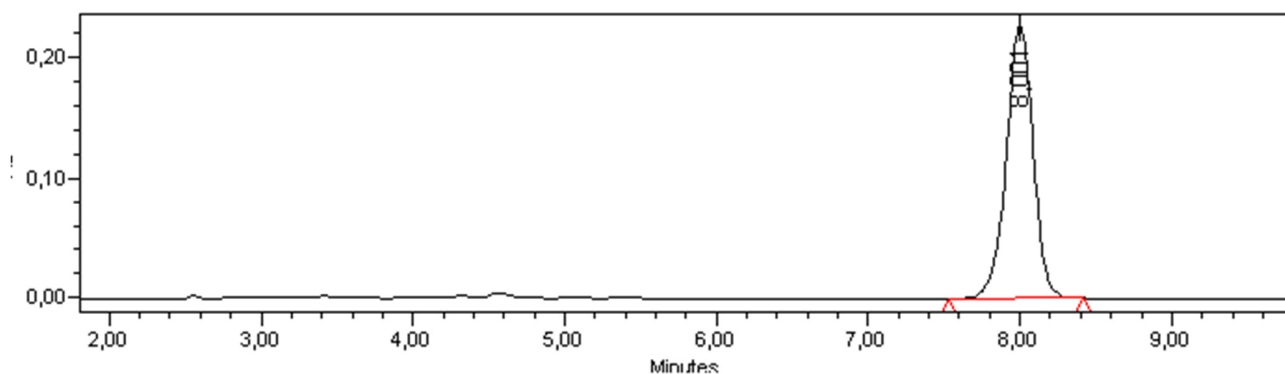
Racemate



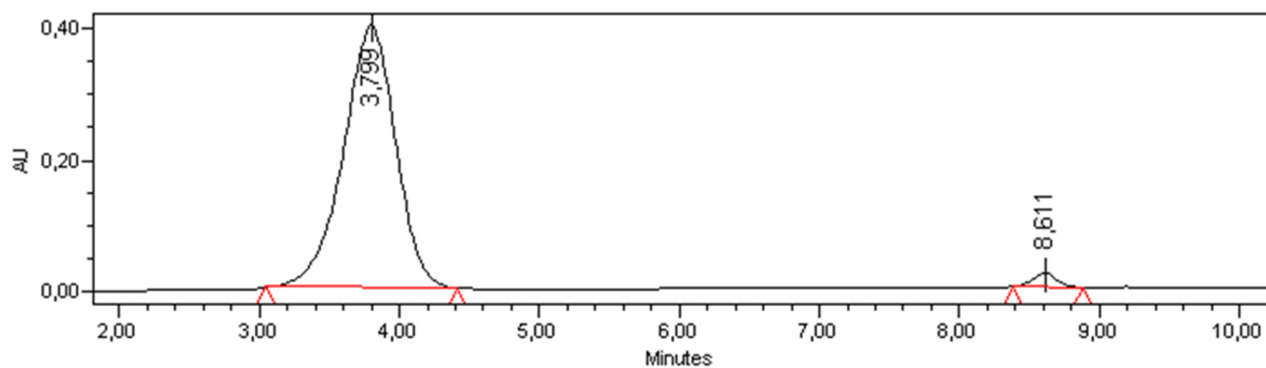
| 1b(OH) Racemic | Retention time (min) | %Area |
|----------------|----------------------|-------|
| (+) | 3.922 | 50.88 |
| (-) | 7.973 | 49.12 |



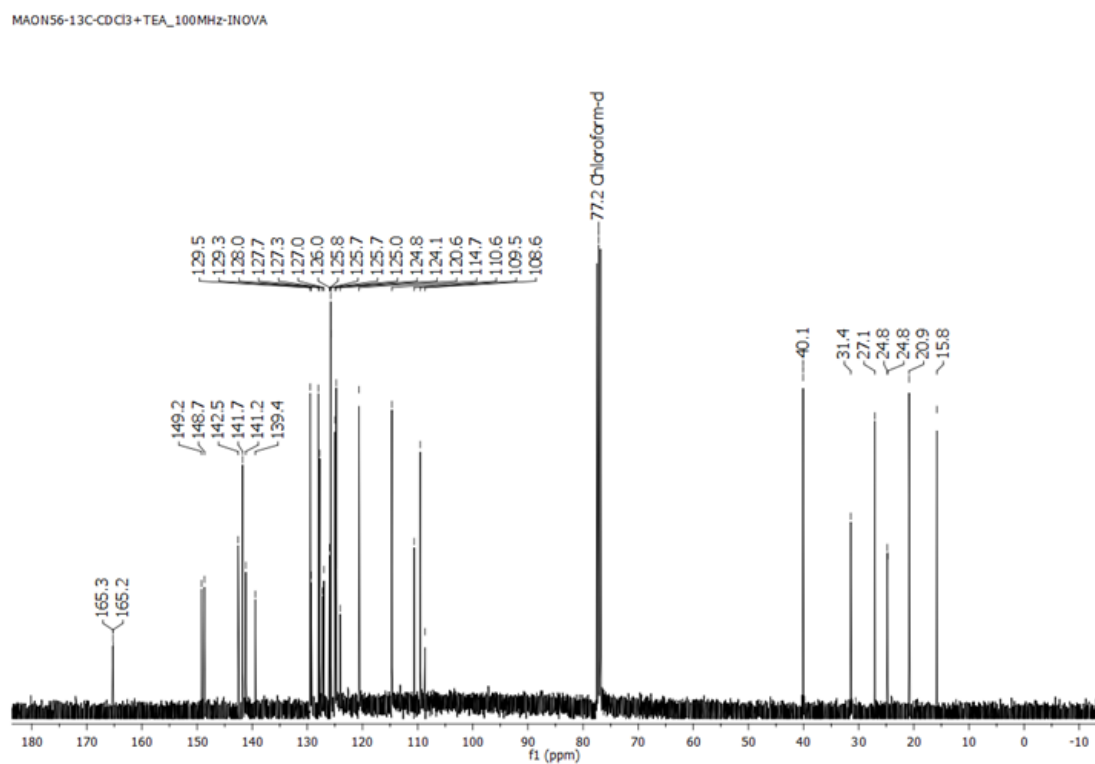
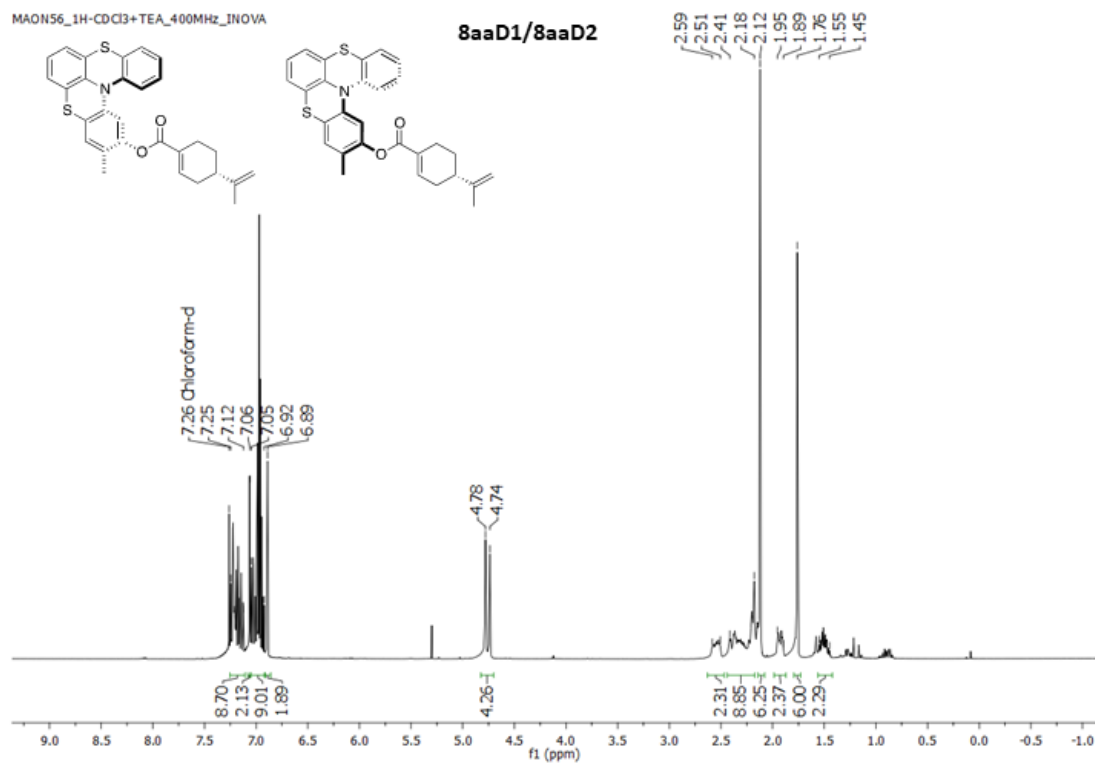
| $[\alpha]_D^{20} -167$ | | |
|---|----------------------|--------|
| (c 0.1, CH ₂ Cl ₂) | Retention time (min) | %Area |
| (M)-1b(OH) (-) | 8.001 | 100.00 |



| $[\alpha]_D^{20} +166$ | | |
|---|----------------------|-------|
| (c 0.1, CH ₂ Cl ₂) | Retention time (min) | %Area |
| (P)-1b(OH) (+) | 3.799 | 97.80 |
| (M)-1b(OH) (-) | 8.611 | 2.20 |

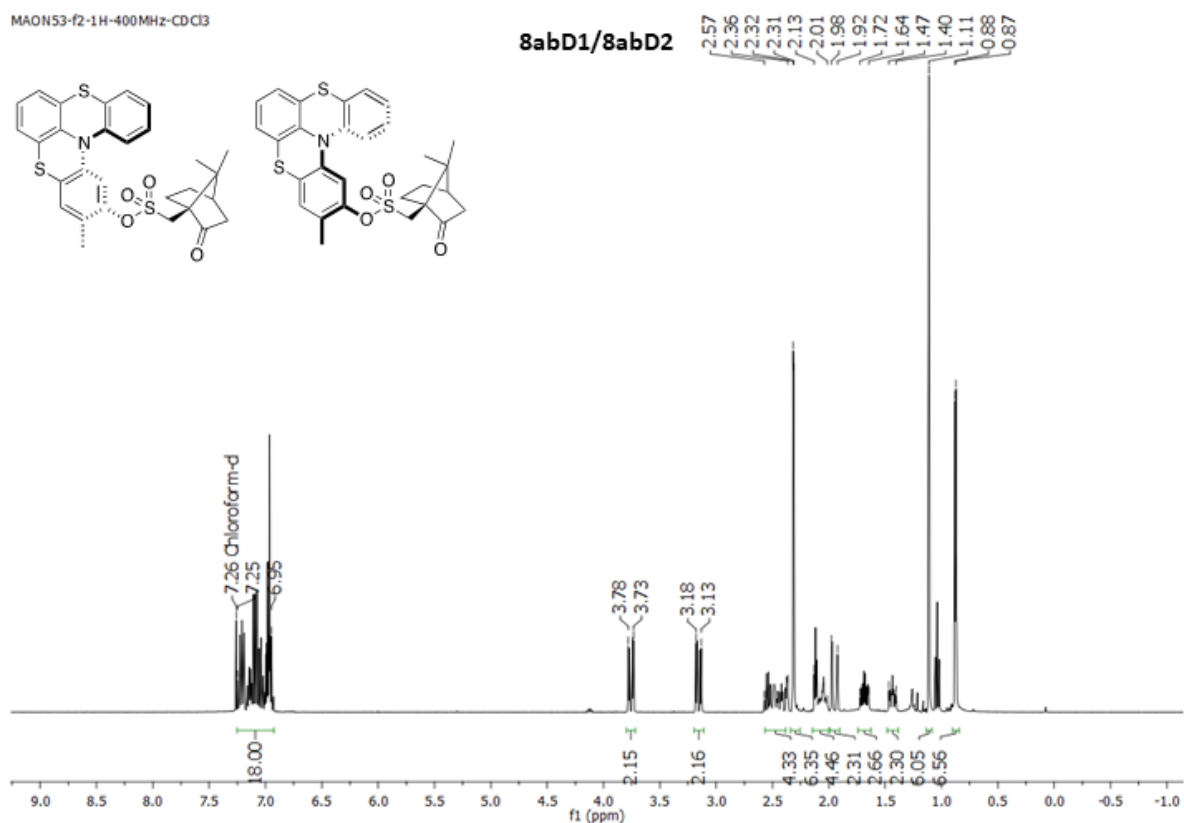


NMR spectra 400 MHz

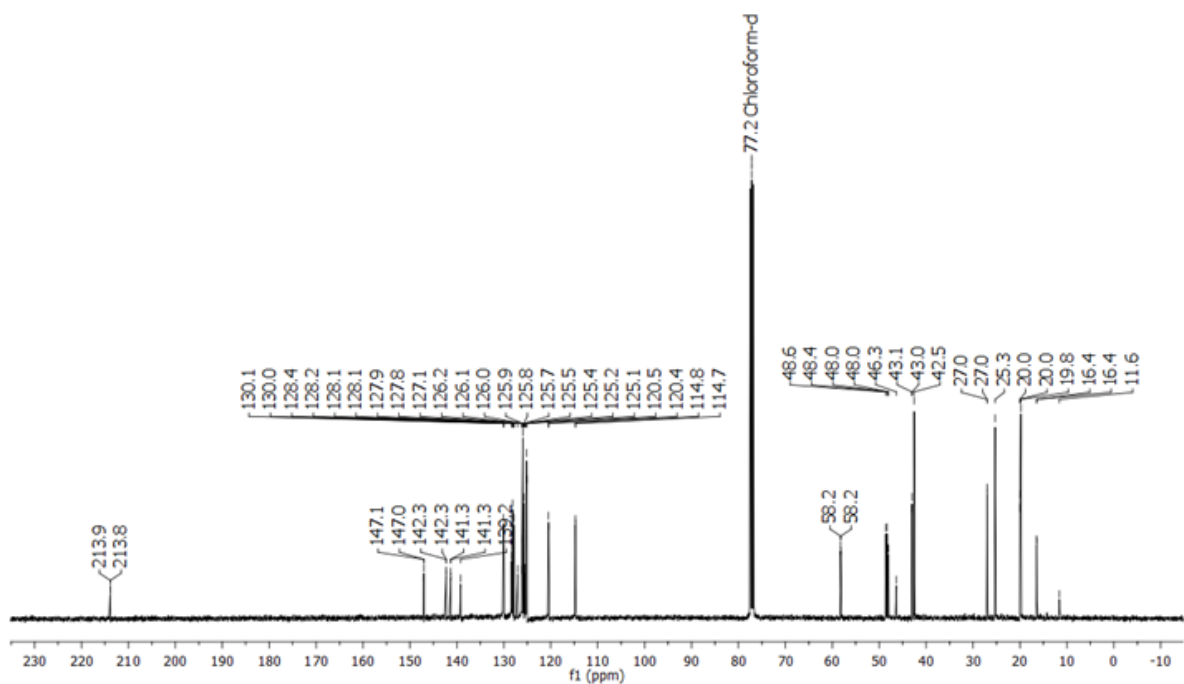


MAON53-f2-1H-400MHz-CDCl3

8abD1/8abD2

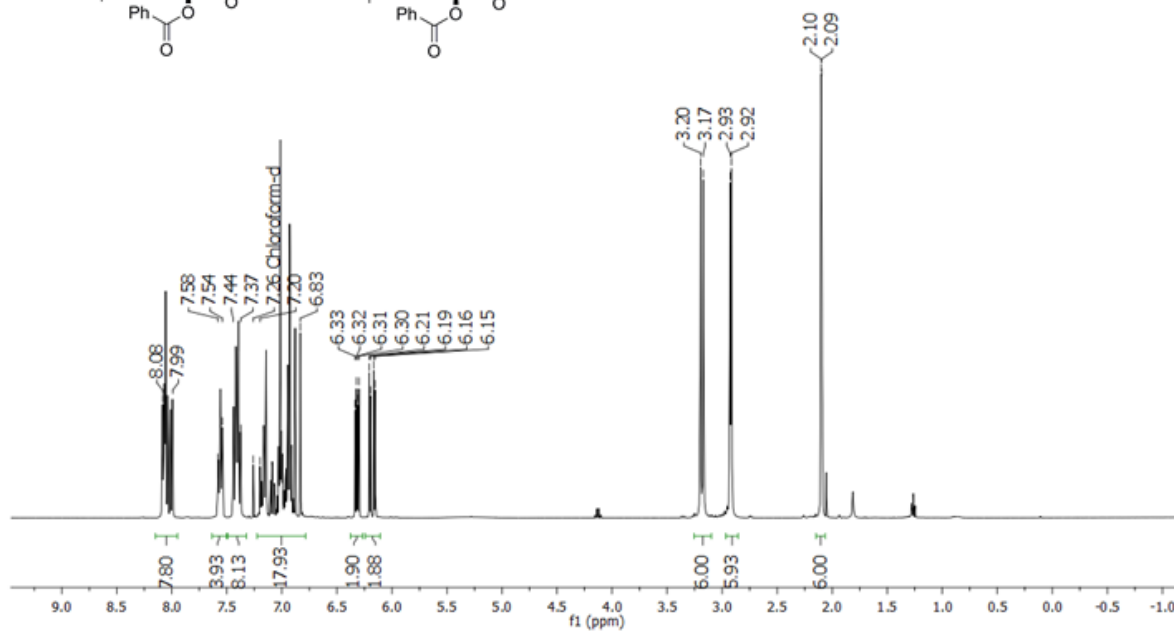
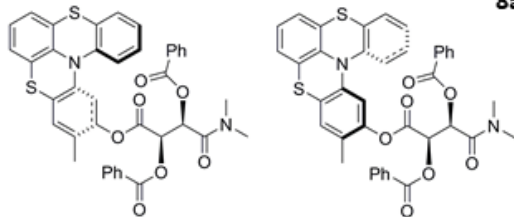


MAON53_f2_28-48_CDCl3_TEA_13C_400MHz

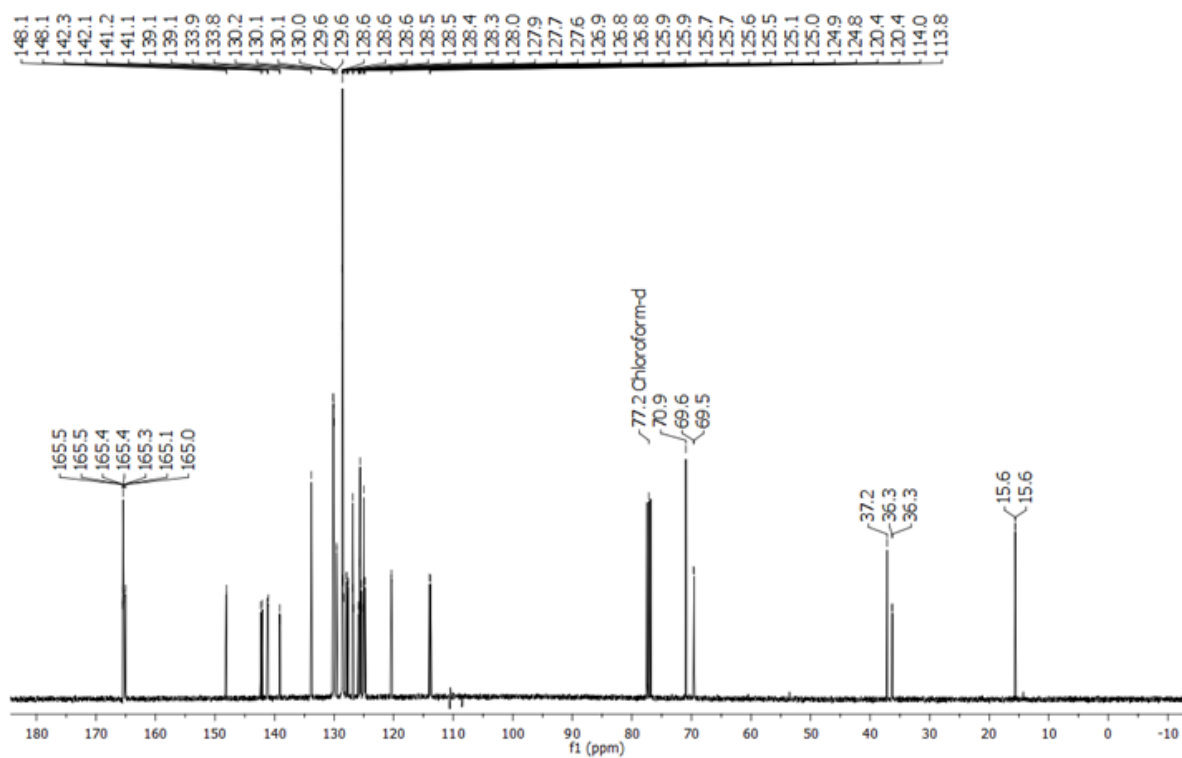


LM215_F2_CDCl3_1HNMR_400MHz

8acD1/8acD2

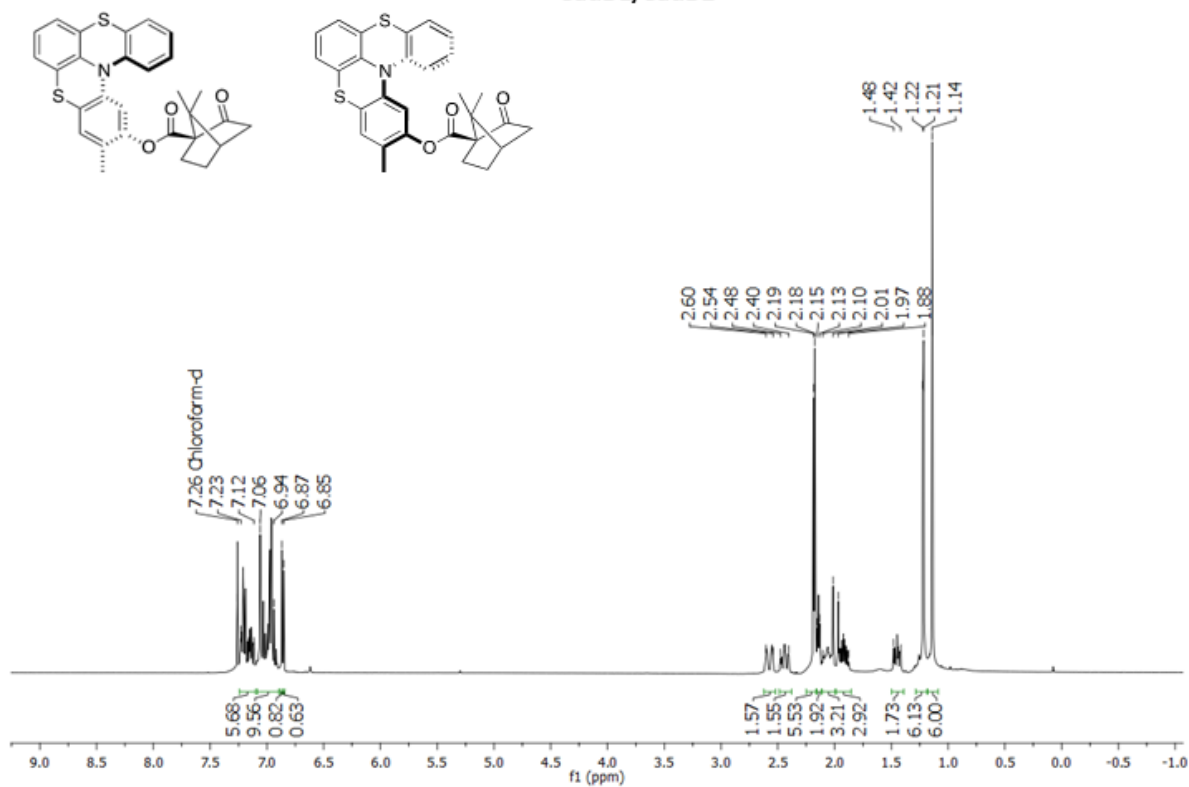


LM215_F2_CDCl3_13CNMR_100MHz

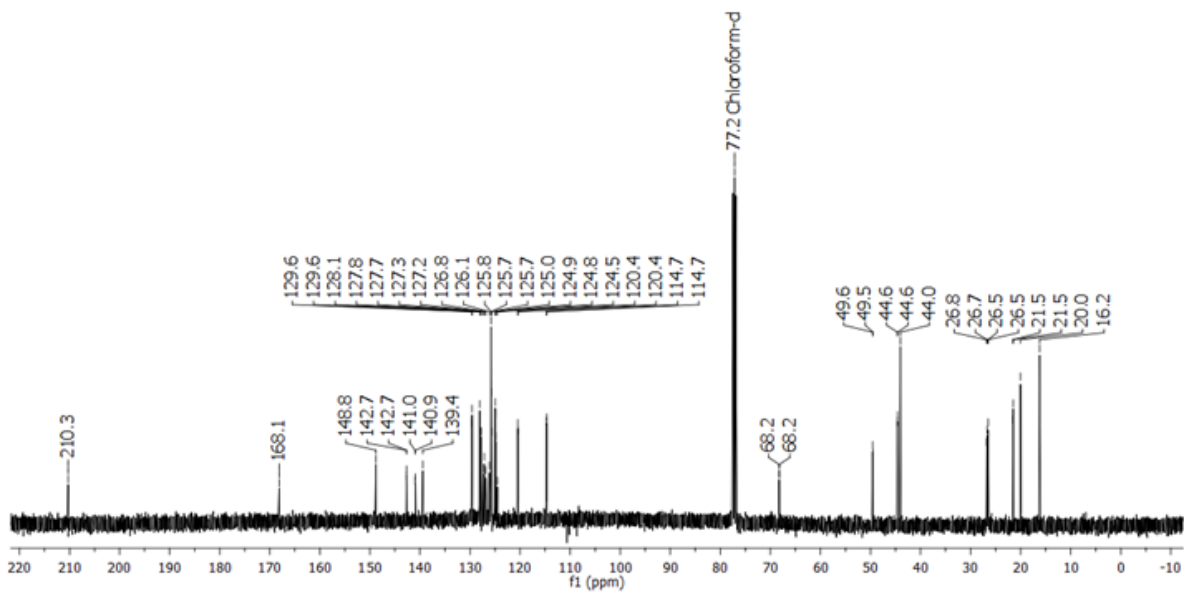


MAON68-f1_1H-CDCl3+TEA_400MHZ-INOVA

8adD1/8adD2

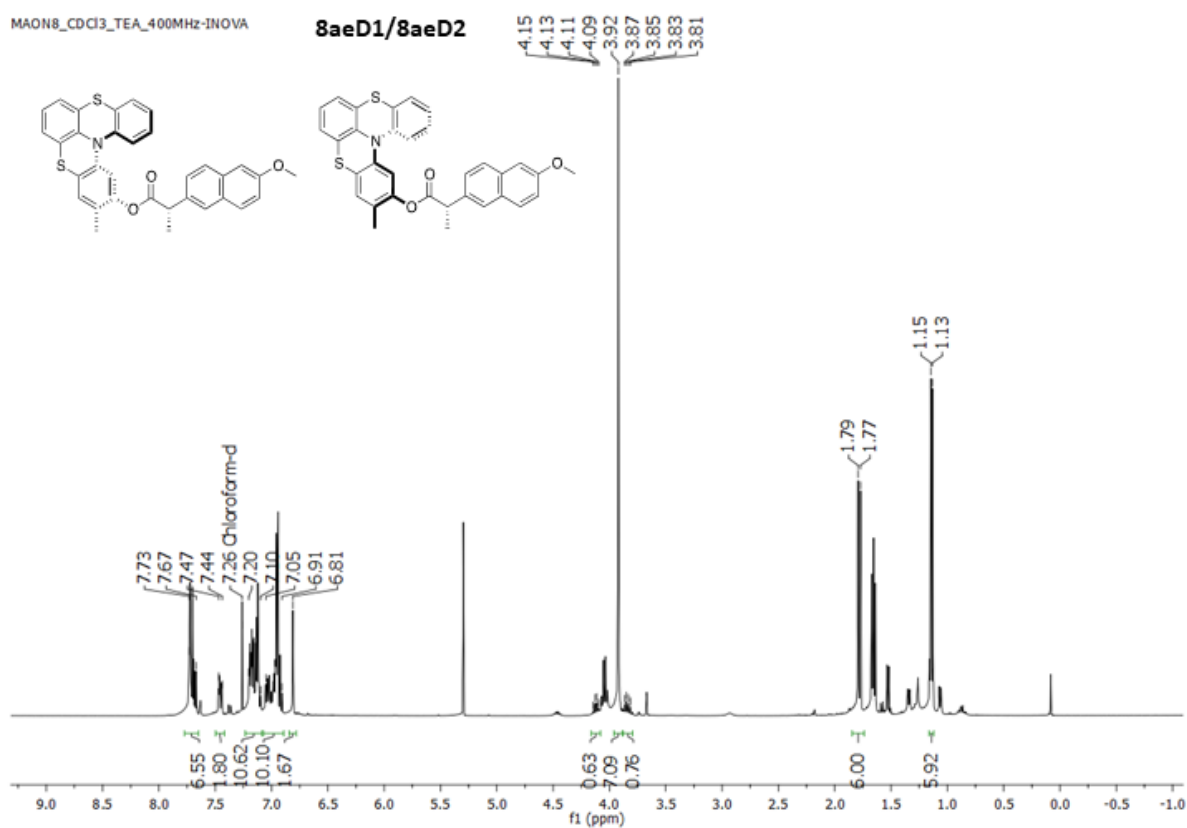


MAON68-f1_13C-CDCl3+TEA_100MHZ-INOVA

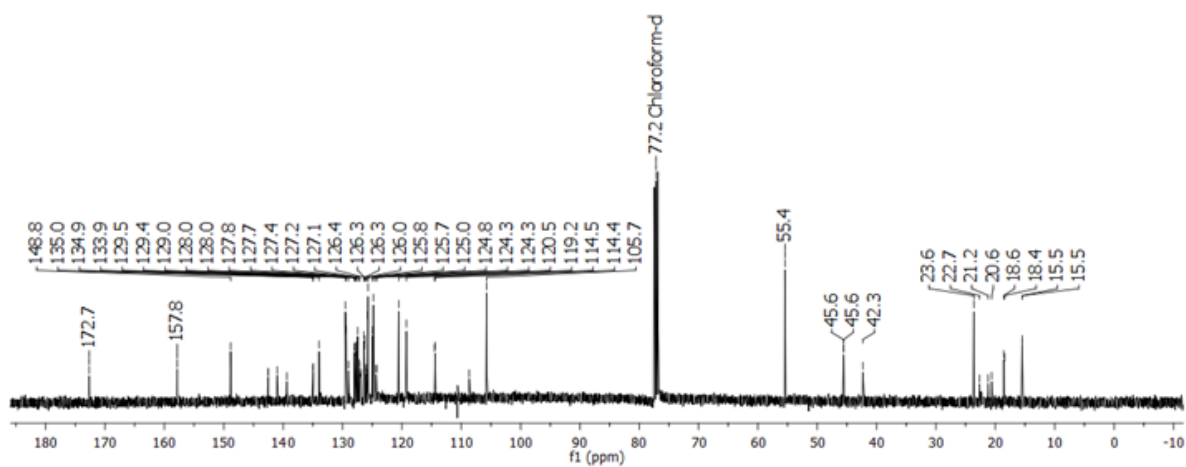


MAON8-CDCl₃-TEA_400MHz-INOVA

8aeD1/8aeD2

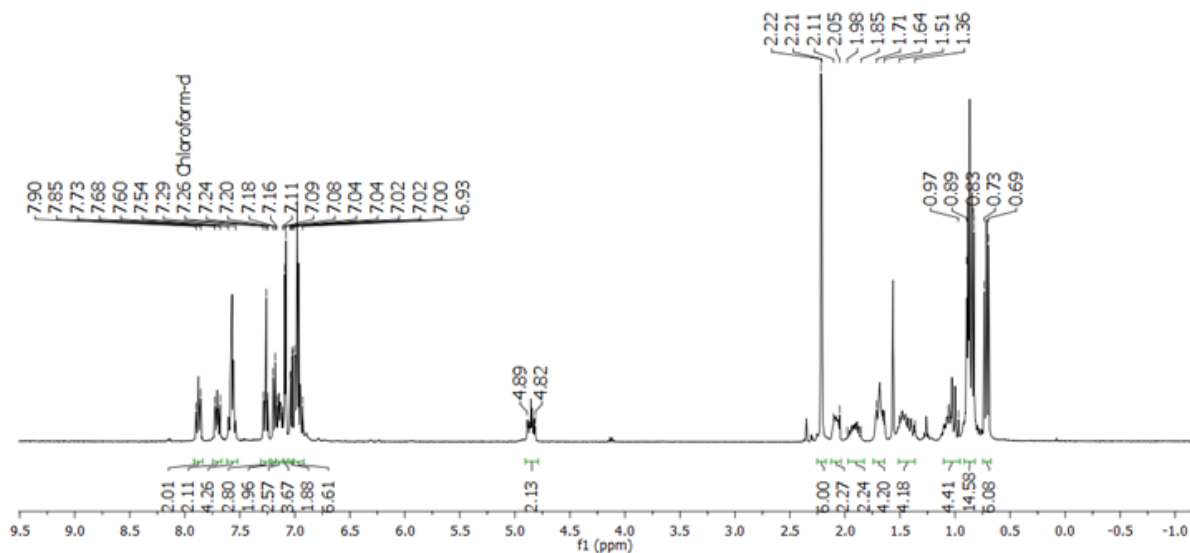
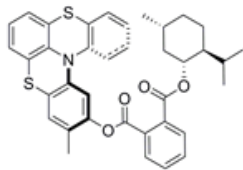
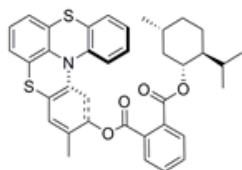


MAON8-CDCl₃+TEA_100MHz-INOVA

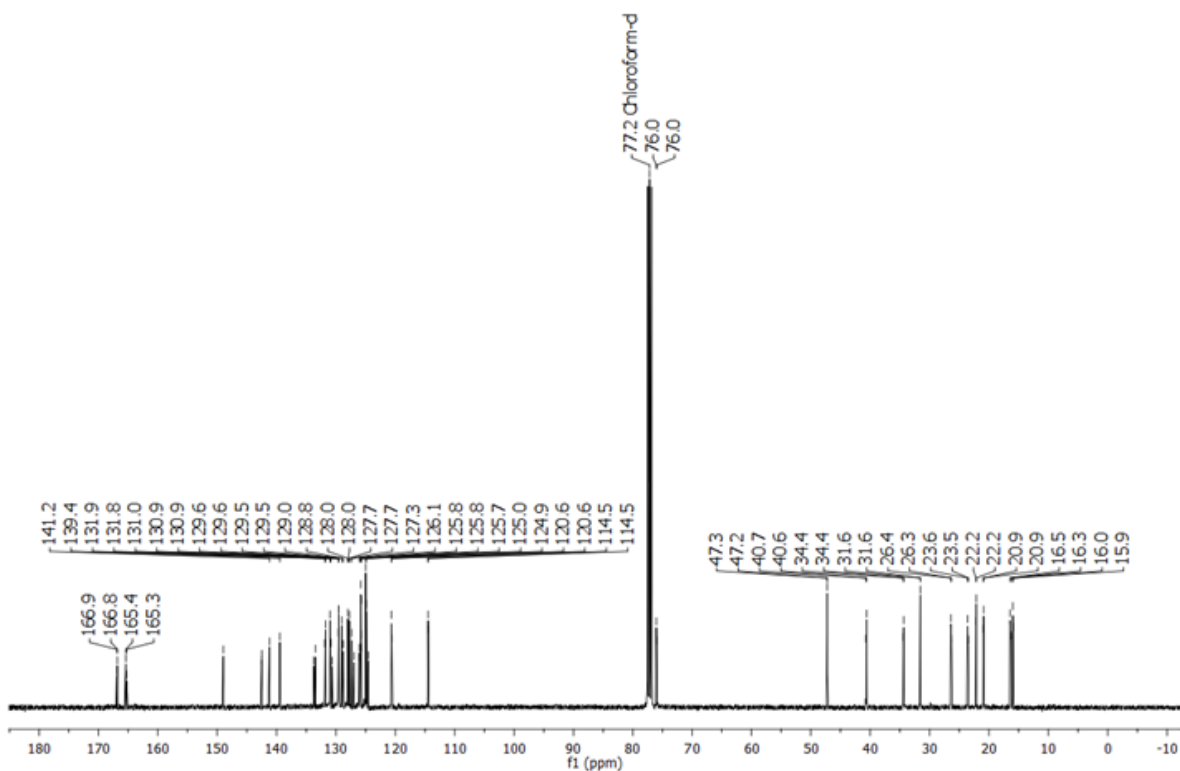


LM219-2col-F3-CDCl₃-1H NMR-400MHz

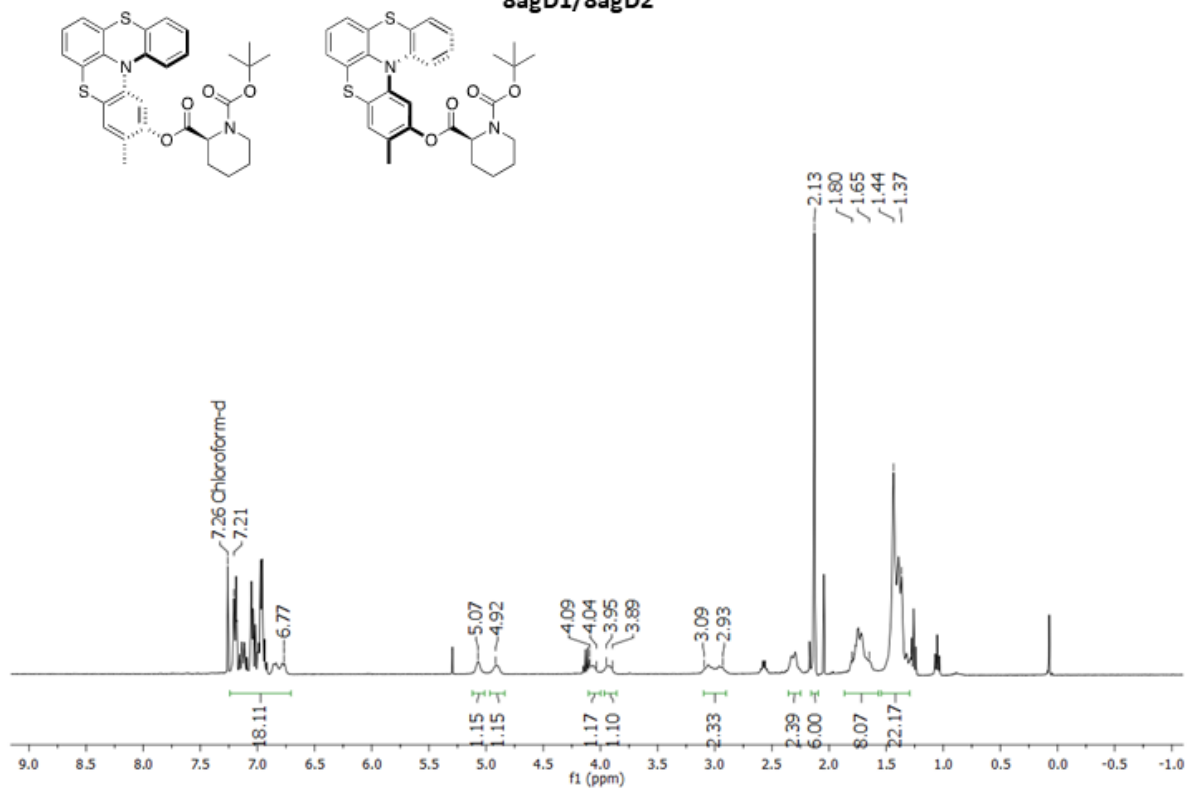
8afD1/8afD2



LM219-2col-F3-CDCl₃-¹³C NMR-100MHz

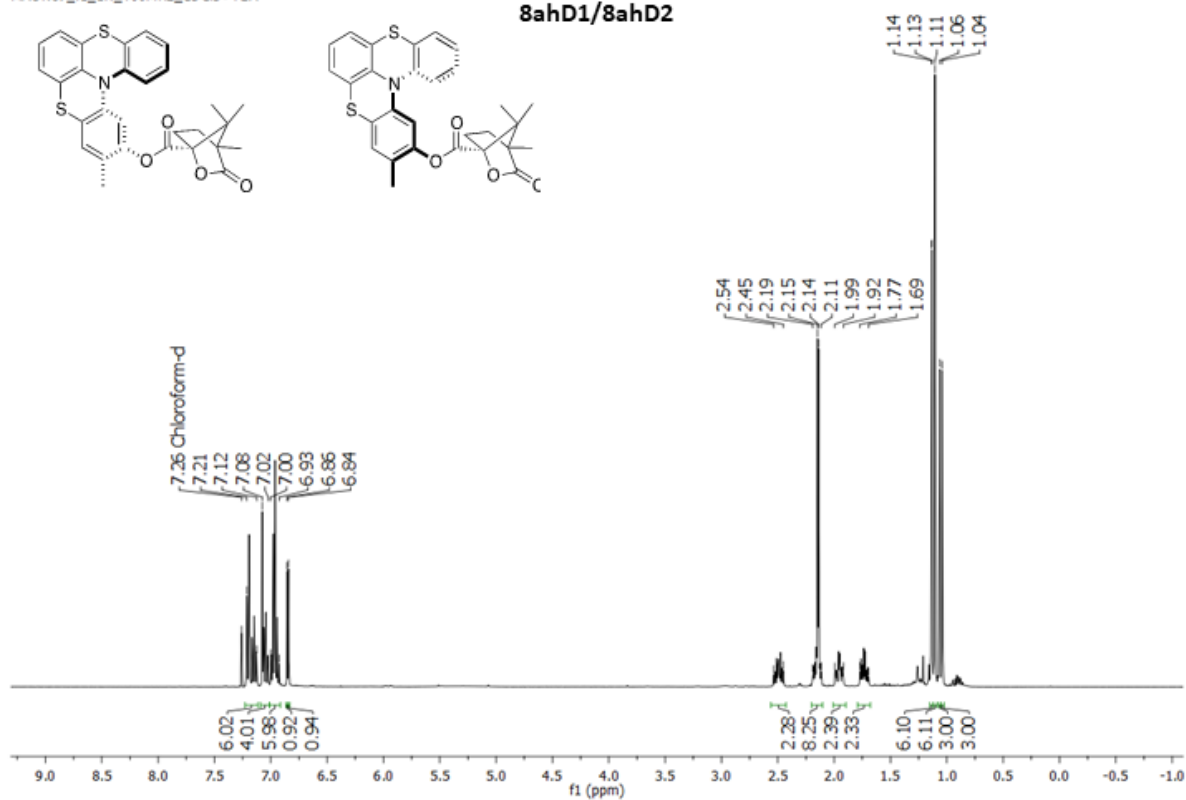


8agD1/8agD2

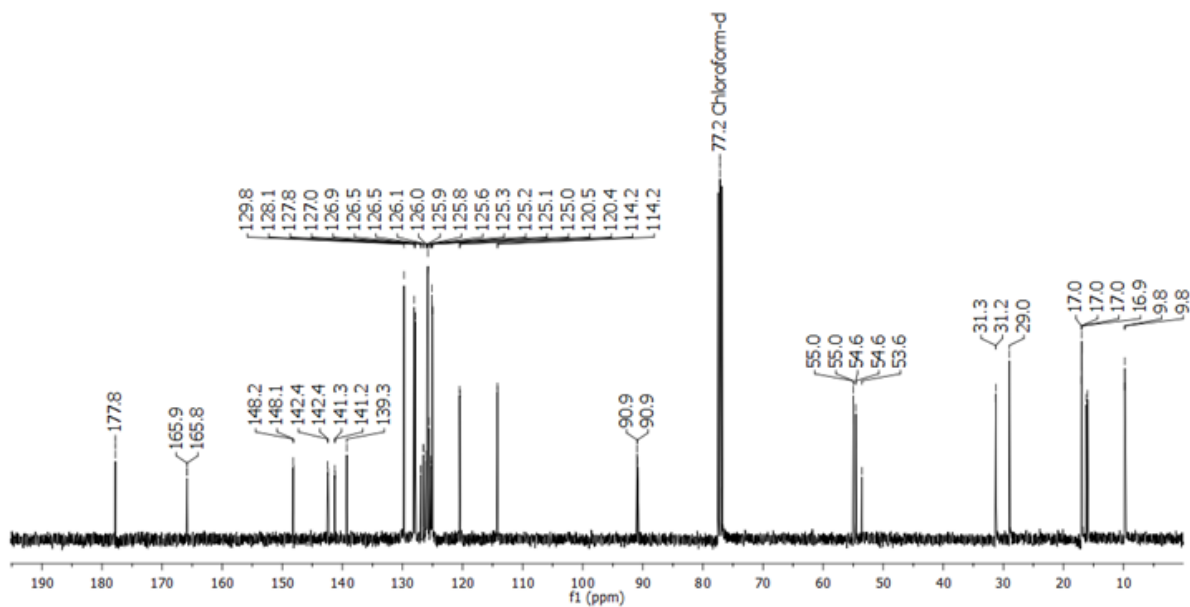


MAON87_f1_1H_400MHz_CDCl3+TEA

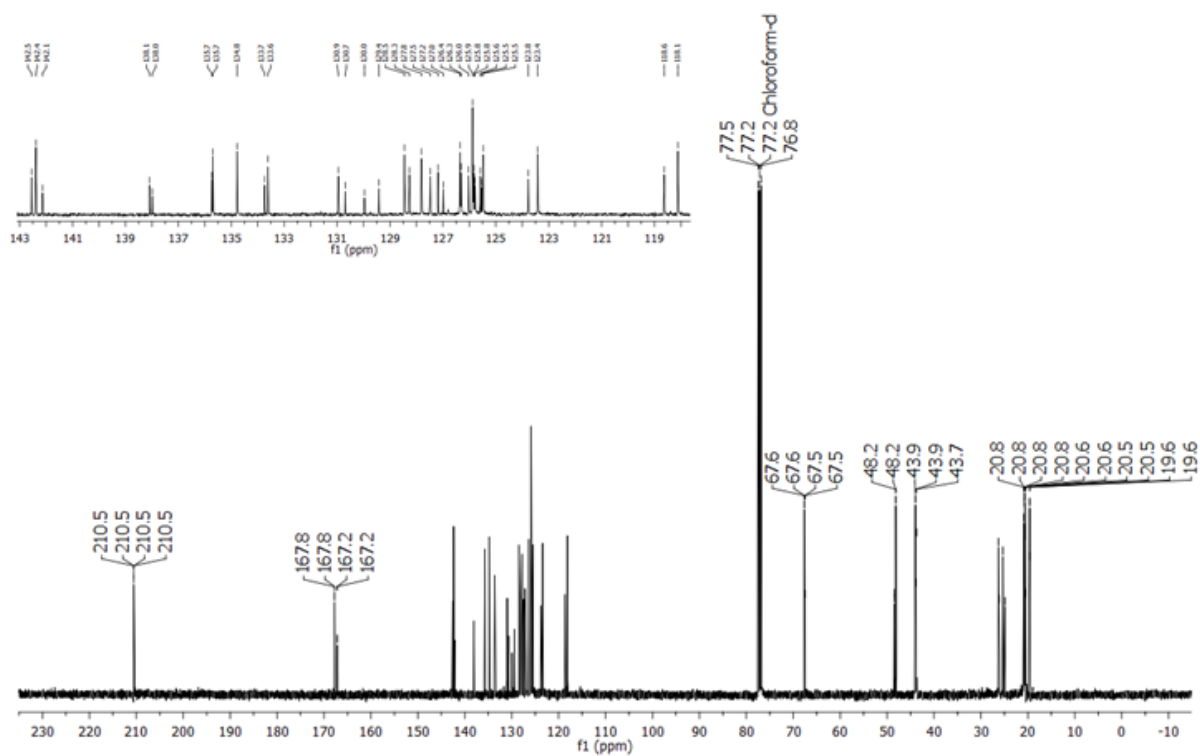
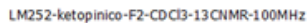
8ahD1/8ahD2



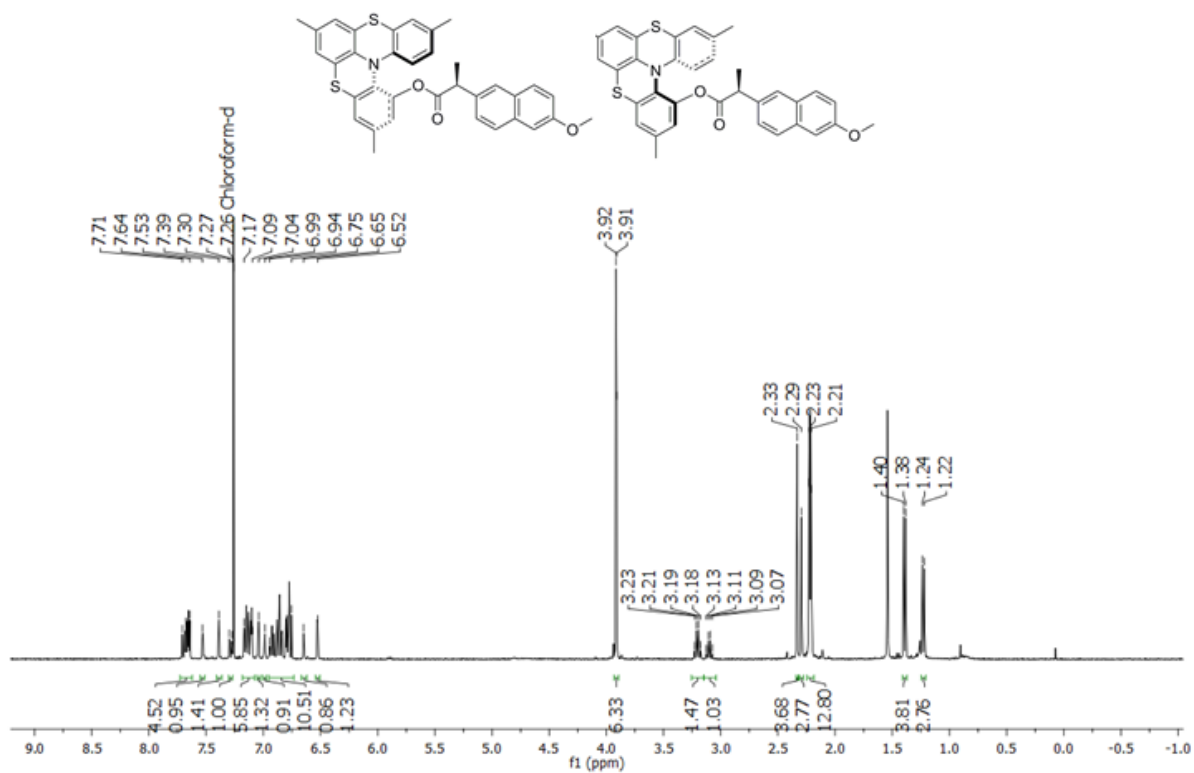
MAON87-f1-13C-100MHz-CDCl3



8bdD1/8bdD2

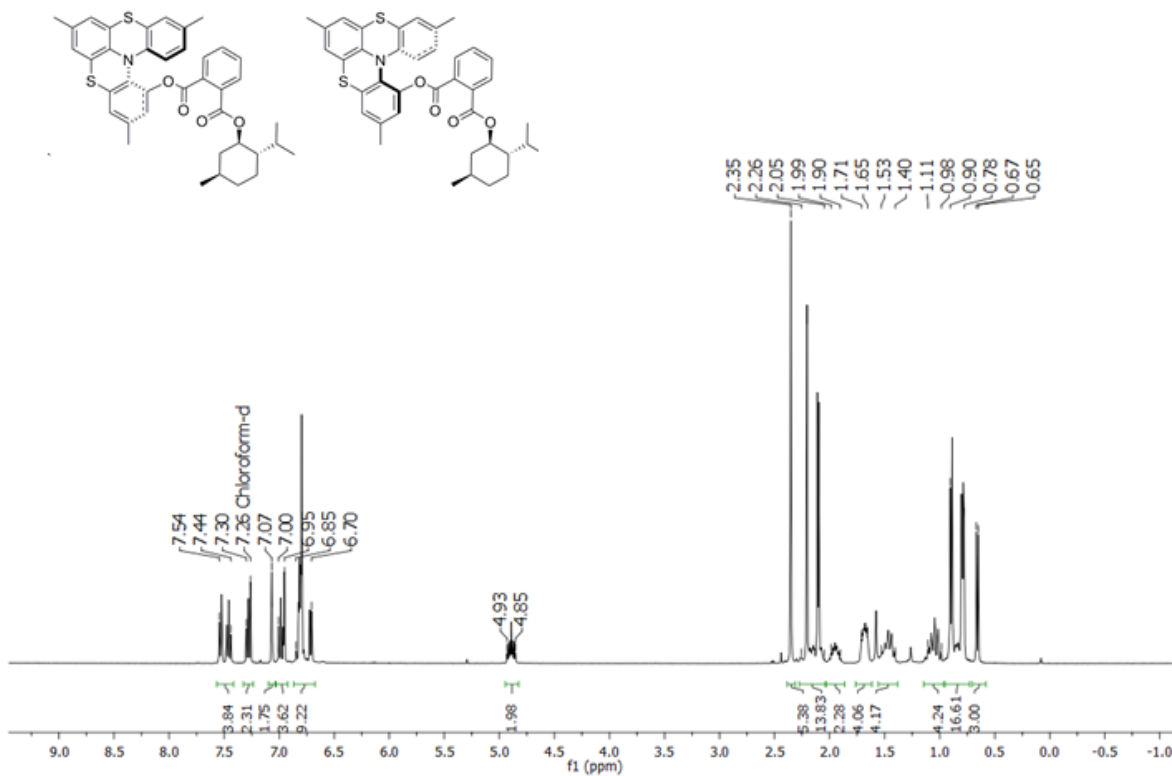


8beD1/8beD2

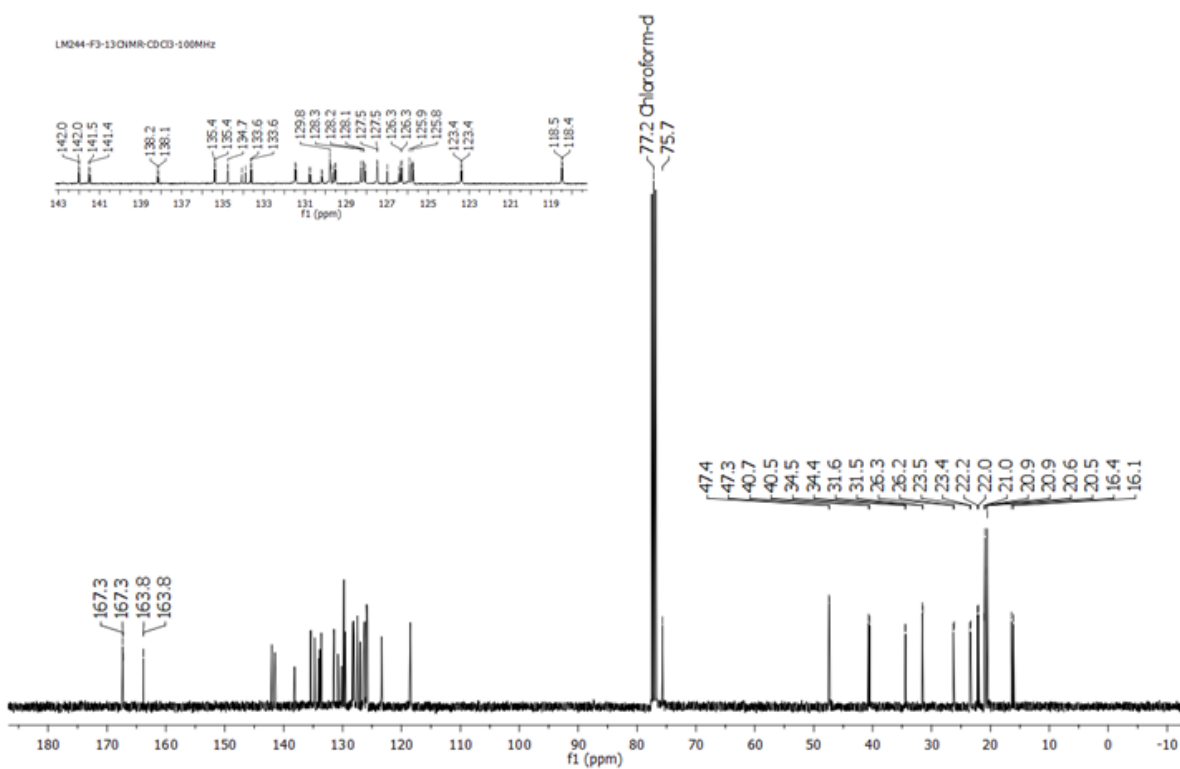


LM244-F3-1H NMR-CDCl3-400MHz

8bfD1/8bfD2

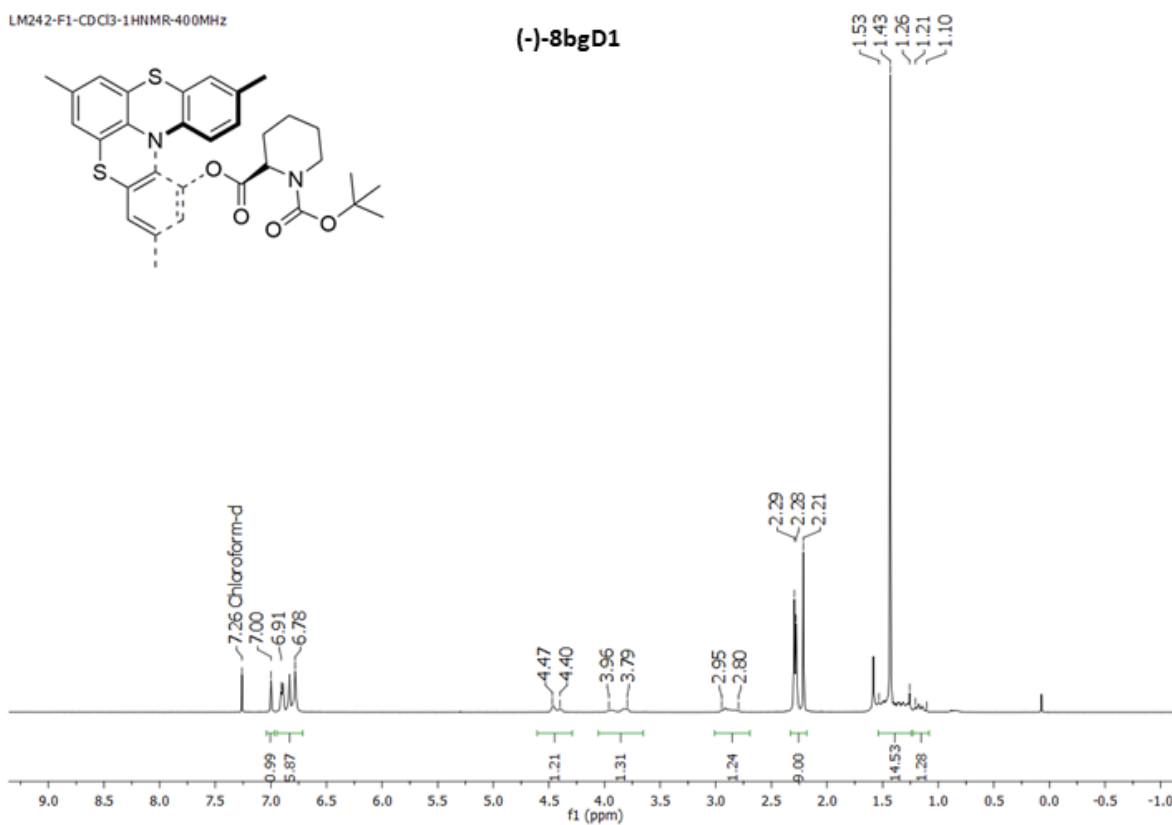


LM244-F3-13C NMR-CDCl3-100MHz

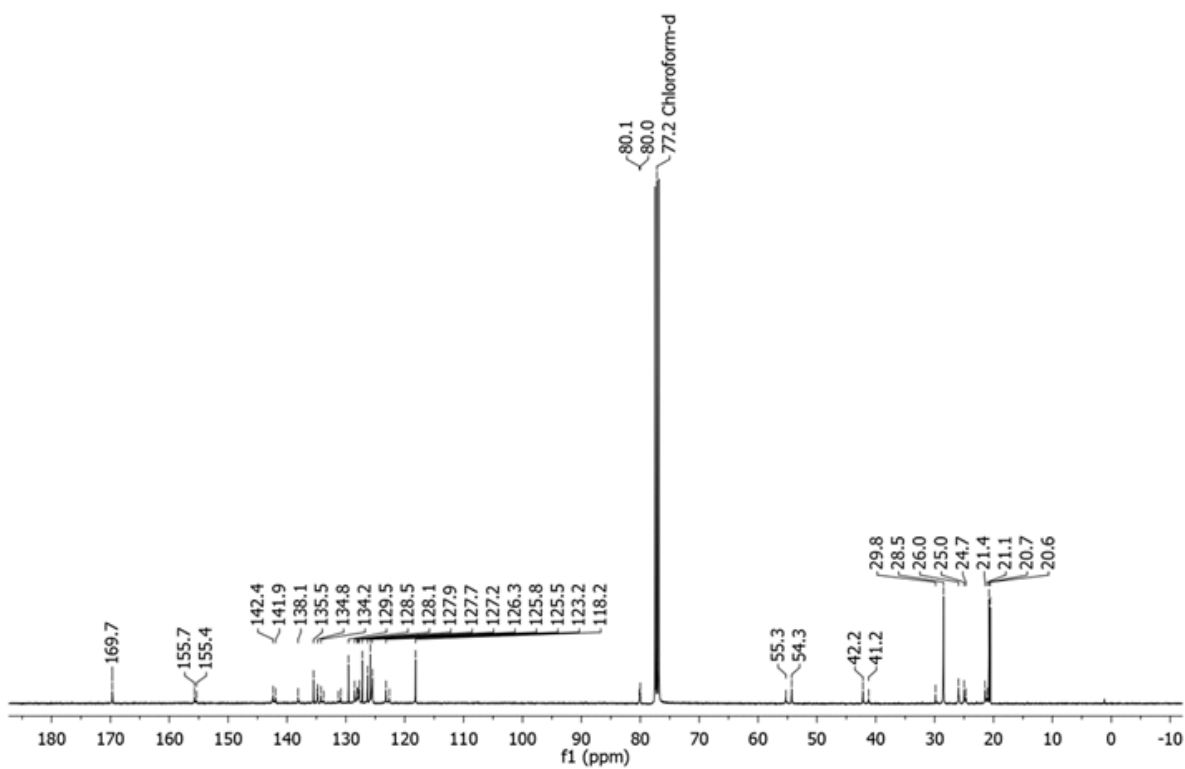


LM242-F1-CDCl3-1H NMR-400MHz

(-)-8bgD1

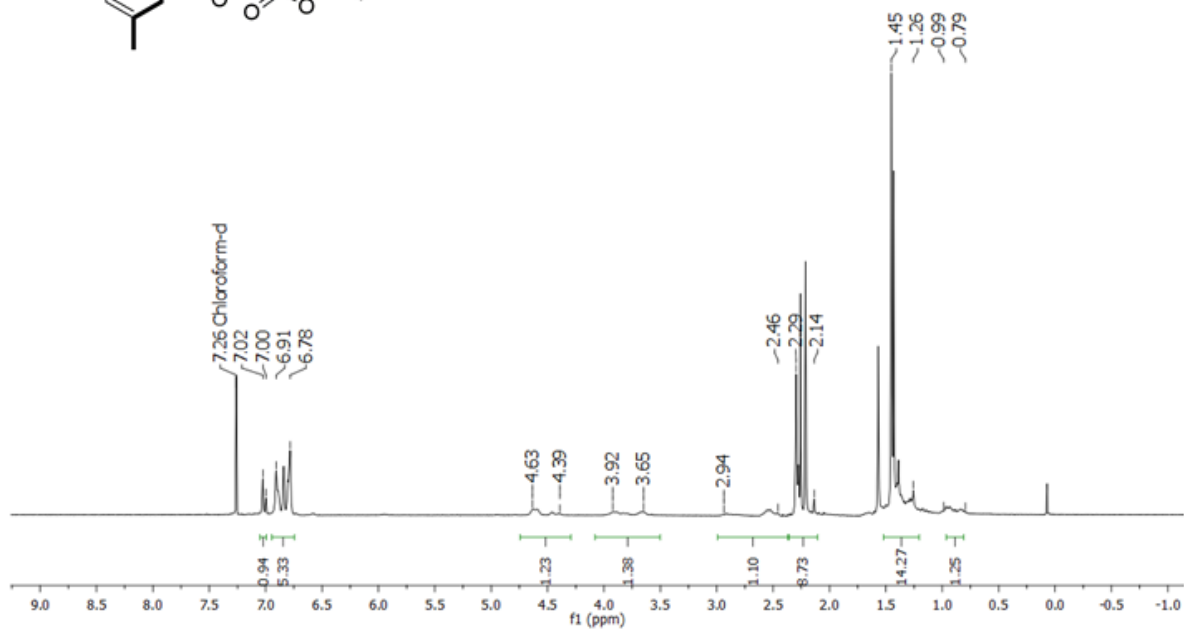
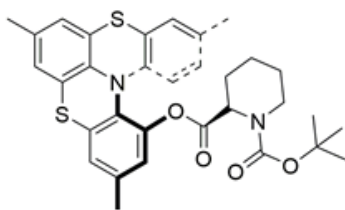


LM242-F1-CDCl3-13C NMR-100MHz-weekend

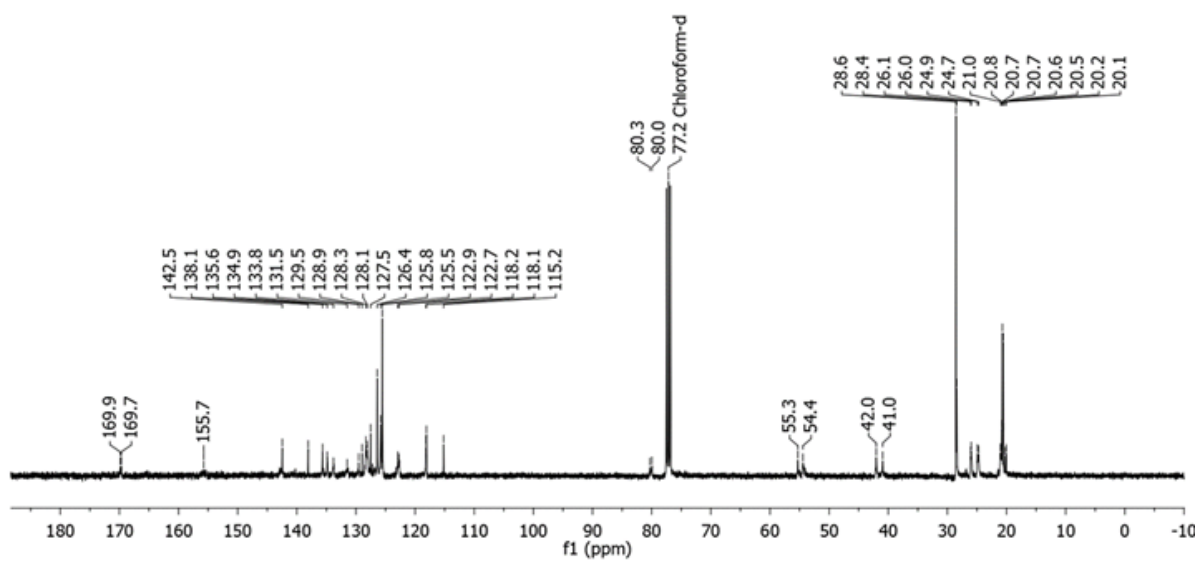


LM242-F2-CDCl3-1H NMR-400MHz

(+)-8bgD2

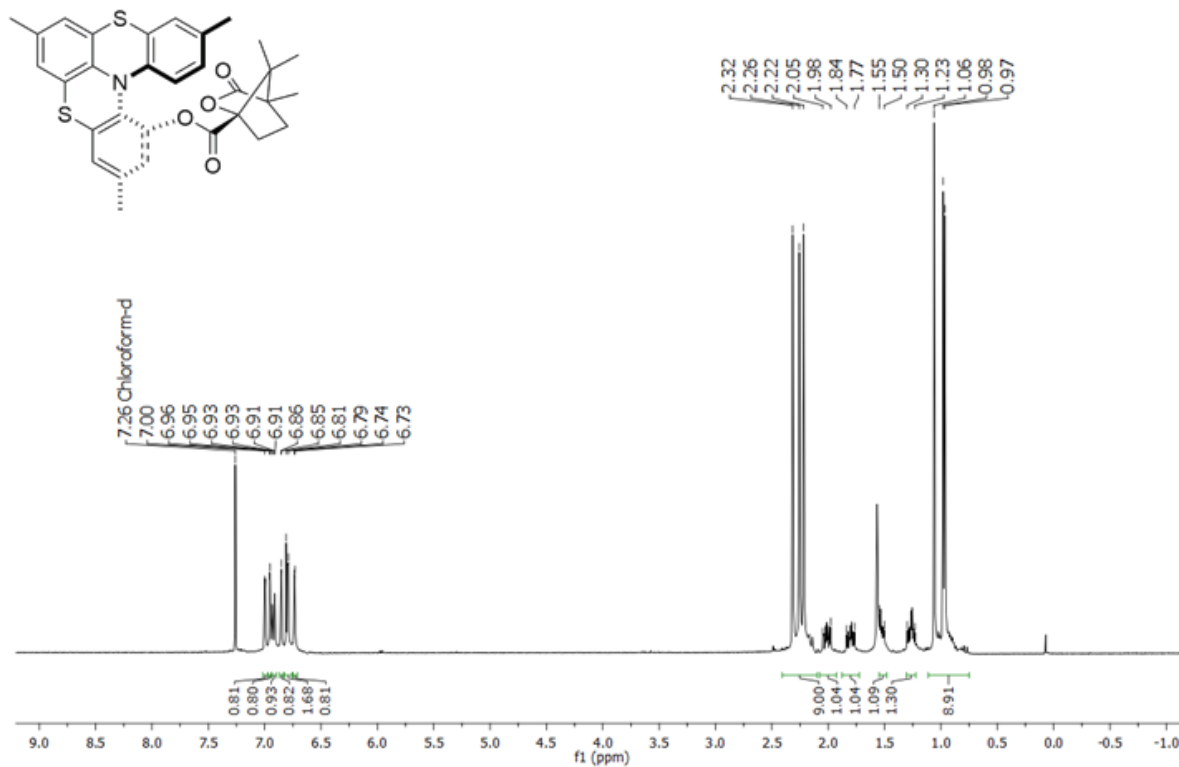


LM325_F3-CDCl3-13CNMR-100MHz

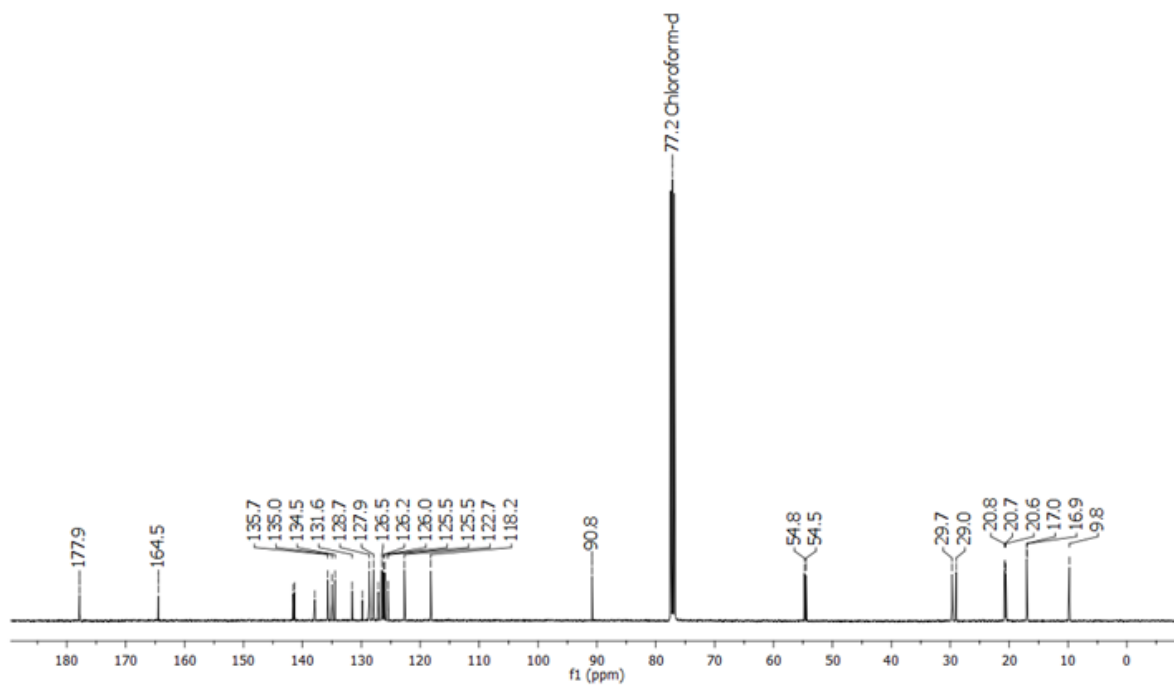


LM251-F3-CDCl₃-1H NMR-400MHz

(-)-8bHD1

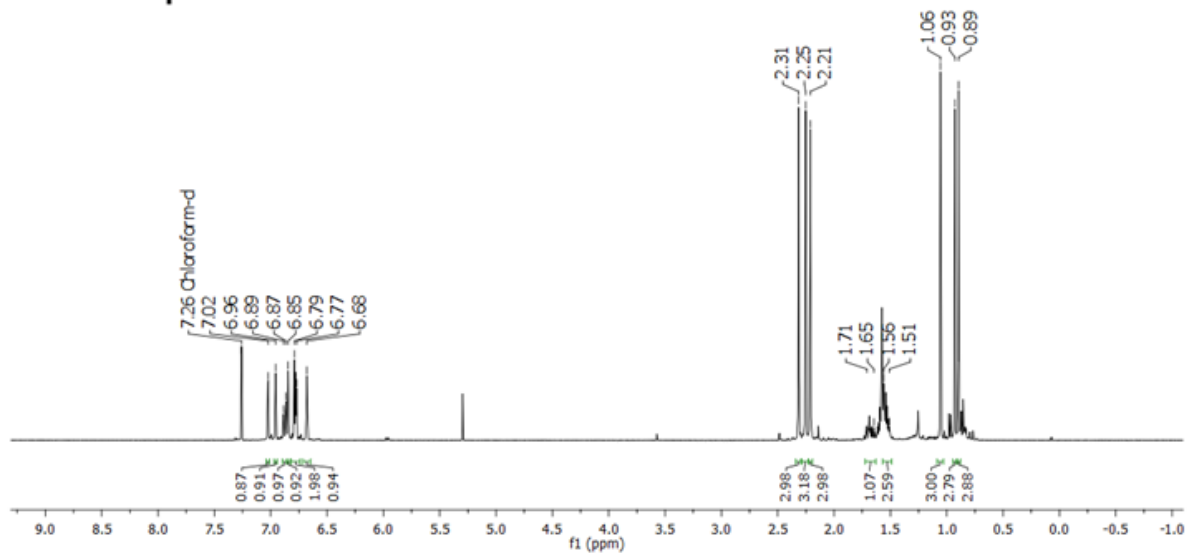
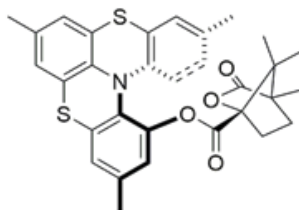


LM251-F3-CDCl₃-¹³C NMR-100MHz

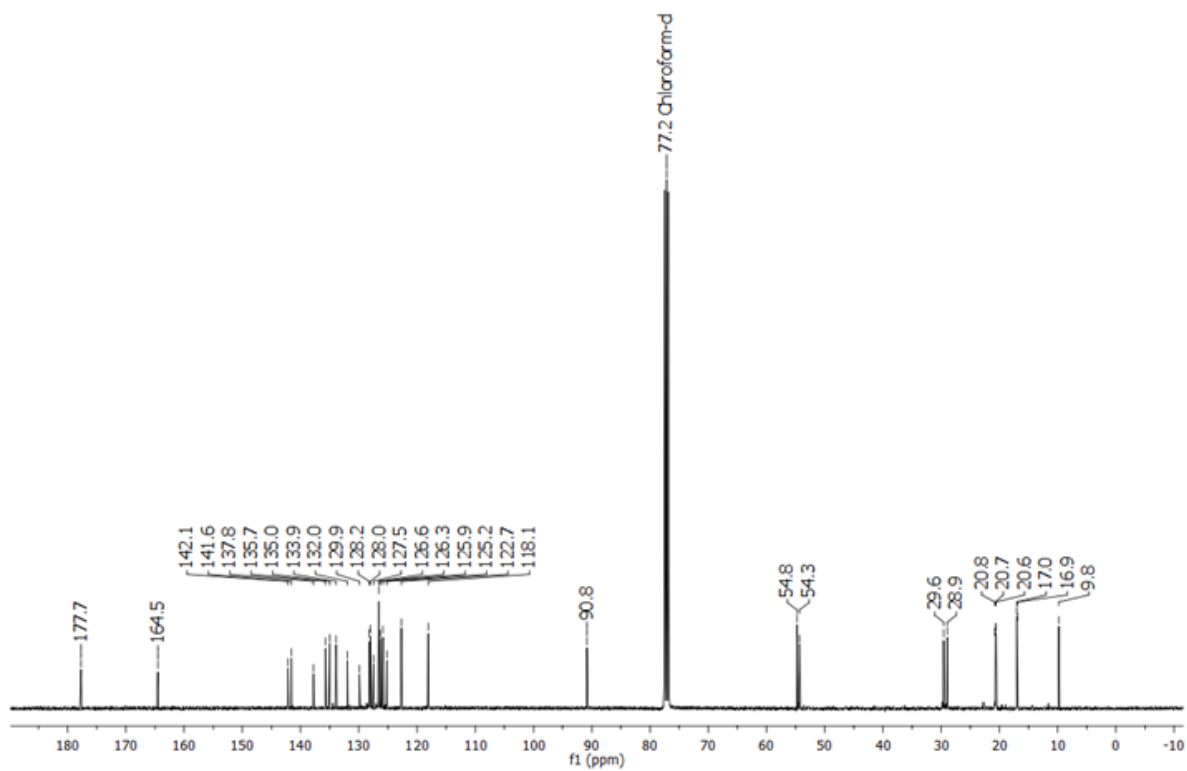


LM251-2diastero-F5-CDCl3-1HNMR-400MHz

(+)-8ahD2



LM251-2diastero-F5-CDCl3-13CNMR-100MHz



DFT calculations of compound 1b(OH)

The 3D-structures of compound **1b(OH)** in its *M* form have been optimized by DFT calculations and thereafter the CD and absorption spectra have been evaluated considering M06 and M06-2X functionals, the solvent has been treated at ieefpcm level. Two conformers have been found with energy and population values reported in the following Table.

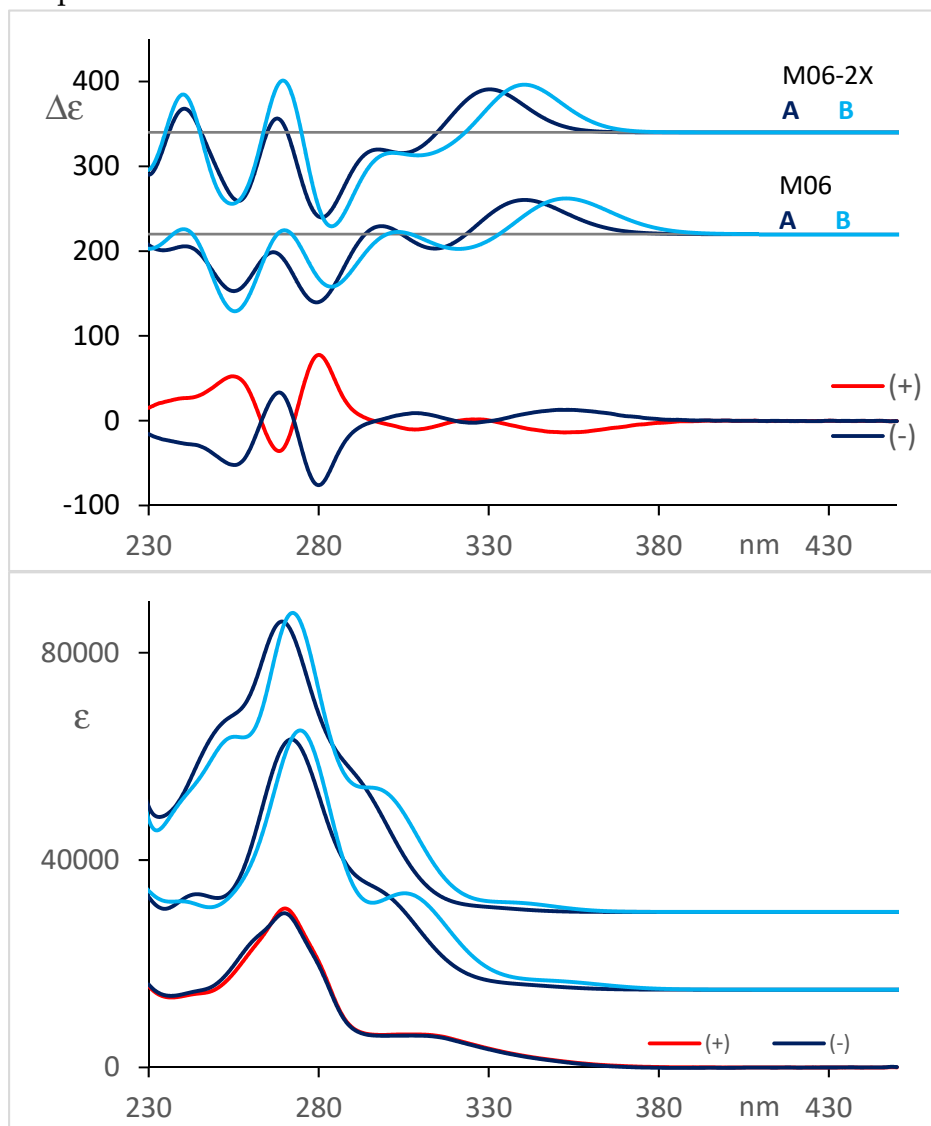
Table S1. Energy and Boltzmann population values for the two conformers reported in Figure 4 of the text

| | Kcal/mole | pop | G Kcal/mole | pop |
|--------------|-----------|-------|----------------|-------|
| M062X | | | | |
| B | 1.38 | 8.8% | 1.26 | 10.6% |
| A | 0.00 | 91.2% | 0.00 | 89.4% |
| M06 | | | | |
| B | 1.06 | 14.3% | 0.83 | 19.7% |
| A | 0.00 | 85.7% | 0.00 | 80.3% |

Spectra have been simulated as superpositions of Gaussian-shaped bands, assuming 0.2 eV-bandwidths. The calculated average spectra are reported in Figure 5 in the text; below we report the calculated spectra of each conformer. A wavelength shift of +4 nm has been applied for the M06 calculation, of +26 nm for the M06-2X calculation: the applied shift has been chosen such as to maximize the similarity index S.I. introduced by as recommended in [Kuppens, T.; Langenaeker, W.; Tollenaere, J. P.; Bultinck, P. J. Phys. Chem. A 2003, 107, 542–553]. The similarity index obtained for the M06 calculated CD spectrum of Figure 5 in the text is 0.81 (0.96 for the absorption spectrum), the similarity index for the M062X CD spectrum is 0.76 (0.98 for the absorption spectrum).

CD and absorption experimental and calculated spectra with two choices of the DFT functional, both conformers are presented considering *M*-1b(OH).

Calculated spectra are presented for the two conformers A and B.



Optimized structures' coordinates.

Conformer A M06

| Center Number | Atomic Number | Atomic Type | Coordinates (Angstroms) | | |
|------------------|------------------|----------------|-------------------------|-----------|-----------|
| | | | X | Y | Z |
| 1 | 6 | 0 | 4.150720 | -1.048962 | 0.283855 |
| 2 | 6 | 0 | 3.234180 | -1.762657 | 1.042086 |
| 3 | 6 | 0 | 1.888342 | -1.437703 | 1.028100 |
| 4 | 6 | 0 | 1.417702 | -0.423713 | 0.186894 |
| 5 | 6 | 0 | 2.347584 | 0.320986 | -0.525347 |
| 6 | 6 | 0 | 3.699366 | 0.022213 | -0.475326 |
| 7 | 6 | 0 | -0.918811 | -1.114251 | -0.184706 |
| 8 | 6 | 0 | -0.620896 | -2.190656 | -1.008074 |
| 9 | 6 | 0 | -2.207839 | -0.995979 | 0.327951 |
| 10 | 6 | 0 | -1.583581 | 1.592994 | 0.670678 |
| 11 | 6 | 0 | -0.377619 | 1.215997 | 0.097420 |
| 12 | 6 | 0 | 0.370311 | 2.174936 | -0.573569 |
| 13 | 1 | 0 | 4.402918 | 0.632165 | -1.029937 |
| 14 | 6 | 0 | -1.588767 | -3.135915 | -1.291380 |
| 15 | 6 | 0 | -3.179824 | -1.929123 | 0.009084 |
| 16 | 1 | 0 | -4.182346 | -1.804876 | 0.403627 |
| 17 | 6 | 0 | -2.881358 | -3.023988 | -0.790769 |
| 18 | 7 | 0 | 0.037050 | -0.129365 | 0.154547 |
| 19 | 6 | 0 | -2.016457 | 2.907536 | 0.613133 |
| 20 | 6 | 0 | -0.040726 | 3.497965 | -0.588630 |
| 21 | 6 | 0 | -1.235222 | 3.882688 | 0.007816 |
| 22 | 1 | 0 | -2.973674 | 3.171929 | 1.047666 |
| 23 | 1 | 0 | 0.569458 | 4.236669 | -1.096493 |
| 24 | 6 | 0 | -1.665197 | 5.314093 | 0.000799 |
| 25 | 16 | 0 | -2.567640 | 0.336456 | 1.449475 |
| 26 | 16 | 0 | 1.801616 | 1.694196 | -1.505806 |
| 27 | 6 | 0 | -3.915216 | -4.060944 | -1.090669 |
| 28 | 1 | 0 | -3.783568 | -4.473807 | -2.091100 |
| 29 | 1 | 0 | -3.851148 | -4.894823 | -0.387683 |
| 30 | 1 | 0 | -4.922994 | -3.652545 | -1.018655 |
| 31 | 6 | 0 | 5.597802 | -1.421440 | 0.297061 |
| 32 | 1 | 0 | 5.906716 | -1.781575 | 1.278591 |
| 33 | 1 | 0 | 5.799375 | -2.222695 | -0.417628 |
| 34 | 1 | 0 | 6.229431 | -0.576186 | 0.025421 |
| 35 | 1 | 0 | -1.339247 | -3.975045 | -1.930210 |
| 36 | 1 | 0 | 0.376907 | -2.284609 | -1.419016 |
| 37 | 1 | 0 | 3.559895 | -2.569173 | 1.688262 |
| 38 | 1 | 0 | -2.750502 | 5.403223 | 0.042647 |
| 39 | 1 | 0 | -1.261942 | 5.847154 | 0.864799 |
| 40 | 1 | 0 | -1.311938 | 5.831129 | -0.891275 |
| 41 | 8 | 0 | 1.061313 | -2.132833 | 1.835604 |
| 42 | 1 | 0 | 0.192224 | -1.713614 | 1.847202 |

Conformer B M06

| Center Number | Atomic Number | Atomic Type | Coordinates (Angstroms) | | |
|------------------|------------------|----------------|-------------------------|-----------|-----------|
| | | | X | Y | Z |
| 1 | 6 | 0 | 4.010156 | -1.468163 | 0.320511 |
| 2 | 6 | 0 | 2.991695 | -2.117694 | 1.006139 |
| 3 | 6 | 0 | 1.682415 | -1.672140 | 0.946212 |
| 4 | 6 | 0 | 1.344250 | -0.585596 | 0.129236 |
| 5 | 6 | 0 | 2.374404 | 0.078100 | -0.523056 |
| 6 | 6 | 0 | 3.691828 | -0.344083 | -0.424048 |
| 7 | 6 | 0 | -1.055886 | -1.007372 | -0.218435 |
| 8 | 6 | 0 | -0.905389 | -2.118249 | -1.036629 |
| 9 | 6 | 0 | -2.309698 | -0.744956 | 0.322736 |
| 10 | 6 | 0 | -1.387677 | 1.759879 | 0.641079 |
| 11 | 6 | 0 | -0.241254 | 1.251040 | 0.044396 |
| 12 | 6 | 0 | 0.623025 | 2.136544 | -0.587274 |
| 13 | 1 | 0 | 4.470123 | 0.211236 | -0.934282 |
| 14 | 6 | 0 | -1.974755 | -2.958934 | -1.278721 |
| 15 | 6 | 0 | -3.385595 | -1.572116 | 0.045624 |
| 16 | 1 | 0 | -4.357066 | -1.333402 | 0.465236 |
| 17 | 6 | 0 | -3.232560 | -2.703619 | -0.743353 |
| 18 | 7 | 0 | 0.013229 | -0.131849 | 0.050046 |
| 19 | 6 | 0 | -1.653487 | 3.119448 | 0.622012 |
| 20 | 6 | 0 | 0.381881 | 3.500480 | -0.560412 |
| 21 | 6 | 0 | -0.759912 | 4.010763 | 0.043462 |
| 22 | 1 | 0 | -2.568914 | 3.486573 | 1.072292 |
| 23 | 1 | 0 | 1.084833 | 4.172592 | -1.040039 |
| 24 | 6 | 0 | -1.013594 | 5.483379 | 0.079667 |
| 25 | 16 | 0 | -2.508405 | 0.631926 | 1.428088 |
| 26 | 16 | 0 | 2.001901 | 1.505333 | -1.508609 |
| 27 | 6 | 0 | -4.381000 | -3.626408 | -0.997352 |
| 28 | 1 | 0 | -4.321770 | -4.068501 | -1.992378 |
| 29 | 1 | 0 | -4.391635 | -4.450448 | -0.279815 |
| 30 | 1 | 0 | -5.336221 | -3.108773 | -0.910007 |
| 31 | 6 | 0 | 5.413980 | -1.974026 | 0.397241 |
| 32 | 1 | 0 | 5.679666 | -2.253961 | 1.417446 |
| 33 | 1 | 0 | 5.542668 | -2.864600 | -0.221967 |
| 34 | 1 | 0 | 6.126085 | -1.226122 | 0.050746 |
| 35 | 1 | 0 | -1.835262 | -3.826521 | -1.913448 |
| 36 | 1 | 0 | 0.063490 | -2.322357 | -1.476517 |
| 37 | 1 | 0 | 3.223617 | -2.975297 | 1.630600 |
| 38 | 1 | 0 | -2.081166 | 5.702584 | 0.091127 |
| 39 | 1 | 0 | -0.580889 | 5.933271 | 0.976213 |
| 40 | 1 | 0 | -0.569460 | 5.984562 | -0.780258 |
| 41 | 8 | 0 | 0.692097 | -2.253257 | 1.655350 |
| 42 | 1 | 0 | 1.049632 | -2.985661 | 2.166051 |

Conformer A M06-2X

| Center Number | Atomic Number | Atomic Type | Coordinates (Angstroms) | | |
|------------------|------------------|----------------|-------------------------|-----------|-----------|
| | | | X | Y | Z |
| 1 | 6 | 0 | 4.186950 | -0.915600 | 0.278361 |
| 2 | 6 | 0 | 3.291055 | -1.676816 | 1.020656 |
| 3 | 6 | 0 | 1.934737 | -1.390975 | 1.012277 |
| 4 | 6 | 0 | 1.432111 | -0.367492 | 0.197243 |
| 5 | 6 | 0 | 2.338260 | 0.417905 | -0.505006 |
| 6 | 6 | 0 | 3.701952 | 0.157605 | -0.460153 |
| 7 | 6 | 0 | -0.876373 | -1.141189 | -0.174951 |
| 8 | 6 | 0 | -0.521342 | -2.217366 | -0.980342 |
| 9 | 6 | 0 | -2.177355 | -1.065548 | 0.320433 |
| 10 | 6 | 0 | -1.654182 | 1.543694 | 0.661784 |
| 11 | 6 | 0 | -0.424374 | 1.213810 | 0.104097 |
| 12 | 6 | 0 | 0.297758 | 2.201559 | -0.559070 |
| 13 | 1 | 0 | 4.384354 | 0.796915 | -1.004886 |
| 14 | 6 | 0 | -1.451268 | -3.202400 | -1.269777 |
| 15 | 6 | 0 | -3.111006 | -2.039764 | -0.005649 |
| 16 | 1 | 0 | -4.121766 | -1.949373 | 0.372604 |
| 17 | 6 | 0 | -2.758338 | -3.131433 | -0.791924 |
| 18 | 7 | 0 | 0.039713 | -0.118334 | 0.174555 |
| 19 | 6 | 0 | -2.136571 | 2.843984 | 0.597628 |
| 20 | 6 | 0 | -0.163524 | 3.510800 | -0.577839 |
| 21 | 6 | 0 | -1.382439 | 3.849766 | 0.003143 |
| 22 | 1 | 0 | -3.107886 | 3.070462 | 1.018305 |
| 23 | 1 | 0 | 0.425150 | 4.271143 | -1.075907 |
| 24 | 6 | 0 | -1.867423 | 5.272657 | -0.009030 |
| 25 | 16 | 0 | -2.594623 | 0.253987 | 1.431082 |
| 26 | 16 | 0 | 1.750685 | 1.775331 | -1.477185 |
| 27 | 6 | 0 | -3.753629 | -4.214994 | -1.101154 |
| 28 | 1 | 0 | -3.622736 | -4.586506 | -2.116518 |
| 29 | 1 | 0 | -3.626177 | -5.060650 | -0.423224 |
| 30 | 1 | 0 | -4.774647 | -3.853700 | -0.993854 |
| 31 | 6 | 0 | 5.653311 | -1.246070 | 0.286937 |
| 32 | 1 | 0 | 5.984049 | -1.530701 | 1.284829 |
| 33 | 1 | 0 | 5.859321 | -2.085819 | -0.378715 |
| 34 | 1 | 0 | 6.248567 | -0.399359 | -0.048762 |
| 35 | 1 | 0 | -1.157921 | -4.038370 | -1.891930 |
| 36 | 1 | 0 | 0.486267 | -2.280219 | -1.369541 |
| 37 | 1 | 0 | 3.638502 | -2.489227 | 1.645442 |
| 38 | 1 | 0 | -2.953278 | 5.316575 | 0.050810 |
| 39 | 1 | 0 | -1.464699 | 5.821851 | 0.843575 |
| 40 | 1 | 0 | -1.547864 | 5.787521 | -0.913521 |
| 41 | 8 | 0 | 1.122853 | -2.137754 | 1.799773 |
| 42 | 1 | 0 | 0.243099 | -1.742075 | 1.832899 |

Conformer B M06-2X

| Center Number | Atomic Number | Atomic Type | Coordinates (Angstroms) | | |
|------------------|------------------|----------------|-------------------------|-----------|-----------|
| | | | X | Y | Z |
| 1 | 6 | 0 | 4.069110 | -1.319315 | 0.312972 |
| 2 | 6 | 0 | 3.071929 | -2.025415 | 0.980000 |
| 3 | 6 | 0 | 1.746727 | -1.622399 | 0.926189 |
| 4 | 6 | 0 | 1.371300 | -0.524122 | 0.138196 |
| 5 | 6 | 0 | 2.377030 | 0.190210 | -0.501834 |
| 6 | 6 | 0 | 3.711577 | -0.188965 | -0.407832 |
| 7 | 6 | 0 | -1.008667 | -1.046466 | -0.207472 |
| 8 | 6 | 0 | -0.796471 | -2.158567 | -1.013774 |
| 9 | 6 | 0 | -2.280486 | -0.831883 | 0.318703 |
| 10 | 6 | 0 | -1.470738 | 1.707925 | 0.640948 |
| 11 | 6 | 0 | -0.294310 | 1.250798 | 0.054795 |
| 12 | 6 | 0 | 0.541416 | 2.174739 | -0.566352 |
| 13 | 1 | 0 | 4.468696 | 0.402915 | -0.905263 |
| 14 | 6 | 0 | -1.829479 | -3.046314 | -1.265402 |
| 15 | 6 | 0 | -3.319141 | -1.706607 | 0.030885 |
| 16 | 1 | 0 | -4.303813 | -1.506465 | 0.435660 |
| 17 | 6 | 0 | -3.106976 | -2.837530 | -0.750459 |
| 18 | 7 | 0 | 0.020691 | -0.122718 | 0.072204 |
| 19 | 6 | 0 | -1.794181 | 3.058085 | 0.622409 |
| 20 | 6 | 0 | 0.241961 | 3.529919 | -0.535759 |
| 21 | 6 | 0 | -0.932371 | 3.989020 | 0.052952 |
| 22 | 1 | 0 | -2.724945 | 3.384408 | 1.069022 |
| 23 | 1 | 0 | 0.925407 | 4.232432 | -0.996307 |
| 24 | 6 | 0 | -1.279096 | 5.452080 | 0.035282 |
| 25 | 16 | 0 | -2.540469 | 0.532217 | 1.421334 |
| 26 | 16 | 0 | 1.951958 | 1.607687 | -1.473891 |
| 27 | 6 | 0 | -4.220582 | -3.813363 | -1.012908 |
| 28 | 1 | 0 | -4.094678 | -4.299243 | -1.979245 |
| 29 | 1 | 0 | -4.238923 | -4.594654 | -0.250914 |
| 30 | 1 | 0 | -5.189654 | -3.317223 | -0.999434 |
| 31 | 6 | 0 | 5.498870 | -1.777675 | 0.382302 |
| 32 | 1 | 0 | 5.773529 | -2.047055 | 1.401619 |
| 33 | 1 | 0 | 5.649980 | -2.660086 | -0.241341 |
| 34 | 1 | 0 | 6.176486 | -1.000624 | 0.034957 |
| 35 | 1 | 0 | -1.642469 | -3.912038 | -1.888156 |
| 36 | 1 | 0 | 0.186860 | -2.327364 | -1.433107 |
| 37 | 1 | 0 | 3.331245 | -2.890654 | 1.579863 |
| 38 | 1 | 0 | -1.929394 | 5.710273 | 0.869039 |
| 39 | 1 | 0 | -0.382538 | 6.067124 | 0.091510 |
| 40 | 1 | 0 | -1.801331 | 5.712562 | -0.886818 |
| 41 | 8 | 0 | 0.767278 | -2.258482 | 1.614733 |
| 42 | 1 | 0 | 1.144116 | -2.990827 | 2.113475 |