

Identifying cysteine residues susceptible to oxidation by photoactivatable atomic oxygen precursors using a proteome-wide analysis.

Ankita Isor^a, Benjamin V. Chartier^b, Masahiro Abo^b, Emily R. Currens^a, Eranthie Weerapana^b, Ryan D. McCulla^{*a}

^a Department of Chemistry, Saint Louis University, 3501 Laclede Ave, Saint Louis, MO 63103, United States

^b Department of Chemistry, Boston College, Chestnut Hill, MA 02467, United States

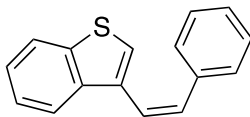
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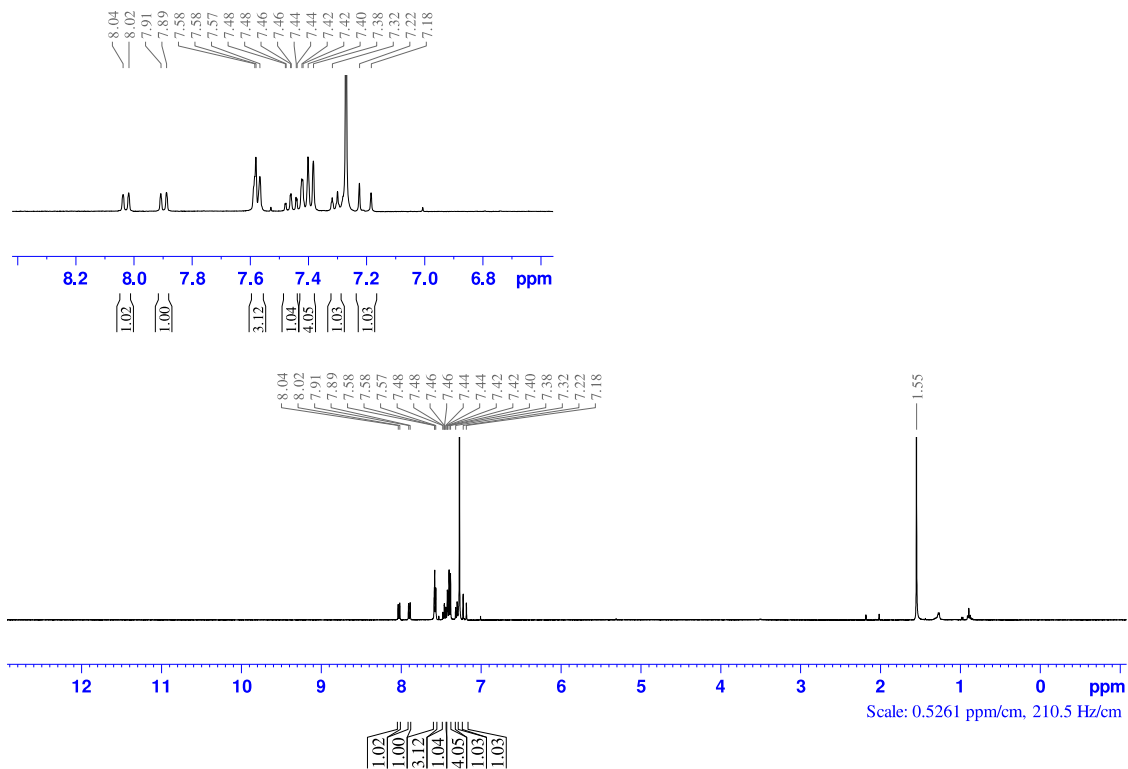
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1. Spectral Data (^1H -NMR, ^{13}C -NMR, HRMS, HPLC Analysis for purity)

3-styrylbenzo[b]thiophene

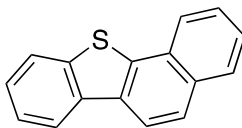


^1H -NMR (in CDCl_3)

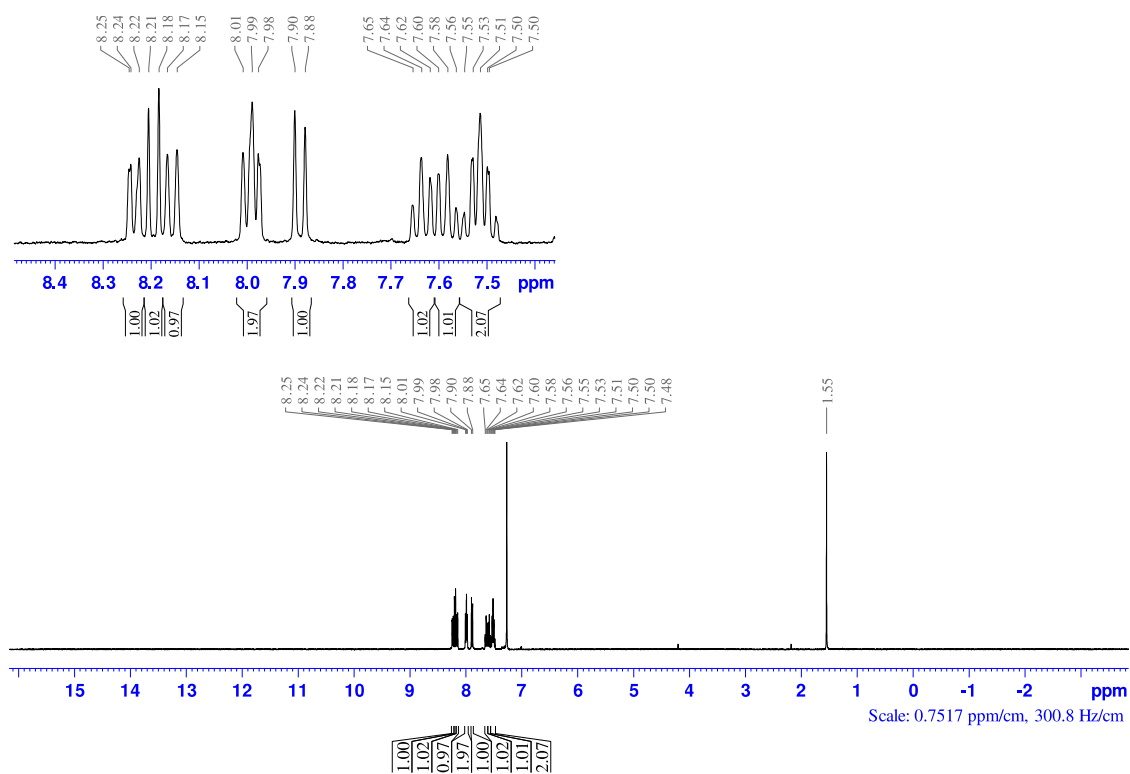


Impurities: Solvent Peak Chloroform (7.27 ppm), Water (1.55 ppm)

Benzo[b]naphtho-[1,2-d]-thiophene

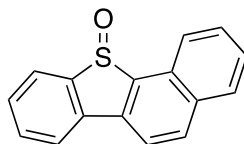


^1H -NMR (in CDCl_3)

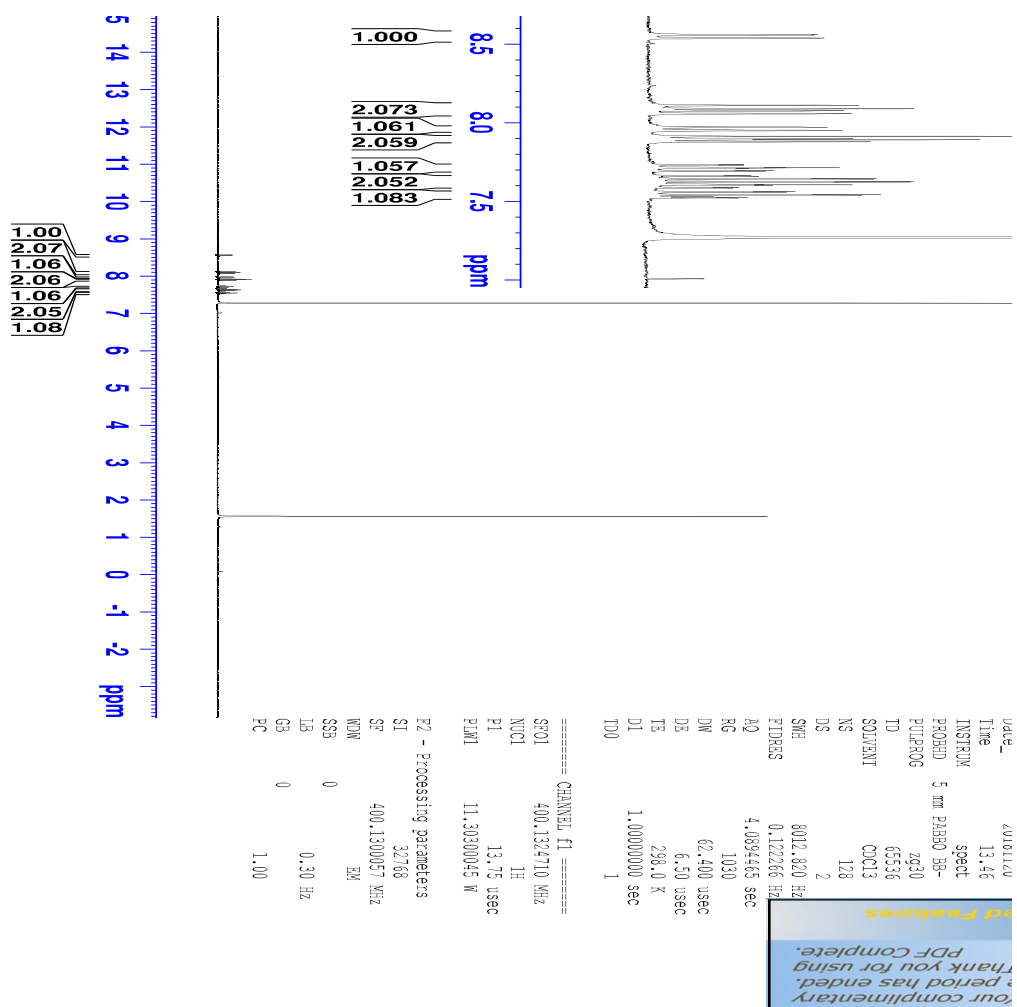


Impurities: Solvent Peak Chloroform (7.27 ppm), Water (1.56 ppm)

Benzo[b]naphtho-[1,2-d]thiophene-S-oxide

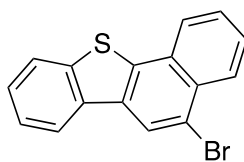


^1H -NMR in CDCl_3

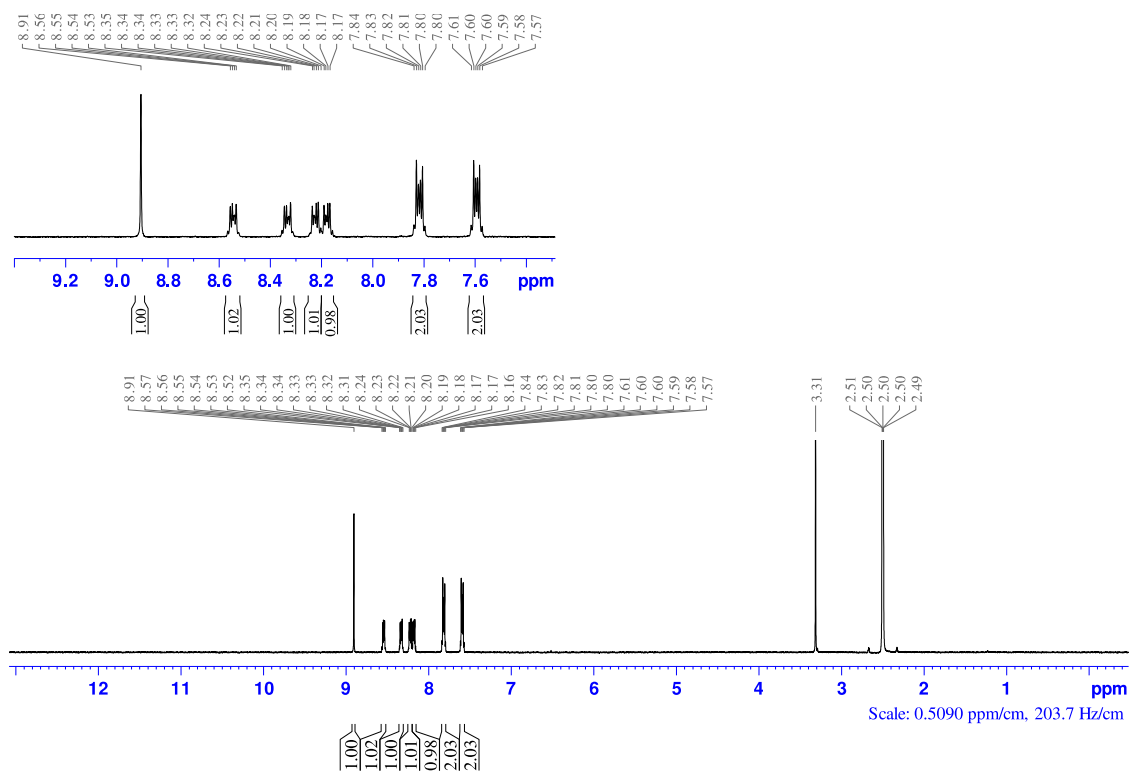


Impurities: Solvent Peak Chloroform (7.27 ppm), Water (1.55 ppm)

5-Bromobenzo[b]naphtho-[1,2-d]thiophene

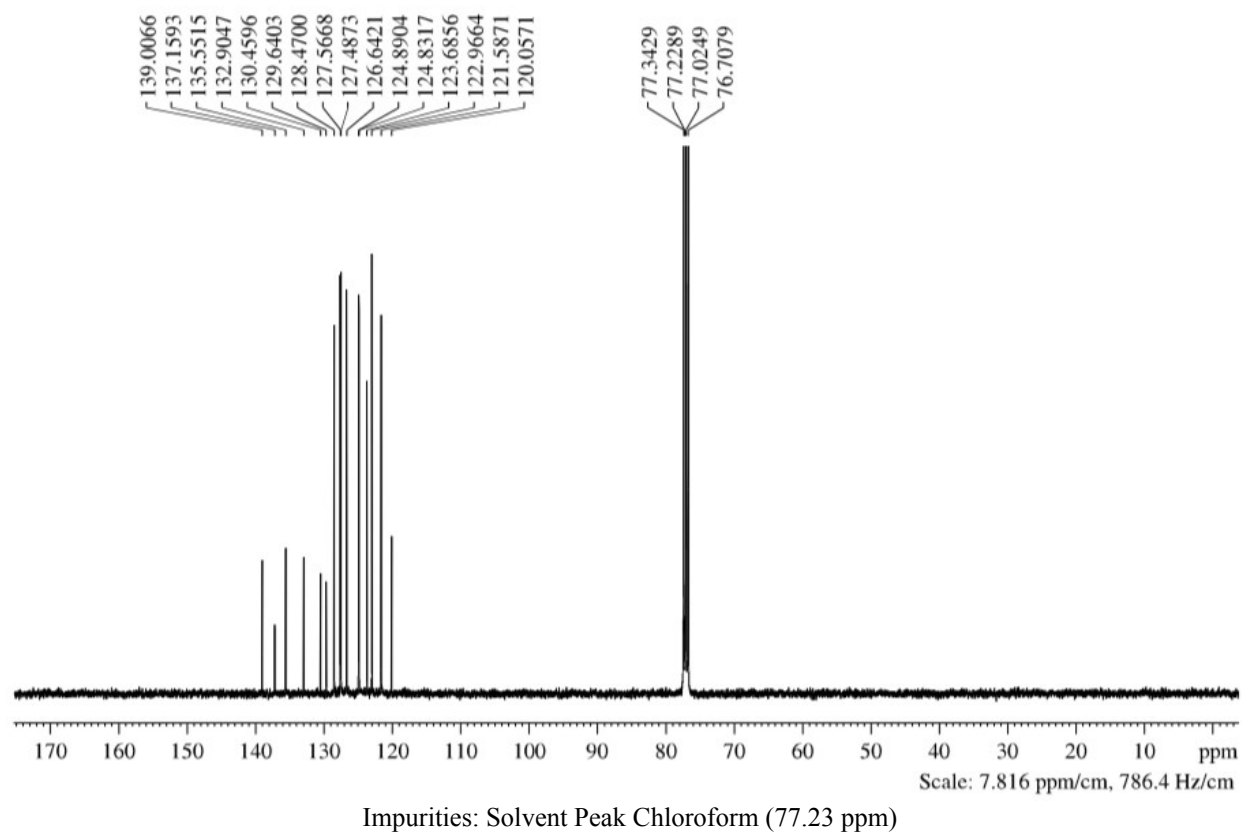


^1H -NMR (in DMSO)

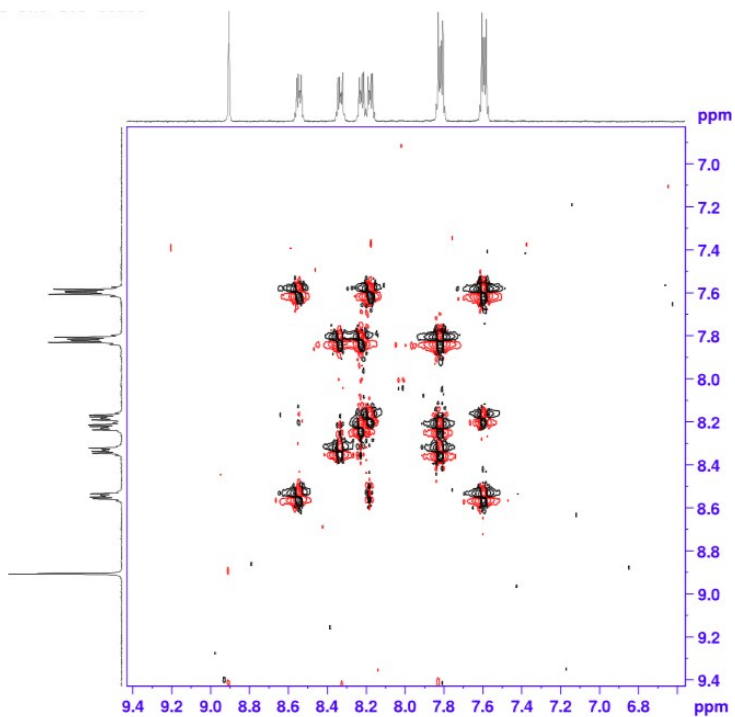


Impurities: Solvent Peak DMSO (2.50 ppm), water (3.31 ppm)

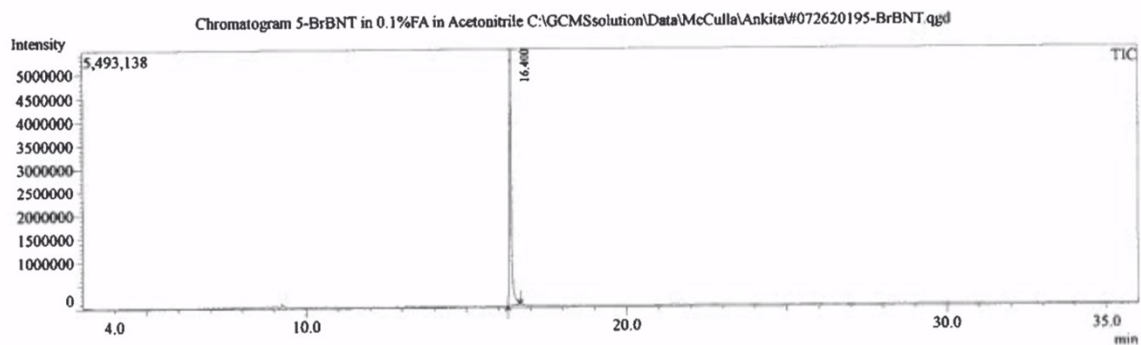
^{13}C -NMR (in CDCl_3)



$\text{COSY } ^1\text{H}$ -NMR (in CDCl_3)

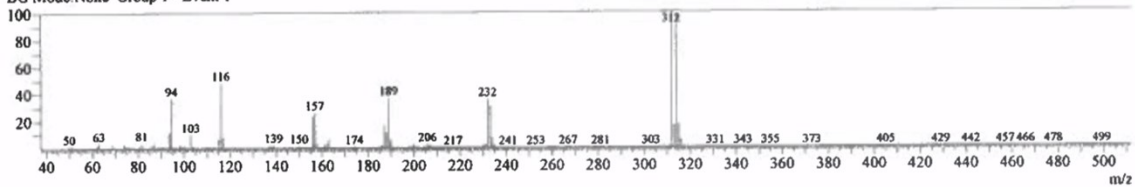


GCMS

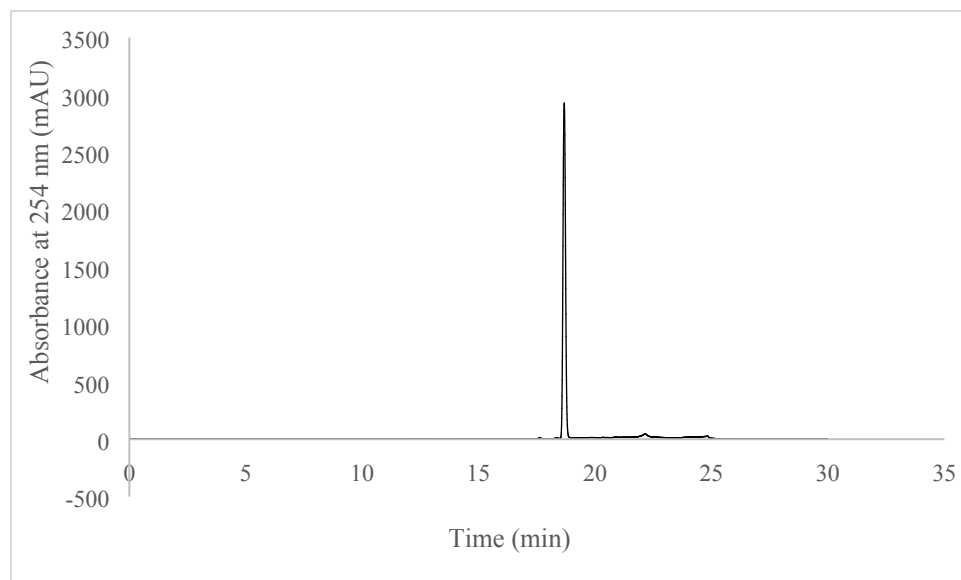


Peak Report TIC							
Peak#	R.Time	I.Time	F.Time	Area	Area%	Height	Height%
1	16.400	16.287	16.697	22711315	100.00	5450028	100.00
				22711315	100.00	5450028	100.00

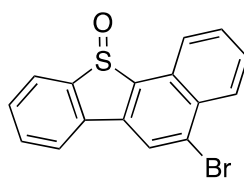
Peak#:1 R.Time 16.400(Scan#:1961)
 MassPeaks:394
 RawMode:Single 16.403(1961)
 BG Mode:None Group 1 - Event 1



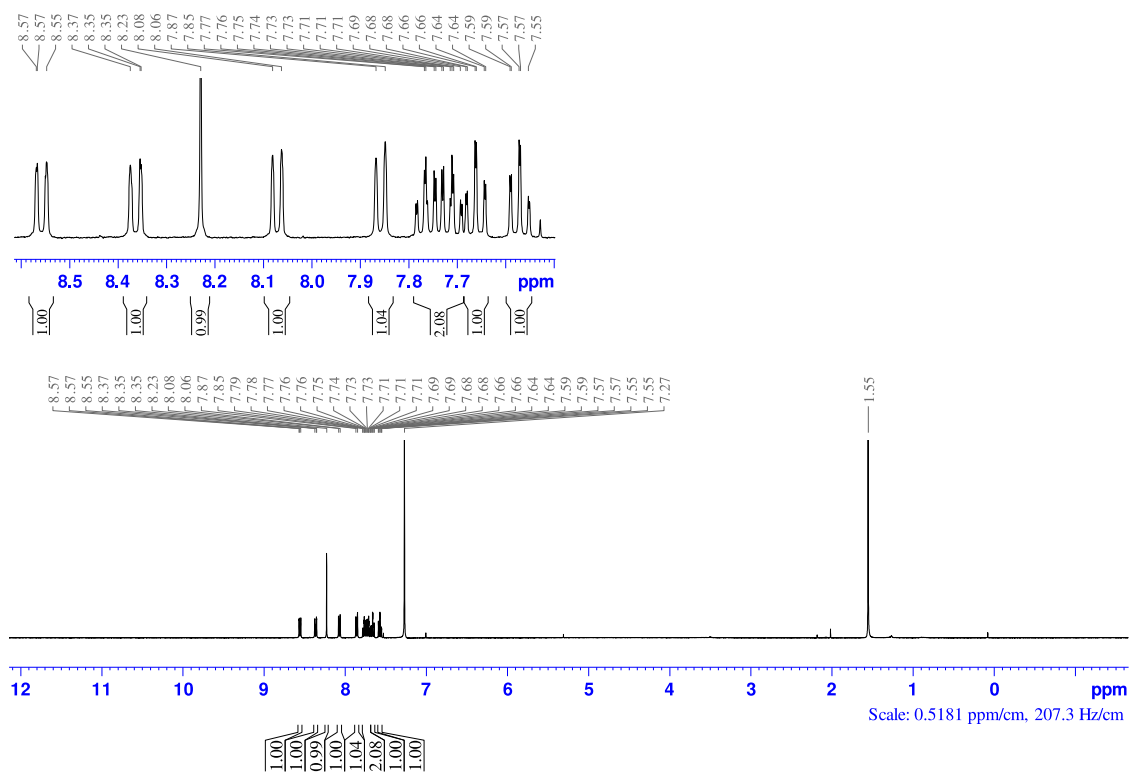
HPLC trace for purity



5-Bromobenzo[b]naphtho-[1,2-d]thiophene-S-oxide

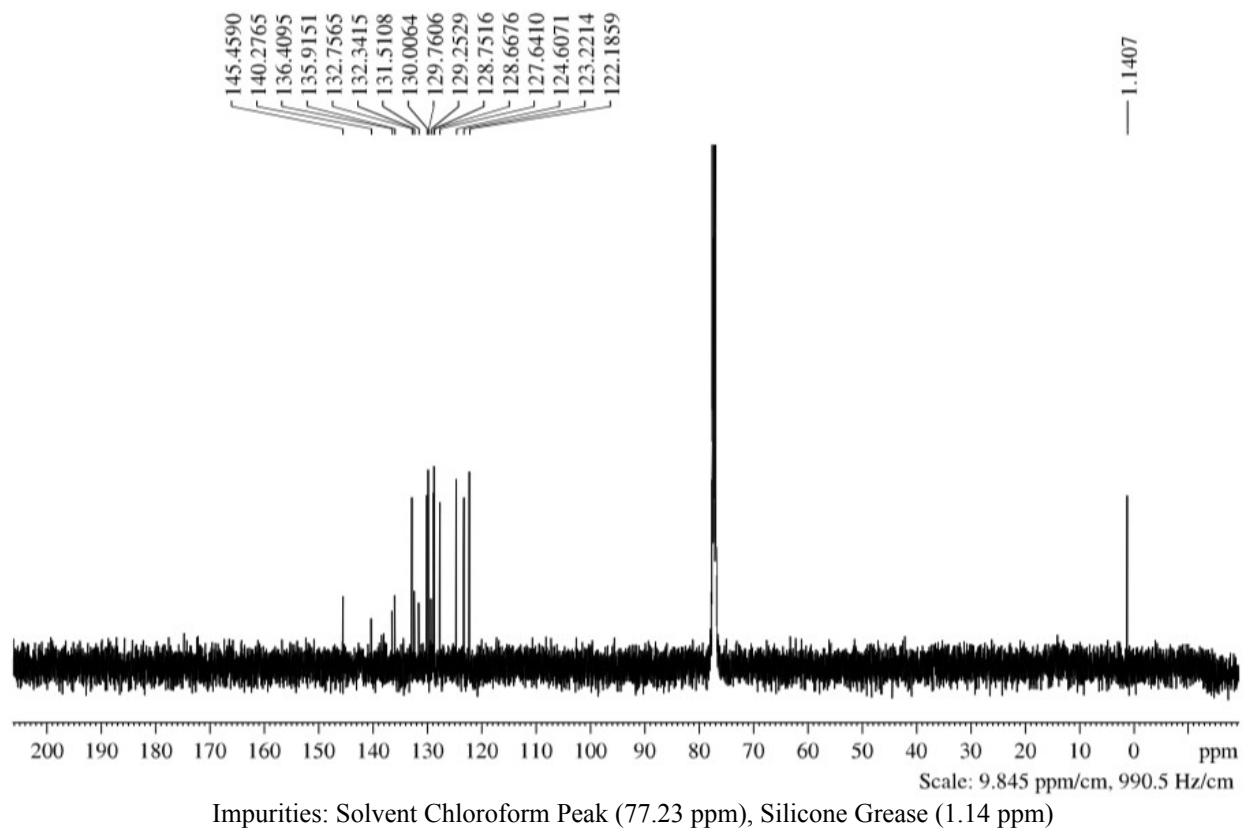


$^1\text{H-NMR}$ in CDCl_3

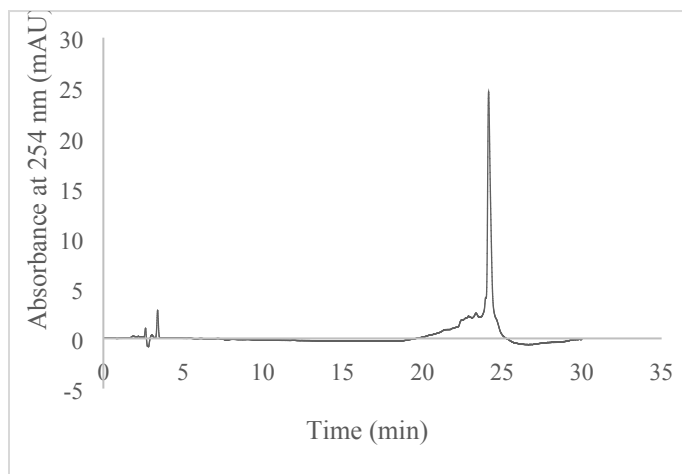


Impurities: Solvent Peak Chloroform (7.27 ppm), Water (1.55 ppm)

^{13}C -NMR (in CDCl_3)

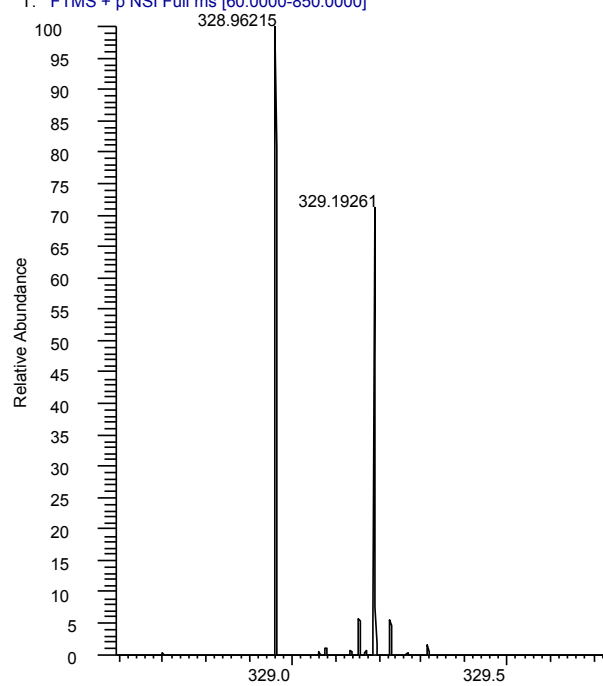


HPLC Analysis for Purity

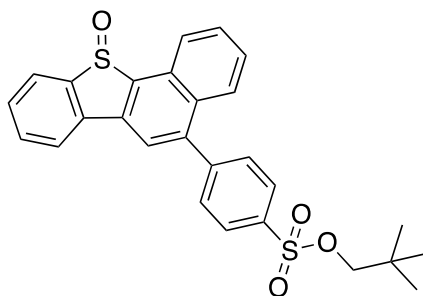


HRMS

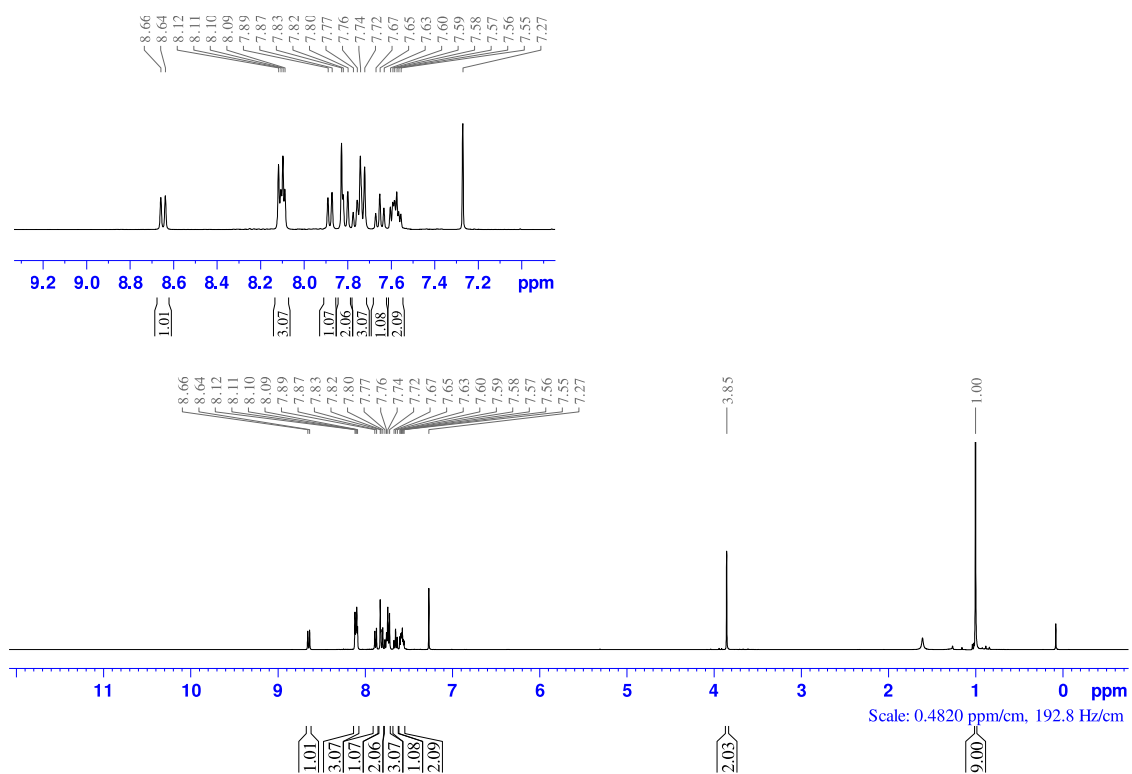
YF2 #49-58 RT: 0.43-0.51 AV: 10 NL: 1.13E6
T: FTMS + p NSI Full ms [60.0000-850.0000]



5-neopentylsulfonatephenylbenzo[b]naphtho-[1,2-d]thiophene-S-oxide

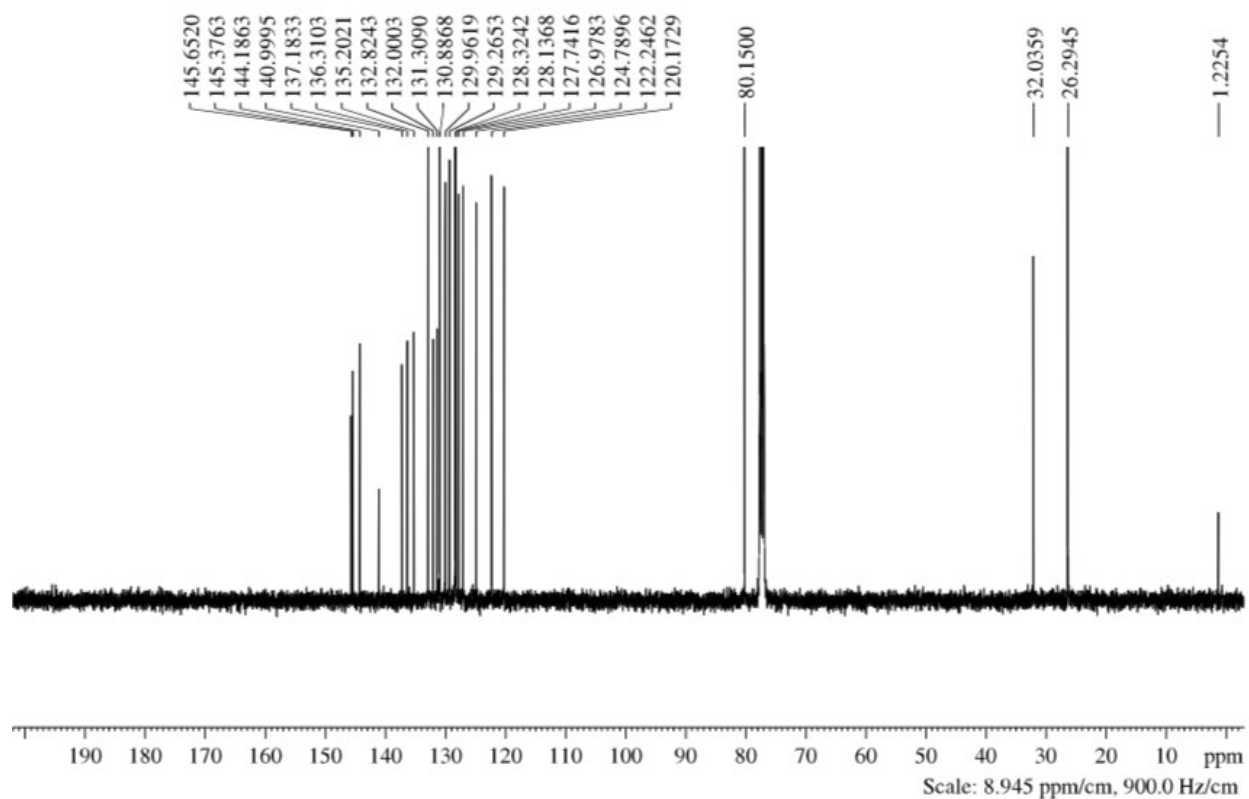


¹H-NMR (in CDCl₃)



Impurities: Solvent Peak Chloroform (7.27 ppm)

^{13}C -NMR (in CDCl_3)

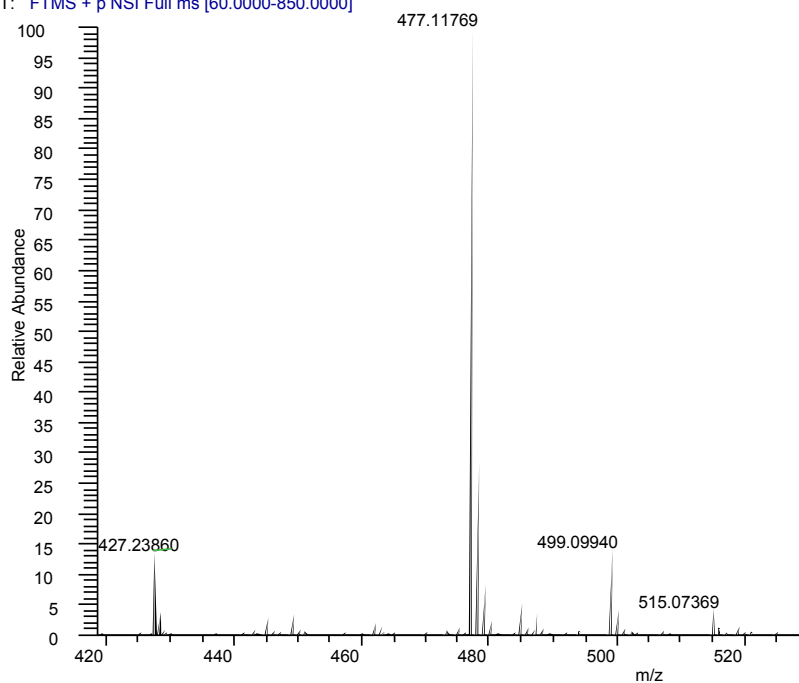


Impurities: Solvent Peak Chloroform (77.23 ppm), Silicone Grease (1.22 ppm)

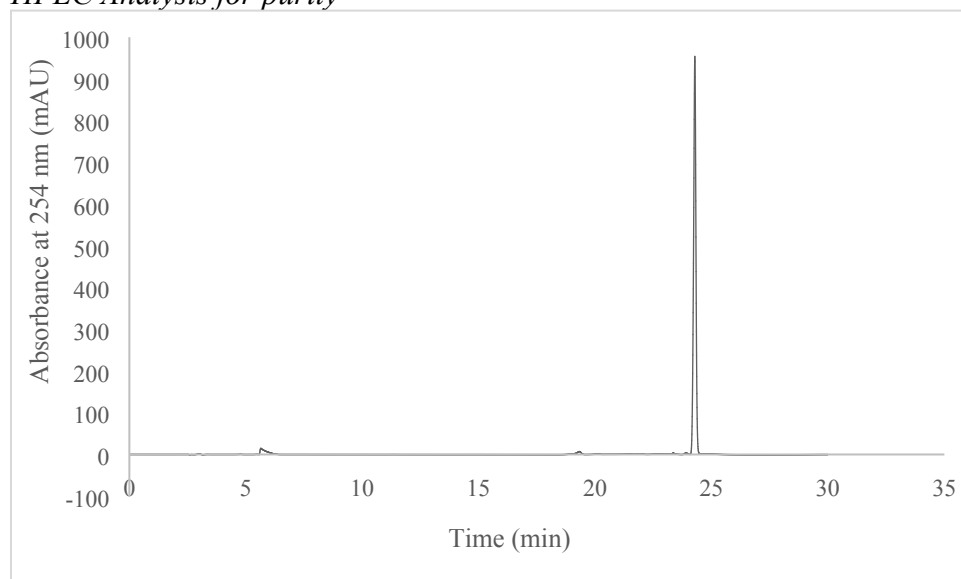
HRMS

YF3 # 1562-1667 RT: 13.63-14.54
T: FTMS + p NSI Full ms [60.0000-850.0000]

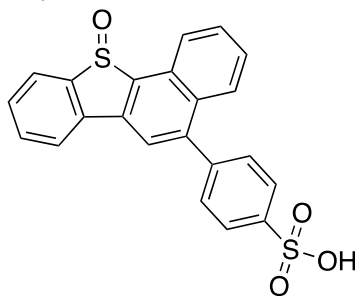
AV: 106 NL: 4.23E7



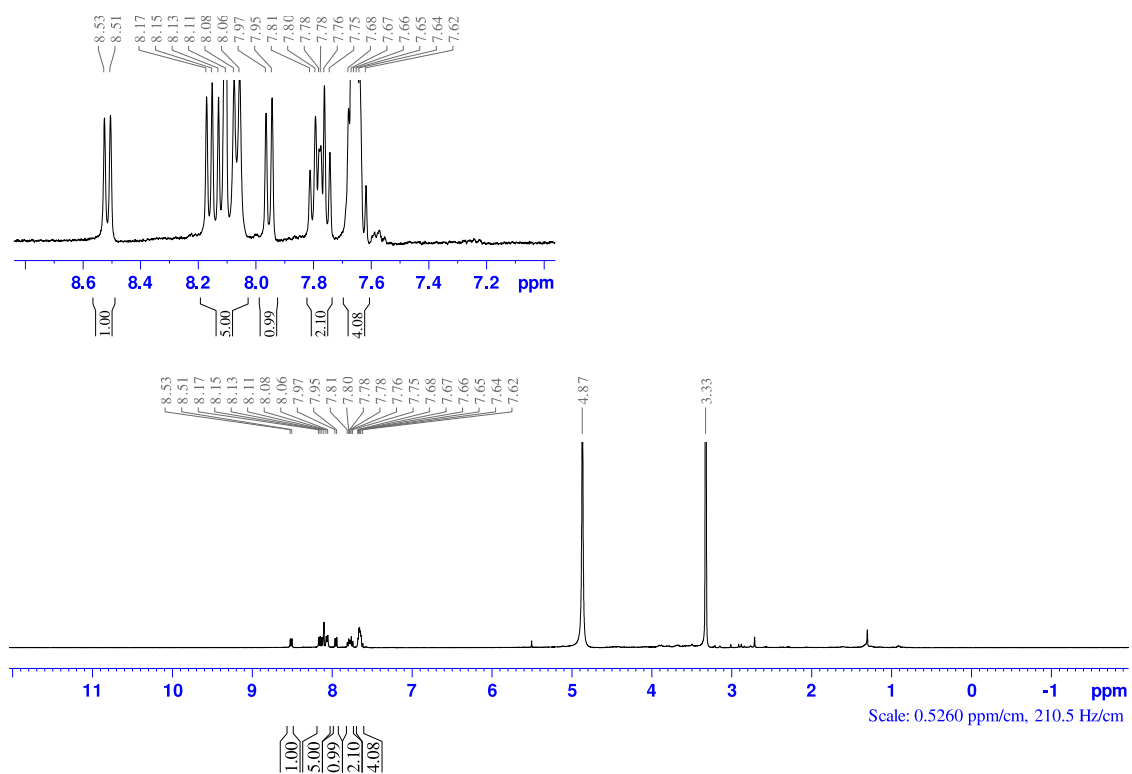
HPLC Analysis for purity



5-phenylbenzo[b]naphtho-[1,2-d]thiophene-S-oxide sulfonic acid

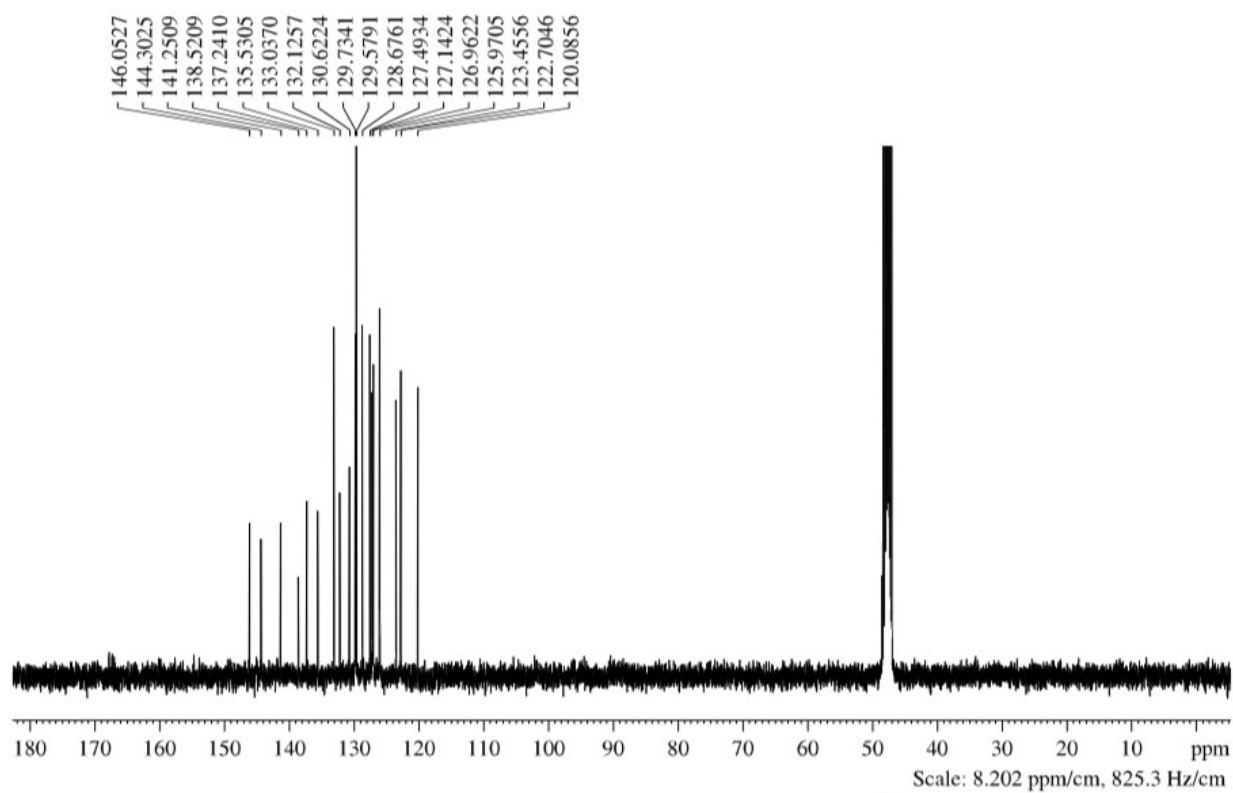


¹H-NMR (in Methanol-D₄)



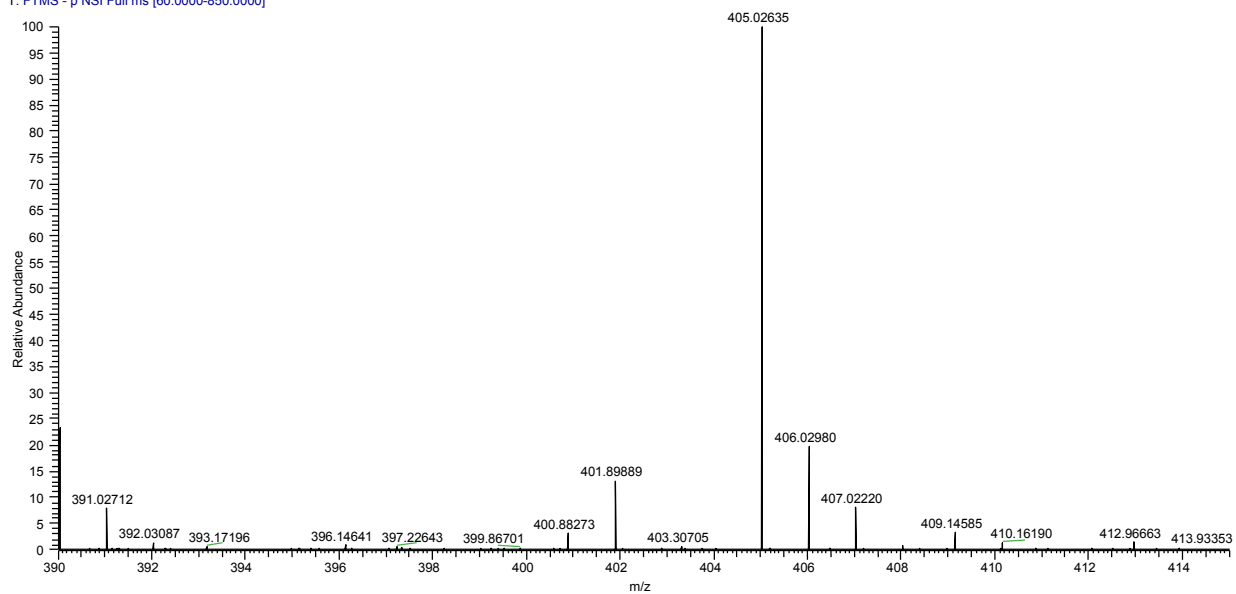
Impurities: Solvent Peak Methanol (3.31 ppm), Water (4.87 ppm)

$^{13}\text{C-NMR}$ (in $\text{Methanol-}D_4$)

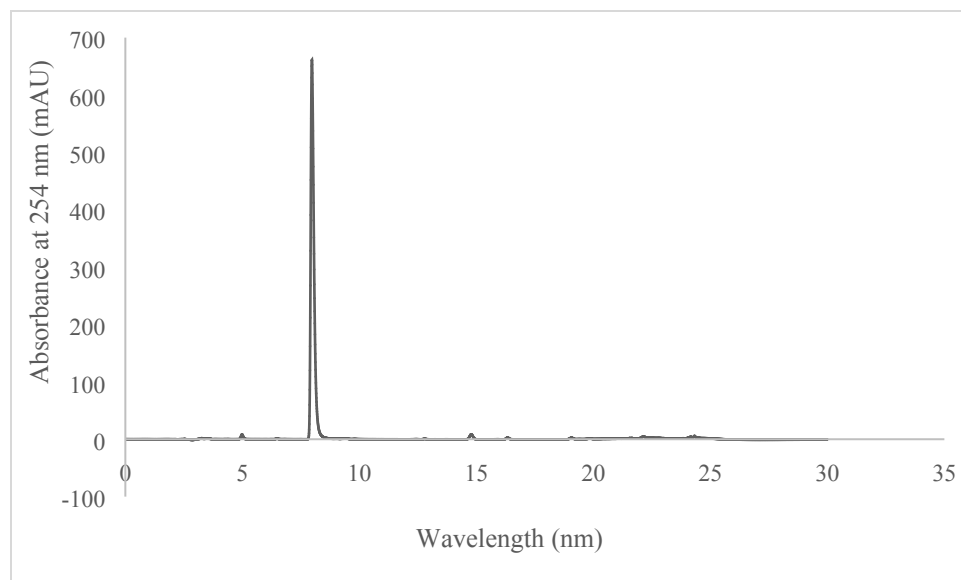


HRMS

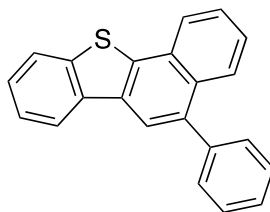
YF5_20190305193153 #336-358 RT: 4.29-4.57 AV: 23 NL: 6.45E3
T: FTMS - p NSI Full ms [60.0000-850.0000]



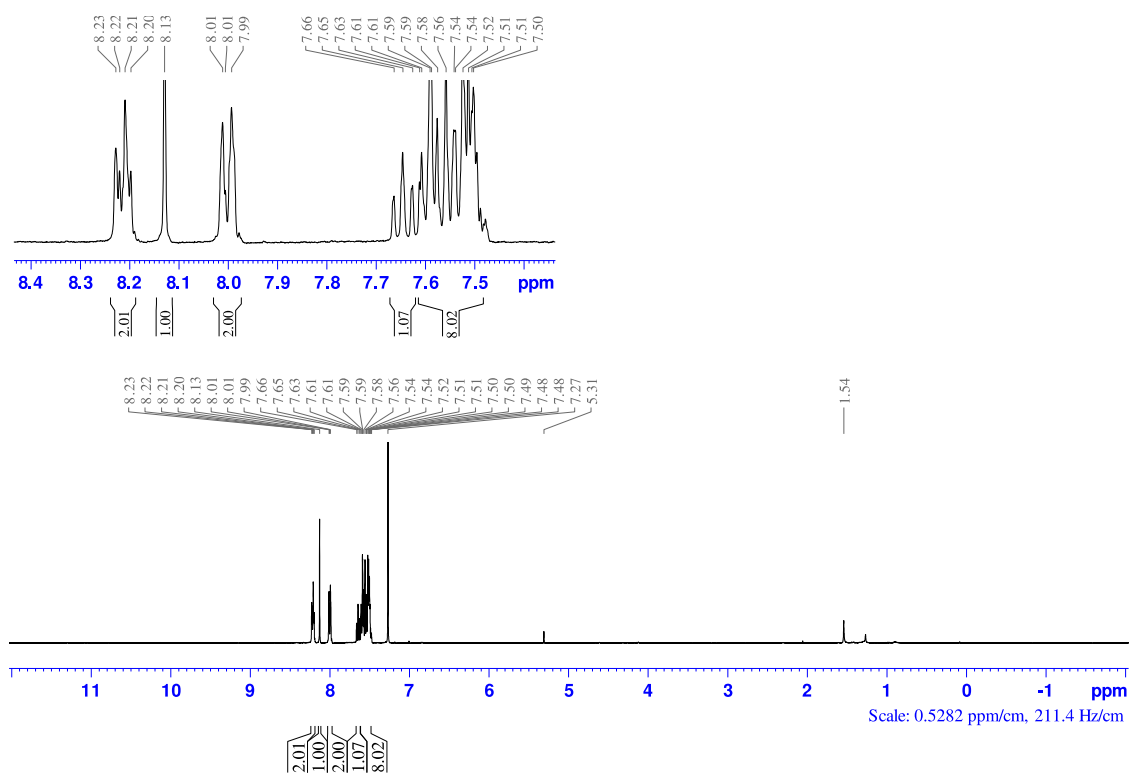
HPLC Analysis for purity



5-phenylbenzo[b]naphtho-[1,2-d]thiophene

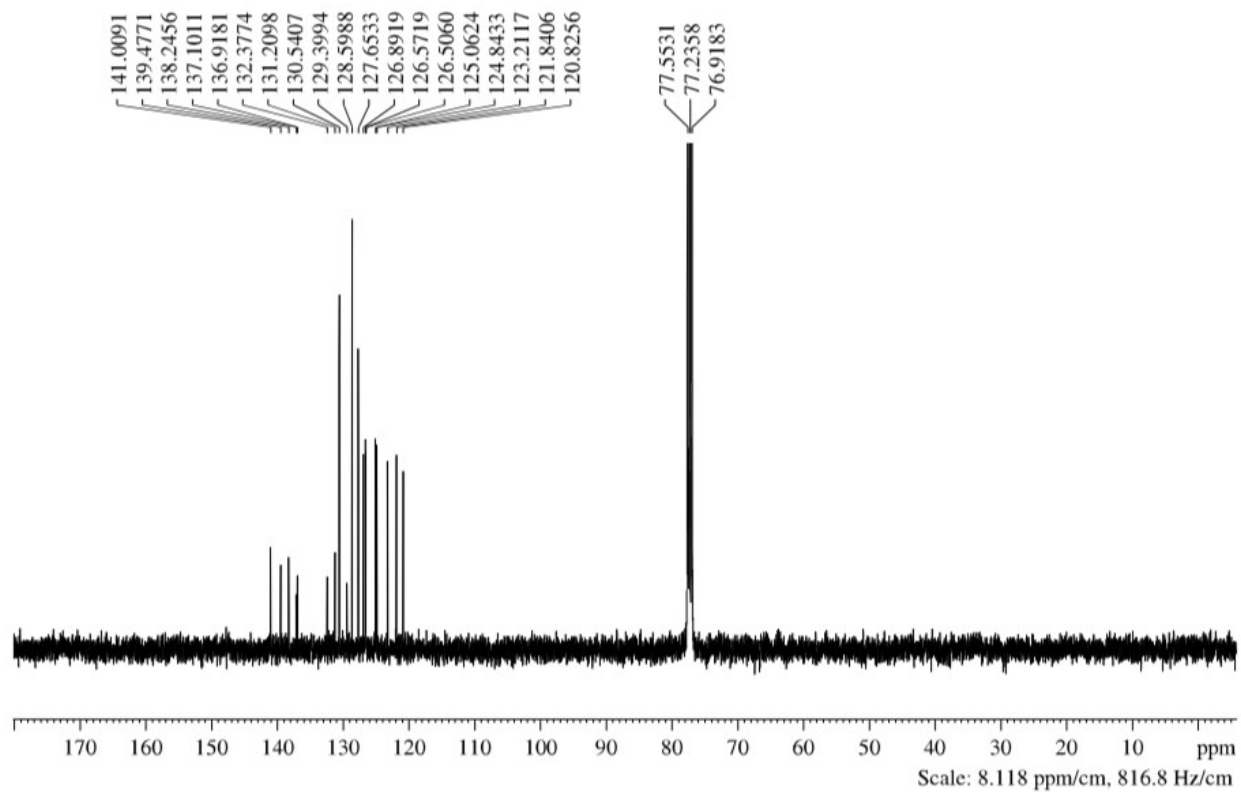


$^1\text{H-NMR}$ (in CDCl_3)



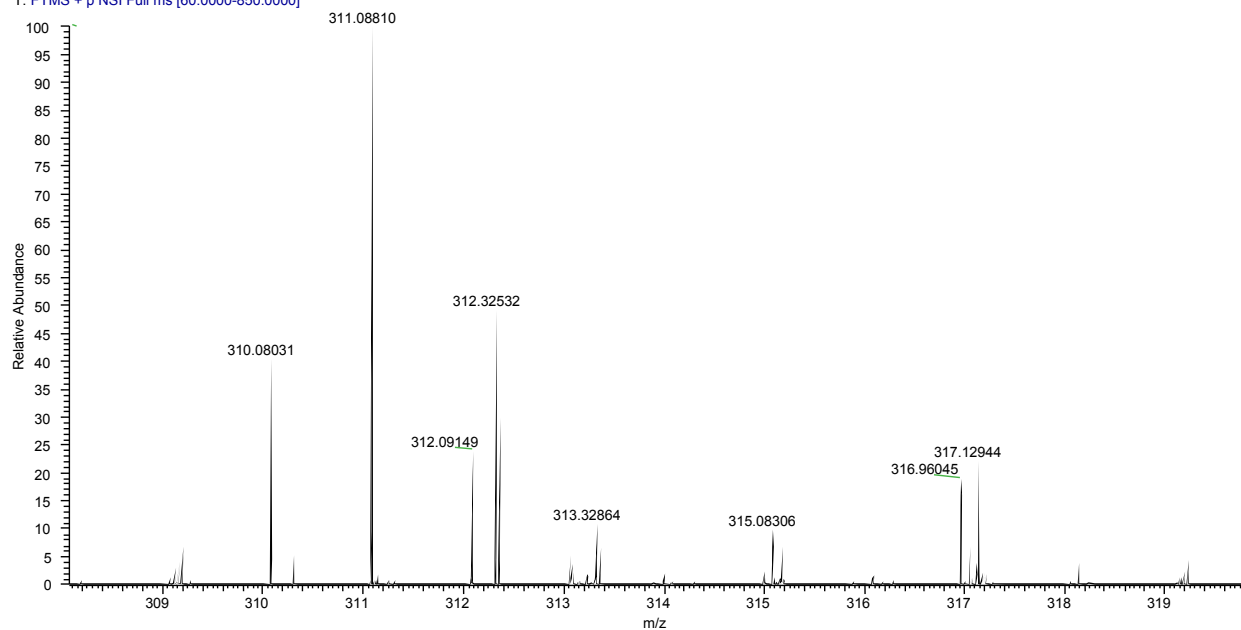
Impurities: Solvent Peak Chloroform (7.27 ppm), Dichloromethane (5.31 ppm), Water (1.54 ppm)

$^{13}\text{C-NMR}$ (in CDCl_3)

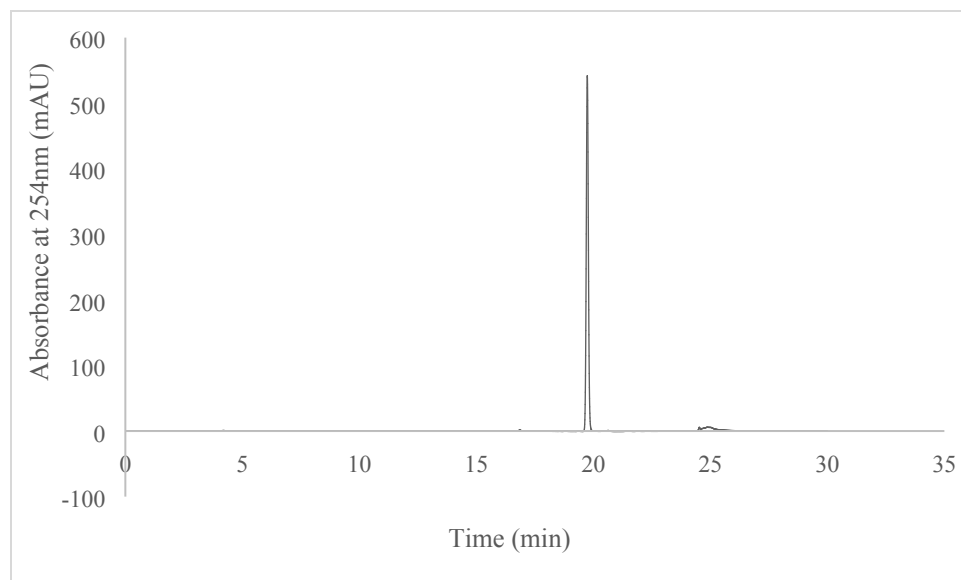


HRMS

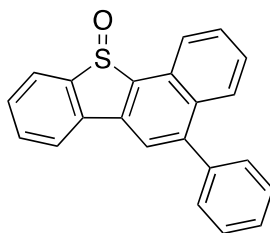
YF6 #1249-1371 RT: 10.90-11.96 AV: 123 NL: 1.63E6
T: FTMS + p NSI Full ms [60.0000-850.0000]



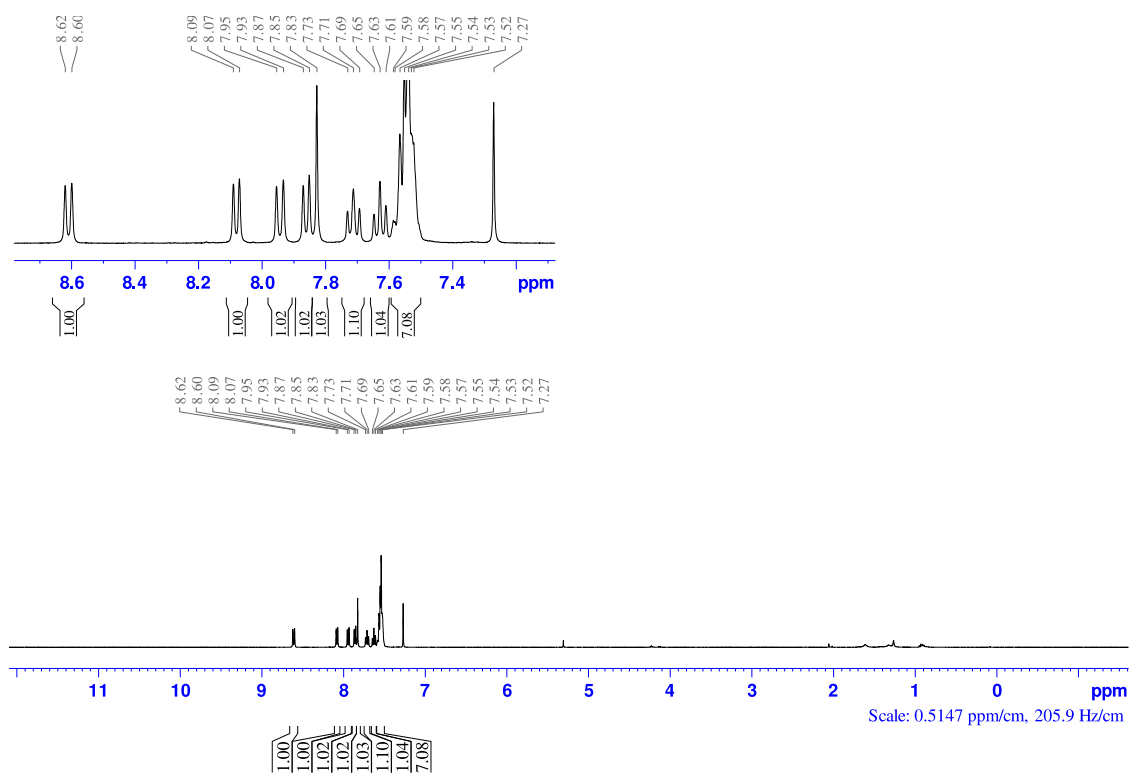
HPLC Signal for purity



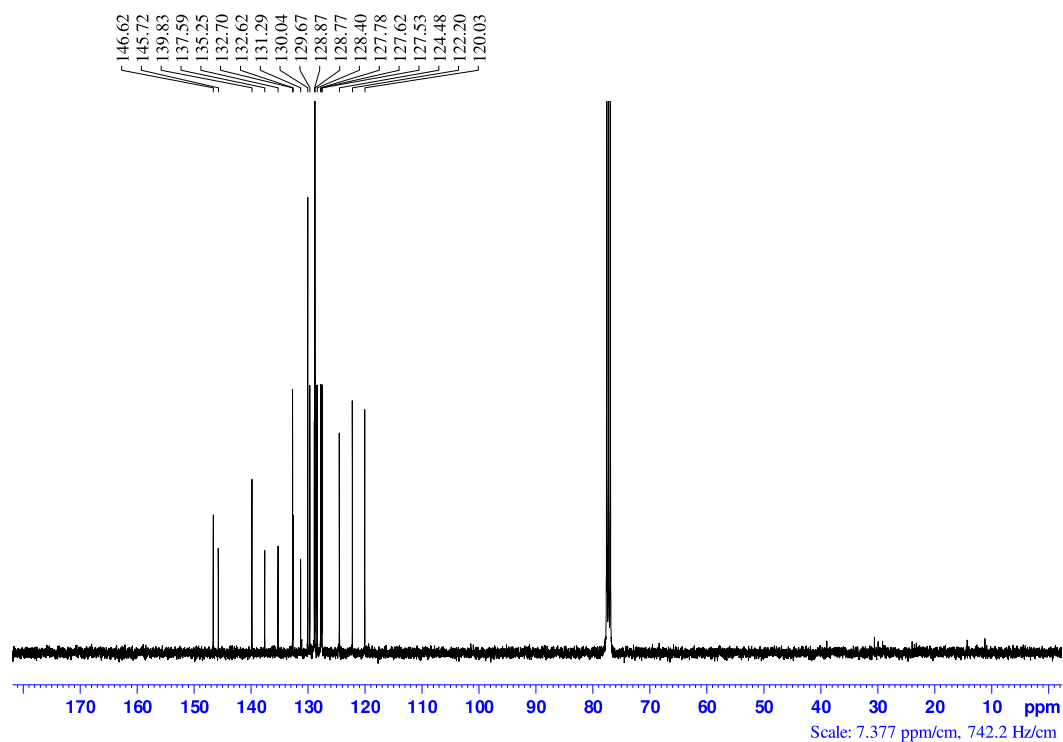
5-phenylbenzo[b]naphtho-[1,2-d]thiophene-S-oxide



$^1\text{H-NMR}$ (in CDCl_3)

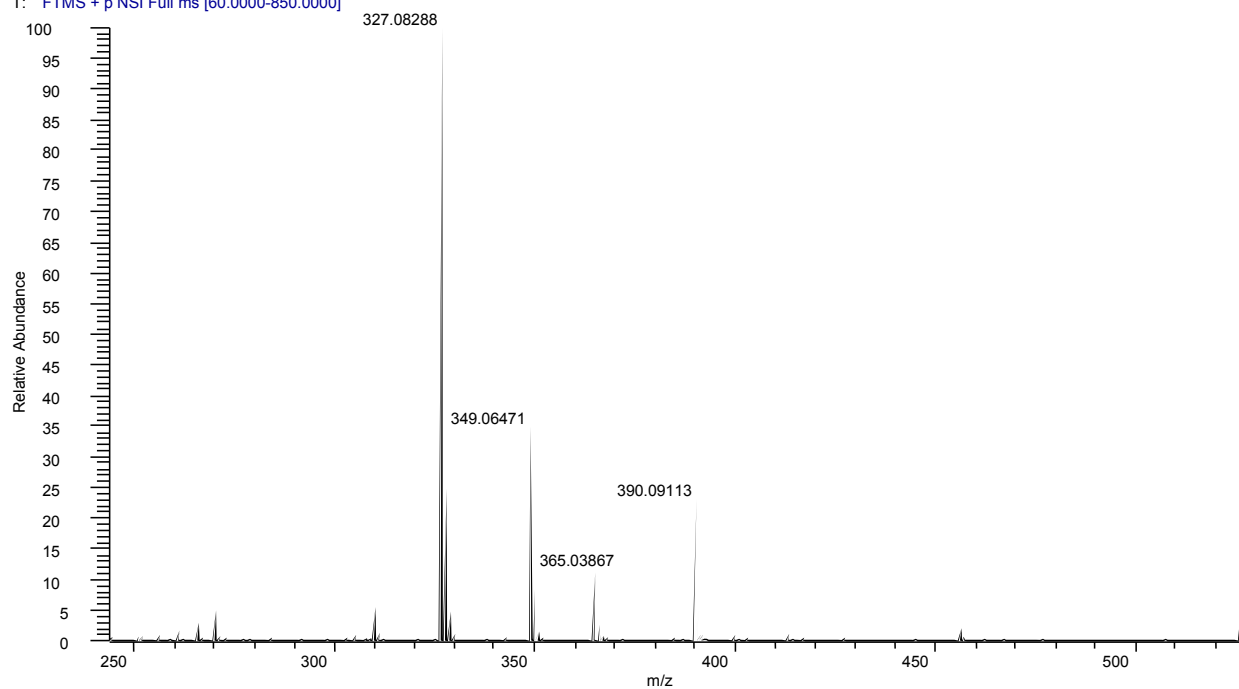


^{13}C -NMR (in CDCl_3)

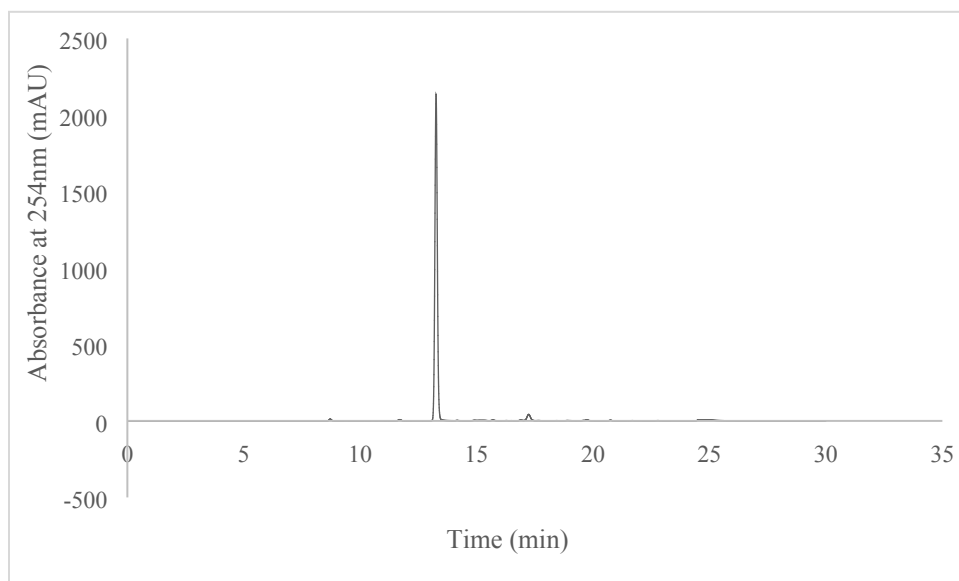


HRMS

YF7 # 1136-1315 RT: 9.91-11.47 AV: 180 NL: 1.51E8
T: FTMS + p NSI Full ms [60.0000-850.0000]

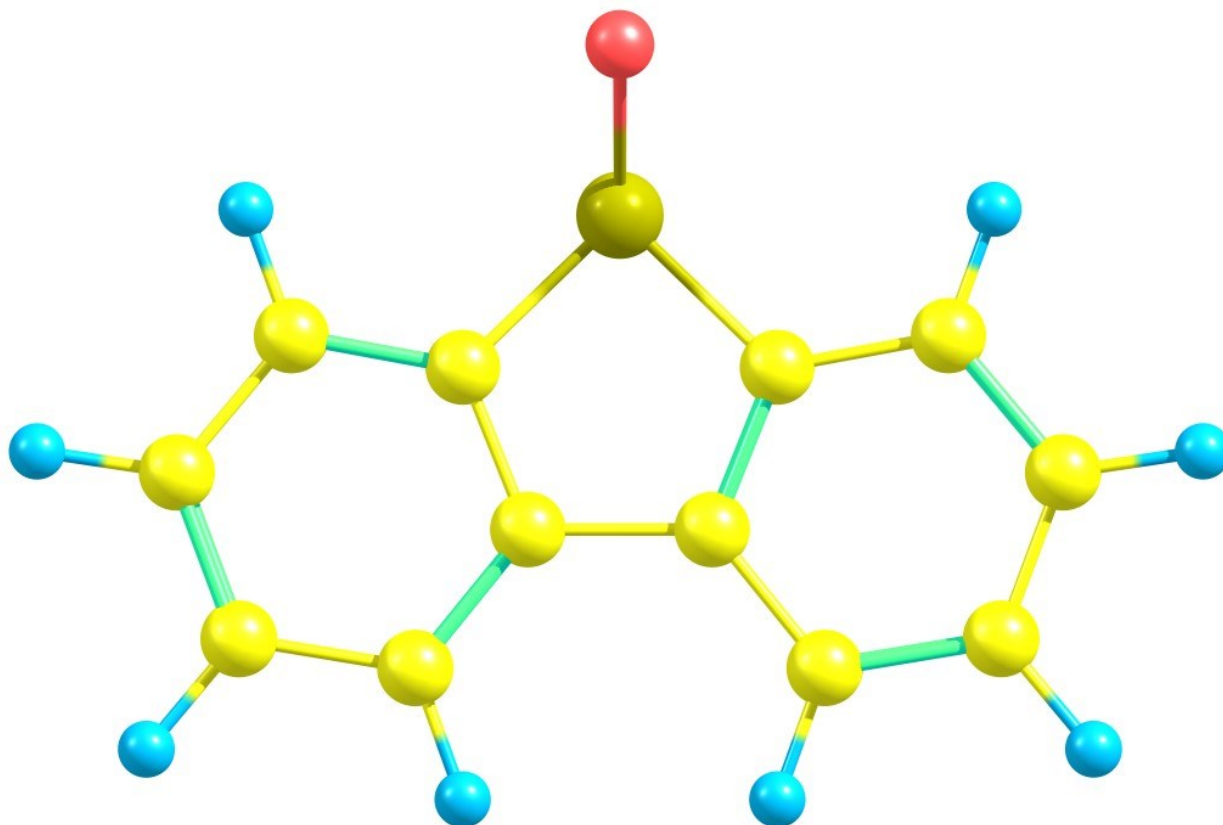


HPLC signal (for purity)



2. Optimized geometry using HSEH1PBE with 6-311G(d,p) basis set¹

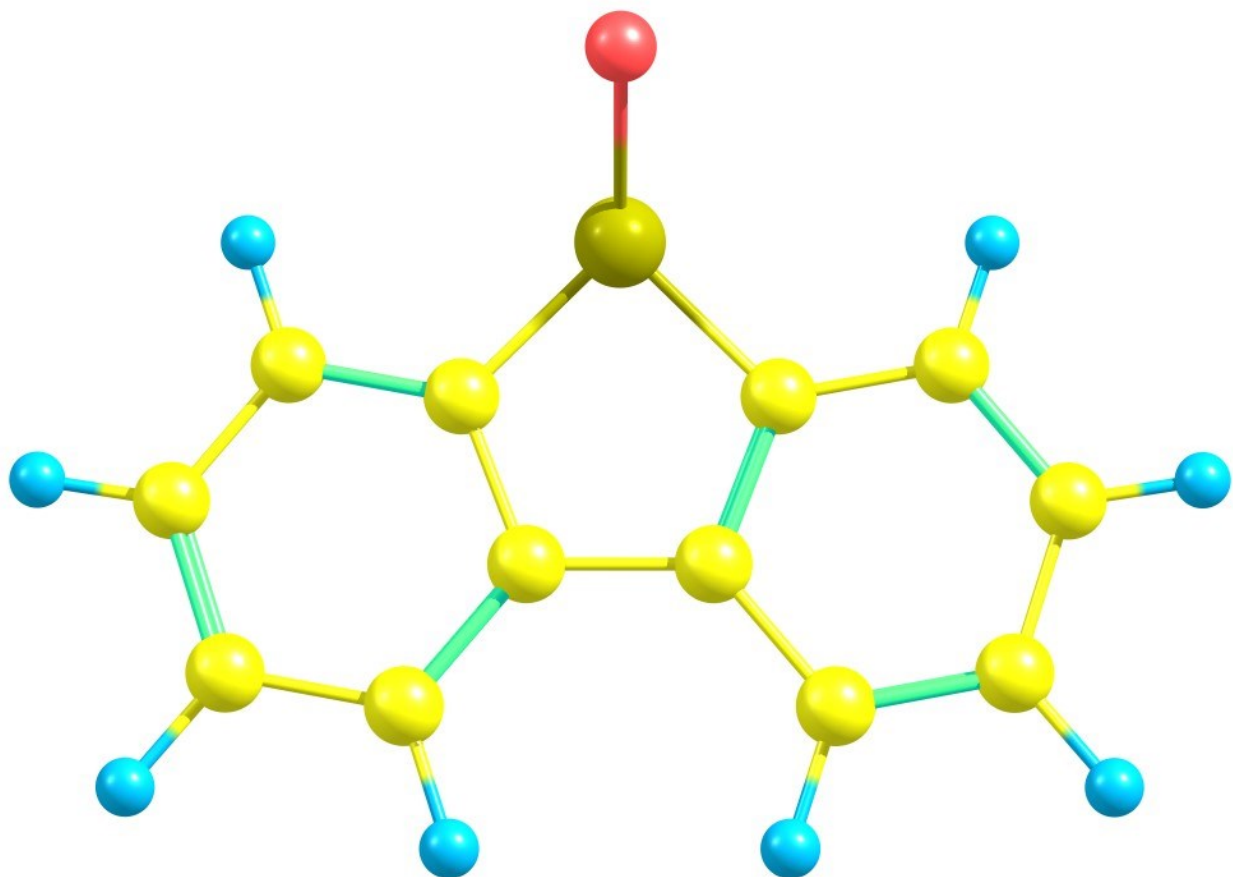
A1) DBTO (Charge = 0, Multiplicity = 1)



Coordinates

C	-0.731700000000	-0.816666000000	-0.048182000000
C	-1.606430000000	-1.895278000000	0.047422000000
C	-1.258377000000	0.476146000000	-0.118410000000
C	-2.975070000000	-1.657584000000	0.098028000000
C	-2.617291000000	0.725185000000	-0.071982000000
C	-3.480052000000	-0.360068000000	0.049740000000
H	-1.227675000000	-2.910790000000	0.094535000000
H	-3.659545000000	-2.494586000000	0.182810000000
H	-2.996259000000	1.740632000000	-0.114454000000
H	-4.550129000000	-0.194346000000	0.105347000000
C	0.731907000000	-0.816534000000	-0.048055000000
C	1.258406000000	0.476386000000	-0.117681000000
C	1.606747000000	-1.895113000000	0.046871000000
C	2.617295000000	0.725525000000	-0.071374000000
C	2.975375000000	-1.657296000000	0.097514000000
C	3.480179000000	-0.359693000000	0.049885000000
H	1.228091000000	-2.910690000000	0.093336000000
H	2.996170000000	1.741026000000	-0.113392000000
H	3.659958000000	-2.494265000000	0.181731000000
H	4.550233000000	-0.193823000000	0.105507000000
S	-0.000111000000	1.745336000000	-0.367871000000
O	-0.000625000000	2.765177000000	0.733481000000

A2) DBTO (Charge = 0, Multiplicity = 3)



Coordinates

C	-0.719779000000	-0.827922000000	-0.005504000000
C	-1.621251000000	-1.874356000000	0.186266000000
C	-1.250351000000	0.472154000000	-0.239358000000
C	-2.985398000000	-1.637048000000	0.122962000000
C	-2.626946000000	0.721998000000	-0.280527000000
C	-3.486456000000	-0.344780000000	-0.116612000000
H	-1.252544000000	-2.876599000000	0.379947000000
H	-3.677884000000	-2.459304000000	0.263373000000
H	-3.003864000000	1.725137000000	-0.446566000000
H	-4.557185000000	-0.183843000000	-0.168966000000
C	0.718323000000	-0.828526000000	-0.005784000000
C	1.250051000000	0.471216000000	-0.238137000000
C	1.618941000000	-1.875996000000	0.184551000000
C	2.626724000000	0.719943000000	-0.279417000000
C	2.983269000000	-1.639813000000	0.121274000000
C	3.485419000000	-0.347780000000	-0.116822000000
H	1.249369000000	-2.878131000000	0.377090000000
H	3.004451000000	1.722952000000	-0.444340000000
H	3.675045000000	-2.462869000000	0.260509000000
H	4.556273000000	-0.187663000000	-0.169031000000
S	0.000393000000	1.680161000000	-0.192722000000

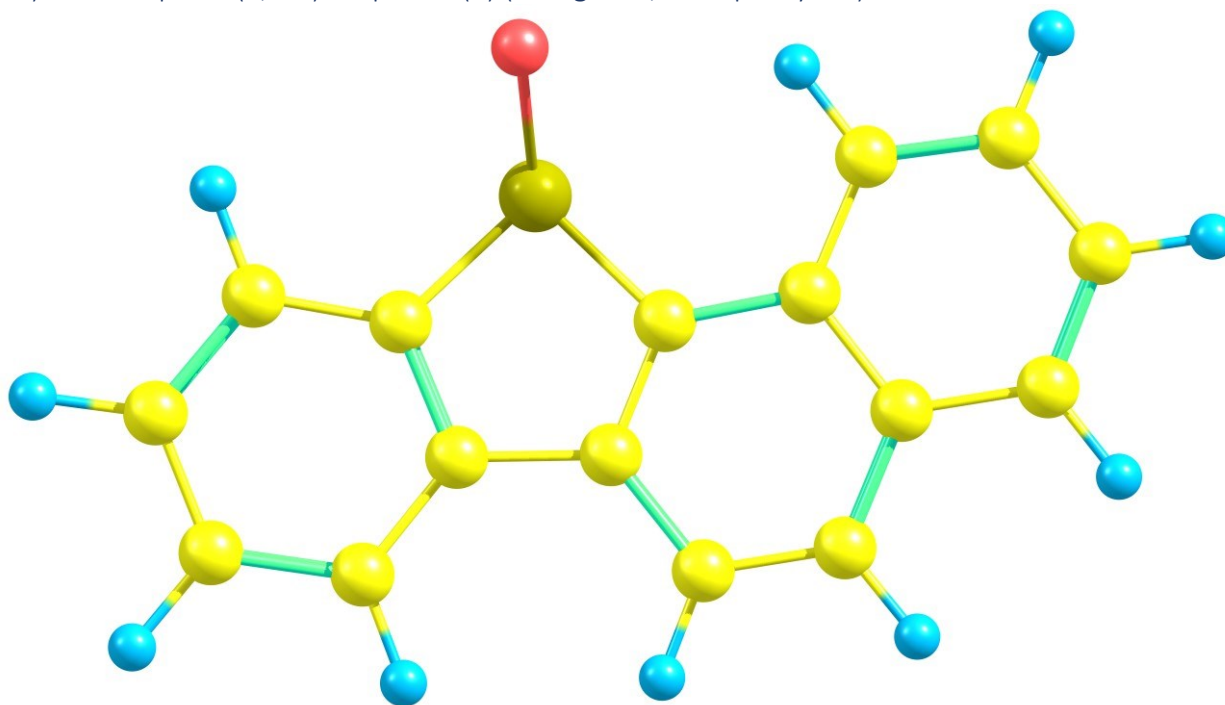
O 0.005597000000 2.832902000000 0.879273000000

A3) Energy calculations for DBTO

Charge	Multiplicity	Sum of electronic and thermal free energies (hartree/particle)
0	1	-934.761157
0	3	-934.664716
Difference		0.096441

In kcal/mol = 0.096441 * 627.5 = 60.5 kcal/mol

B1) Benzonaphtho(1,2-d)thiophene (1) (Charge = 0, Multiplicity = 1)

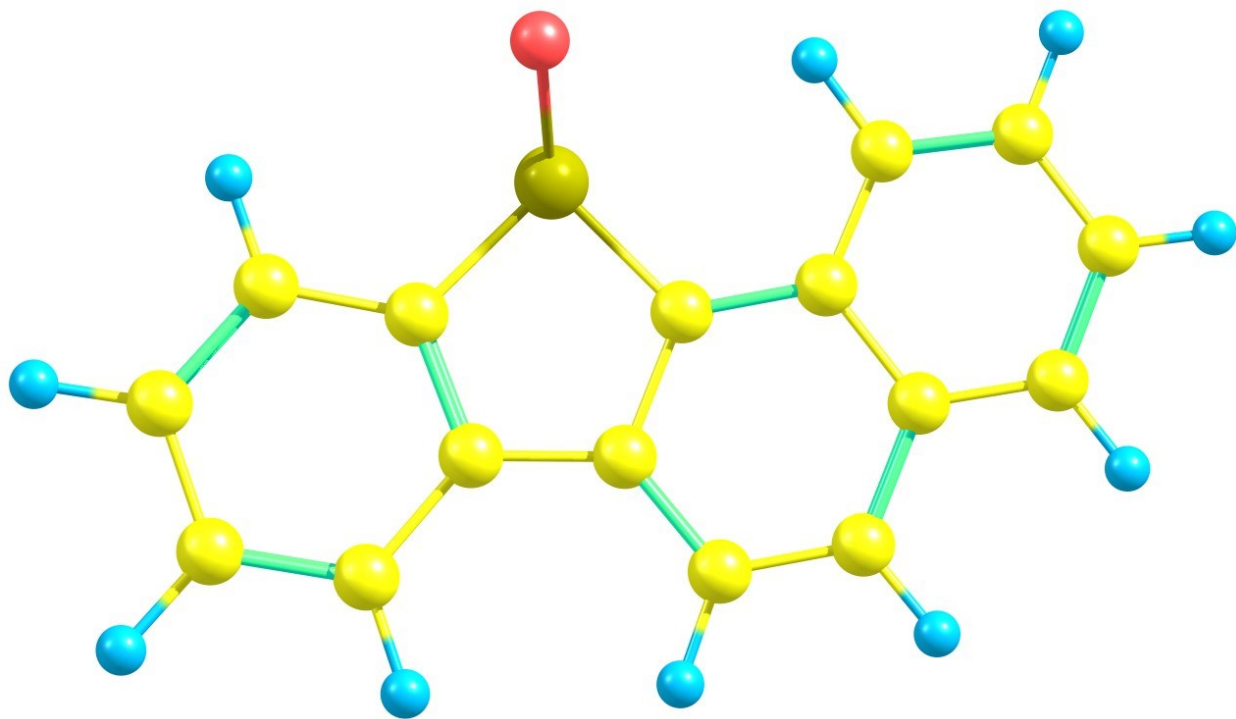


Coordinates

C	4.449124000000	-0.139866000000	0.079334000000
C	3.414694000000	-1.063889000000	-0.045211000000
C	4.172974000000	1.224903000000	0.115541000000
H	3.613451000000	-2.129556000000	-0.079739000000
H	4.989911000000	1.933052000000	0.201479000000
C	2.120539000000	-0.584657000000	-0.101733000000
C	2.865174000000	1.693474000000	0.052566000000
H	2.665767000000	2.759090000000	0.090638000000
C	1.820284000000	0.778864000000	-0.041231000000
H	5.474553000000	-0.485661000000	0.146398000000
C	-0.360257000000	-0.143294000000	-0.080319000000

C	-1.770376000000	-0.157411000000	-0.044344000000
C	0.378530000000	1.023337000000	-0.046404000000
C	-2.429565000000	1.108109000000	0.019307000000
C	-0.287619000000	2.266607000000	0.005474000000
H	0.281452000000	3.189665000000	0.025512000000
C	-1.656321000000	2.296855000000	0.034499000000
H	-2.176879000000	3.248512000000	0.075940000000
S	0.660114000000	-1.597359000000	-0.380926000000
C	-2.542267000000	-1.343258000000	-0.039728000000
C	-3.910854000000	-1.273984000000	0.003270000000
C	-4.567128000000	-0.027082000000	0.052915000000
C	-3.841789000000	1.136420000000	0.065218000000
H	-2.036537000000	-2.302574000000	-0.041494000000
H	-4.495542000000	-2.187391000000	0.008826000000
H	-5.650556000000	0.007728000000	0.088105000000
H	-4.343689000000	2.097880000000	0.112141000000
O	0.498171000000	-2.667973000000	0.661510000000

B2) Benzonaphtho(1,2-d)thiophene (1) (Charge = 0, Multiplicity = 3)



Coordinates

C	4.435600000000	-0.193406000000	-0.015848000000
C	3.377998000000	-1.086275000000	-0.149364000000
C	4.189729000000	1.184191000000	0.107719000000
H	3.557277000000	-2.152494000000	-0.237755000000
H	5.027162000000	1.865381000000	0.212641000000

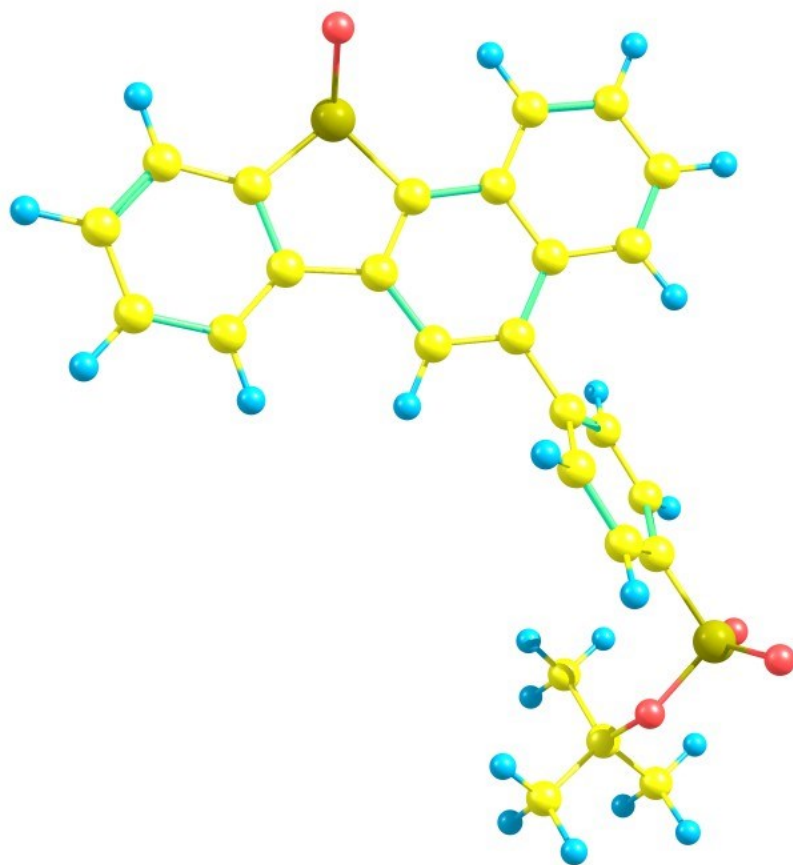
C	2.088948000000	-0.587301000000	-0.159666000000
C	2.904244000000	1.681366000000	0.106485000000
H	2.728523000000	2.745672000000	0.220586000000
C	1.815803000000	0.794440000000	-0.022329000000
H	5.455311000000	-0.561239000000	-0.008044000000
C	-0.364166000000	-0.164391000000	-0.143513000000
C	-1.768213000000	-0.164300000000	-0.065490000000
C	0.426769000000	1.077282000000	-0.026909000000
C	-2.425733000000	1.110529000000	0.007370000000
C	-0.245085000000	2.288557000000	0.053790000000
H	0.311623000000	3.217916000000	0.115505000000
C	-1.628638000000	2.316160000000	0.055896000000
H	-2.150385000000	3.265090000000	0.117709000000
S	0.628847000000	-1.633159000000	-0.337110000000
C	-2.548308000000	-1.335147000000	-0.057861000000
C	-3.947847000000	-1.263918000000	-0.033793000000
C	-4.576363000000	-0.040239000000	0.007333000000
C	-3.806384000000	1.147702000000	0.035602000000
H	-2.052267000000	-2.299056000000	-0.039037000000
H	-4.528115000000	-2.179638000000	-0.034881000000
H	-5.658307000000	0.023051000000	0.030146000000
H	-4.309115000000	2.108870000000	0.080286000000
O	0.498326000000	-2.561813000000	0.842510000000

B3) Energy calculations for 1

Charge	Multiplicity	Sum of electronic and thermal free energies (hartree/particle)
0	1	-1088.226958
0	3	-1088.146953
Difference		0.080005

in kcal/mol = 0.080005 * 627.5 = 50.2 kcal/mol

C1) 5-neopentylsulfonatephenylbenzonaphtho(1,2-d)thiophene (2) (Charge = 0, Multiplicity = 1)

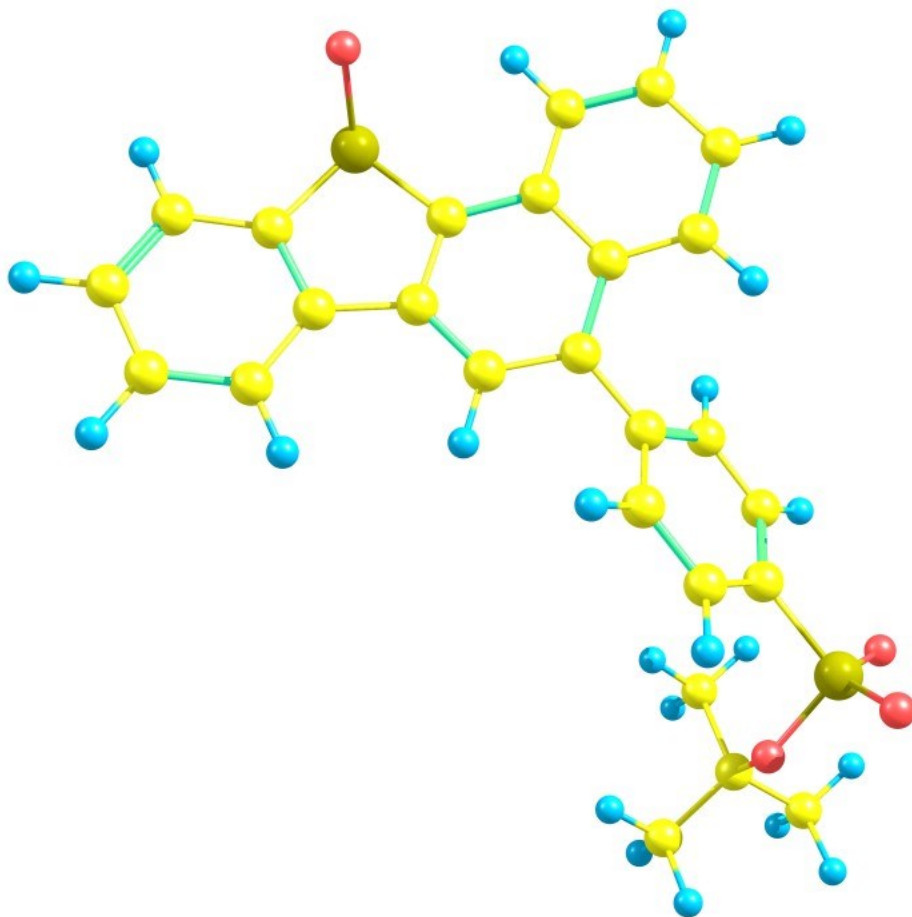


Coordinates

C	5.750362000000	3.846545000000	0.208670000000
C	6.136166000000	2.537916000000	-0.070331000000
C	4.405831000000	4.160240000000	0.392965000000
H	7.178540000000	2.278839000000	-0.220821000000
H	4.120139000000	5.185900000000	0.599155000000
C	5.151719000000	1.570811000000	-0.128501000000
C	3.424671000000	3.176902000000	0.326313000000
H	2.382682000000	3.435769000000	0.481446000000
C	3.800311000000	1.860069000000	0.077788000000
H	6.501401000000	4.625148000000	0.280912000000
C	3.709387000000	-0.493978000000	-0.162660000000
C	3.125008000000	-1.776787000000	-0.179918000000
C	2.972826000000	0.655012000000	0.037164000000
C	1.707172000000	-1.863463000000	-0.006990000000
C	1.575890000000	0.562579000000	0.181898000000
H	0.982195000000	1.463763000000	0.294467000000
C	0.943017000000	-0.660366000000	0.140770000000
S	5.426011000000	-0.146878000000	-0.587508000000
C	3.888510000000	-2.960519000000	-0.314538000000
C	3.277178000000	-4.186332000000	-0.281734000000

C	1.885464000000	-4.281511000000	-0.091945000000
C	1.121588000000	-3.150376000000	0.047729000000
H	4.966370000000	-2.882707000000	-0.404671000000
H	3.870511000000	-5.088658000000	-0.379972000000
H	1.415372000000	-5.257476000000	-0.040880000000
H	0.055531000000	-3.239181000000	0.219584000000
O	6.415996000000	-0.827377000000	0.314821000000
C	-0.535389000000	-0.691133000000	0.243254000000
C	-1.313603000000	-1.259194000000	-0.769122000000
C	-2.699318000000	-1.240017000000	-0.695028000000
C	-3.308838000000	-0.652481000000	0.404633000000
C	-2.559194000000	-0.092055000000	1.432493000000
C	-1.176695000000	-0.107711000000	1.341696000000
H	-0.827813000000	-1.705644000000	-1.629840000000
H	-3.310412000000	-1.676223000000	-1.476580000000
H	-3.053949000000	0.338129000000	2.295496000000
H	-0.579349000000	0.315689000000	2.141986000000
S	-5.082084000000	-0.667338000000	0.533177000000
O	-5.433945000000	-1.129435000000	1.852877000000
O	-5.617034000000	-1.305228000000	-0.654259000000
O	-5.440071000000	0.900394000000	0.547929000000
C	-5.837533000000	1.698858000000	-0.634441000000
C	-7.239950000000	1.292170000000	-1.054485000000
H	-7.257628000000	0.267688000000	-1.426689000000
H	-7.924818000000	1.372654000000	-0.207214000000
H	-7.590287000000	1.960208000000	-1.846486000000
C	-5.815667000000	3.110075000000	-0.073714000000
H	-6.130293000000	3.818445000000	-0.844383000000
H	-6.496119000000	3.192397000000	0.776117000000
H	-4.809717000000	3.376219000000	0.258261000000
C	-4.829764000000	1.543108000000	-1.761457000000
H	-5.084698000000	2.239603000000	-2.564545000000
H	-3.819487000000	1.776561000000	-1.416401000000
H	-4.845474000000	0.533323000000	-2.173652000000

C2) 5-neopentylsulfonatephenylbenzonaphtho(1,2-d)thiophene (2) (Charge = 0, Multiplicity = 3)



Coordinates

C	5.811281000000	-3.773514000000	-0.180725000000
C	6.113189000000	-2.484979000000	0.244148000000
C	4.515811000000	-4.094013000000	-0.604728000000
H	7.114652000000	-2.223397000000	0.568344000000
H	4.297662000000	-5.104225000000	-0.933193000000
C	5.105463000000	-1.537588000000	0.234321000000
C	3.515288000000	-3.140057000000	-0.621056000000
H	2.521405000000	-3.396718000000	-0.971876000000
C	3.802126000000	-1.832504000000	-0.202952000000
H	6.582809000000	-4.535143000000	-0.182854000000
C	3.670929000000	0.502063000000	0.283766000000
C	3.080026000000	1.783370000000	0.260196000000
C	2.942219000000	-0.683794000000	-0.171794000000
C	1.686627000000	1.878358000000	-0.072554000000
C	1.611189000000	-0.577786000000	-0.461891000000
H	1.042866000000	-1.467286000000	-0.712293000000
C	0.922786000000	0.654094000000	-0.330950000000
S	5.342872000000	0.163308000000	0.792374000000
C	3.817819000000	2.955822000000	0.490966000000

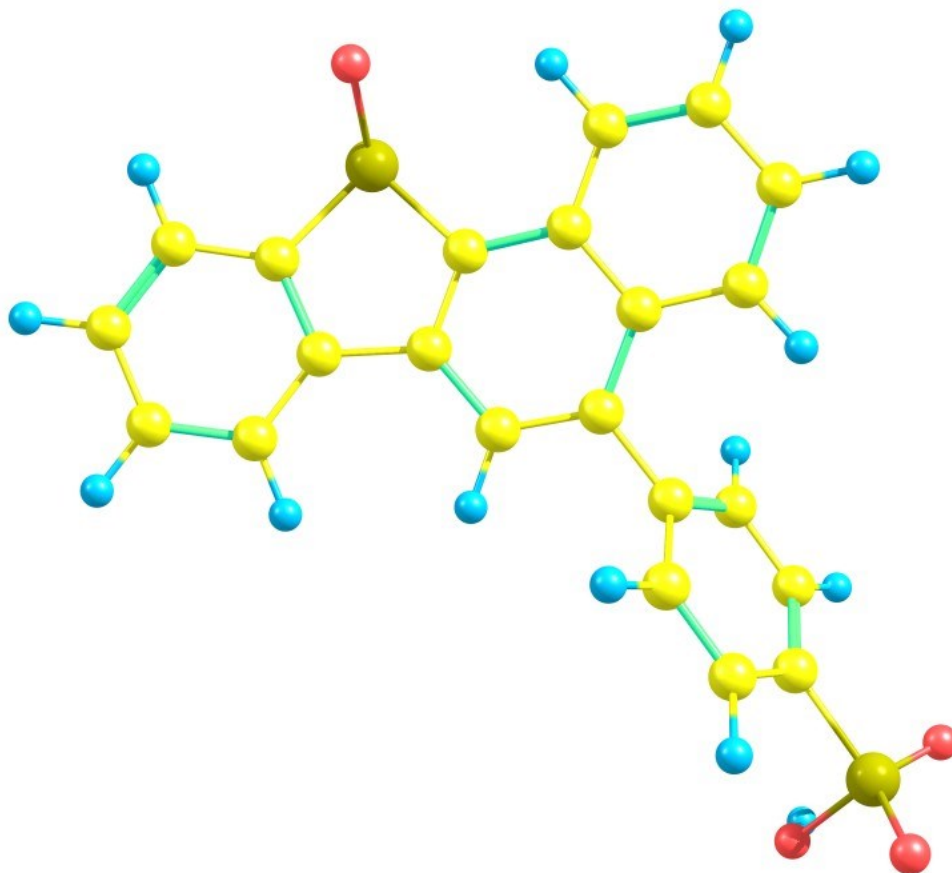
C	3.215630000000	4.215169000000	0.400945000000
C	1.890647000000	4.312019000000	0.042104000000
C	1.138891000000	3.143677000000	-0.210517000000
H	4.879127000000	2.875795000000	0.698106000000
H	3.804375000000	5.105722000000	0.588741000000
H	1.415506000000	5.280800000000	-0.061943000000
H	0.110603000000	3.248415000000	-0.534815000000
O	6.369319000000	0.864464000000	-0.057589000000
C	-0.528845000000	0.651111000000	-0.434183000000
C	-1.334942000000	1.422689000000	0.425044000000
C	-2.716221000000	1.365373000000	0.356882000000
C	-3.319657000000	0.542343000000	-0.586812000000
C	-2.553792000000	-0.224396000000	-1.463022000000
C	-1.177066000000	-0.177279000000	-1.374488000000
H	-0.866821000000	2.040037000000	1.182618000000
H	-3.334648000000	1.947603000000	1.030079000000
H	-3.038877000000	-0.836527000000	-2.214550000000
H	-0.582809000000	-0.754085000000	-2.074368000000
S	-5.087899000000	0.495485000000	-0.712808000000
O	-5.446079000000	0.622675000000	-2.104084000000
O	-5.638975000000	1.386112000000	0.290578000000
O	-5.423534000000	-1.037883000000	-0.356441000000
C	-5.782947000000	-1.542612000000	0.987647000000
C	-7.193366000000	-1.085948000000	1.321477000000
H	-7.238460000000	-0.002743000000	1.436338000000
H	-7.883227000000	-1.386745000000	0.529621000000
H	-7.515597000000	-1.552922000000	2.256578000000
C	-5.723797000000	-3.045939000000	0.779161000000
H	-6.013817000000	-3.557696000000	1.700435000000
H	-6.406869000000	-3.346069000000	-0.017942000000
H	-4.712671000000	-3.357190000000	0.507621000000
C	-4.771168000000	-1.095630000000	2.030205000000
H	-4.989895000000	-1.599469000000	2.975327000000
H	-3.755427000000	-1.362619000000	1.728399000000
H	-4.824031000000	-0.019465000000	2.201705000000

C3) Energy calculations for 2

Charge	Multiplicity	Sum of electronic and thermal free energies (hartree/particle)
0	1	-2099.540992
0	3	-2099.464567
Difference		0.076425

in kcal/mol = 0.076425 * 627.5 = 47.9 kcal/mol

D1) 5-phenylsulfonicacidbenzonaphtho(1,2-d)thiophene (3) (Charge = 0, Multiplicity = 1)

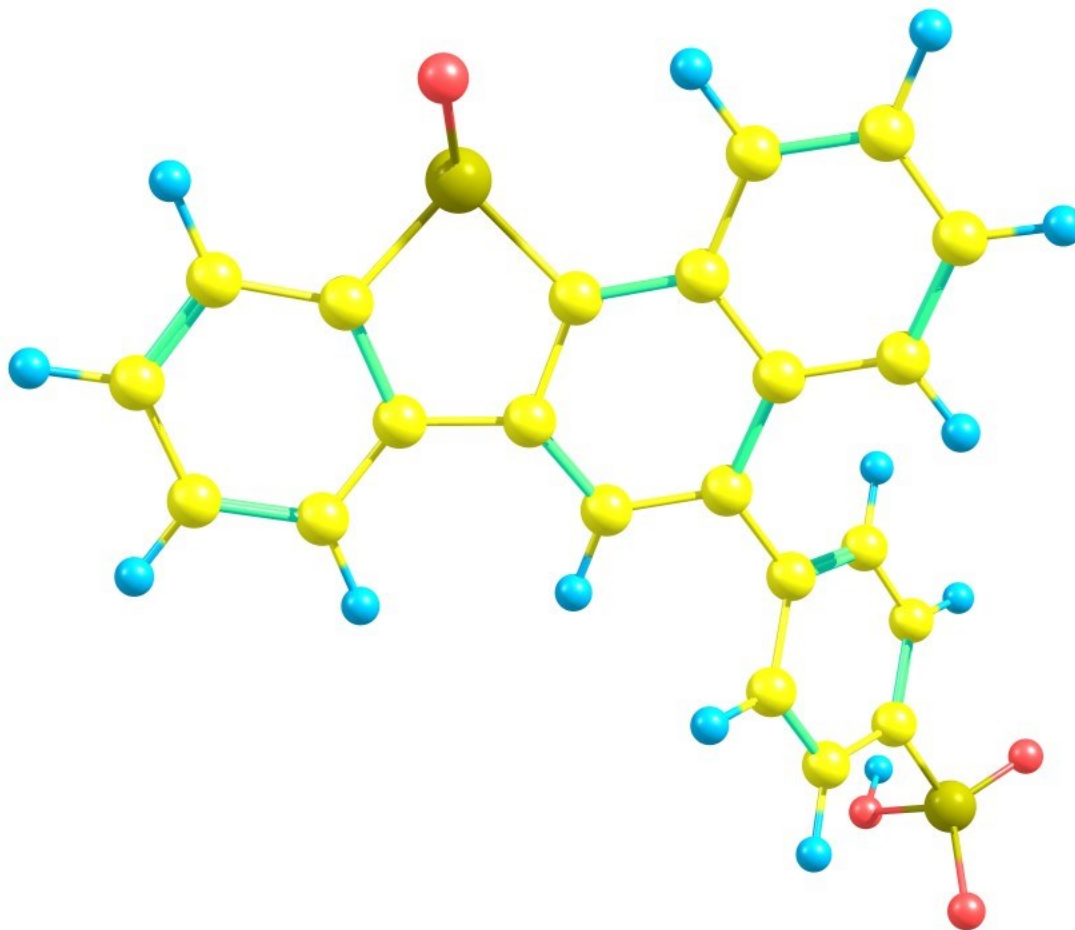


Coordinates

C	-5.415741000000	-3.348146000000	-0.013607000000
C	-5.606061000000	-1.974196000000	-0.136391000000
C	-4.129140000000	-3.878096000000	0.050442000000
H	-6.601292000000	-1.546818000000	-0.191964000000
H	-3.997100000000	-4.951176000000	0.134851000000
C	-4.486876000000	-1.165255000000	-0.162725000000
C	-3.011005000000	-3.051825000000	0.017360000000
H	-2.015786000000	-3.479342000000	0.077106000000
C	-3.188553000000	-1.674561000000	-0.074597000000
H	-6.274476000000	-4.008540000000	0.030359000000
C	-2.744490000000	0.651446000000	-0.085671000000
C	-1.970294000000	1.827094000000	-0.011688000000
C	-2.186654000000	-0.609557000000	-0.048775000000
C	-0.549660000000	1.683095000000	0.077950000000
C	-0.786906000000	-0.742225000000	0.015382000000
H	-0.334860000000	-1.728492000000	-0.000839000000
C	0.024407000000	0.370486000000	0.056354000000
S	-4.512325000000	0.613145000000	-0.432834000000
C	-2.547692000000	3.118497000000	0.022377000000

C	-1.753702000000	4.229185000000	0.137872000000
C	-0.356451000000	4.095641000000	0.246071000000
C	0.229530000000	2.855202000000	0.223060000000
H	-3.627805000000	3.210972000000	-0.004680000000
H	-2.204739000000	5.214801000000	0.168978000000
H	0.260441000000	4.979730000000	0.363906000000
H	1.303489000000	2.766171000000	0.335241000000
O	-5.339914000000	1.335845000000	0.591900000000
C	1.492525000000	0.167660000000	0.067081000000
C	2.301942000000	0.724089000000	-0.928932000000
C	3.668065000000	0.489129000000	-0.946409000000
C	4.227544000000	-0.305773000000	0.046495000000
C	3.450577000000	-0.862347000000	1.054266000000
C	2.084907000000	-0.623534000000	1.056288000000
H	1.852078000000	1.335290000000	-1.703509000000
H	4.301336000000	0.927542000000	-1.709011000000
H	3.917506000000	-1.453707000000	1.833095000000
H	1.467805000000	-1.036631000000	1.846686000000
S	5.970372000000	-0.624640000000	0.024500000000
O	6.407328000000	-0.887967000000	1.369539000000
O	6.612646000000	0.350910000000	-0.831259000000
O	6.051862000000	-2.051902000000	-0.749228000000
H	6.291642000000	-1.877039000000	-1.667696000000

D2) 5-neopentylsulfonatephenylbenzonaphtho(1,2-d)thiophene (3) (Charge = 0, Multiplicity = 3)



Coordinates

C	5.423909000000	-3.322629000000	0.063777000000
C	5.568074000000	-1.960933000000	0.303720000000
C	4.161104000000	-3.865412000000	-0.201493000000
H	6.542007000000	-1.527942000000	0.504410000000
H	4.066906000000	-4.929859000000	-0.385909000000
C	4.436734000000	-1.166302000000	0.269806000000
C	3.035529000000	-3.063217000000	-0.243379000000
H	2.066412000000	-3.493538000000	-0.472102000000
C	3.161971000000	-1.686251000000	-0.012400000000
H	6.294420000000	-3.968527000000	0.083935000000
C	2.734619000000	0.655431000000	0.185825000000
C	1.971095000000	1.833555000000	0.039057000000
C	2.151296000000	-0.666017000000	-0.050205000000
C	0.560847000000	1.698705000000	-0.190779000000
C	0.803361000000	-0.777259000000	-0.237877000000
H	0.351543000000	-1.759967000000	-0.319742000000
C	-0.041980000000	0.362210000000	-0.209663000000
S	4.462855000000	0.607957000000	0.607276000000

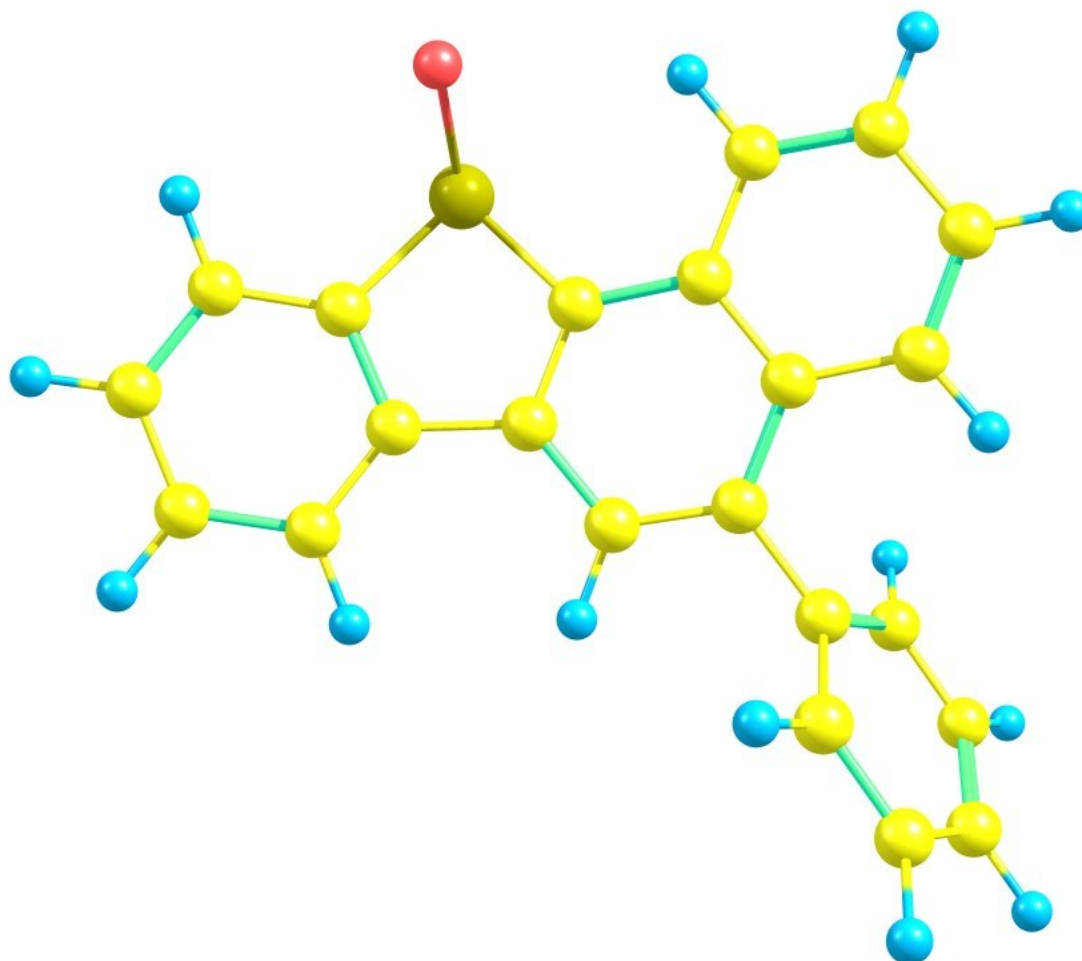
C	2.552417000000	3.111805000000	0.046615000000
C	1.777926000000	4.255913000000	-0.171423000000
C	0.433911000000	4.127550000000	-0.438605000000
C	-0.162785000000	2.848459000000	-0.464708000000
H	3.625074000000	3.199171000000	0.178680000000
H	2.247889000000	5.232550000000	-0.155684000000
H	-0.175092000000	5.000956000000	-0.641938000000
H	-1.212744000000	2.771404000000	-0.720275000000
O	5.333316000000	1.329950000000	-0.386637000000
C	-1.480594000000	0.158414000000	-0.172438000000
C	-2.322116000000	0.957013000000	0.629045000000
C	-3.681770000000	0.718546000000	0.701698000000
C	-4.232703000000	-0.320013000000	-0.043888000000
C	-3.434938000000	-1.124266000000	-0.854247000000
C	-2.075502000000	-0.889042000000	-0.907124000000
H	-1.890130000000	1.746094000000	1.232774000000
H	-4.322417000000	1.334888000000	1.321865000000
H	-3.889641000000	-1.905170000000	-1.452472000000
H	-1.461831000000	-1.494551000000	-1.564412000000
S	-5.968121000000	-0.635599000000	0.047692000000
O	-6.393937000000	-1.241341000000	-1.186343000000
O	-6.623078000000	0.522518000000	0.620044000000
O	-6.056878000000	-1.819575000000	1.159936000000
H	-6.275962000000	-1.413201000000	2.007507000000

D3) Energy calculations for 3

Charge	Multiplicity	Sum of electronic and thermal free energies (hartree/particle)
0	1	-1942.539267
0	3	-1942.462598
Difference		0.076669

in kcal/mol = 0.076669 * 627.5 = 48.1 kcal/mol

E1) 5-phenylbenzonaptho(1,2-d)thiophene (Charge = 0, Multiplicity = 1)

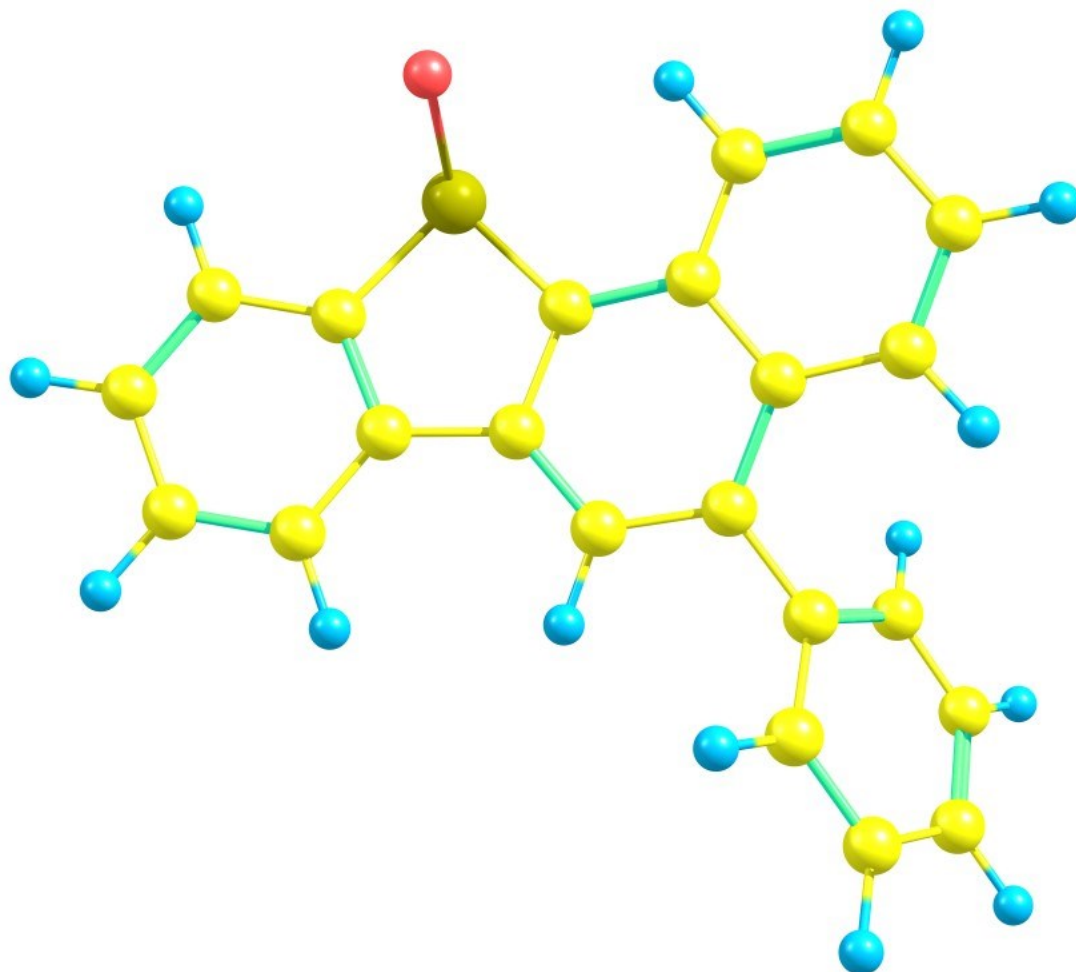


Coordinates

C	4.761393000000	-2.339283000000	0.007195000000
C	4.576375000000	-0.962362000000	0.103801000000
C	3.663876000000	-3.196446000000	-0.020698000000
H	5.420412000000	-0.281754000000	0.131222000000
H	3.823876000000	-4.267208000000	-0.084841000000
C	3.281453000000	-0.483395000000	0.141994000000
C	2.365355000000	-2.700250000000	0.022974000000
H	1.520665000000	-3.380201000000	-0.008699000000
C	2.166768000000	-1.324350000000	0.089029000000
H	5.765491000000	-2.745183000000	-0.044180000000
C	1.112754000000	0.795231000000	0.074938000000
C	0.046829000000	1.715269000000	0.007950000000
C	0.914260000000	-0.569710000000	0.068296000000
C	-1.282564000000	1.188516000000	-0.046850000000
C	-0.397815000000	-1.078094000000	0.036648000000
H	-0.565585000000	-2.149476000000	0.071679000000
C	-1.482947000000	-0.230051000000	-0.001791000000
S	2.831376000000	1.242569000000	0.378173000000

C	0.250111000000	3.114368000000	-0.051104000000
C	-0.818350000000	3.965707000000	-0.155841000000
C	-2.128699000000	3.455573000000	-0.227482000000
C	-2.353977000000	2.102983000000	-0.180806000000
H	1.264744000000	3.497255000000	-0.050075000000
H	-0.653224000000	5.036401000000	-0.205372000000
H	-2.965402000000	4.137328000000	-0.335132000000
H	-3.364035000000	1.719624000000	-0.261385000000
O	3.415953000000	2.132514000000	-0.683028000000
C	-2.843586000000	-0.819952000000	0.002924000000
C	-3.761319000000	-0.511174000000	1.011723000000
C	-5.016166000000	-1.104946000000	1.031382000000
C	-5.376762000000	-2.012703000000	0.041754000000
C	-4.472721000000	-2.326932000000	-0.965386000000
C	-3.215044000000	-1.737109000000	-0.983194000000
H	-3.478081000000	0.185715000000	1.793675000000
H	-5.713114000000	-0.862004000000	1.826422000000
H	-6.358478000000	-2.473584000000	0.056683000000
H	-4.747985000000	-3.030648000000	-1.743721000000
H	-2.514253000000	-1.972836000000	-1.777553000000

E2) 5-phenylbenzonaptho(1,2-d)thiophene (Charge = 0, Multiplicity = 3)



Coordinates

C	4.767703000000	-2.302470000000	0.071133000000
C	4.545712000000	-0.940606000000	0.236967000000
C	3.688406000000	-3.178069000000	-0.108467000000
H	5.373181000000	-0.251686000000	0.368666000000
H	3.878777000000	-4.238064000000	-0.236845000000
C	3.241810000000	-0.478735000000	0.219761000000
C	2.388377000000	-2.711066000000	-0.134273000000
H	1.563868000000	-3.397902000000	-0.293512000000
C	2.143292000000	-1.337687000000	0.024763000000
H	5.780316000000	-2.689604000000	0.081541000000
C	1.108696000000	0.809750000000	0.160108000000
C	0.046870000000	1.729454000000	0.033899000000
C	0.897770000000	-0.632943000000	-0.001692000000
C	-1.284716000000	1.208574000000	-0.105316000000
C	-0.382008000000	-1.107514000000	-0.112571000000
H	-0.559218000000	-2.177622000000	-0.145104000000
C	-1.496822000000	-0.241302000000	-0.082025000000
S	2.808424000000	1.258893000000	0.444629000000

C	0.254891000000	3.117876000000	-0.011577000000
C	-0.816304000000	4.001663000000	-0.183323000000
C	-2.088323000000	3.504768000000	-0.349726000000
C	-2.312699000000	2.109890000000	-0.328088000000
H	1.268763000000	3.497979000000	0.044075000000
H	-0.631650000000	5.069581000000	-0.208939000000
H	-2.926006000000	4.174070000000	-0.510259000000
H	-3.315918000000	1.740513000000	-0.502904000000
O	3.385910000000	2.097608000000	-0.666488000000
C	-2.833751000000	-0.826850000000	-0.012858000000
C	-3.815770000000	-0.322361000000	0.858602000000
C	-5.060830000000	-0.922892000000	0.954462000000
C	-5.368215000000	-2.035614000000	0.177291000000
C	-4.409559000000	-2.548941000000	-0.690702000000
C	-3.159072000000	-1.958912000000	-0.780249000000
H	-3.577314000000	0.524062000000	1.492928000000
H	-5.794259000000	-0.526163000000	1.648473000000
H	-6.345321000000	-2.500338000000	0.249615000000
H	-4.641182000000	-3.410758000000	-1.307584000000
H	-2.428811000000	-2.353307000000	-1.479031000000

E3) Energy calculations for 6

Charge	Multiplicity	Sum of electronic and thermal free energies (hartree/particle)
0	1	-1319.002595
0	3	-1318.925184
Difference		0.077411

in kcal/mol = $0.077411 \times 627.5 = 48.6$ kcal/mol

3. Figures

S1. isoTOP-ABPP workflow scheme

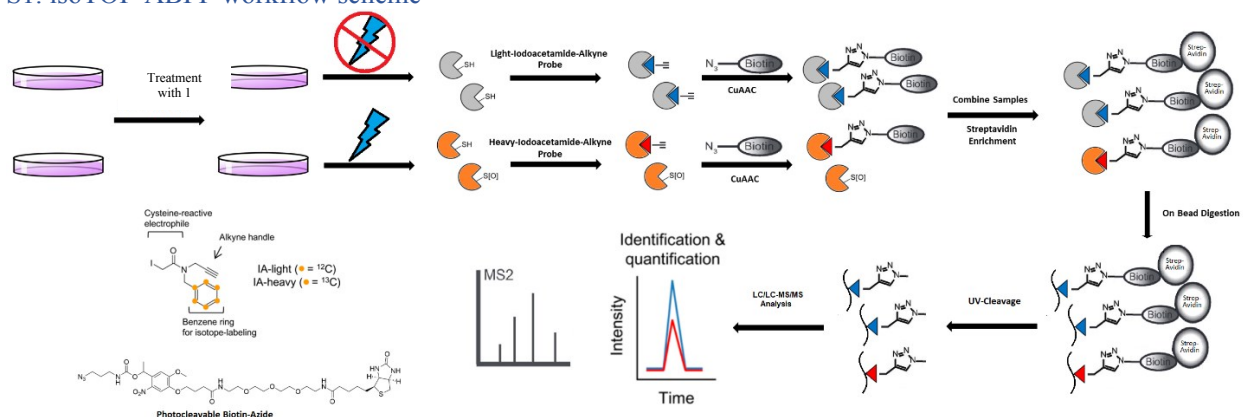


Table S1 – Distribution of peptide sequences based on cysteine reactivity and amino acid length

Cysteine reactivity	Sequence criteria	Single cysteine		Two cysteines			
		Fragment A	Fragment B	Fragment A	Fragment B	Fragment C	Fragment D
Two-fold reduction in labeling (n=461)	Sequence number less than 4	176	131	15	12	12	12
	Sequences	258	303	12	15	15	15
Increase in labeling (n=200)	Sequence number less than 4	58	61	2	2	2	2
	Sequences	138	135	2	2	2	2
No difference in labeling (n=80)	Sequences number less than 4	27	27	x	x	x	x
	Sequences	53	53	x	x	x	x

Table S2 - Summary of ten peptides with the highest reduction in labelling as a result of treatment with 1 and irradiation

Protein	Peptide Sequence	Log ₂ (R _{L:H})
TK1 Thymidine kinase, cytosolic	YHSVC*R	4.32
DCAFDDDB1- and CUL4-associated factor 7	VPC*TPVAR	4.32
ILF3 Interleukin enhancer-binding factor 3	VLEC *LASGIVMPDGSGIYDPC*EK	4.32
PKM Pyruvate kinase isozymes M1/M2	PVIC*ATQMLESNIK	4.32
LRPPRC Leucine-rich PPR motif containing protein, mitochondrial	AILQENG*LSDSMDMFSQAGLR	4.32
ANKHD1 Ankyrin repeat and KH domain-containing protein 1	DTVSLHQQC*SHR	4.32
TUBB Tubulin beta chain	VSDTVVEPYNATLSVHQLVENTDETYC*IDNEALYDIC*FR	3.22
LRRC40 Leucine-rich repeat-containing protein 40	NLSEVPQC*VWR	2.92
TUBB Tubulin beta chain	LTTPTYGDLNHLVSATMSGVTTC*LR	2.80
RRP8 Ribosomal RNA-processing protein 8	NPVHC*FDLASLDPR	2.75

Table S3- Summary of ten peptides with the least difference in labeling as a result of treatment with 1 and irradiation

Protein	Peptide sequence	Log ₂ (R _{L:H})
HSPA4 Heat shock 70 kDa protein 4	FDEVLVNHFC*EEFGK	-0.0097
EIF3CL Eukaryotic translation initiation factor 3 subunit	C*LEEFELLGK	-0.0048
SEC13 Protein SEC13 homolog	FASGGC*DNLIK	-0.0048

SF3A1 Splicing factor 3A subunit 1	EVLDQVC*YR	-0.0048
SMU1 WD40 repeat-containing protein SMU1	NPEHFVVC*NR	0.0000
HNRNPL Heterogeneous nuclear ribonucleoprotein L	ASLNGADIYSGC*CTLK	0.0000
ILVBL Acetolactate synthase-like protein	EQVPSLGSNVAC*GLAYTDYHK	0.0048
FLNB Filamin-B	C*LATGPGIASTVK	0.0048
IARS Isoleucine--tRNA ligase, cytoplasmic	NNDLC*YWVPELVR	0.0048
ANLN Actin-binding protein anillin	SC*EGQNPELLPK	0.0072

Table S4 - Summary of ten peptides with the lowest reduction in labelling (increase in labelling) as a result of treatment with 1 and irradiation

Protein	Peptide sequence	Log ₂ (R _{L:H})
LRRC47 Leucine-rich repeat-containing protein 47	MVSGC*QTR	-6.06
RAP1B Ras-related protein Rap-1b	C*DLEDER	-2.60
RAP1B Ras-related protein Rap-1b	QVEVDAQQC*MLEILDTAGTEQFTAMR	-2.51
CDC42 Cell division control protein 42 homolog	YVEC*SALTQK	-2.35
SDHA Succinate dehydrogenase	TLNEADC*ATVPPAIR	-2.25
ECHS1 Enoyl-CoA hydratase, mitochondrial	IC*PVETLVEEAIQC*AEK	-2.08
RAB8A Ras-related protein Rab-8A	C*DVNDK	-1.94
RAB10 Ras-related protein Rab-10	C*DMDDK	-1.94
RAB21 Ras-related protein Rab-21	GIEELFLDLC*K	-1.91
MDH2 Malate dehydrogenase, mitochondrial	SQETEC*TYFSTPLLLGK	-1.84

Table S5 - Solvent accessible surface area of sulfur atoms in cysteines within peptides with highest reduction in labeling

ID	Description	Cysteine position	PDB file ID	Chain	Cysteine	Solvent accessibility for Sulfur (Å ²)
Q12906	ILF3 Interleukin enhancer-binding factor 3	278 295	4ATB 4(Q9Z1X4)	B	278	10.29
				D		10.42
				B	295	2.83
				D		1.54
P04183	TK1 Thymidine kinase, cytosolic	185	1XBT ⁵	A	185	2.7
				B		5.28
				C		3.73
				D		3.35
				E		4.5

				F		4.25
				G		4.5
				H		4.25
P14618	PKM Pyruvate kinase isozymes M1/M2	326	3BJT ⁶	A B C D	326	0
			1ZJH ⁷	A	325	0
P07437	TUBB Tubulin beta chain	201 211	5N5N ⁸	A B E F	203	3.09
				C D		2.7
				A B	213	0.39
		C D		0.26		
		E F		0.52		
		239		A B C D E F	241	0
O43159	RRP8 Ribosomal RNA-processing protein 8	2ZFU ⁹	A	332	0.16	
			B		0	

Table S6 – Solvent accessible surface area of sulfur atoms in cysteines within peptides that exhibited least reactivity with O(³P)

ID	Description	Cysteine position	PDB file ID	Chain	Cysteine	Solvent accessibility for Sulfur (Å ²)
B5ME19	EIF3CL Eukaryotic translation initiation factor 3 subunit	79	6ZP4 (Q99613) ¹⁰	C	79	0.77
P55735	SEC13 Protein SEC13 homolog	187	3BG1 ¹¹	A	187	3.35
				D		2.83
				E		2.96
				H		2.57
Q15459	SF3A1 Splicing factor 3A subunit 1	244	6FF7 ¹²	p	244	65.11
Q2TAY7	SMU1 WD40 repeat-containing protein SMU1	416	5O9Z ¹³	L	416	0
P14866	HNRNPL Heterogeneous nuclear ribonucleoprotein L	260	2MQP (FILQ48) ¹⁴	A	257	40.79
O75369	FLNB Filamin-B	1617	2DMB ¹⁵	A	14	1.03

Table S7 - Solvent accessible surface area of sulfur atoms in cysteines within peptides that exhibited an increase in labeling

ID	Description	Cysteine position	PDB file ID	Chain	Cysteine	Solvent accessibility for Sulfur (Å ²)
P61224	RAP1B Ras-related protein Rap-1b	118	4HDO ¹⁶	B	118	7.08
P61224	RAP1B Ras-related protein Rap-1b	51	4HDO ¹⁶	B	51	0.77
P60953	CDC42 Cell division control protein 42 homolog	157	2NGR ¹⁷	A	157	0.13
P31040	SDHA Succinate dehydrogenase	654	6VAX ¹⁸	A,C	654	0.64
P30084	ECHS1 Enoyl-CoA hydratase, mitochondrial	213		A B C D E	213	0
				F		0.39
				A		13.12
		225	2HW5 ¹⁹	B	225	11.32
				C		14.03
				D		14.8
				E		16.6
				F		13.51
		123	6RIR ²⁰	A	123	6.18
				B		5.53
P61026	RAB10 Ras-related protein Rab-10	124	5LPN ²¹	A	124	1.42
				C		10.42
Q9UL25	RAB21 Ras-related protein Rab-21	177	1Z0I ²²	A	177	0
P40926	MDH2 Malate dehydrogenase, mitochondrial	285	4WLE ²³	A B C D	285	0

Appendix

Supporting Information excel workbook with protein analysis

Sheet 1 - CysAnn_L-UV_H+UV_LHratios – $\text{Log}_2(R_{L:H})$ for 1 treatment of HeLa cells in UV

Sheet 2 - Control_L&H_noUV_LHratios - $\text{Log}_2(R_{L:H})$ for 1 treatment of HeLa cells in No-UV

Sheet 3 – Seq (two-fold red in label) – peptides with $\text{Log}_2(R_{L:H}) > 1$ in Sheet 1

Sheet 4 – 200 seq (increase in label) – top 200 peptides with highest increase in labeling ($\text{Log}_2(R_{L:H}) \leq 0$)

Sheet 5 - Seq (no reactivity) - peptides with $-0.05 < \text{Log}_2(R_{L:H}) < 0.05$

Sheet 6 - Segmentation for 461 peptides – Location and sub-cellular location of peptides in Sheet 3

Sheet 7 – Mem count increase label – Location and sub-cellular location of peptides in Sheet 4

Sheet 8 – Segmentation of no reactive seq - Location and sub-cellular location of peptides in Sheet 5

Sheet 9 – 2.9% increase in label mem loc – Peptides with highest increase in labeling segmented by location

Sheet 10 - WebLogo (oxidized) - Fragments for peptides in Sheet 3

Sheet 11 - WebLogo (Not Oxidized) – Fragments for peptides in Sheet 4

Sheet 12 – WebLogo (No reactivity) – Fragments for peptides in Sheet 5

Sheet 13 – Cysteine SA structure – Details on crystal structures used to determine Cys solvent accessibility

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