

Catalytic Enantioselective Synthesis of 3,4-Unsubstituted Thiochromenes through Sulfa-Michael/Julia-Kocienski Olefination Cascade Reaction

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SUPPORTING INFORMATION

- | | |
|---|------|
| 1. Optimization of catalyst and reaction conditions | S-2 |
| 2. NMR Spectra and HPLC Chromatograms | S-4 |
| 3. Detection of reactive intermediates | S-63 |
| 4. Single crystal X-ray diffraction analysis of 3ai and 3ak | S-65 |

Optimization of catalyst and reaction conditions:

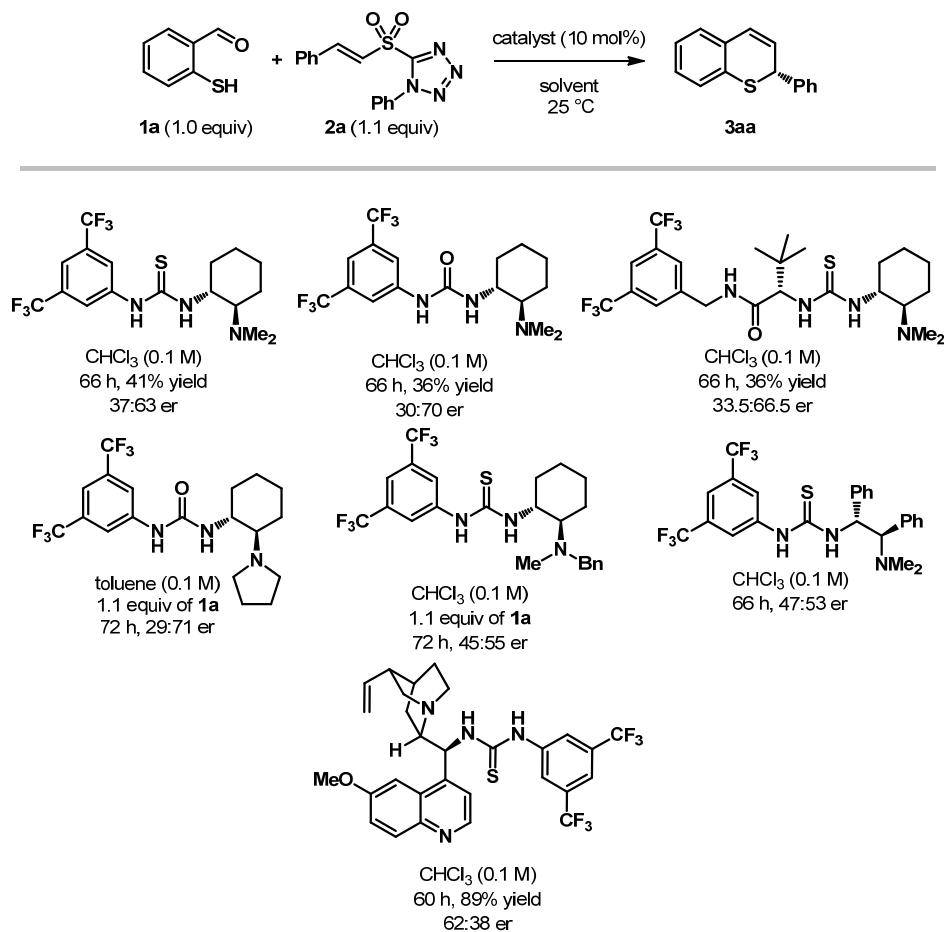


Figure S1. Bifunctional catalyst screening

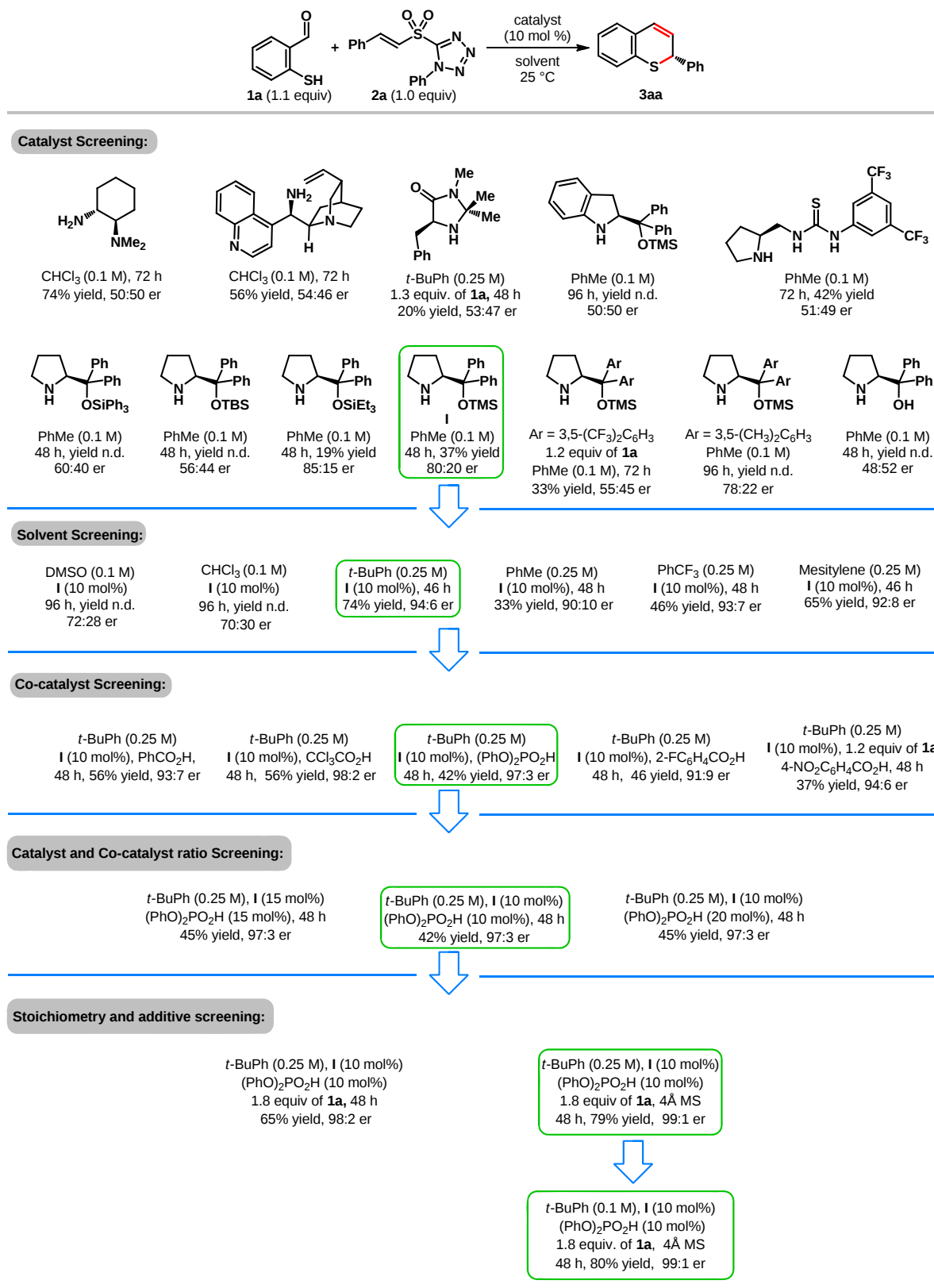
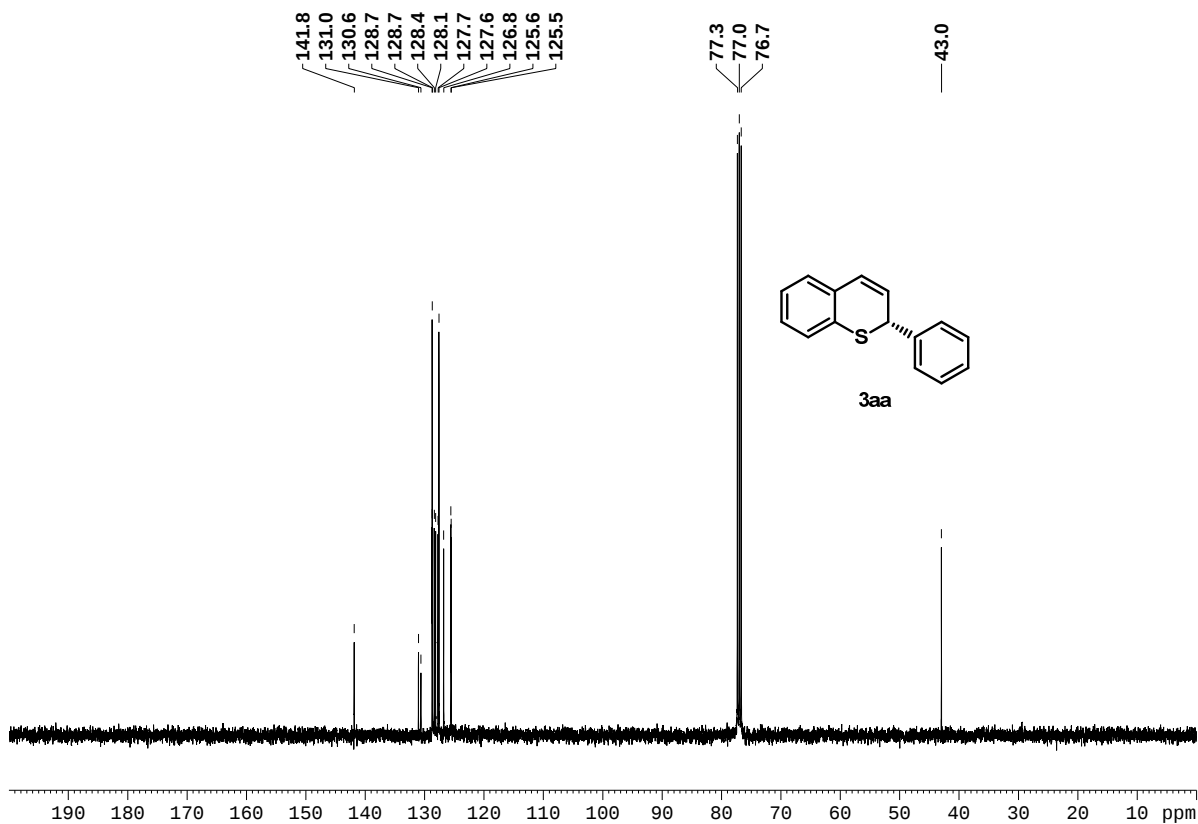
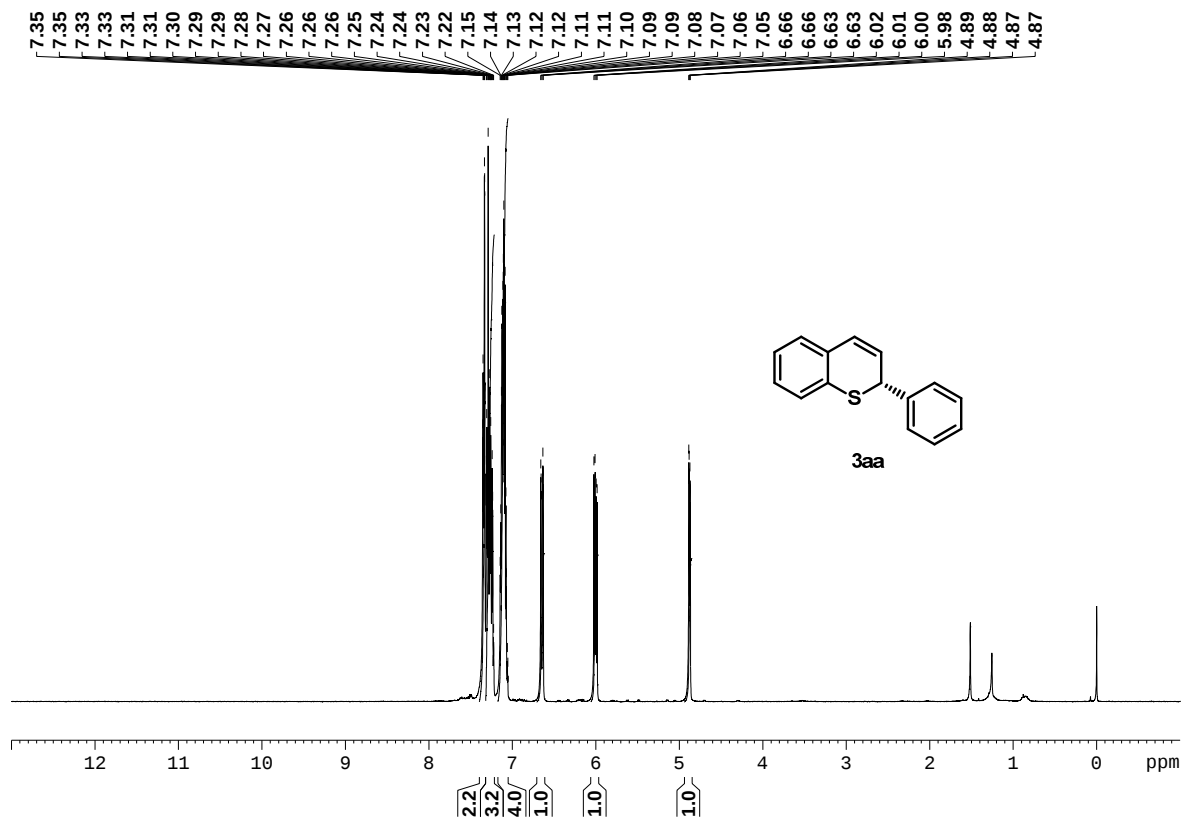
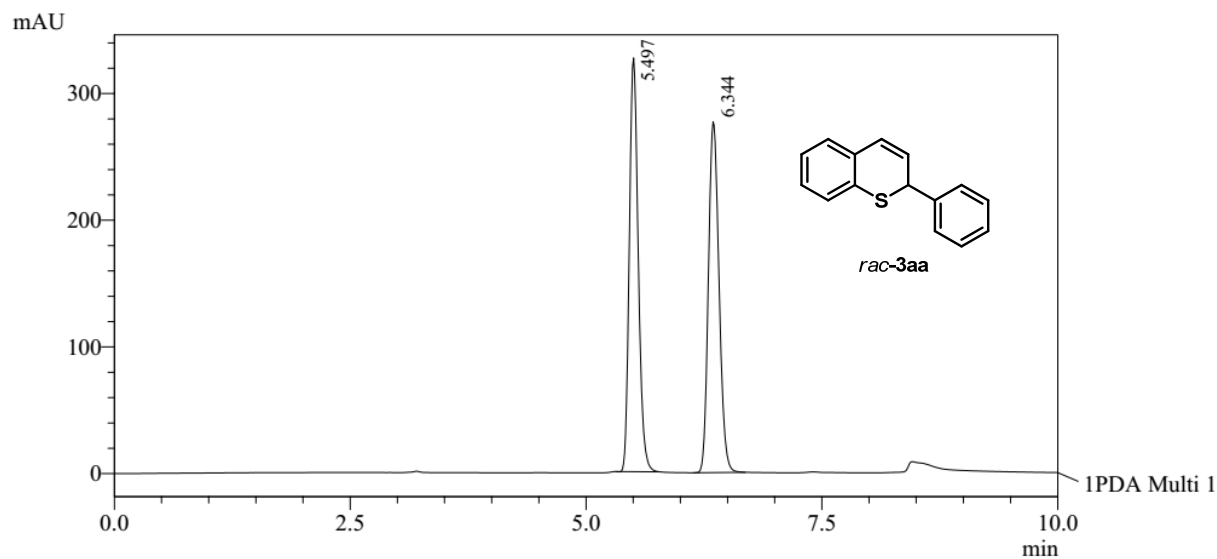


Figure S2. Optimization of reaction conditions with amine catalyst



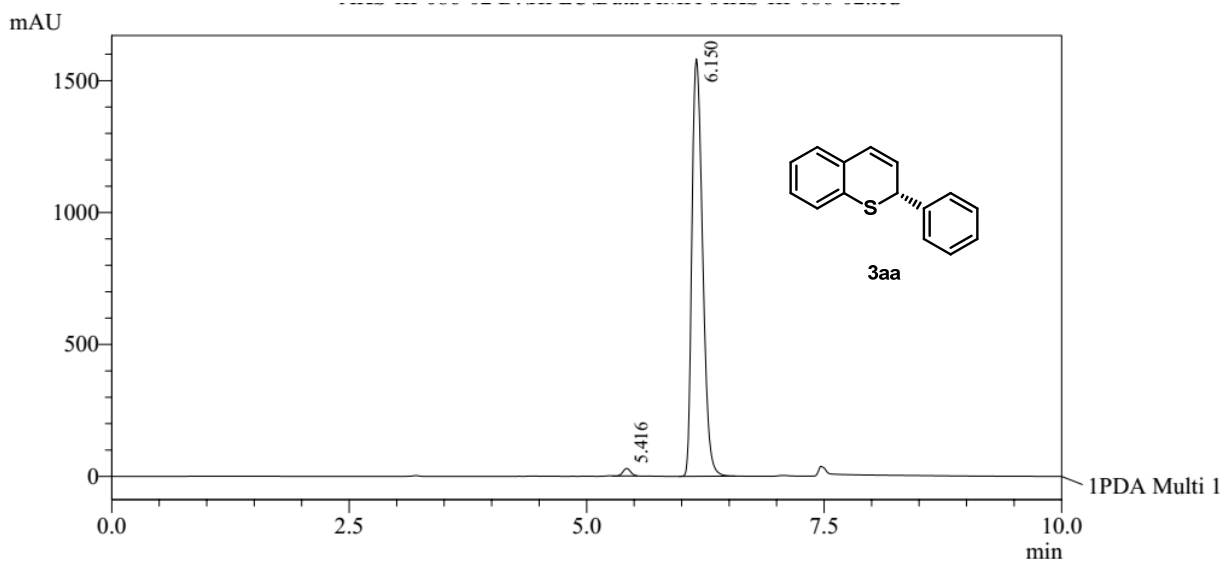


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.497	2145121	49.809
2	6.344	2161532	50.191
Total		4306652	100.000

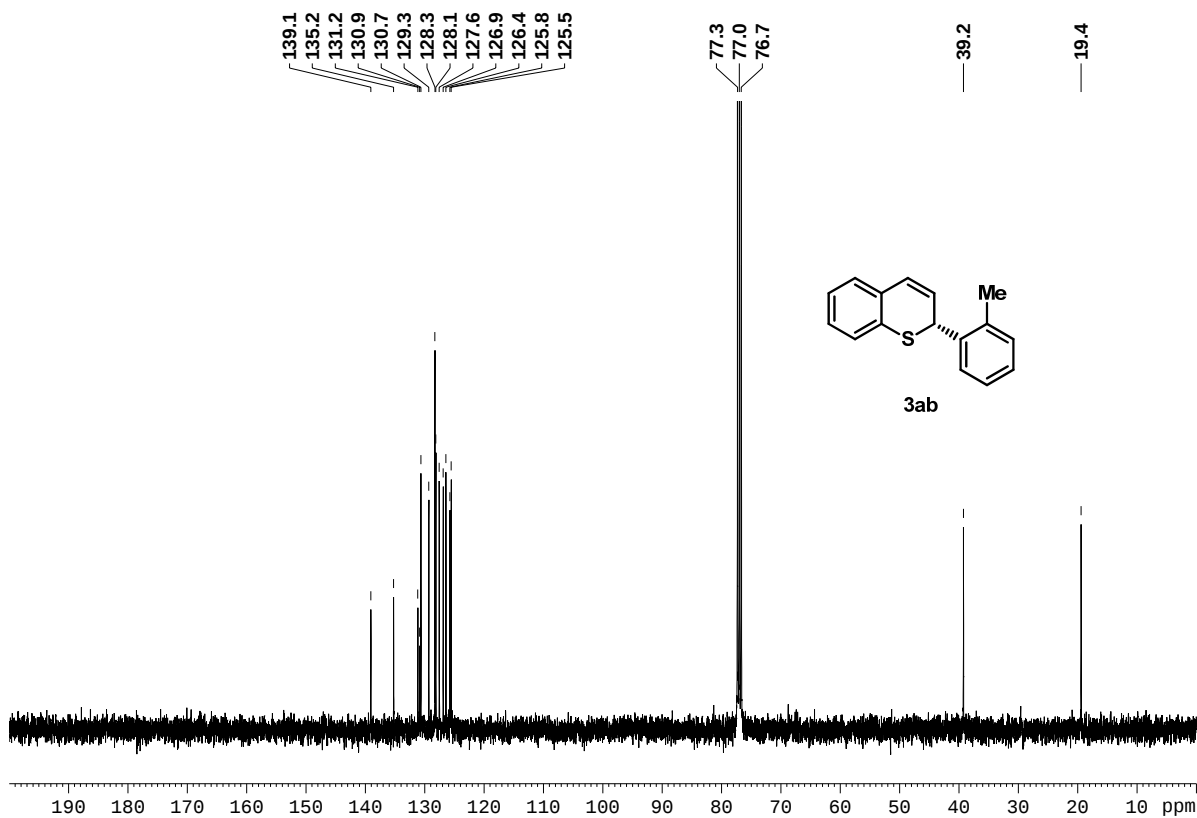
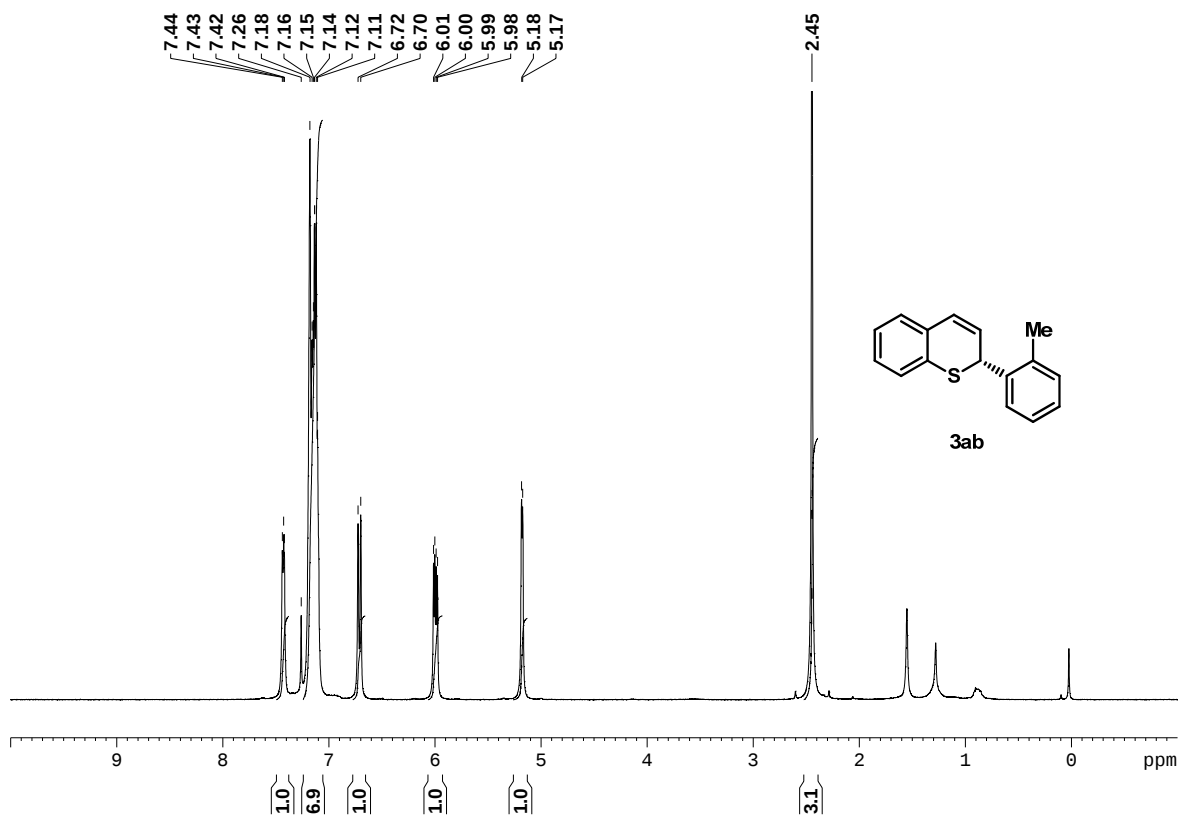
Daicel Chiralpak IF column (80:20 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

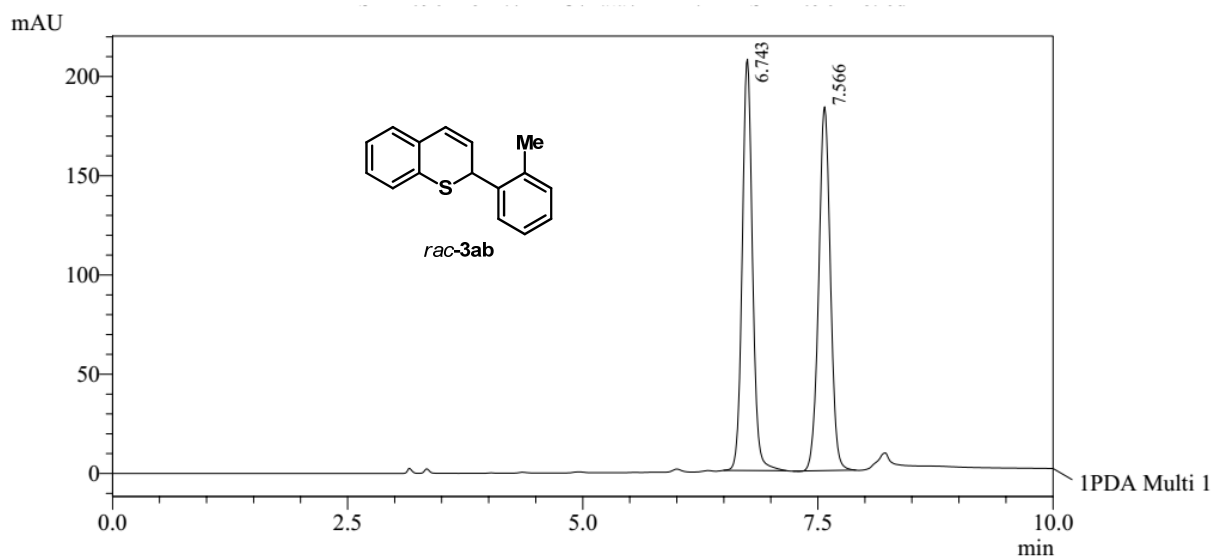


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.416	155202	1.234
2	6.150	12426083	98.766
Total		12581286	100.000





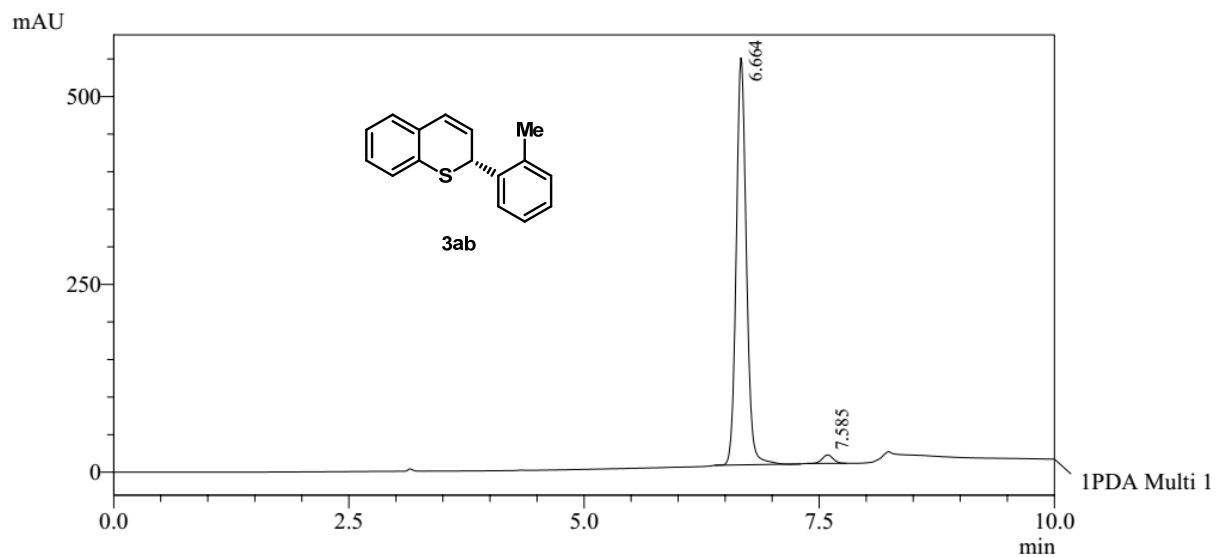
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.743	1573025	49.808
2	7.566	1585129	50.192
Total		3158154	100.000

Daicel Chiralpak IB column (80:20 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

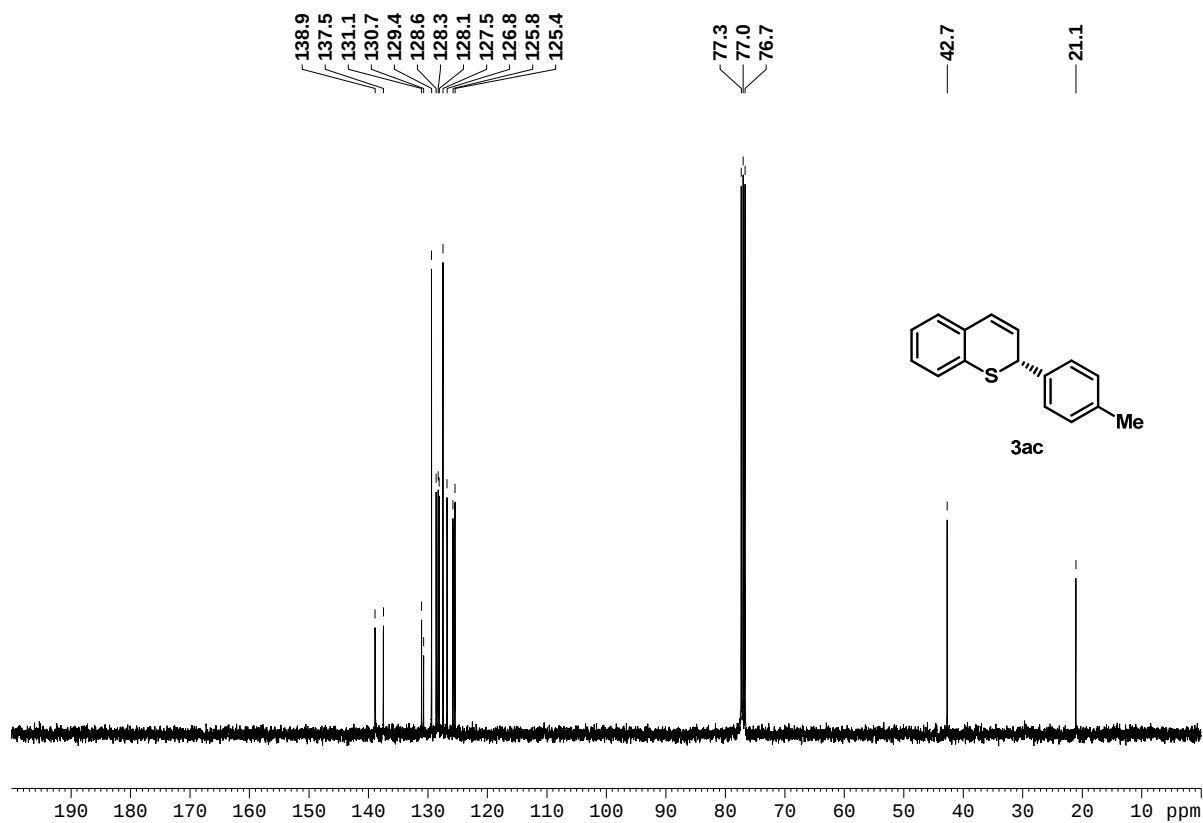
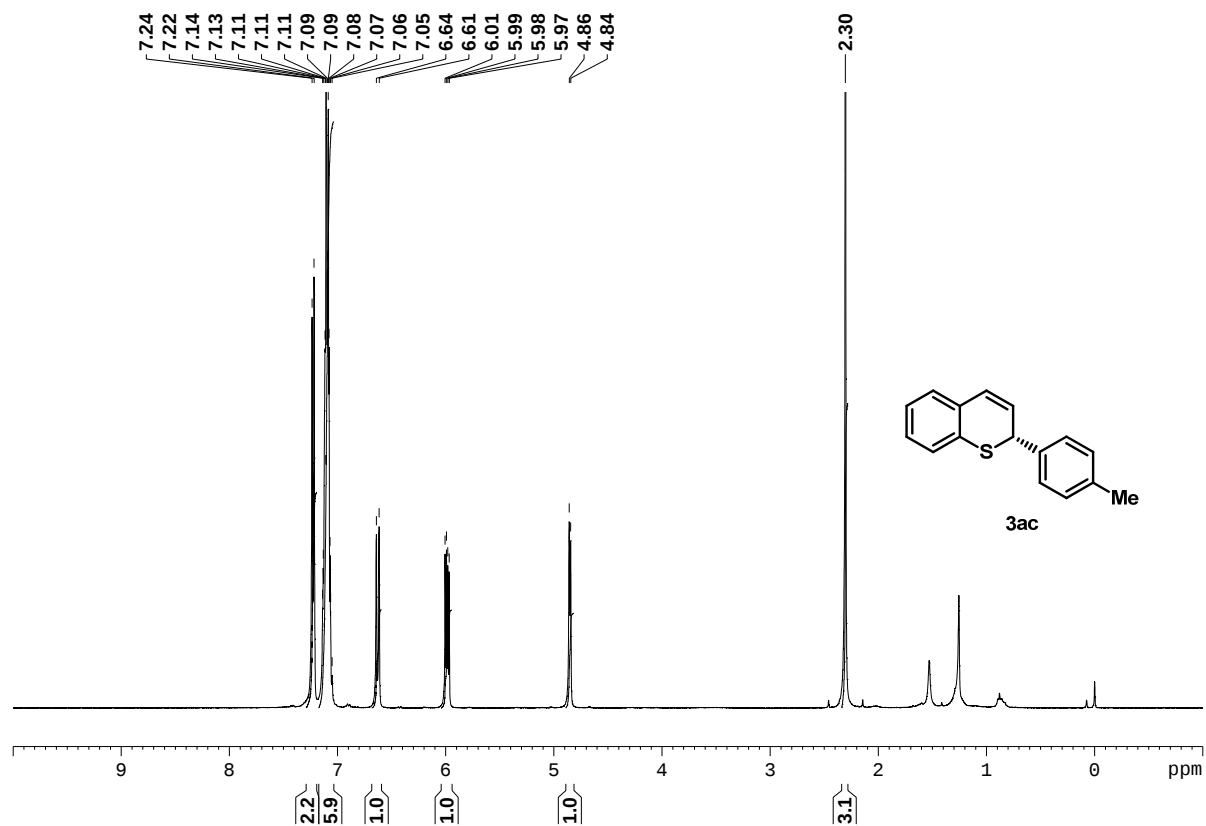


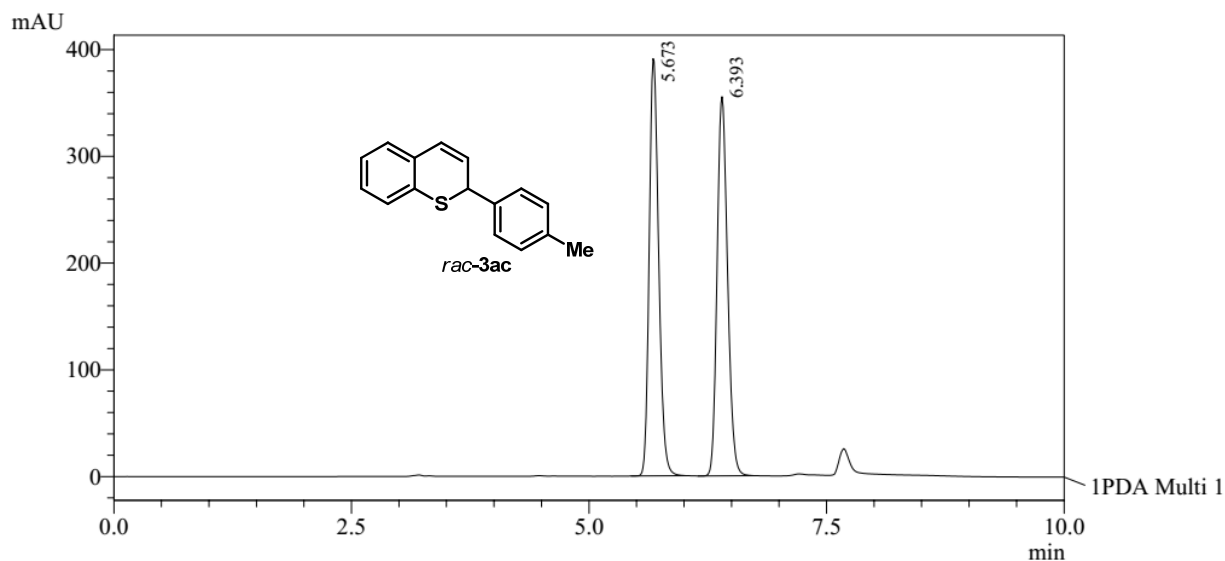
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.664	4060821	97.816
2	7.585	90676	2.184
Total		4151498	100.000





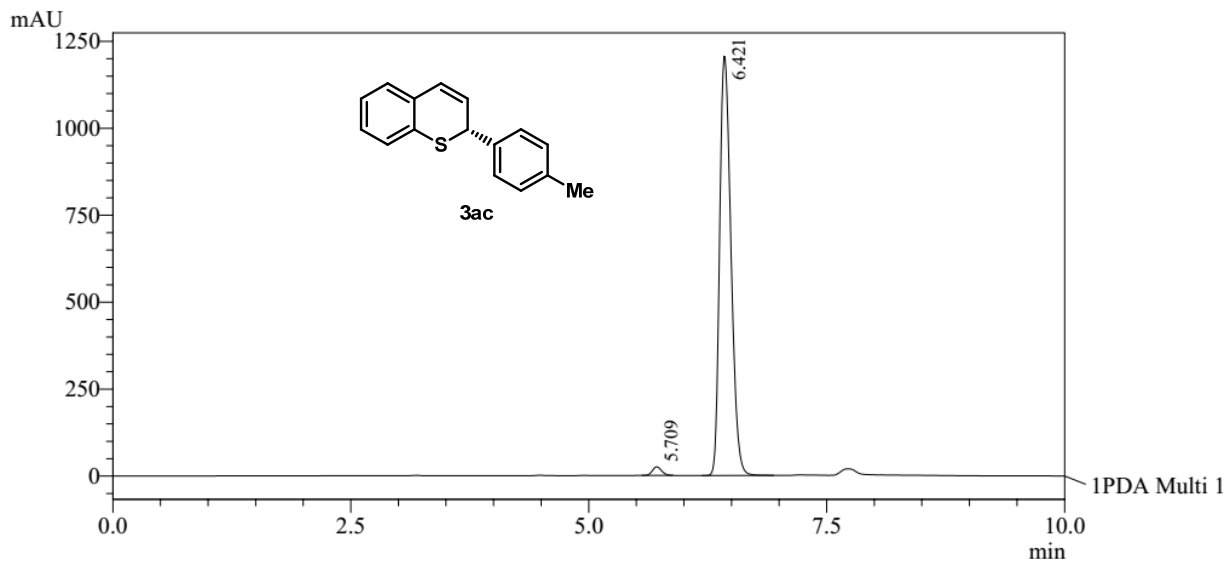
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.673	2714428	49.958
2	6.393	2718979	50.042
Total		5433406	100.000

Daicel Chiralpak IF column (80:20 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

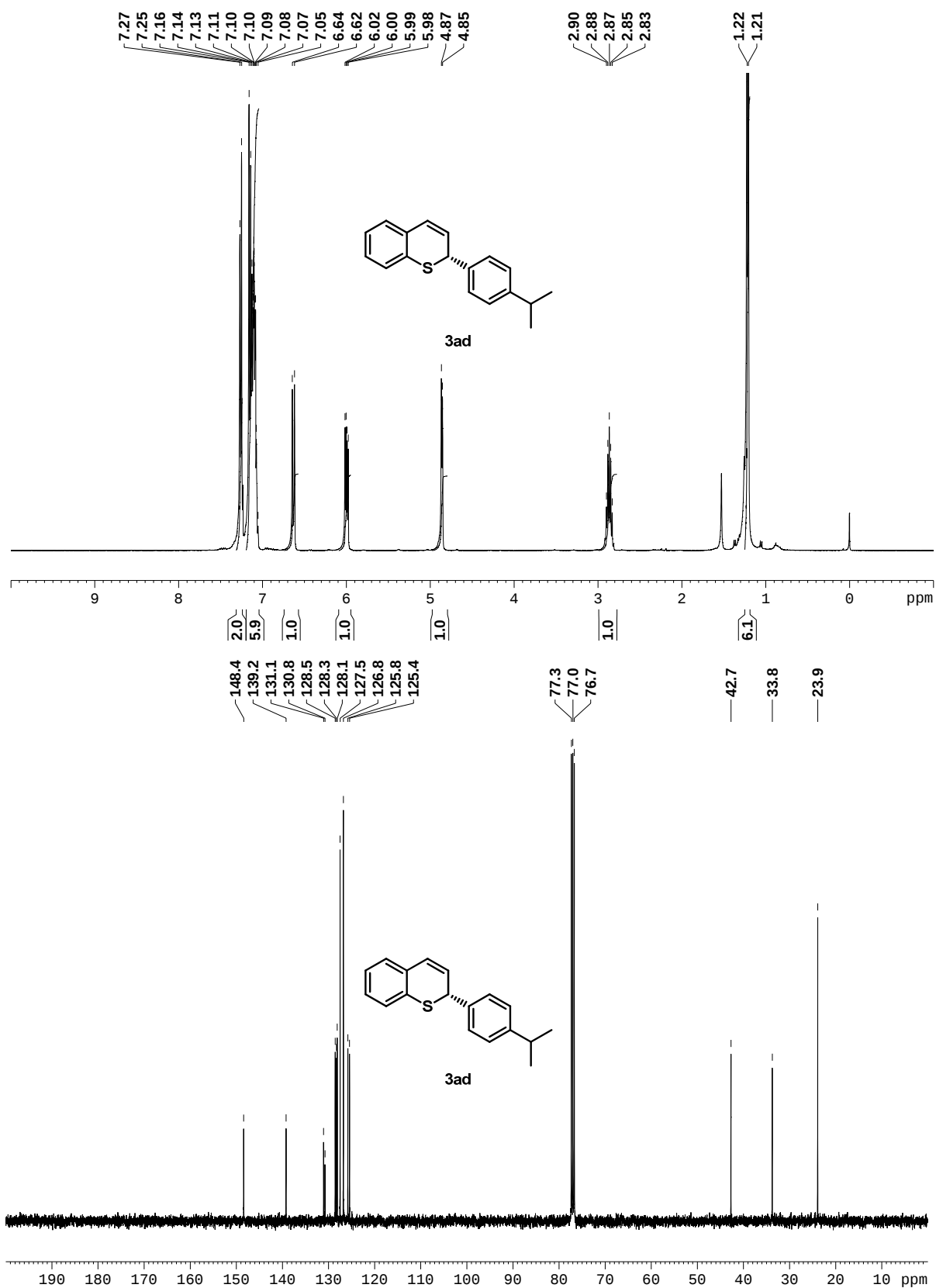


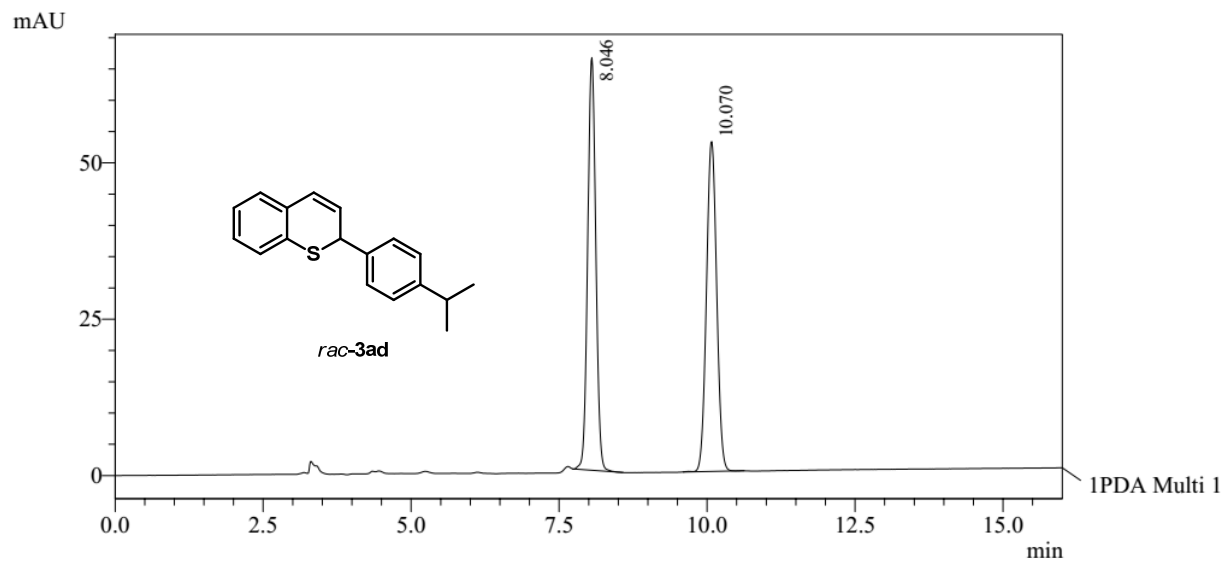
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.709	165267	1.627
2	6.421	9993421	98.373
Total		10158688	100.000





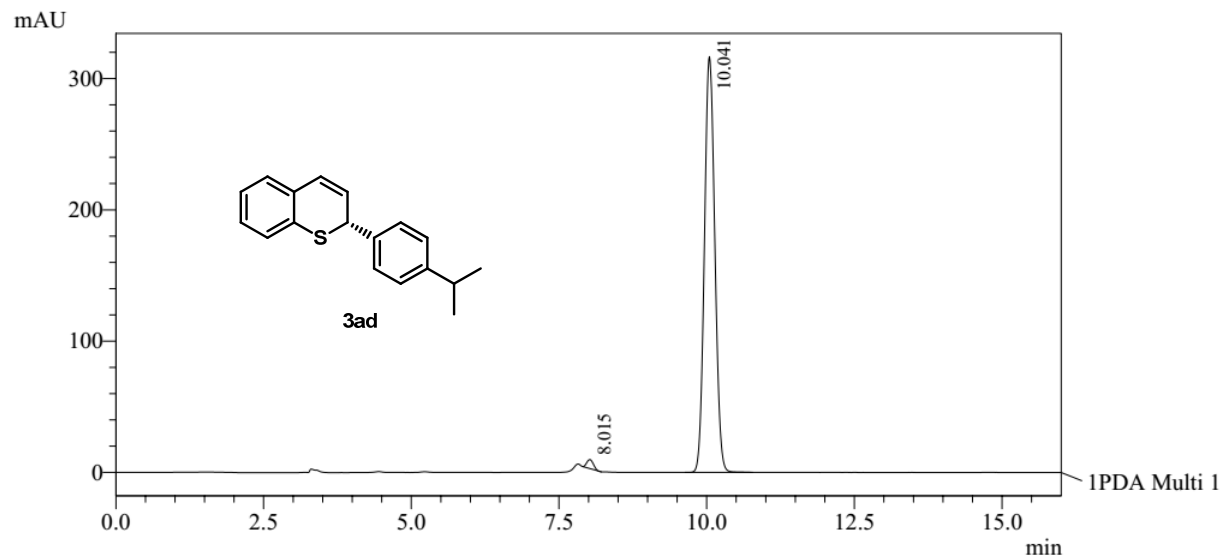
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.046	628147	49.892
2	10.070	630872	50.108
Total		1259019	100.000

Phenomenex Cellulose-3 column (97:3 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

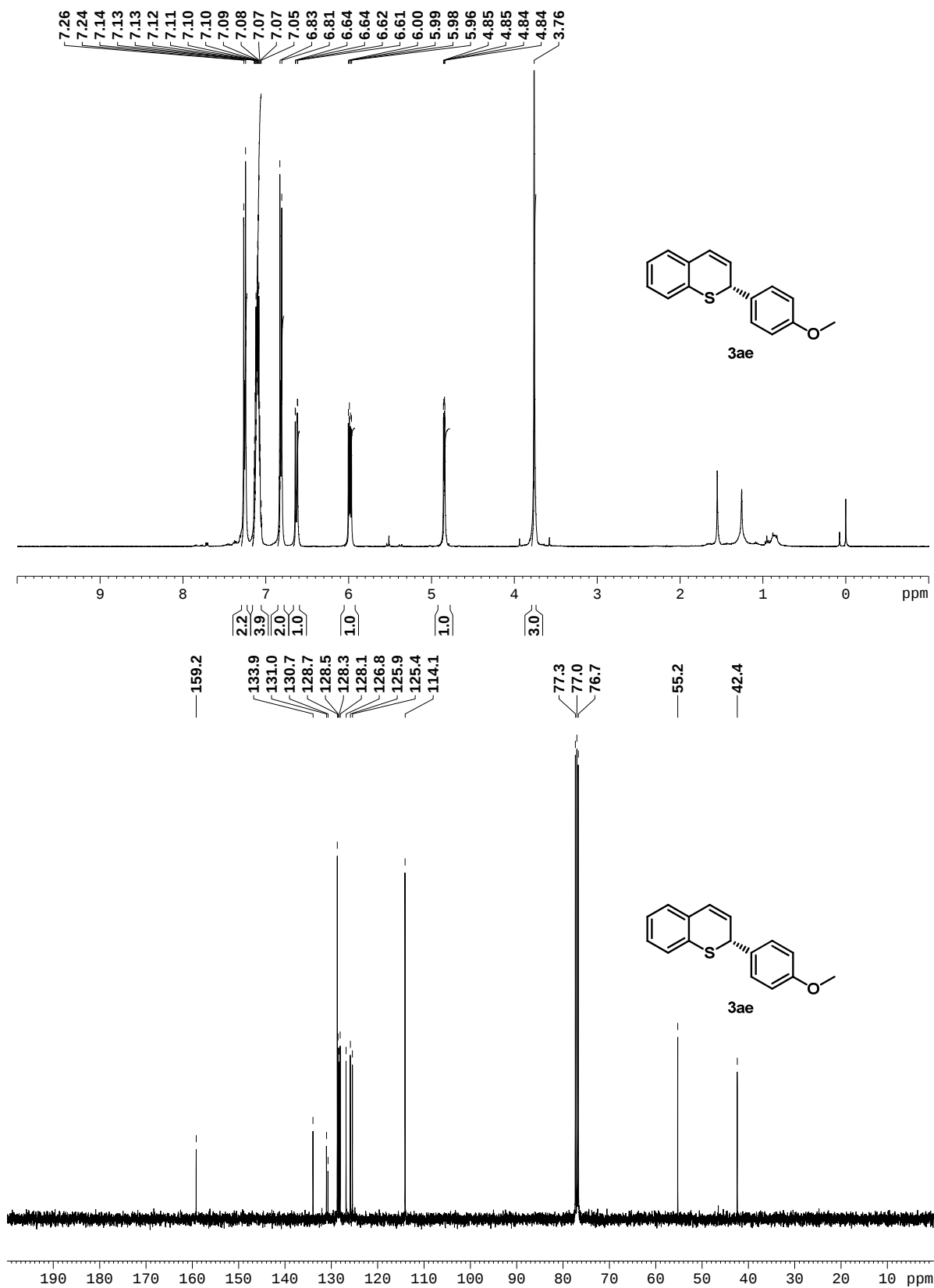


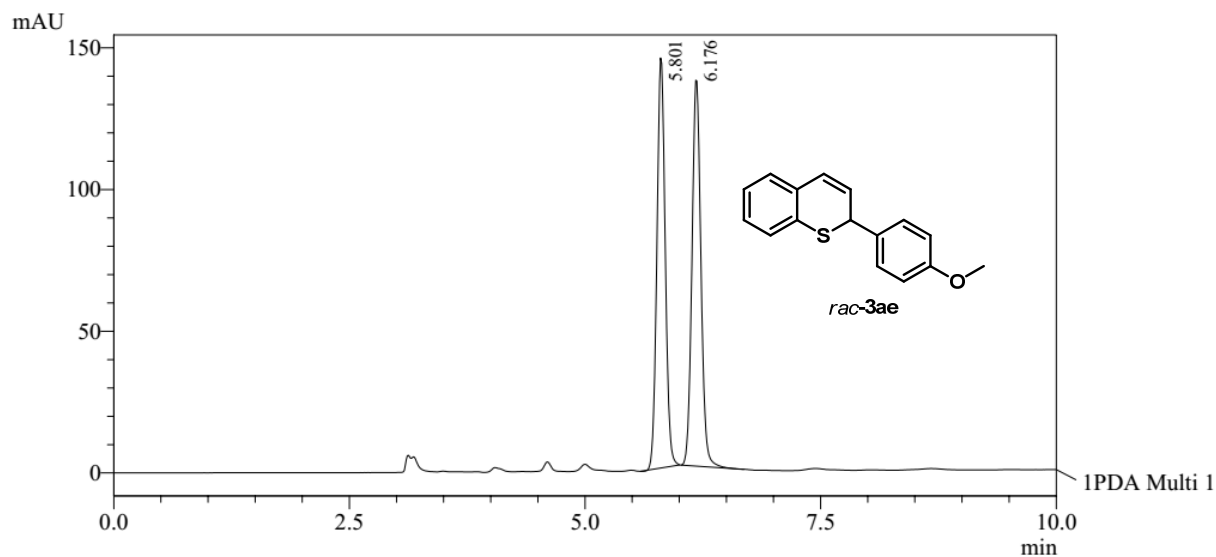
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.015	50141	1.280
2	10.041	3866619	98.720
Total		3916761	100.000





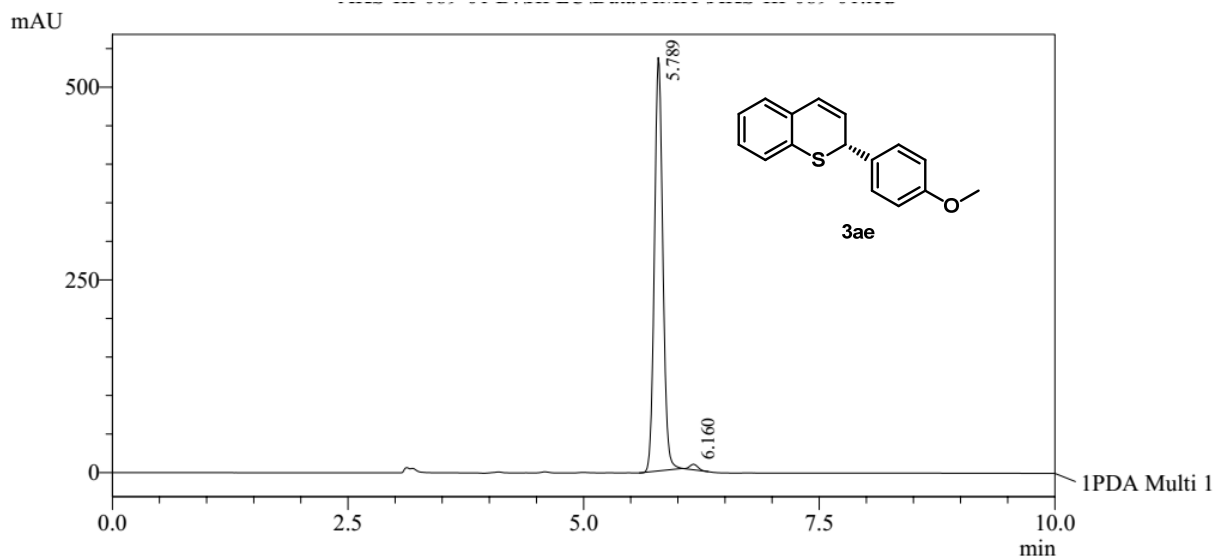
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.801	887248	49.571
2	6.176	902611	50.429
Total		1789858	100.000

Daicel Chiralpak IB column (95:5 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

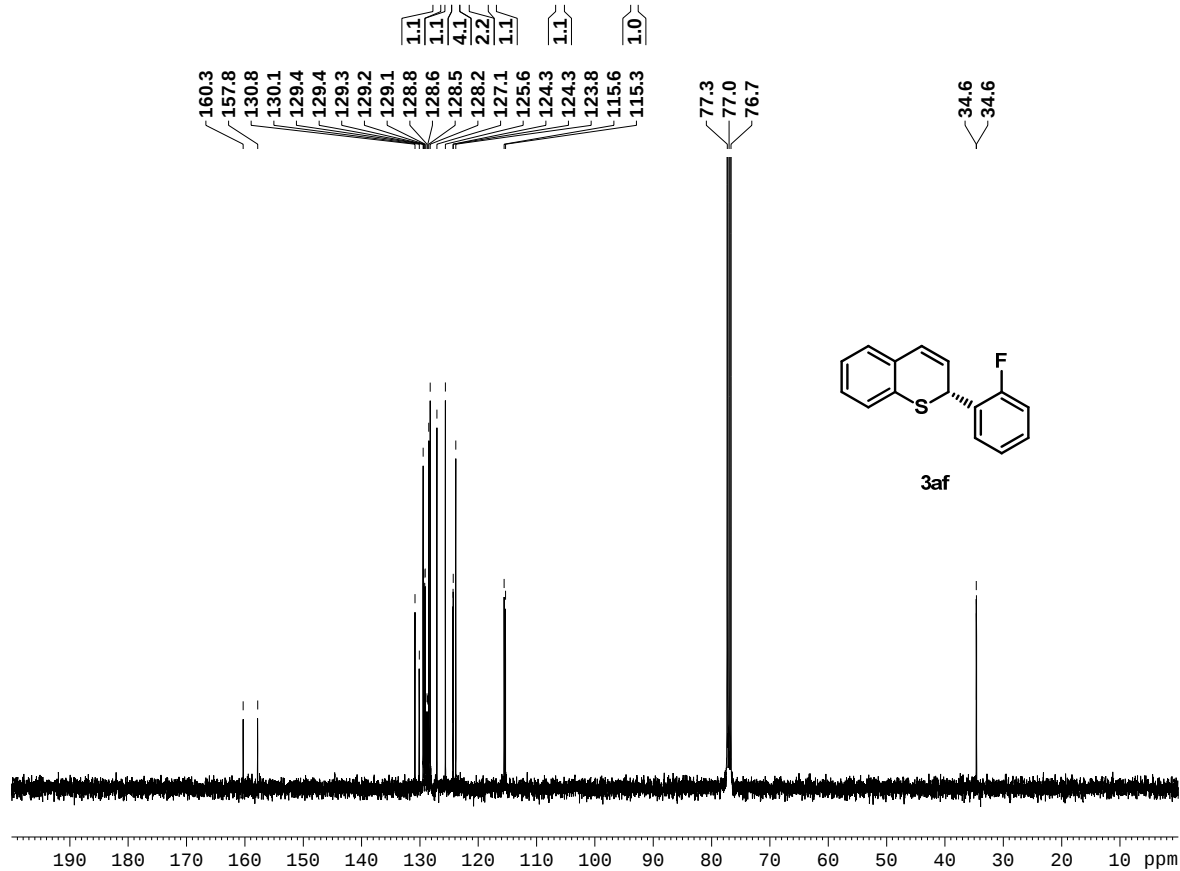
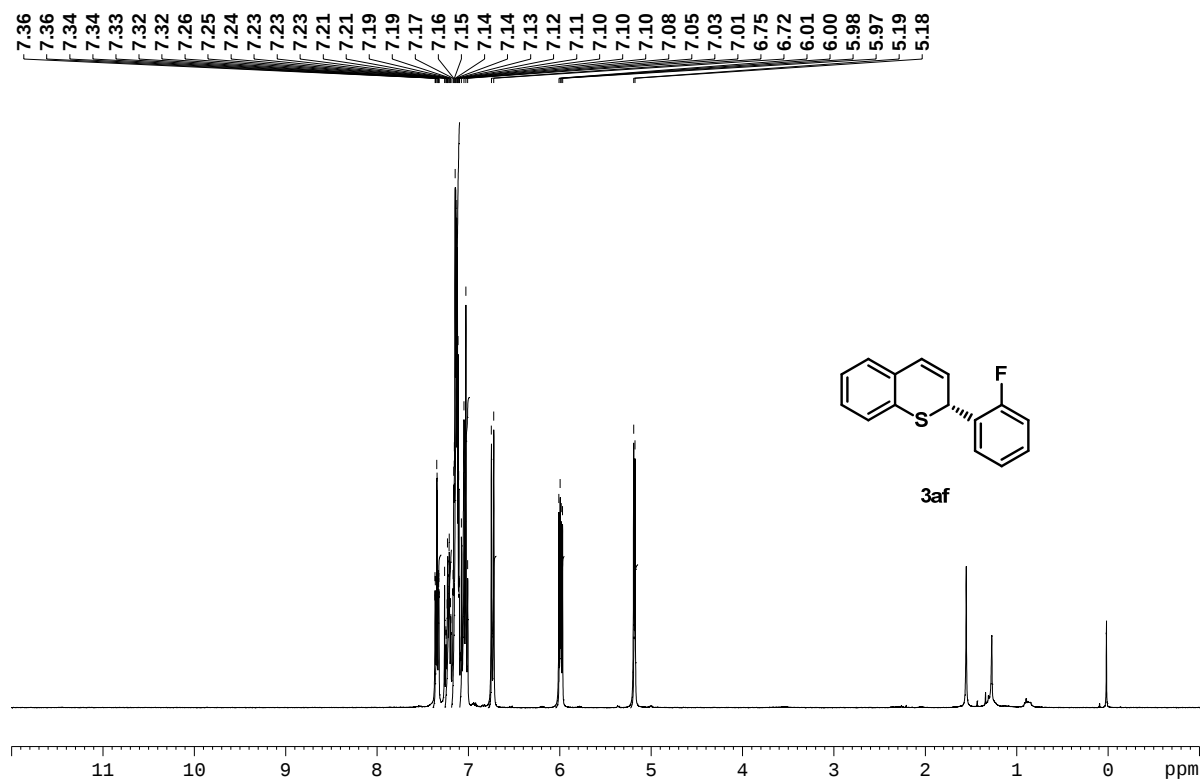


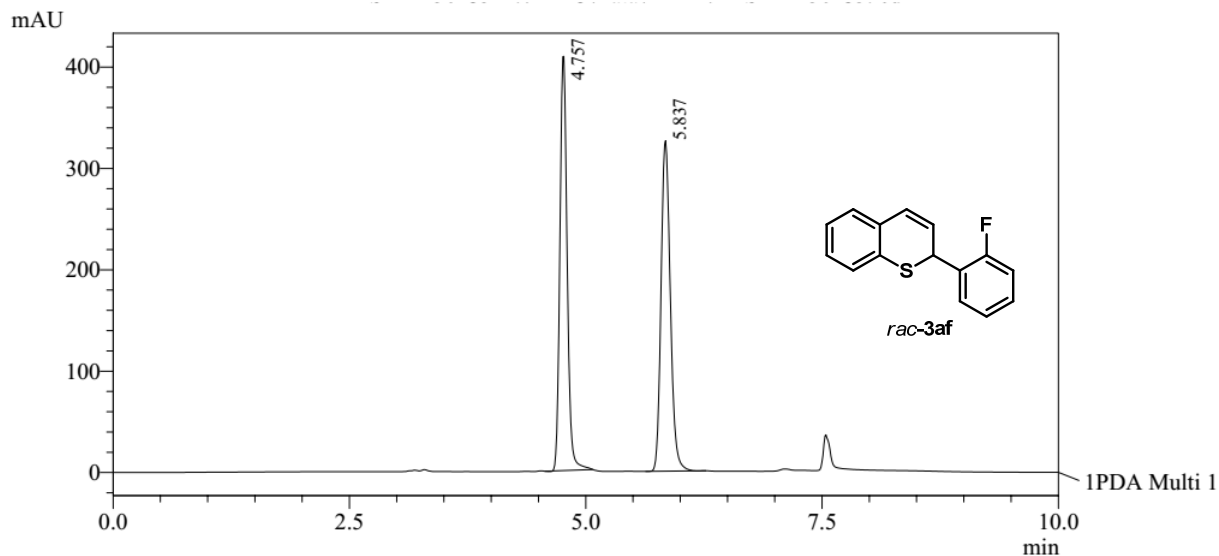
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.789	3321021	98.744
2	6.160	42228	1.256
Total		3363250	100.000





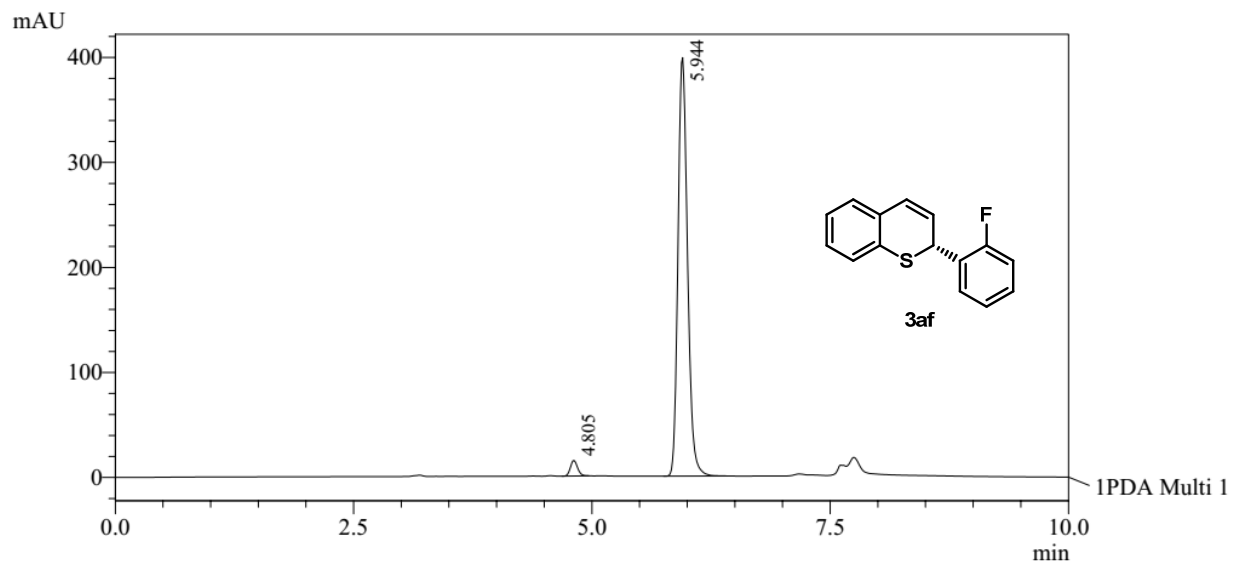
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.757	2190085	49.593
2	5.837	2226006	50.407
Total		4416091	100.000

Daicel Chiralpak IB column (80:20 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

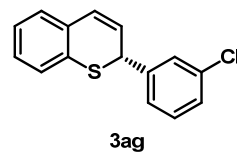
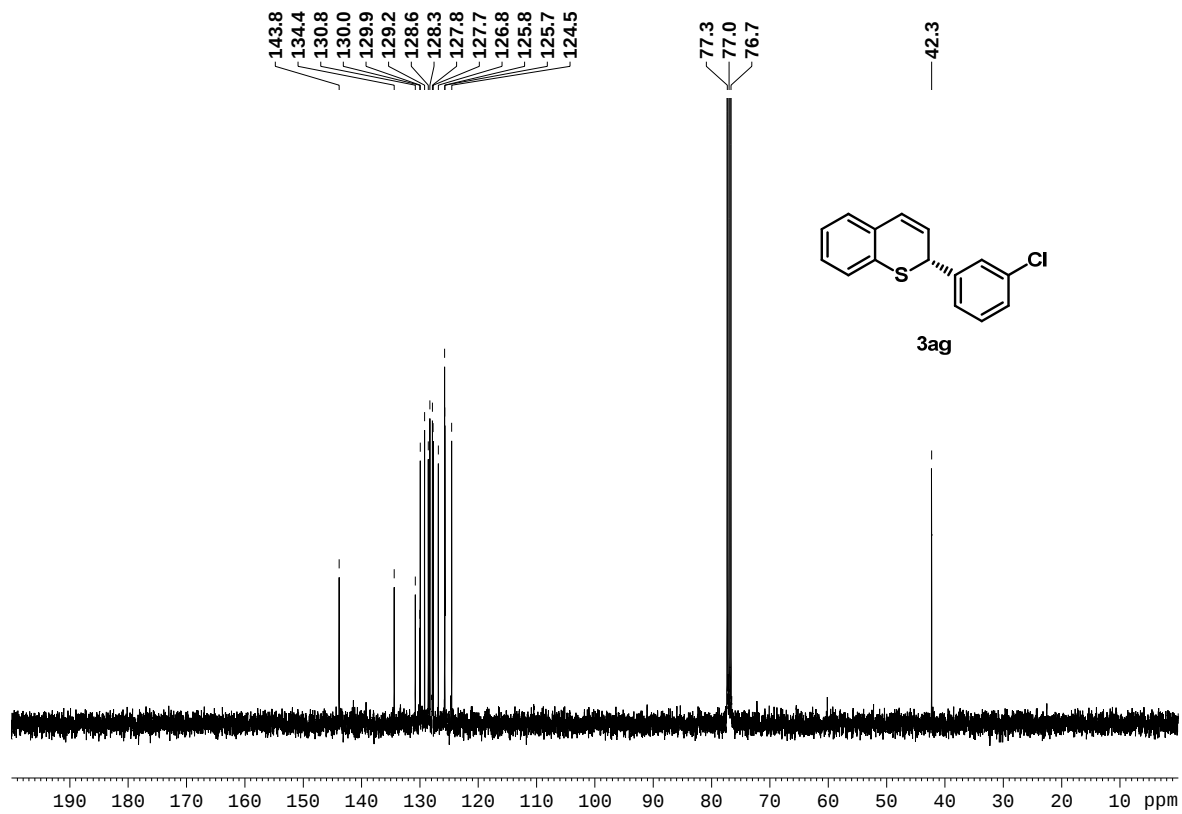
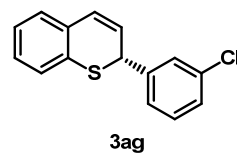
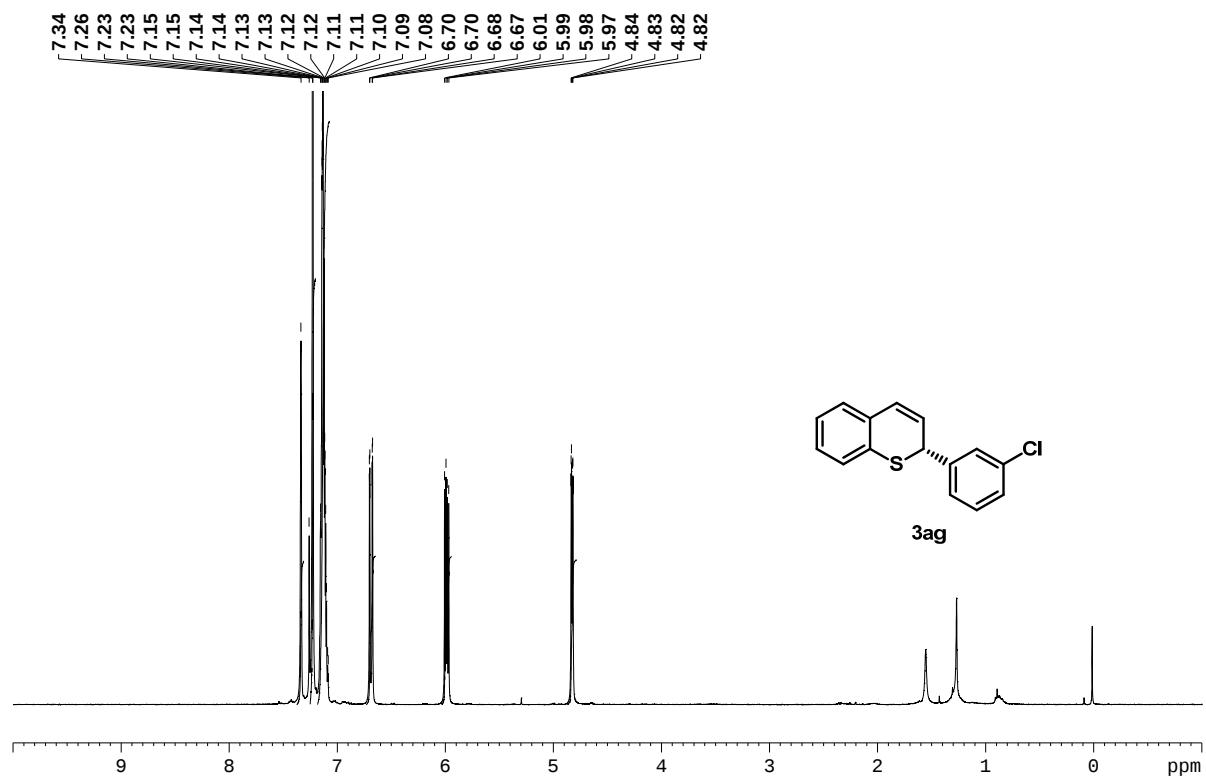


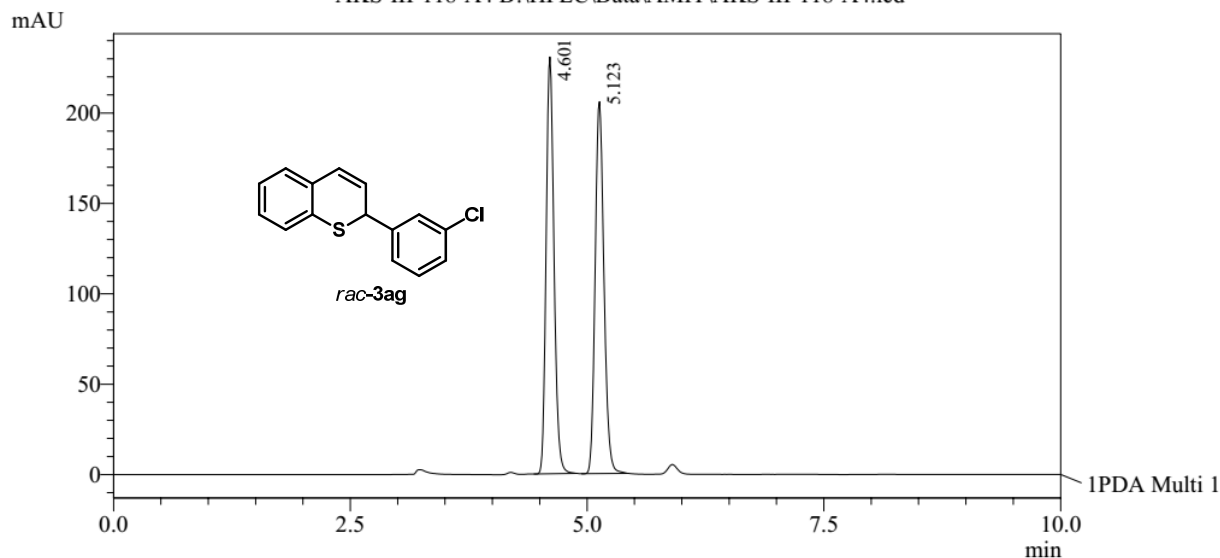
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.805	77757	2.725
2	5.944	2776051	97.275
Total		2853807	100.000





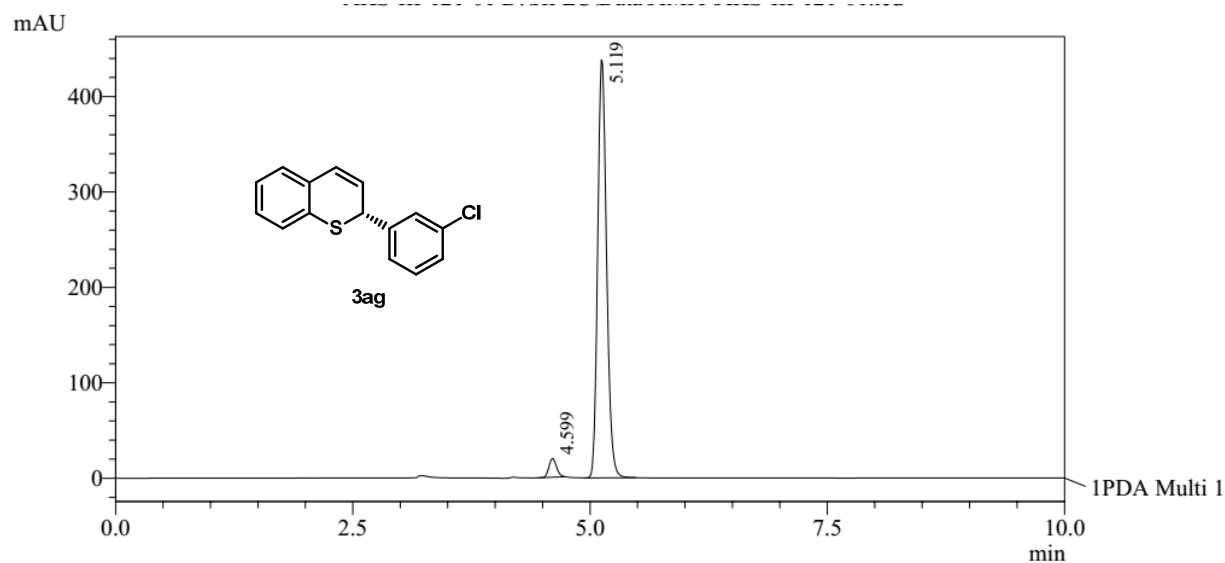
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.601	1329571	50.039
2	5.123	1327523	49.961
Total		2657093	100.000

Daicel Chiralpak IF column (97:3 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

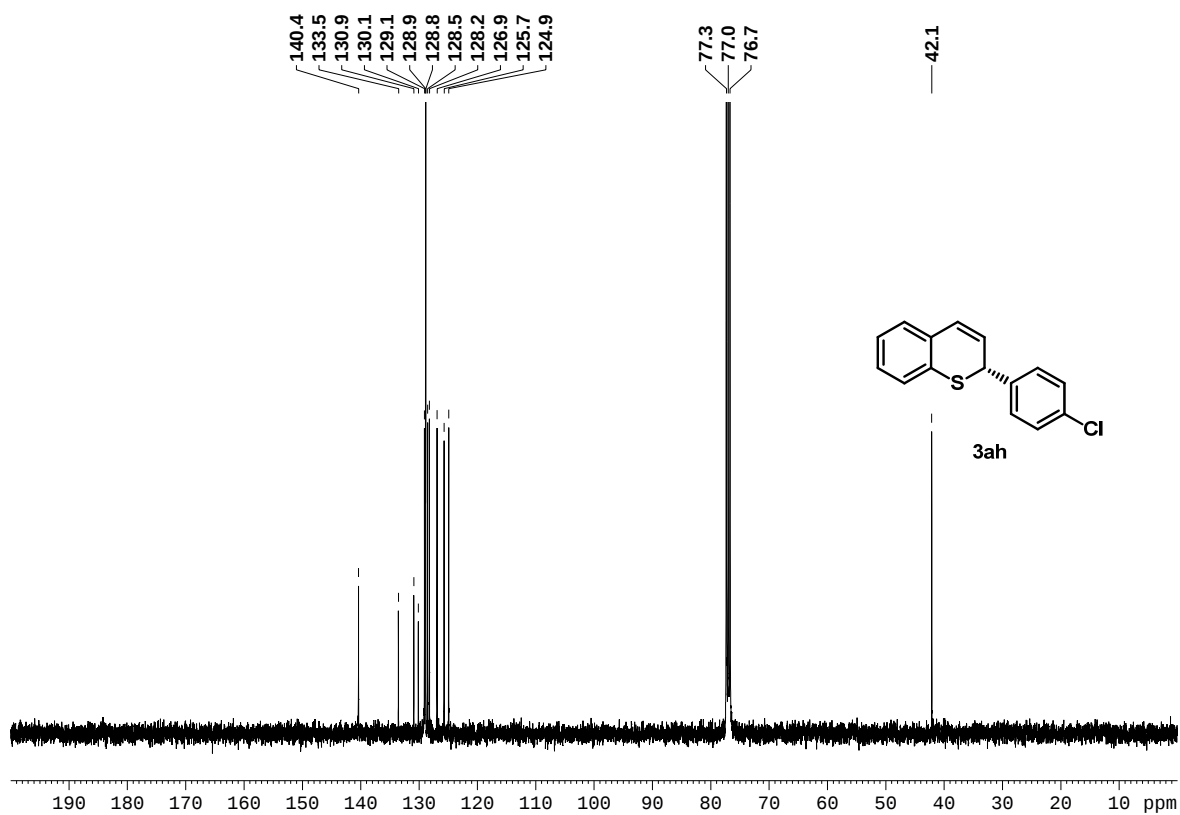
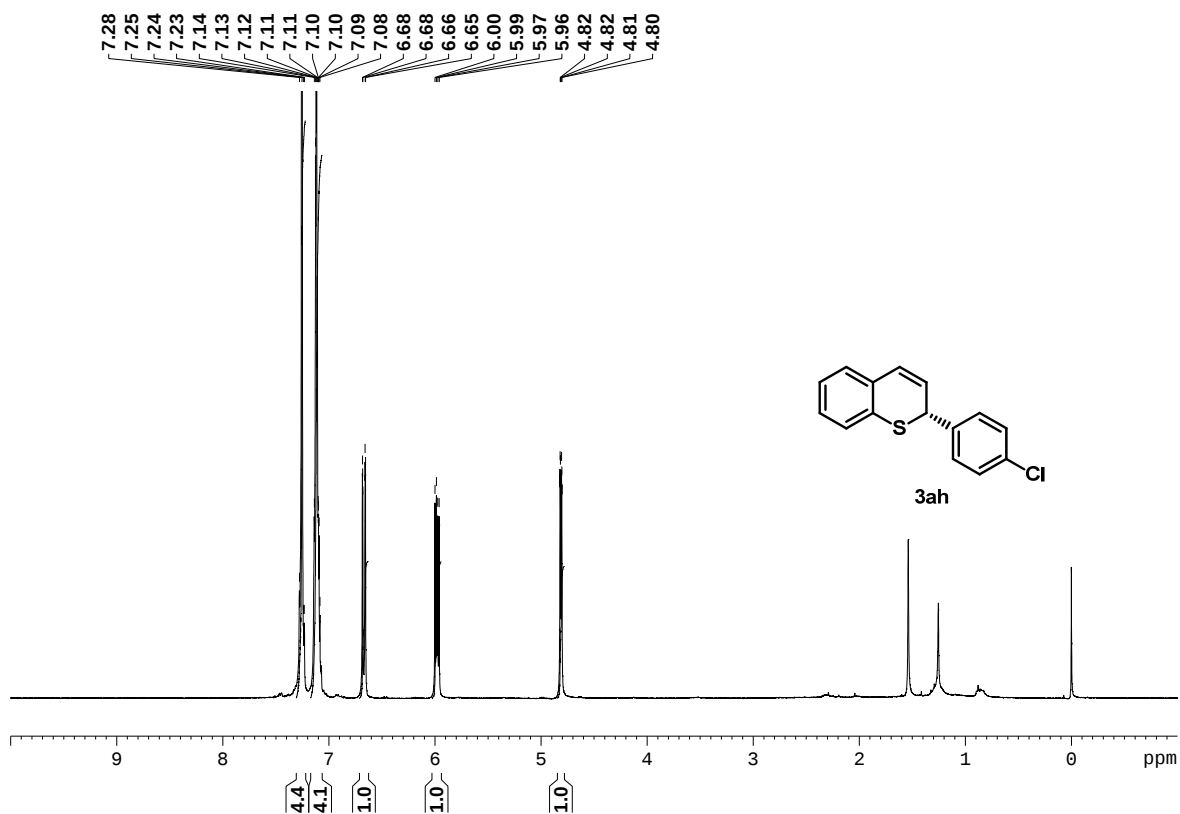


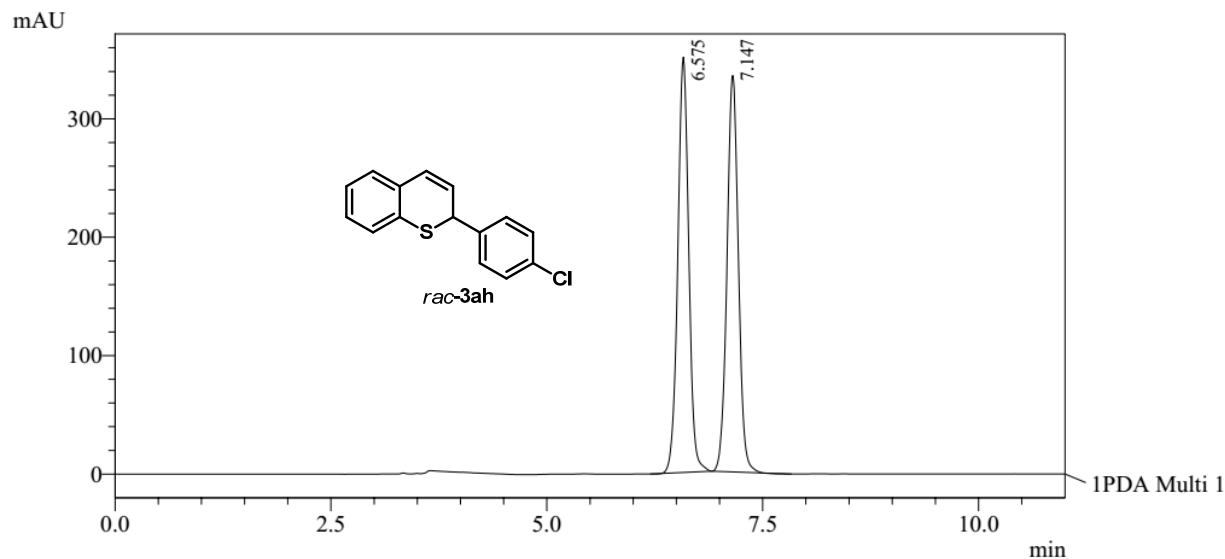
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.599	109244	3.686
2	5.119	2854644	96.314
Total		2963888	100.000





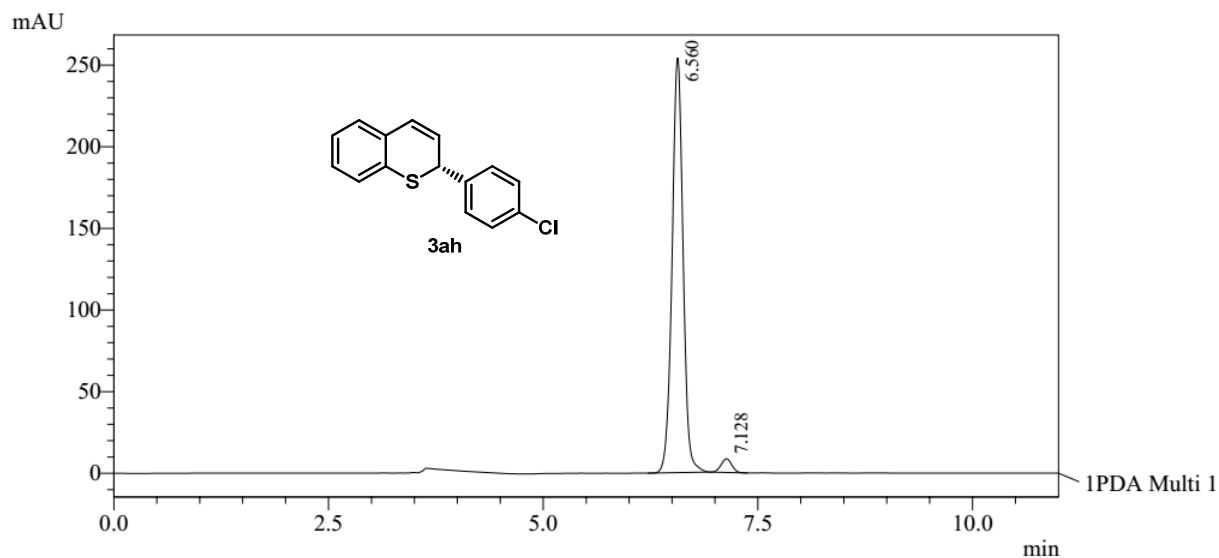
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.575	3089189	50.179
2	7.147	3067186	49.821
Total		6156375	100.000

Daicel Chiralpak IB column (99:1 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

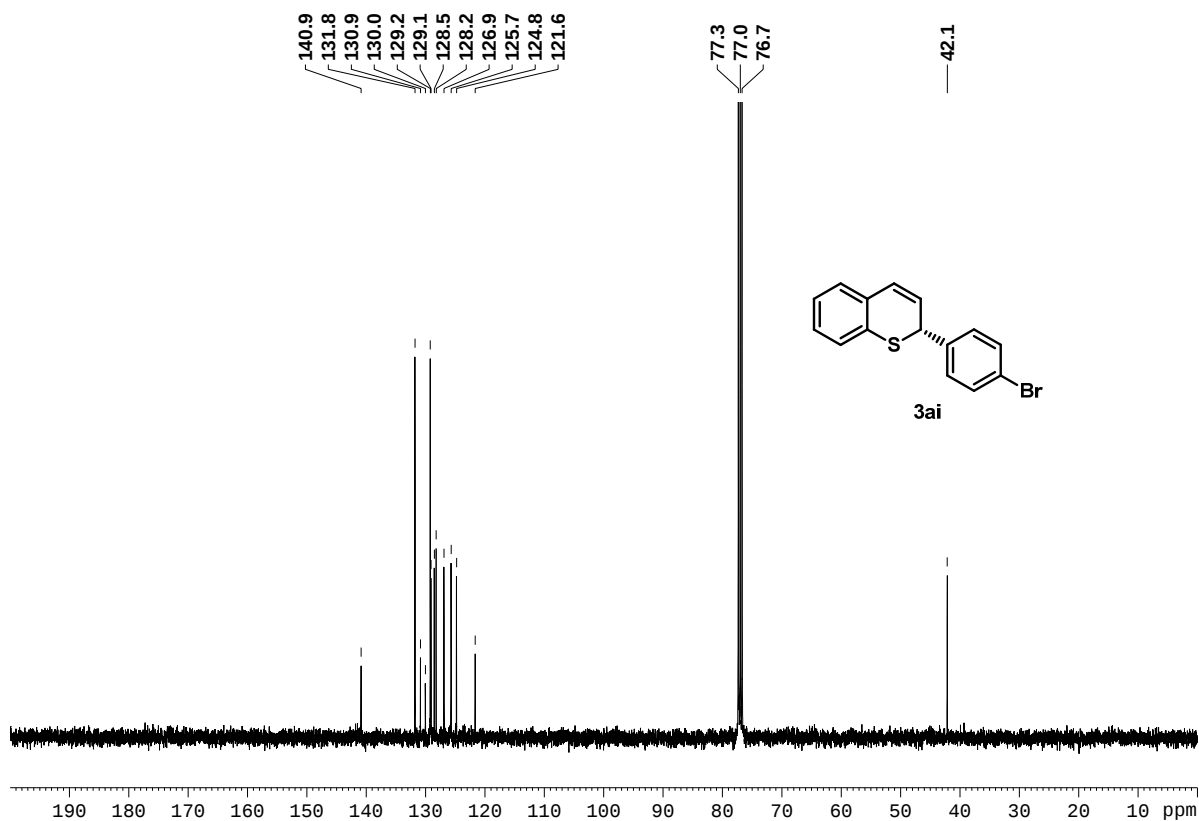
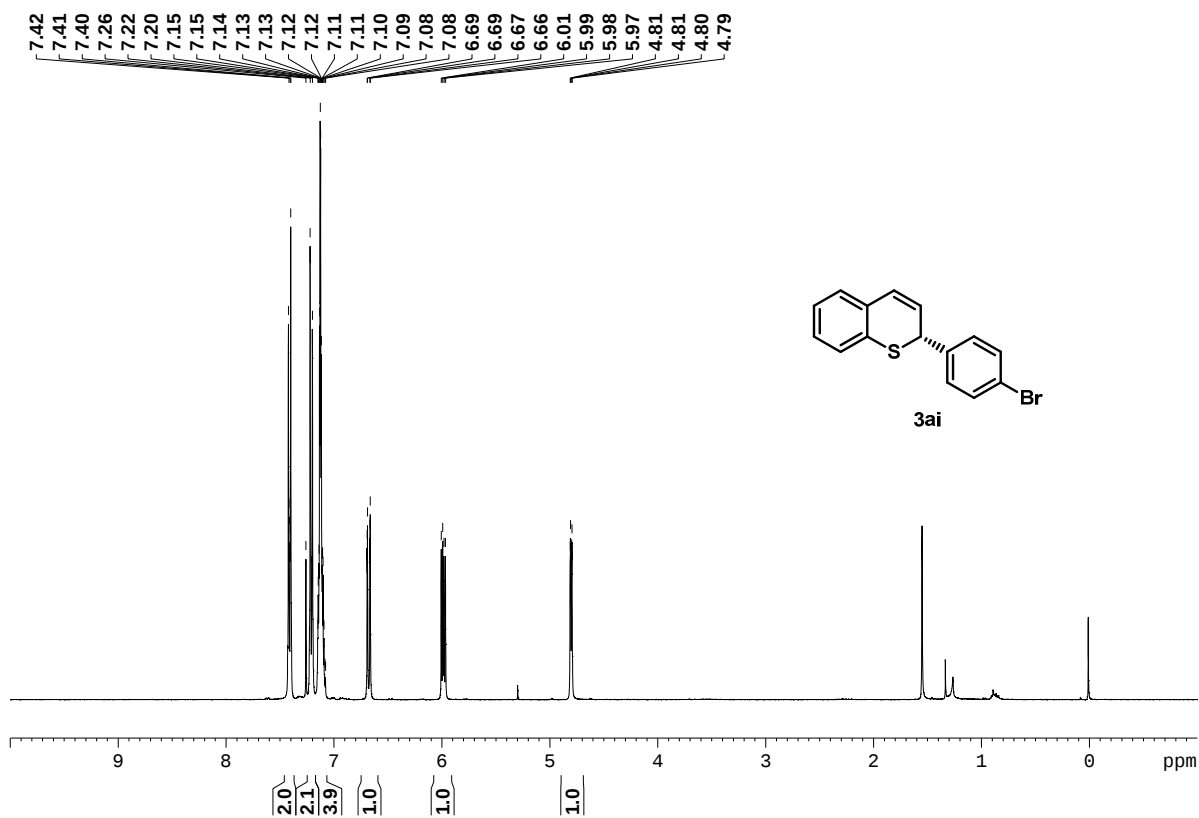


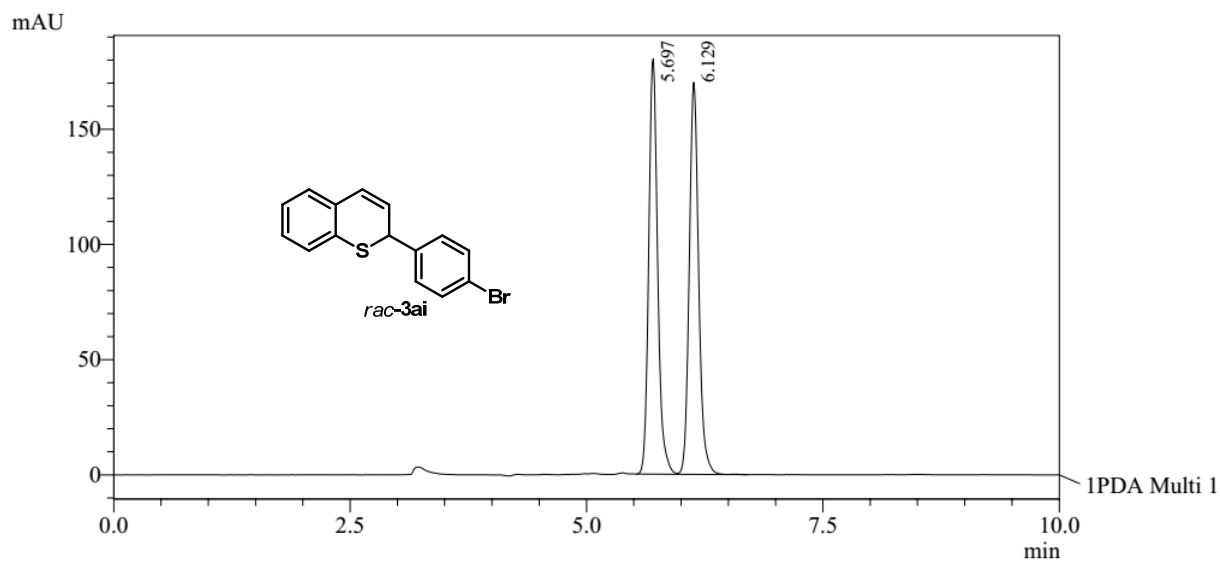
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.560	2262710	96.869
2	7.128	73136	3.131
Total		2335846	100.000





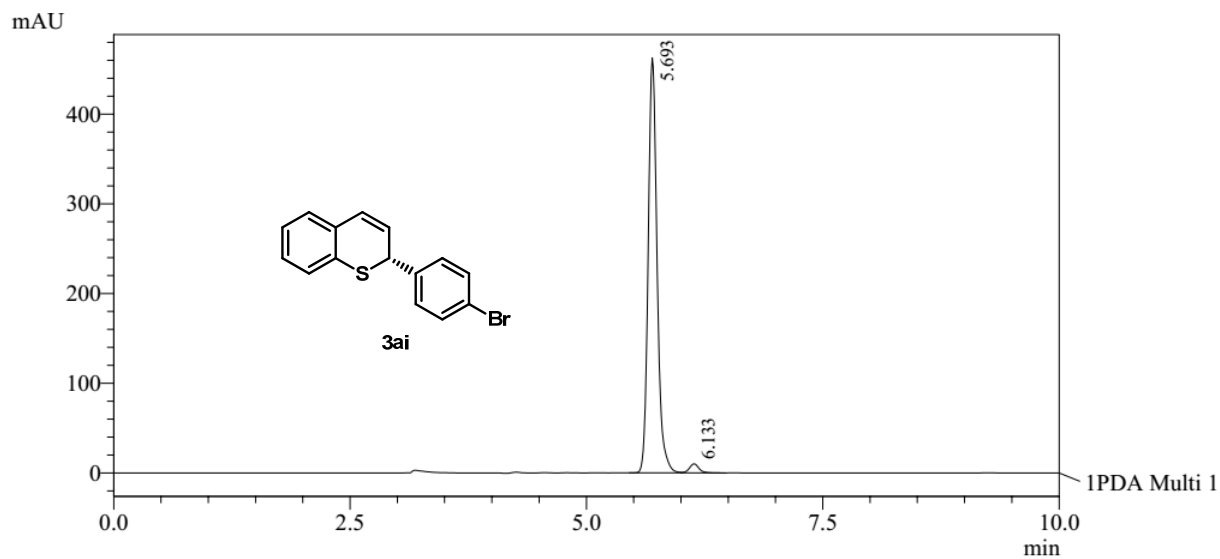
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.697	1183268	50.168
2	6.129	1175337	49.832
Total		2358605	100.000

Daicel Chiralpak IB column (97:3 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

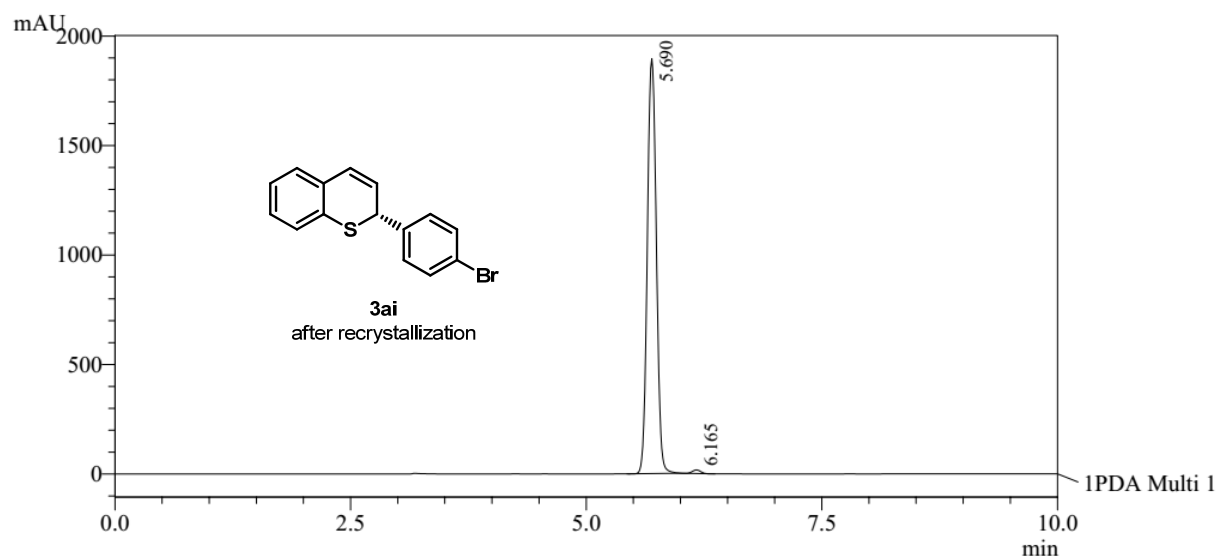


1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.693	3028182	97.713
2	6.133	70887	2.287
Total		3099069	100.000

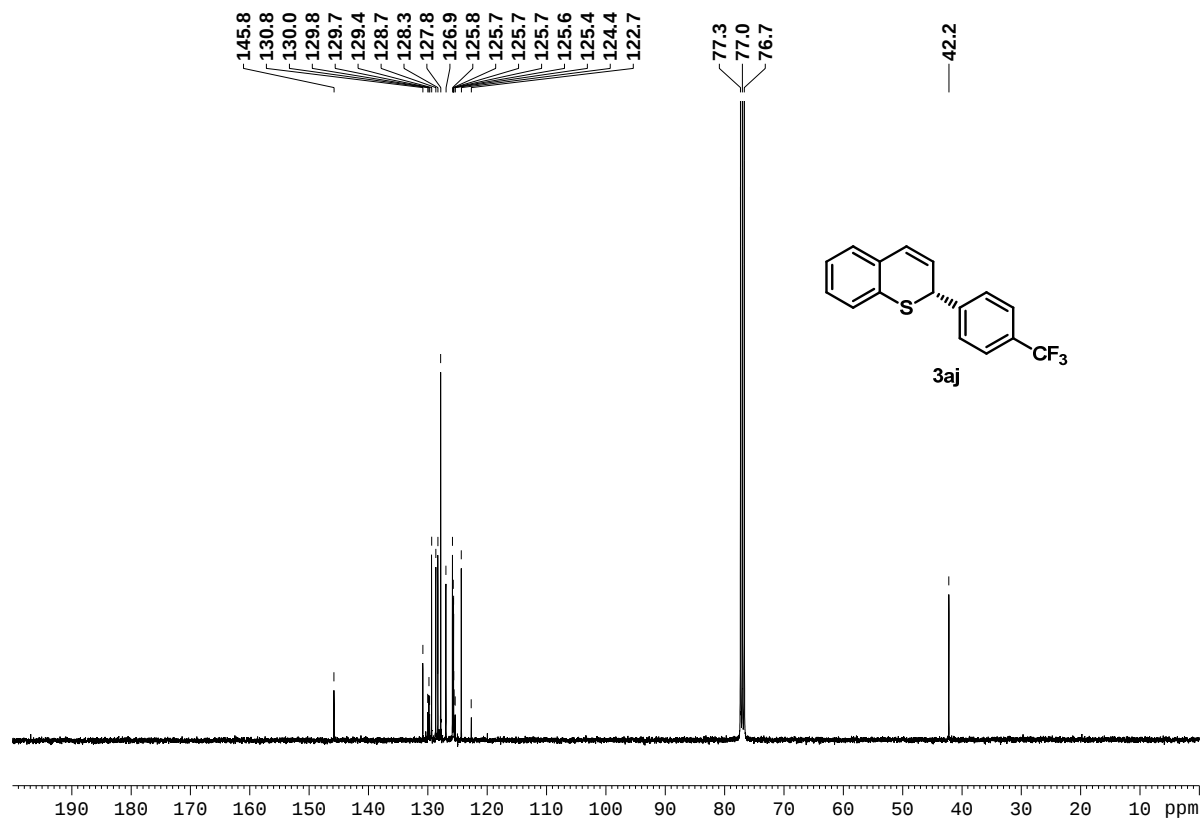
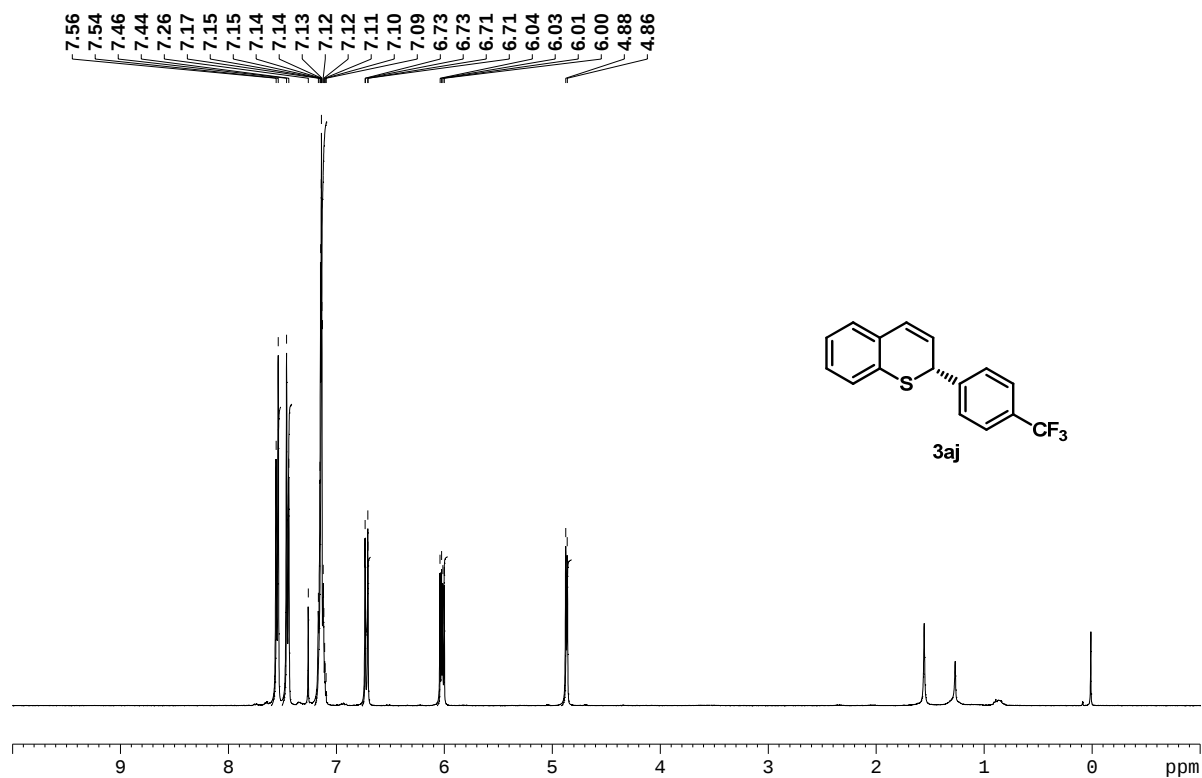


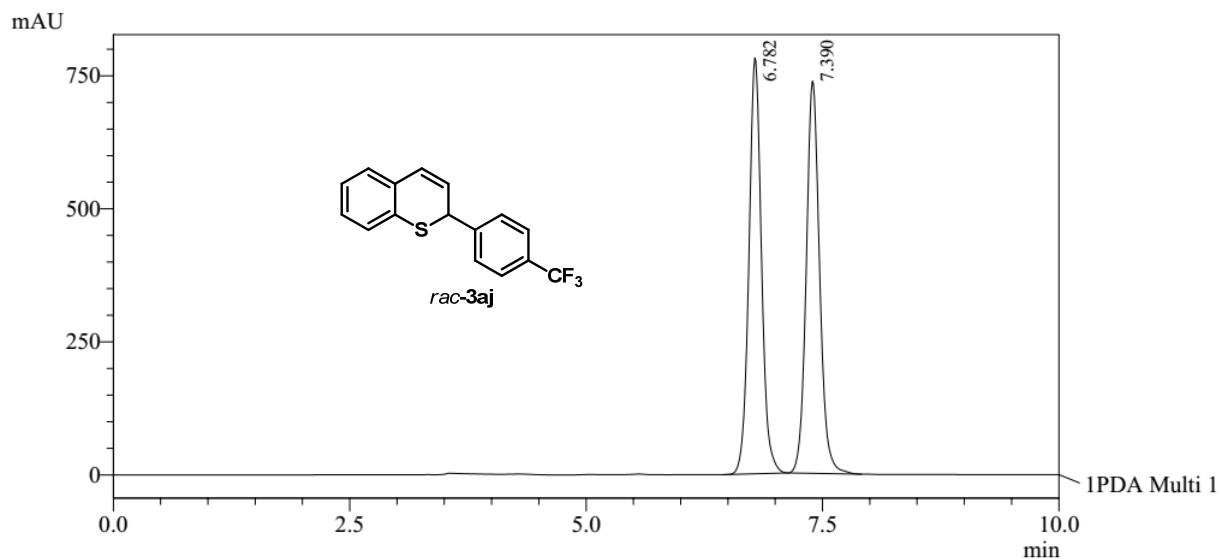
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.690	12595395	99.234
2	6.165	97188	0.766
Total		12692583	100.000





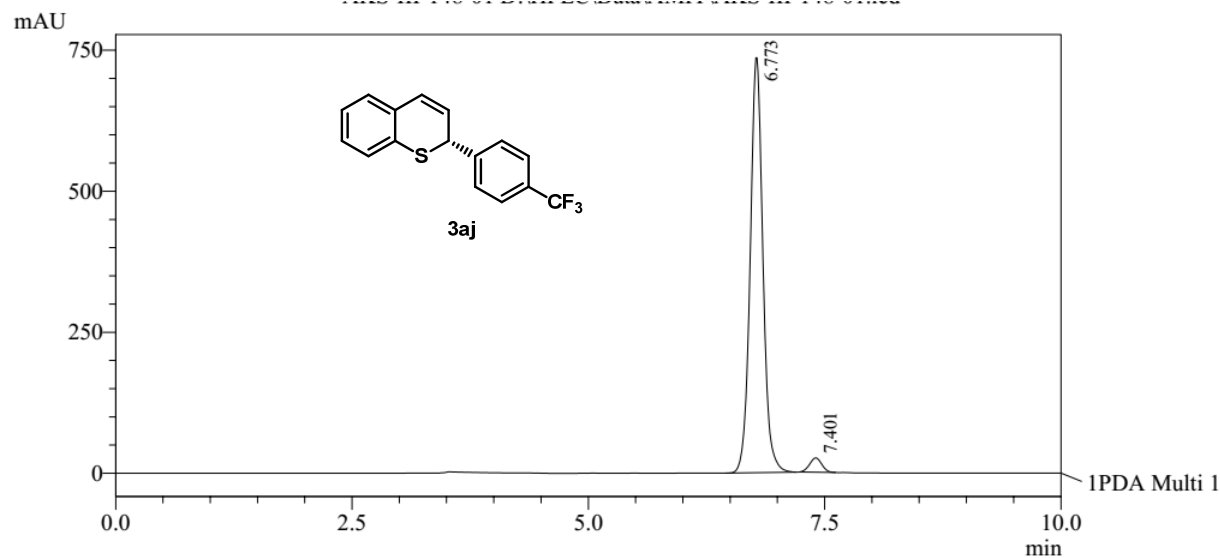
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.782	7206786	49.884
2	7.390	7240318	50.116
Total		14447104	100.000

Daicel Chiralpak IB column (99:1 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

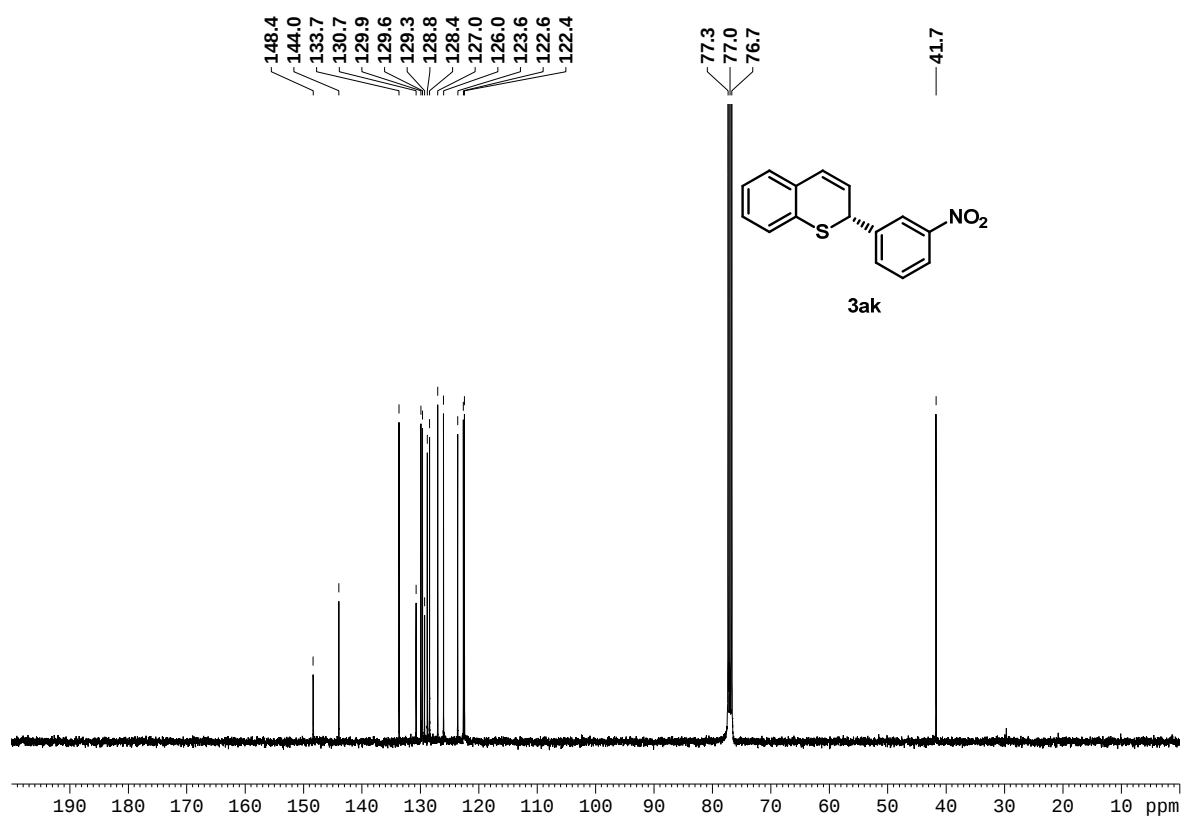
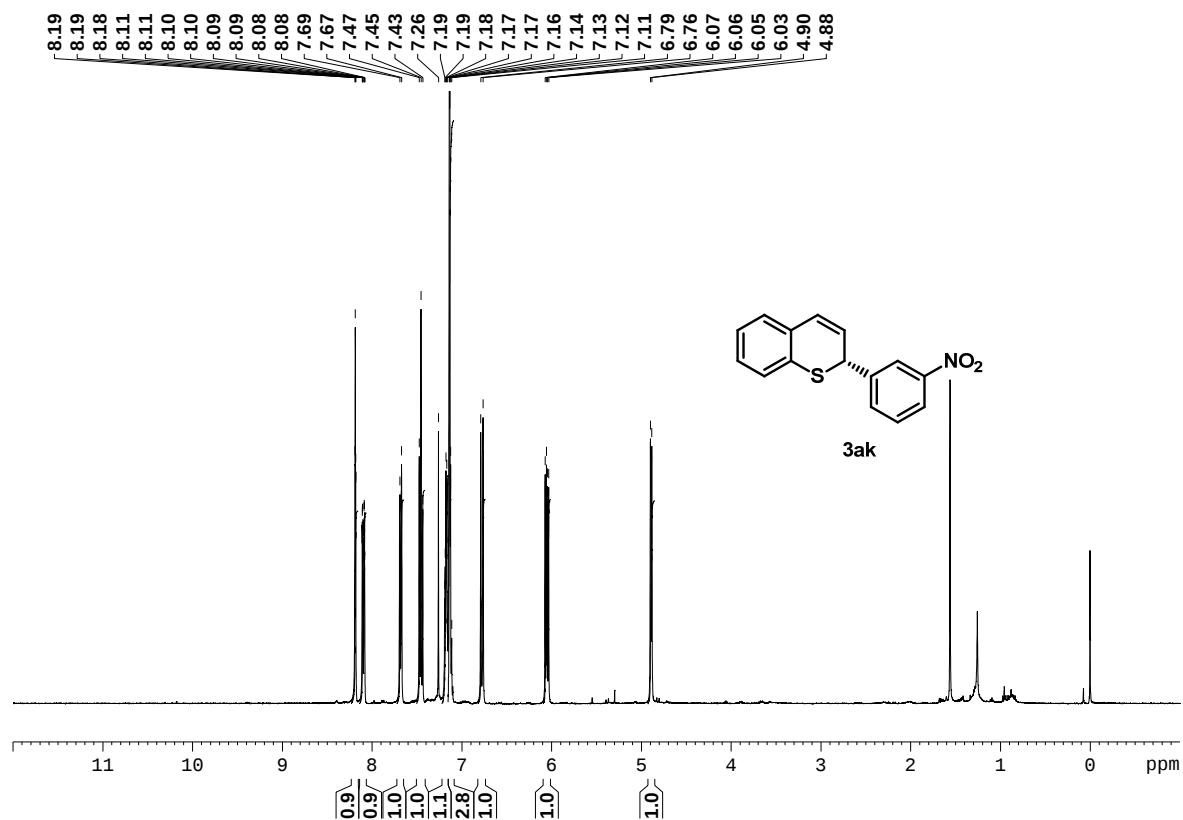


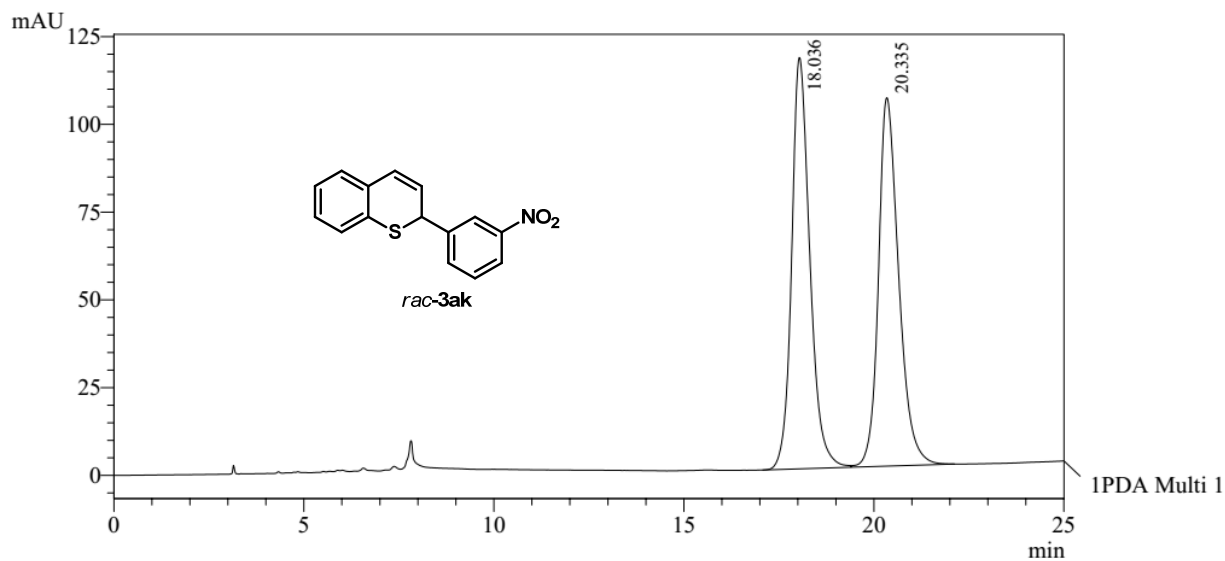
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.773	6880430	96.826
2	7.401	225575	3.174
Total		7106005	100.000



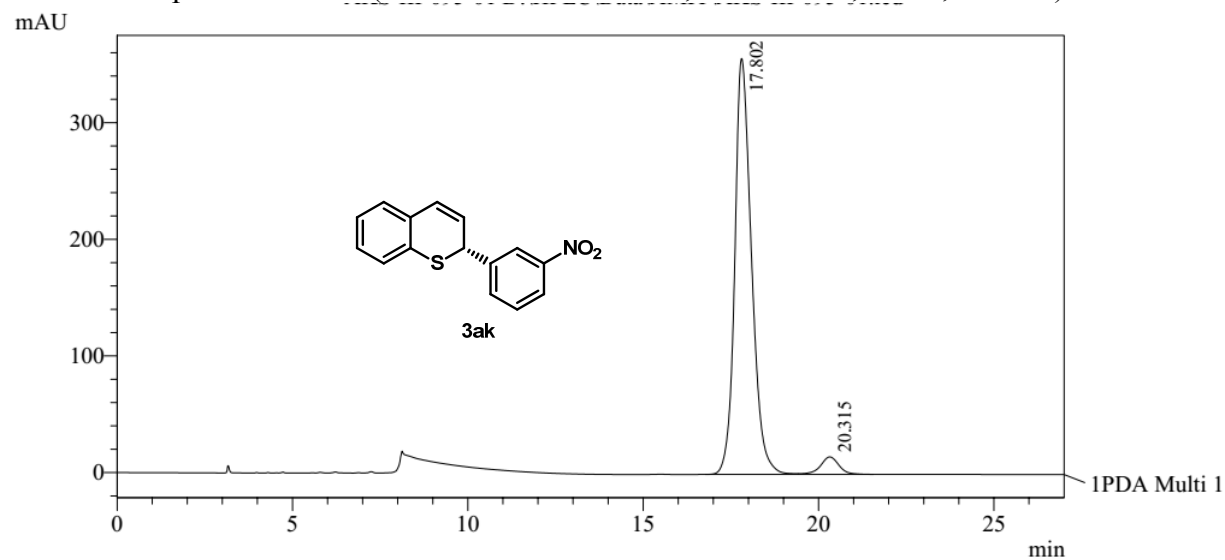


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	18.036	3904147	50.076
2	20.335	3892280	49.924
Total		7796426	100.000

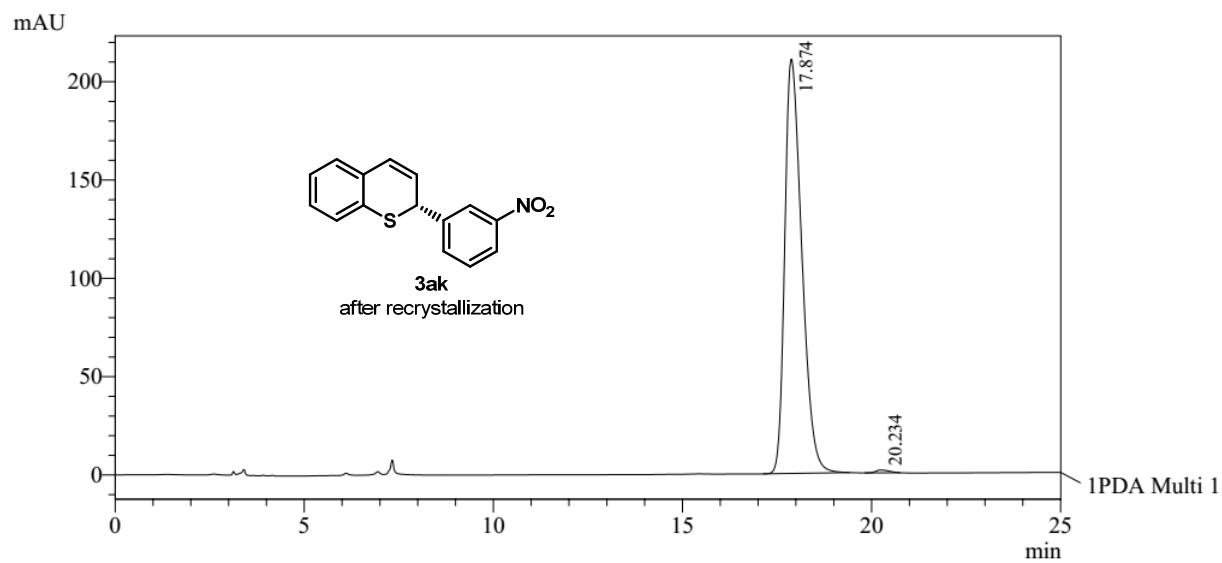
Daicel Chiralpak IB column (80:20 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)



PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	17.802	11838986	95.565
2	20.315	549399	4.435
Total		12388386	100.000

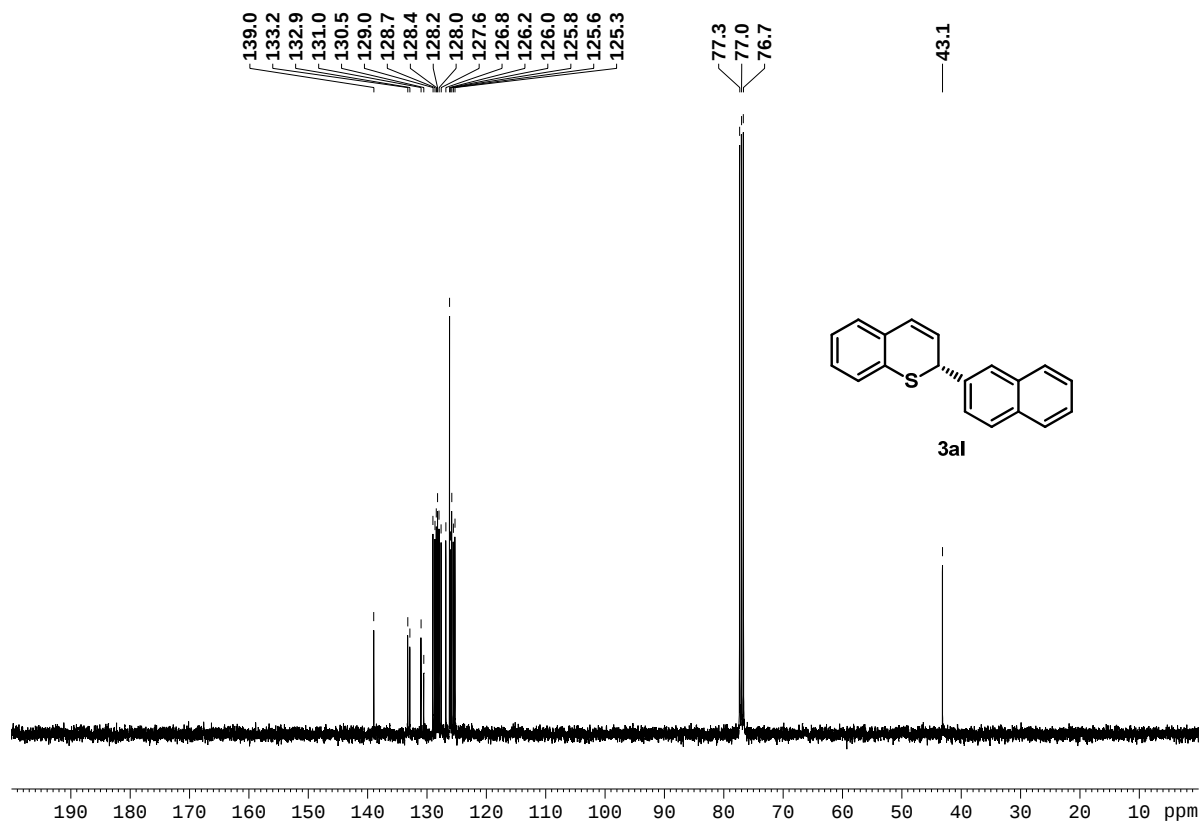
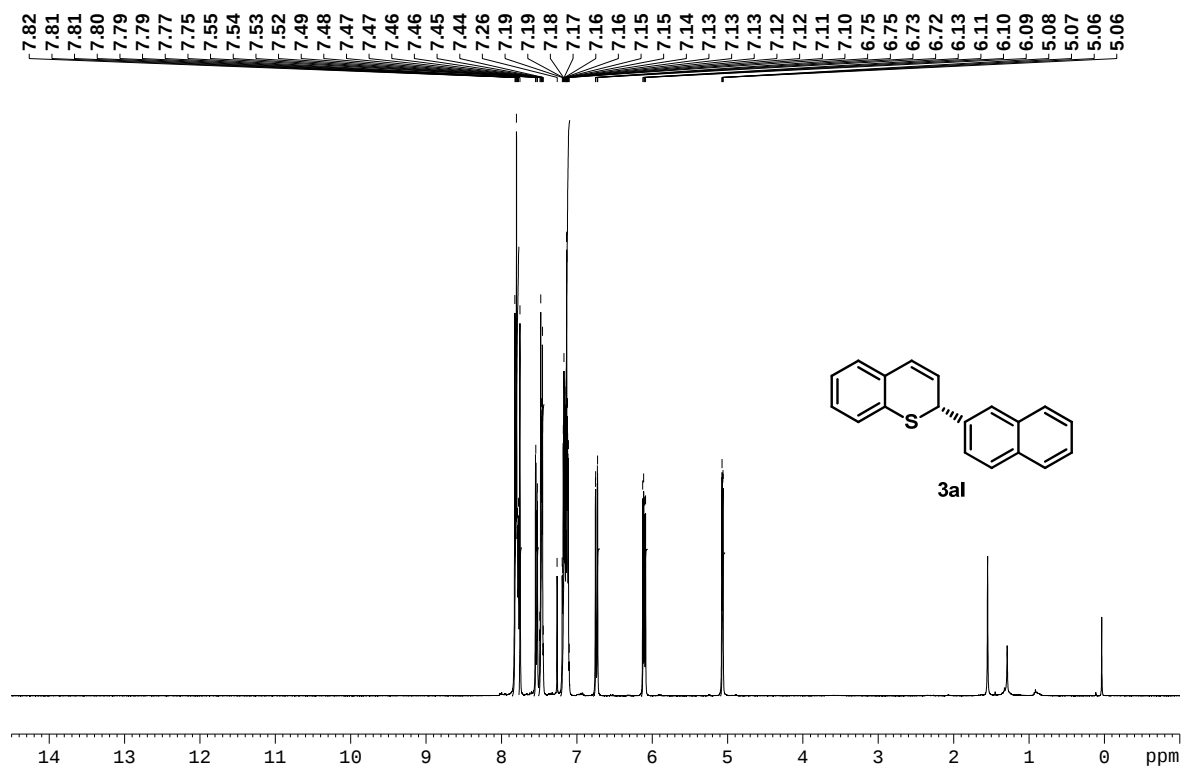


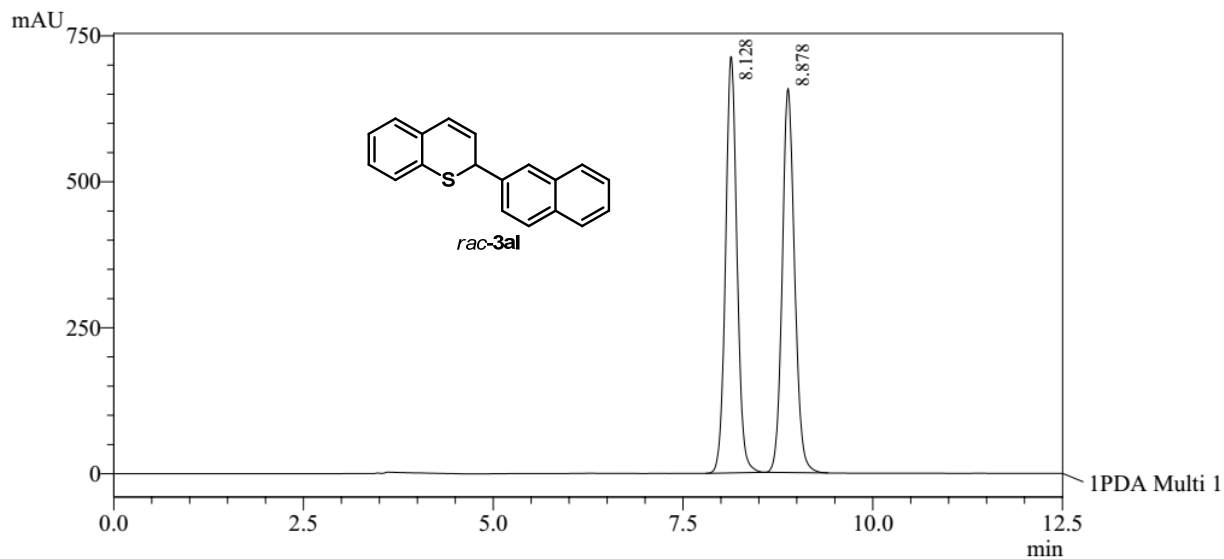
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	17.874	6614035	99.488
2	20.234	34060	0.512
Total		6648096	100.000





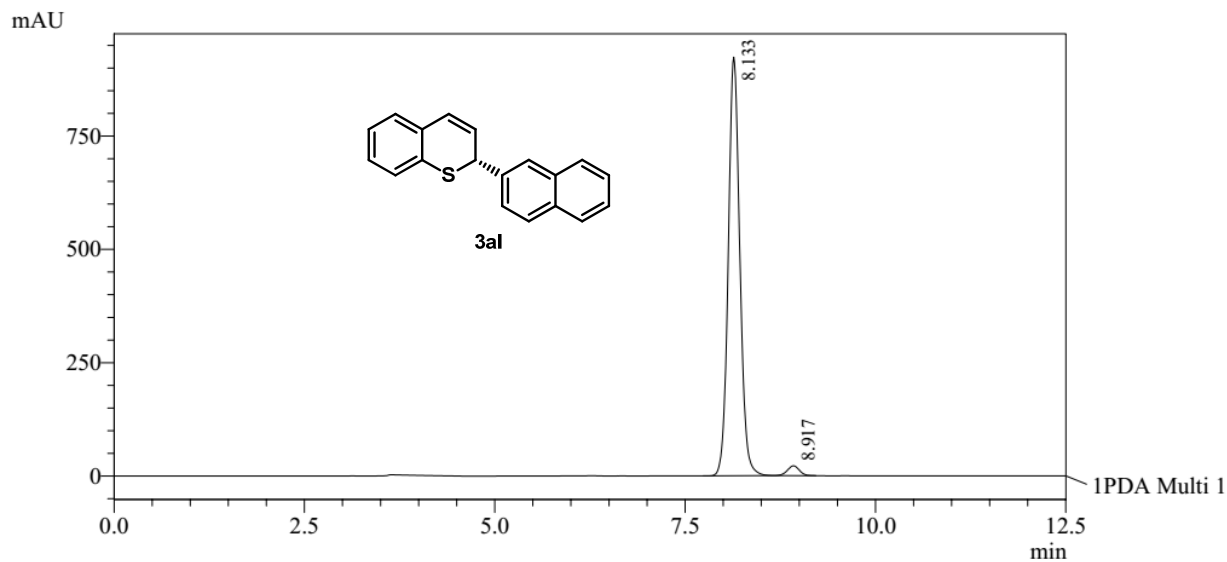
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.128	7490967	50.046
2	8.878	7477099	49.954
Total		14968065	100.000

Daicel Chiralpak IB column (99:1 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

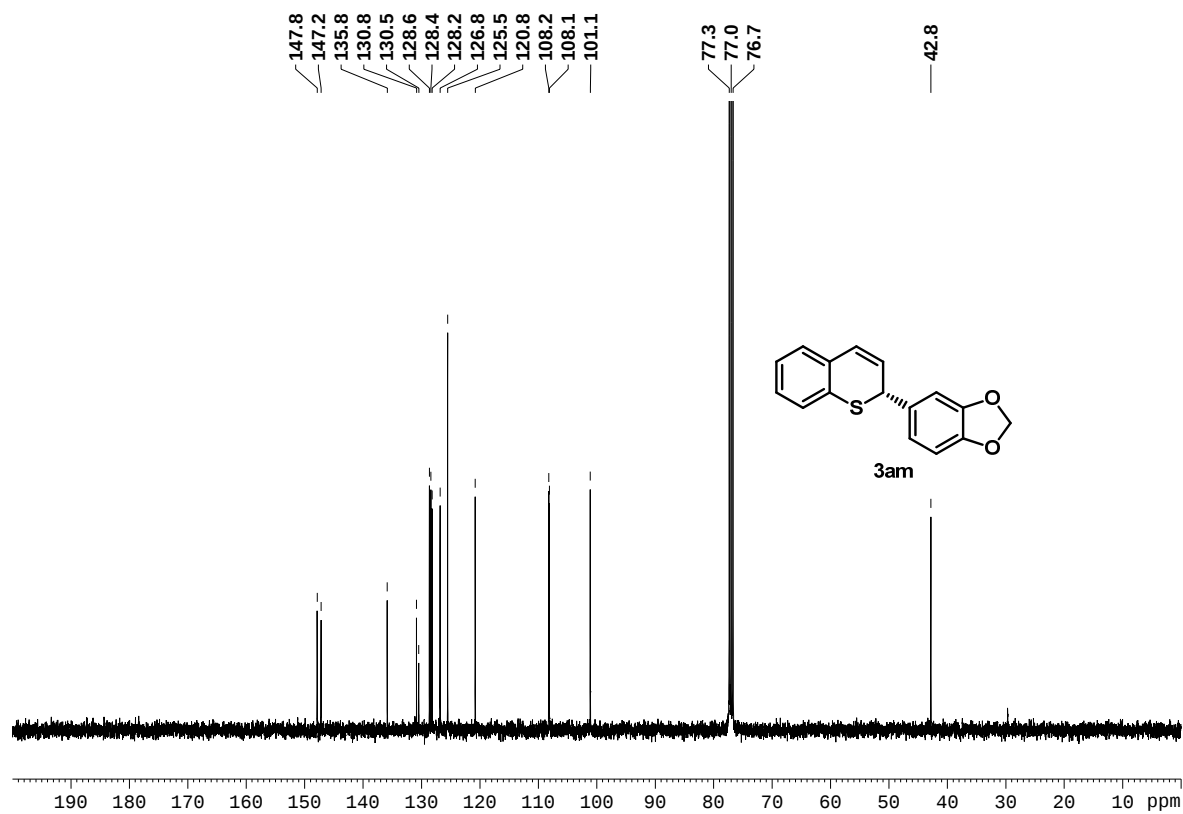
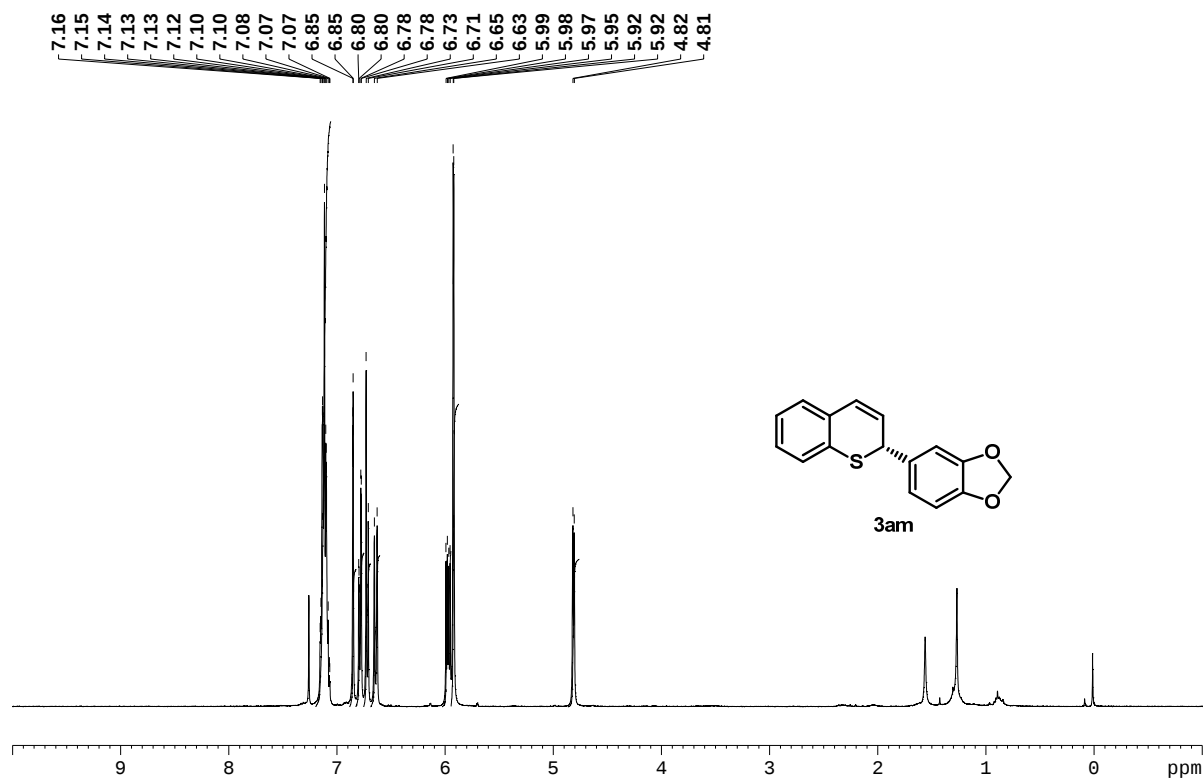


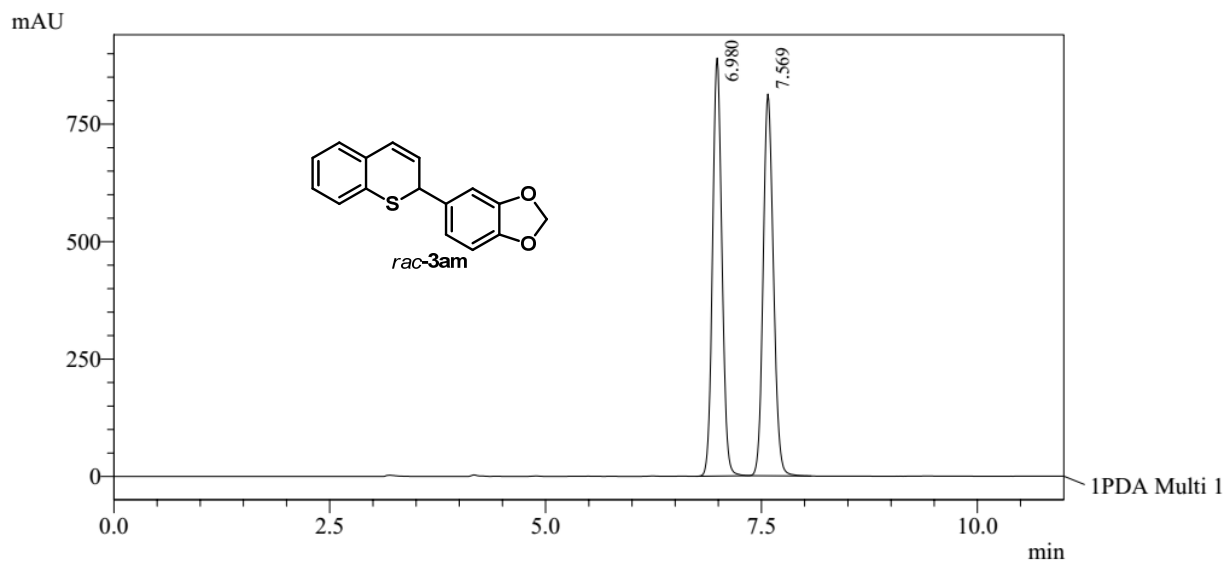
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.133	9744526	97.637
2	8.917	235822	2.363
Total		9980348	100.000





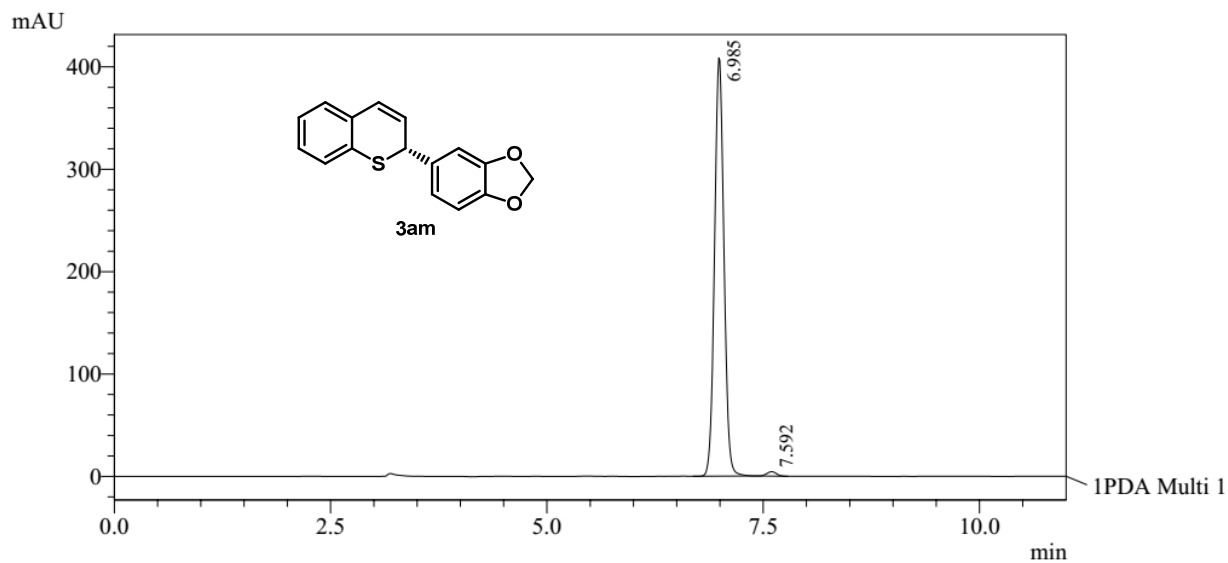
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.980	6719067	50.003
2	7.569	6718310	49.997
Total		13437376	100.000

Daicel Chiralpak IB column (97:3 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

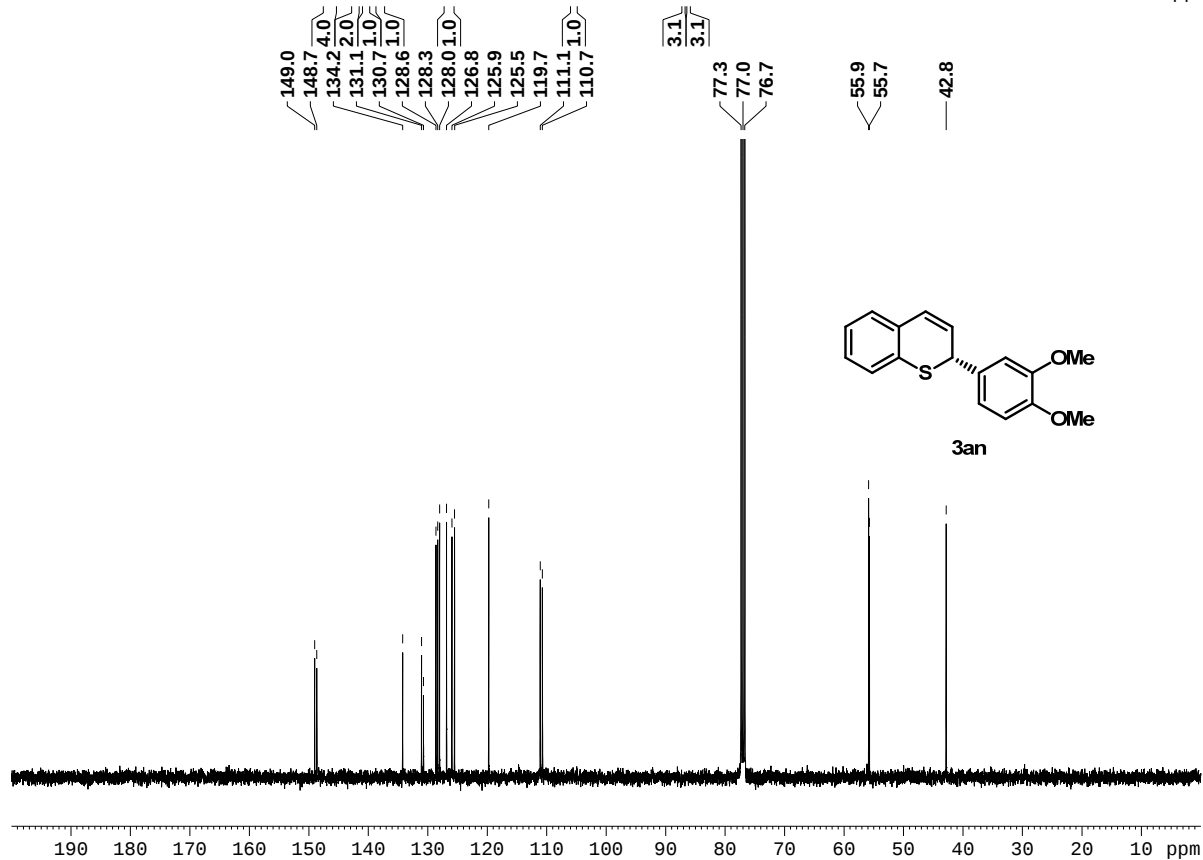


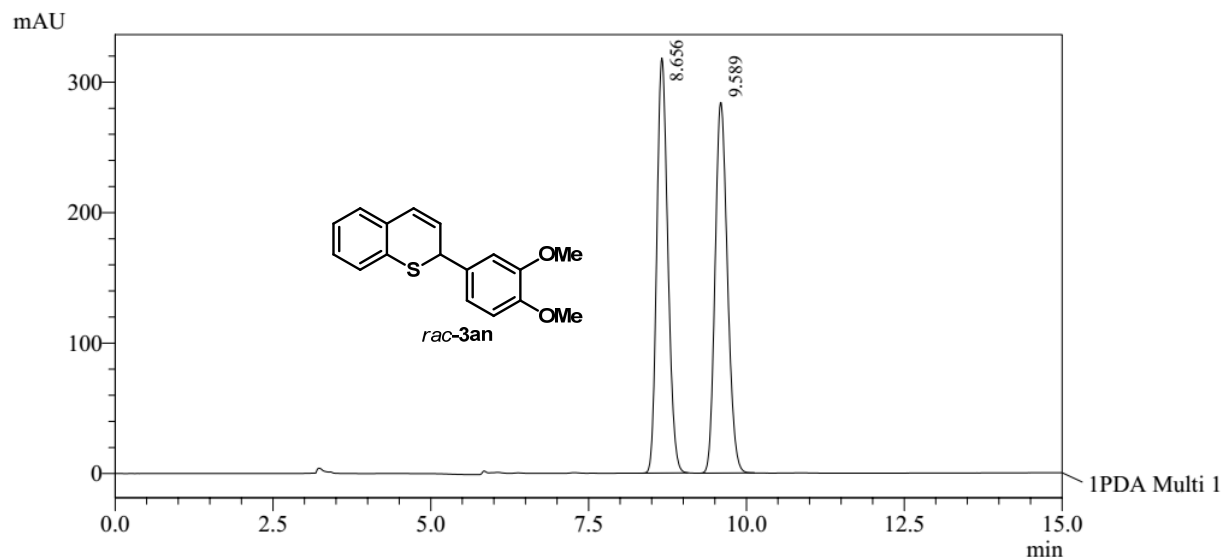
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.985	3086647	98.923
2	7.592	33601	1.077
Total		3120247	100.000





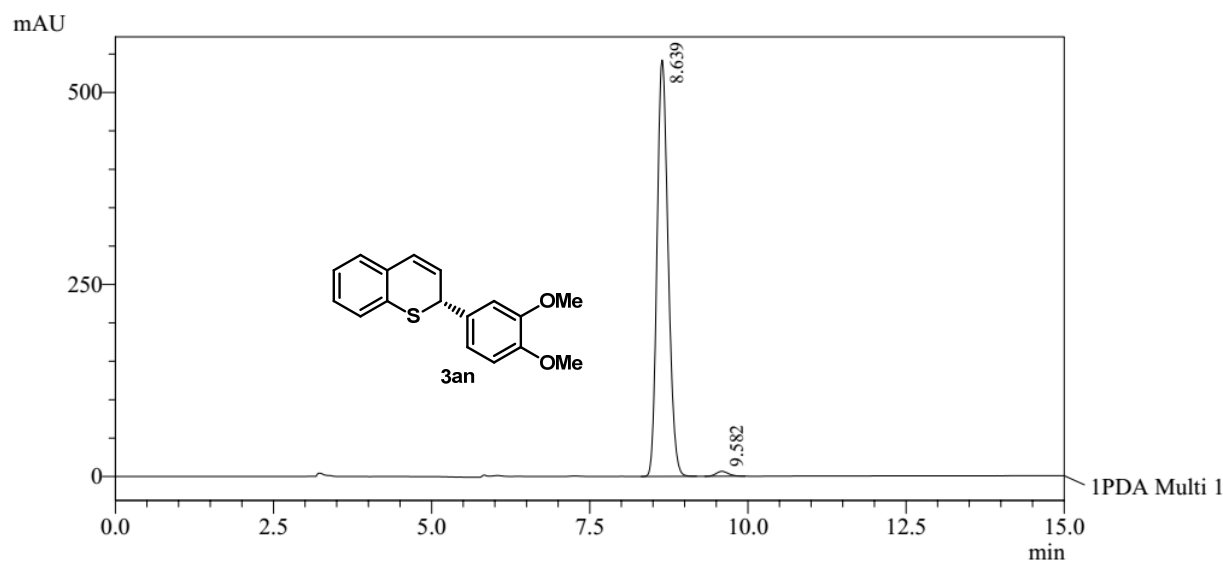
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.656	3877631	50.006
2	9.589	3876675	49.994
Total		7754306	100.000

Daicel Chiralpak AS-H column (97:03 *n*-Hexane/EtOH, 1.0 mL/min, 20 °C, 254 nm)

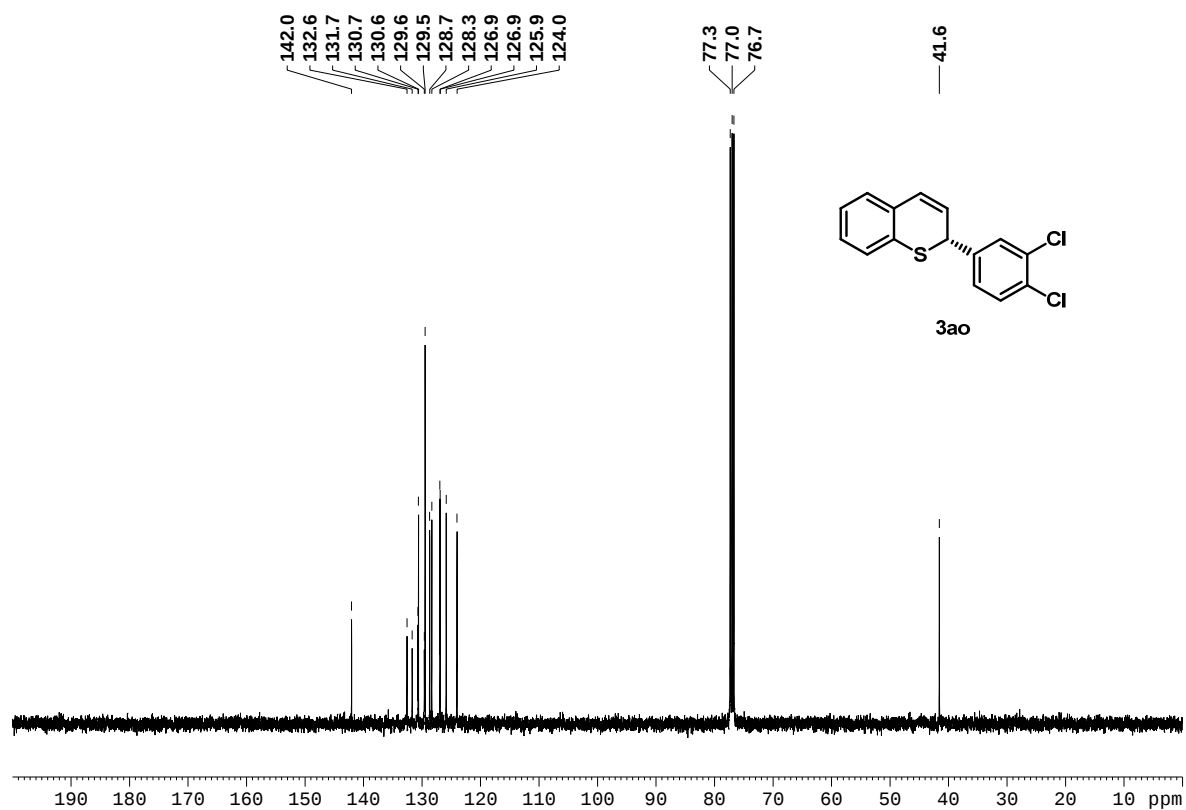
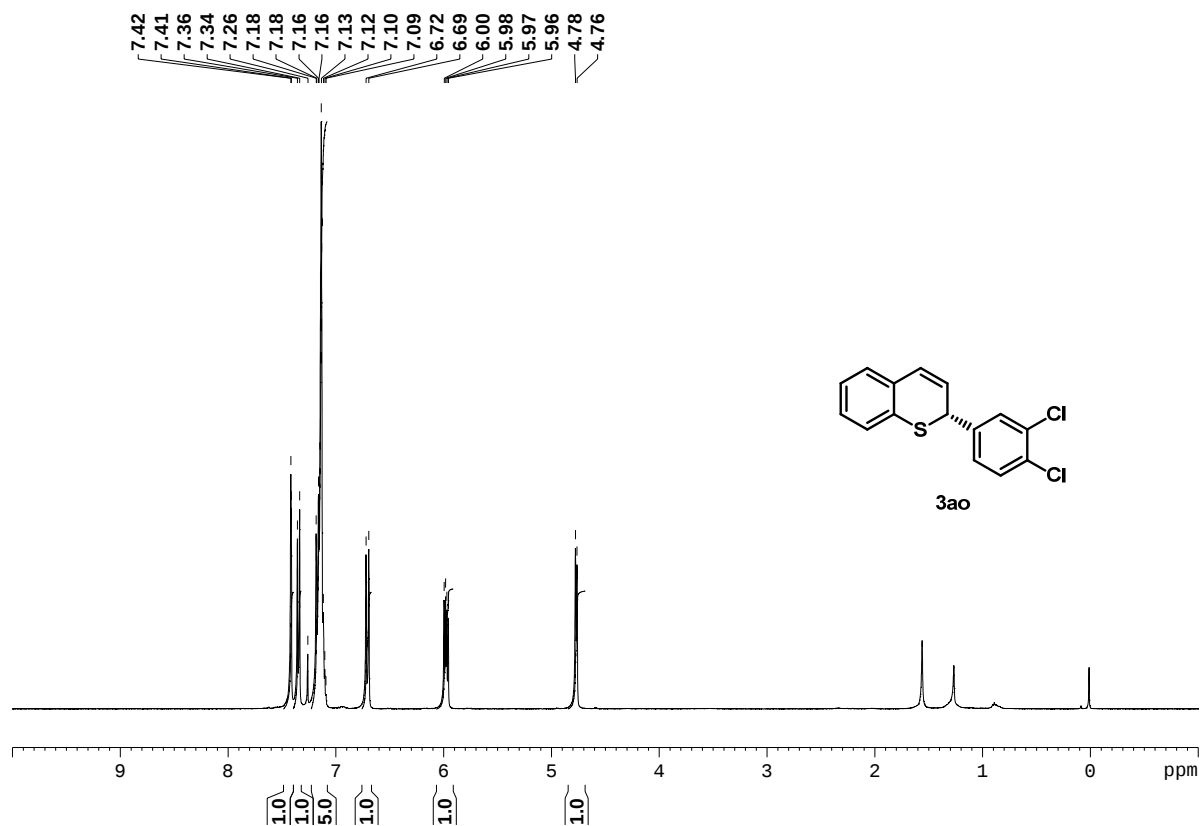


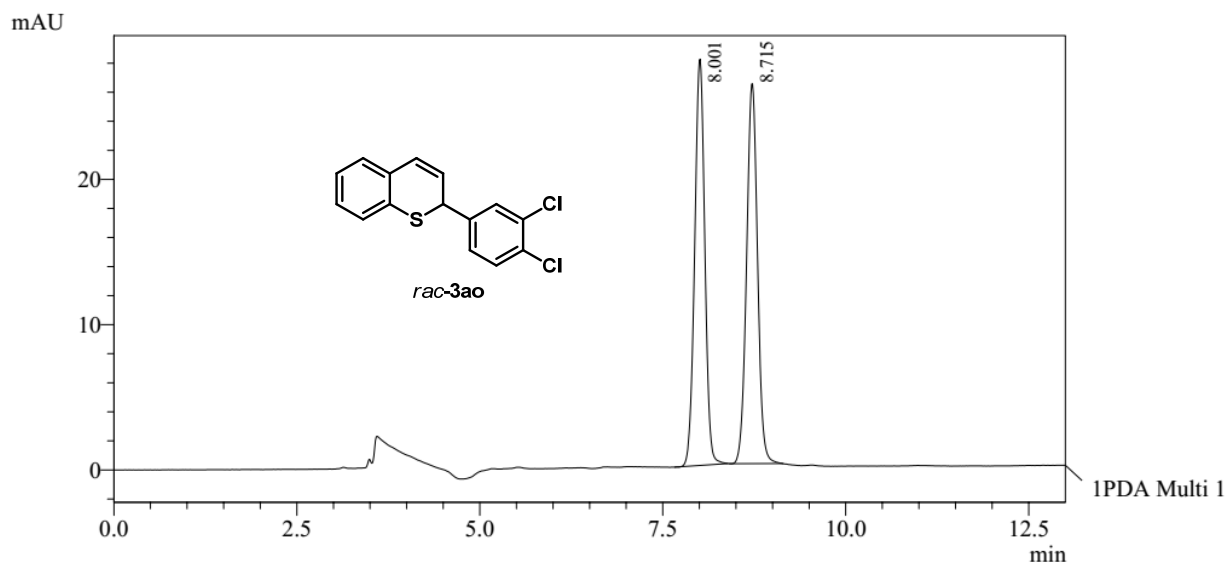
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.639	6624300	98.718
2	9.582	86000	1.282
Total		6710299	100.000



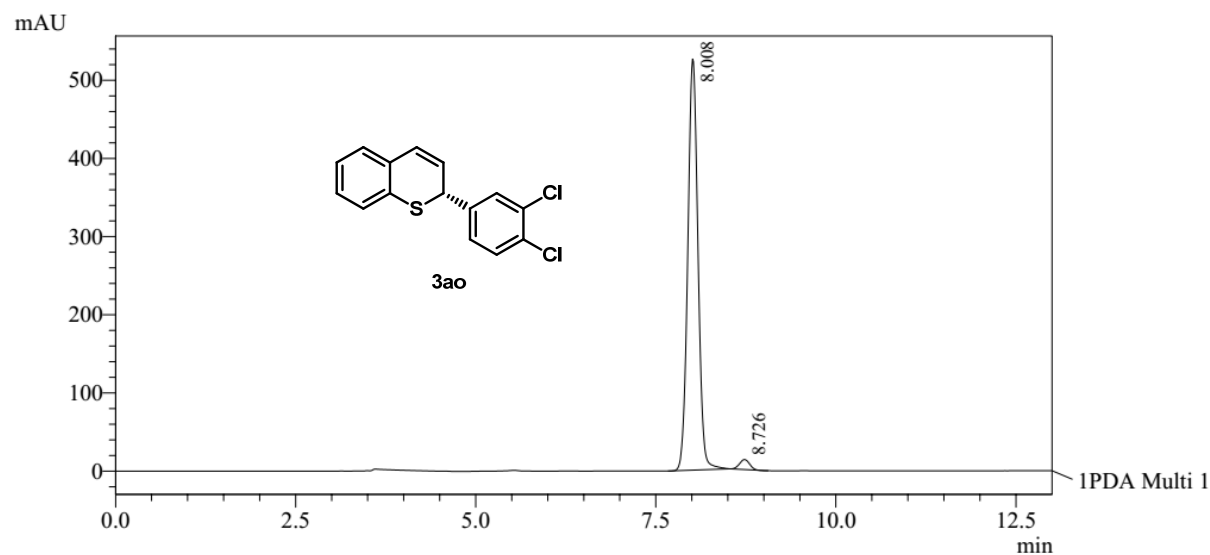


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.001	267922	49.732
2	8.715	270805	50.268
Total		538727	100.000

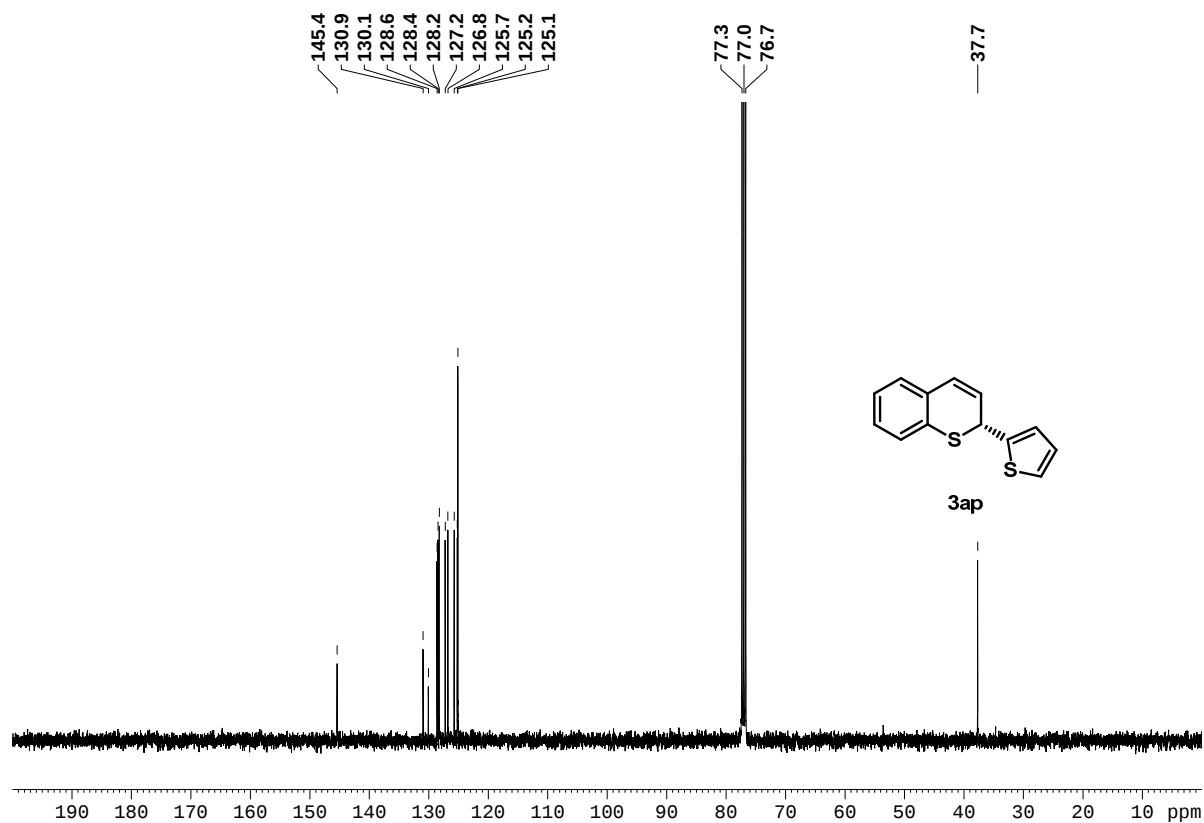
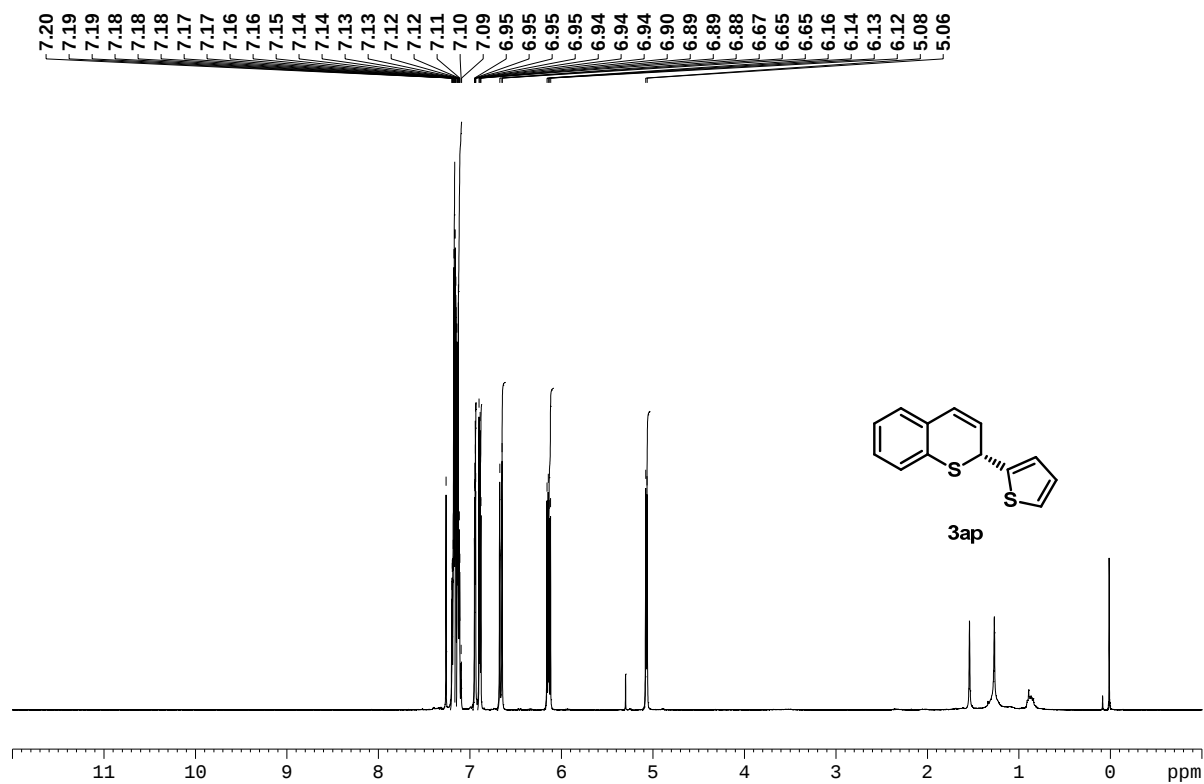
Daicel Chiralpak IB column (99:1 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

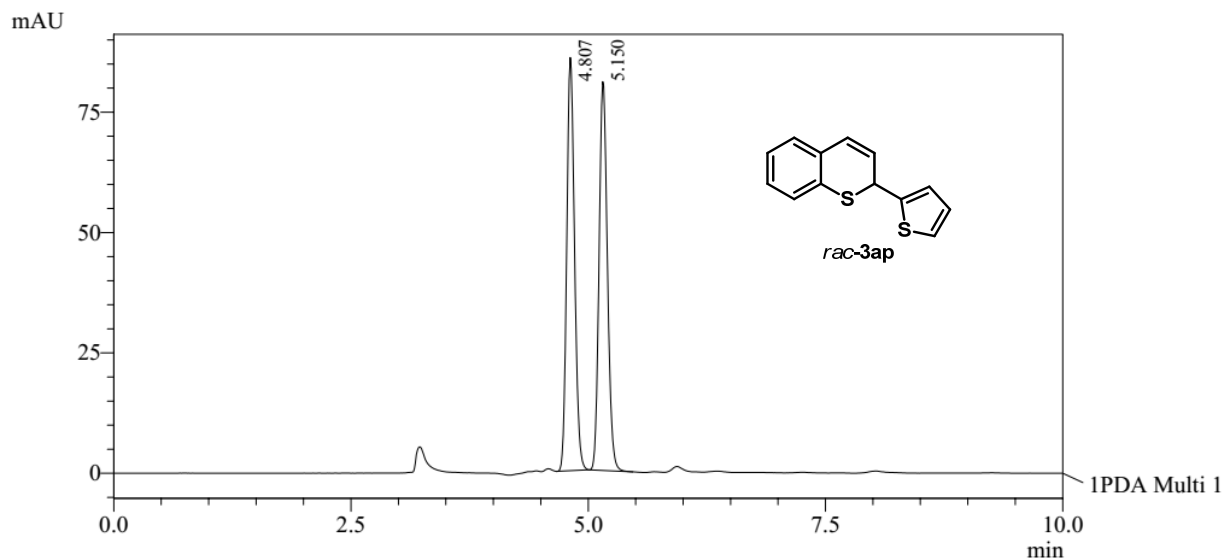


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	8.008	5139749	97.641
2	8.726	124174	2.359
Total		5263923	100.000





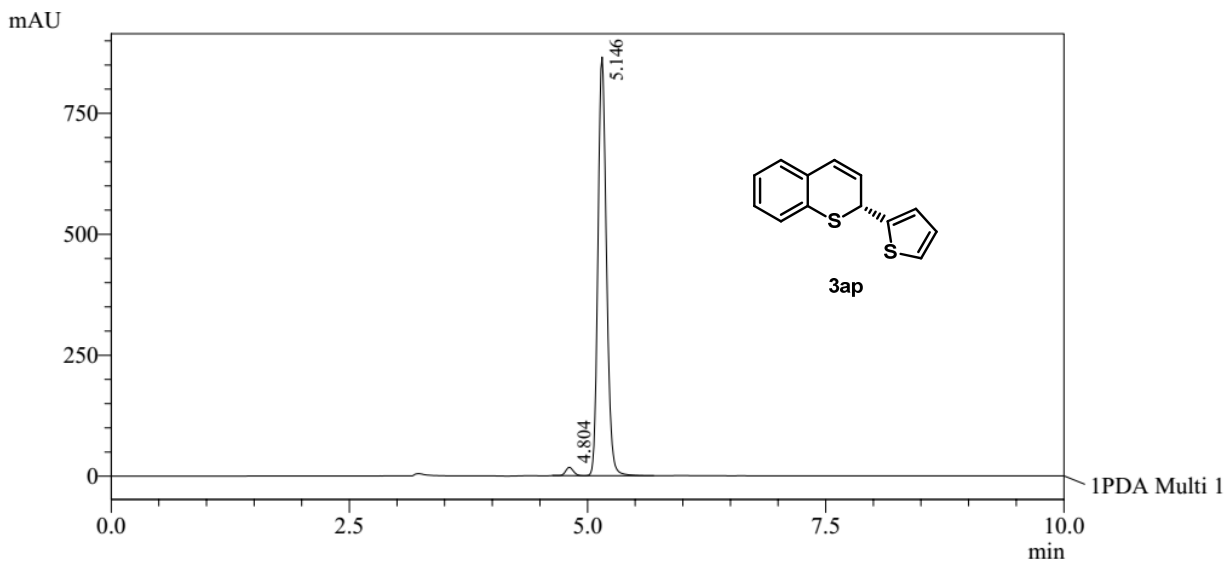
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.807	491875	49.851
2	5.150	494824	50.149
Total		986700	100.000

Daicel Chiralpak IF column (97:3 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

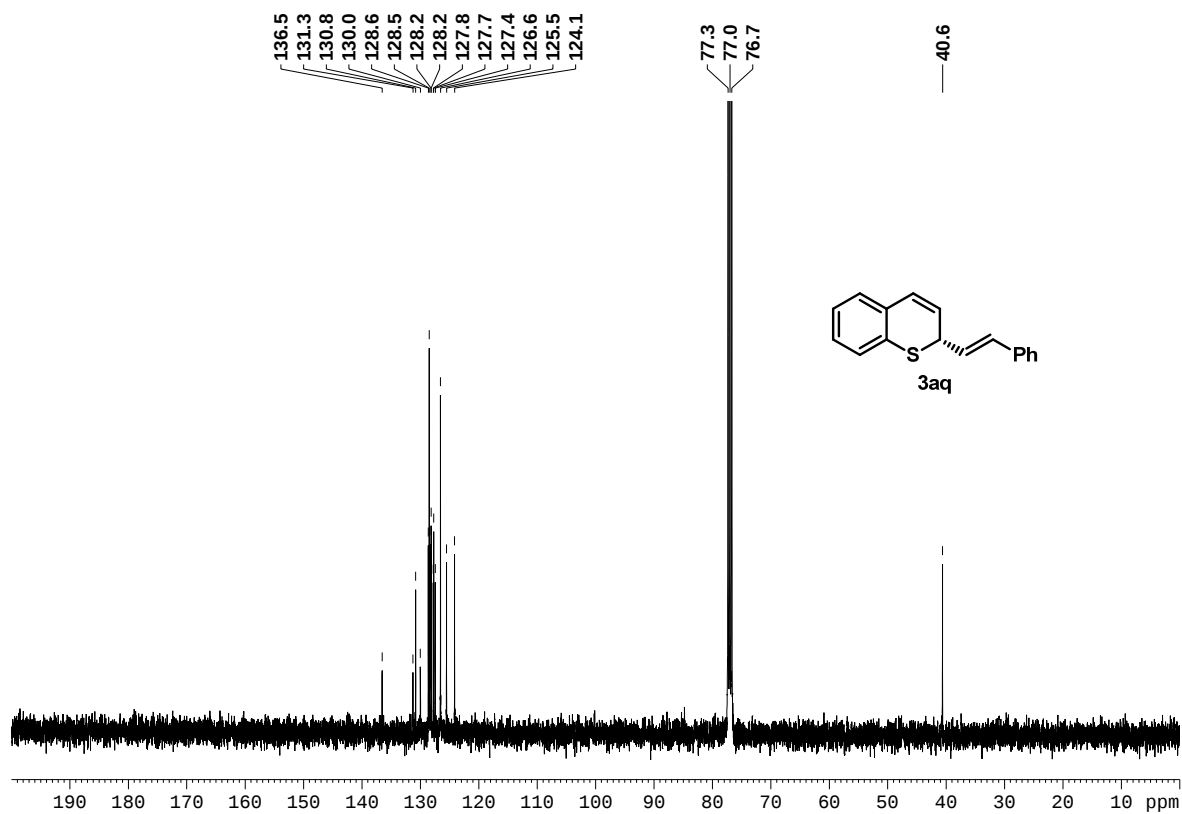
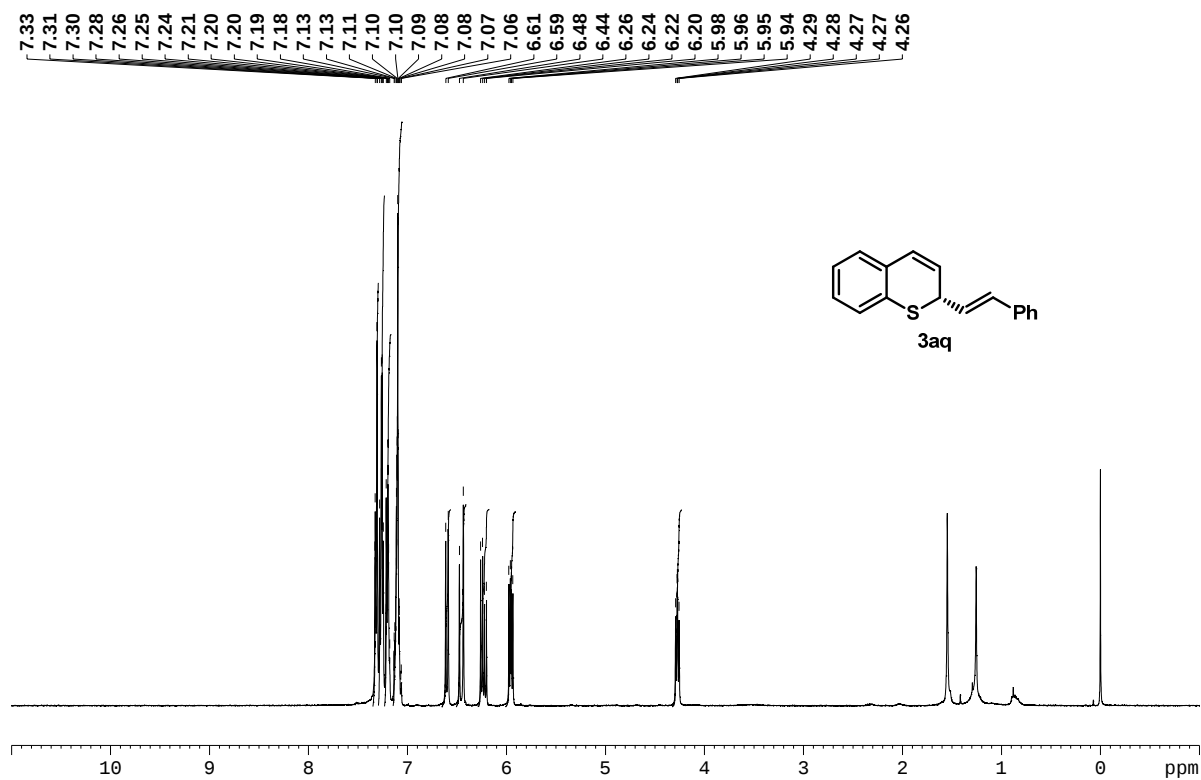


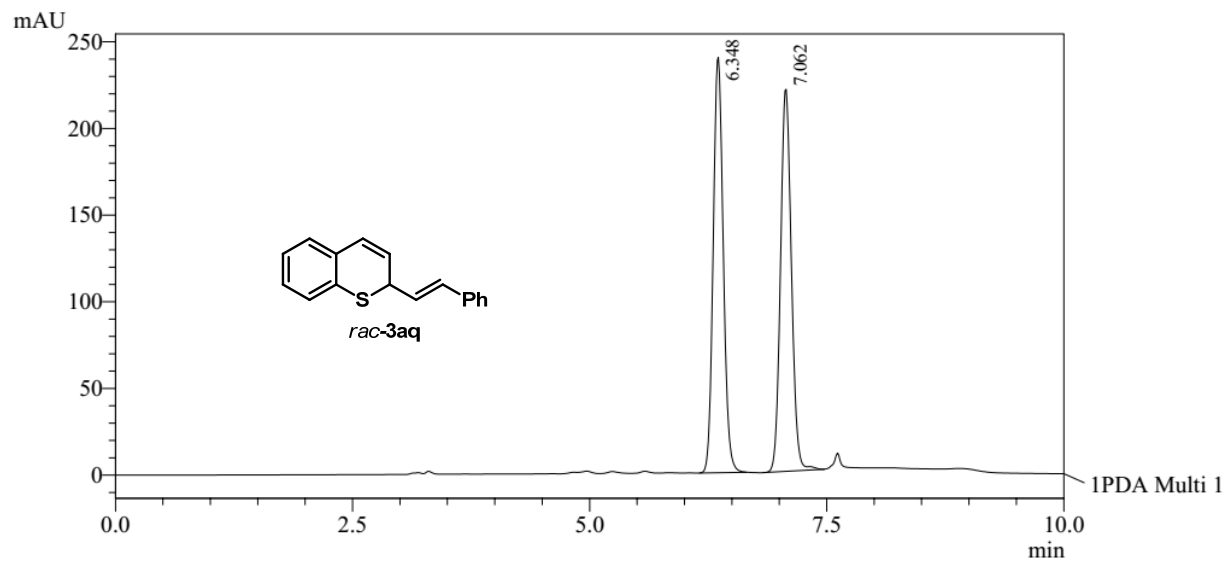
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.804	97052	1.771
2	5.146	5382797	98.229
Total		5479849	100.000





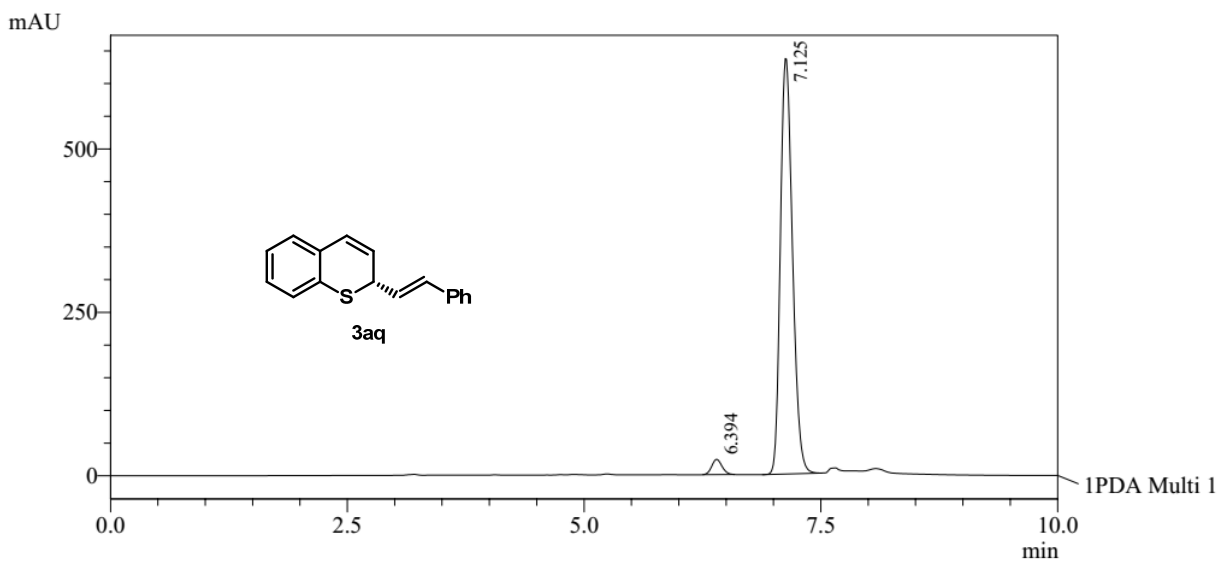
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.348	1766375	50.056
2	7.062	1762432	49.944
Total		3528807	100.000

Daicel Chiralpak IF column (80:20 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

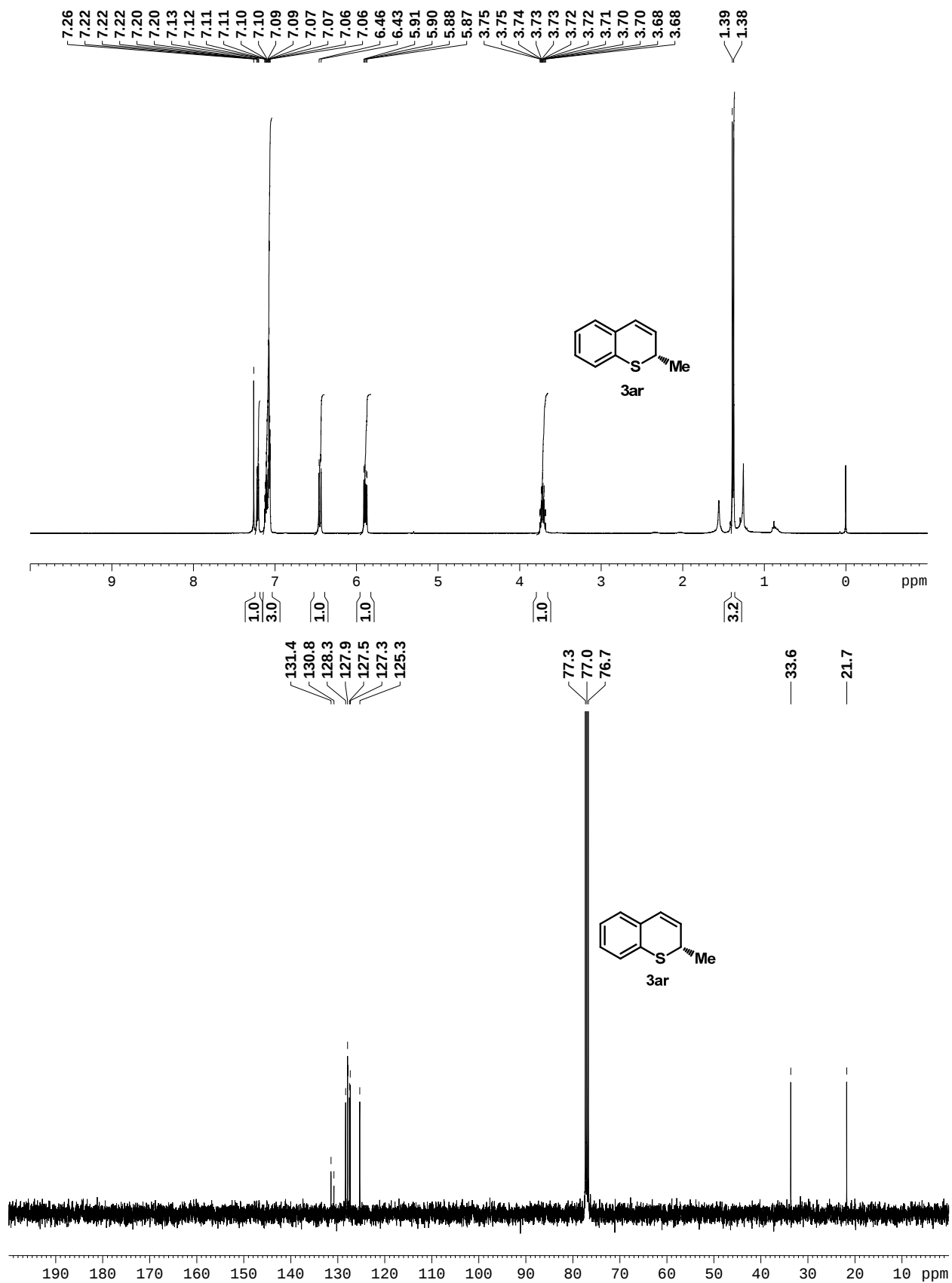


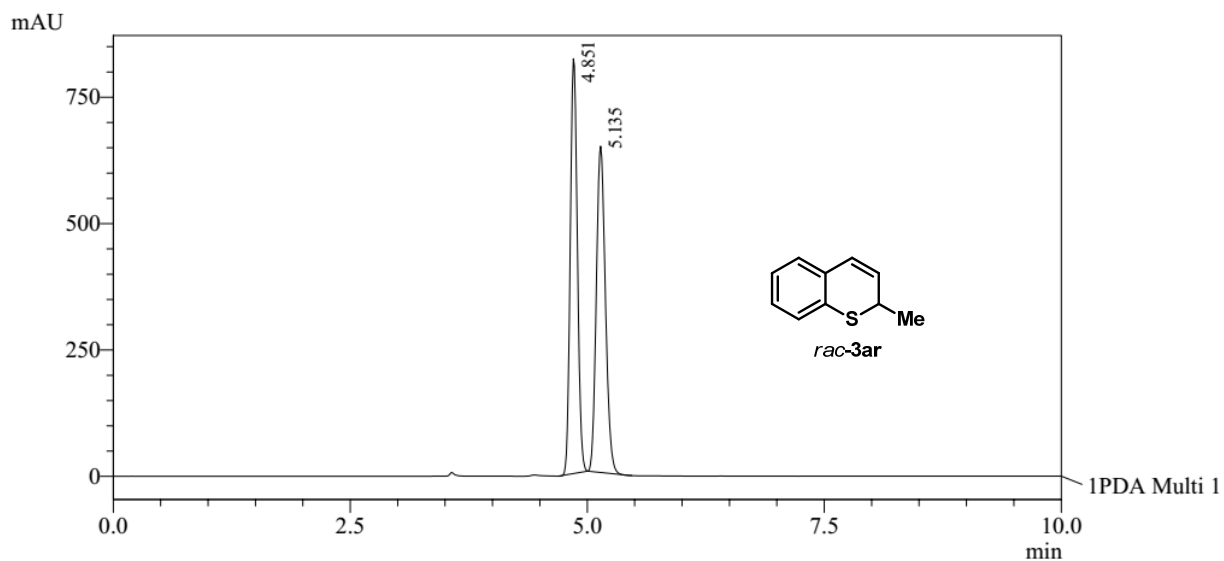
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.394	176644	3.025
2	7.125	5663627	96.975
Total		5840271	100.000





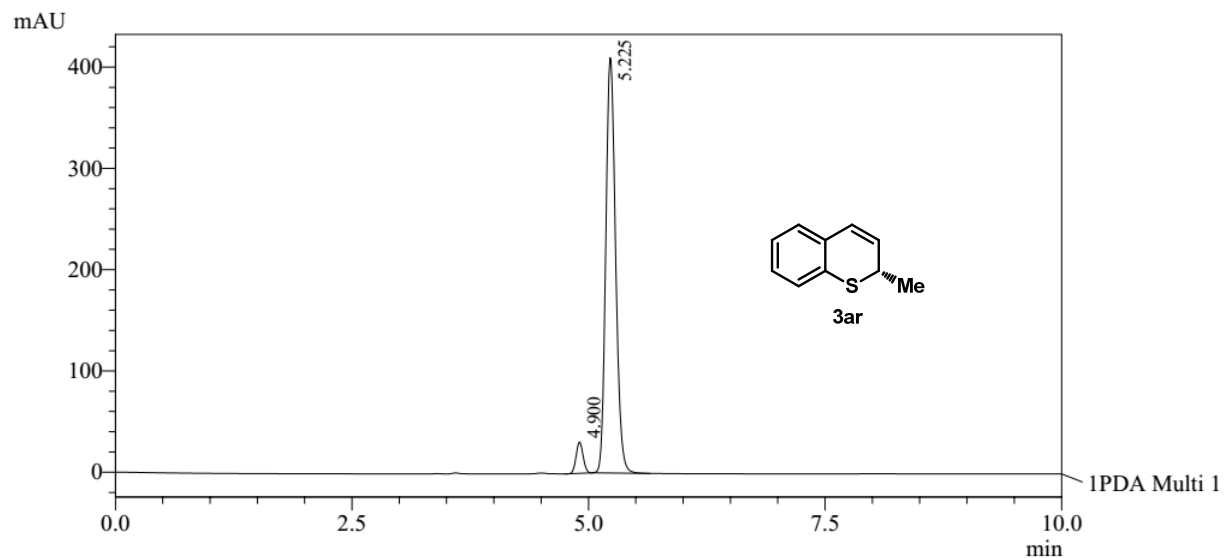
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.851	4282567	49.866
2	5.135	4305588	50.134
Total		8588156	100.000

Daicel Chiralpak ID column (95:5 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

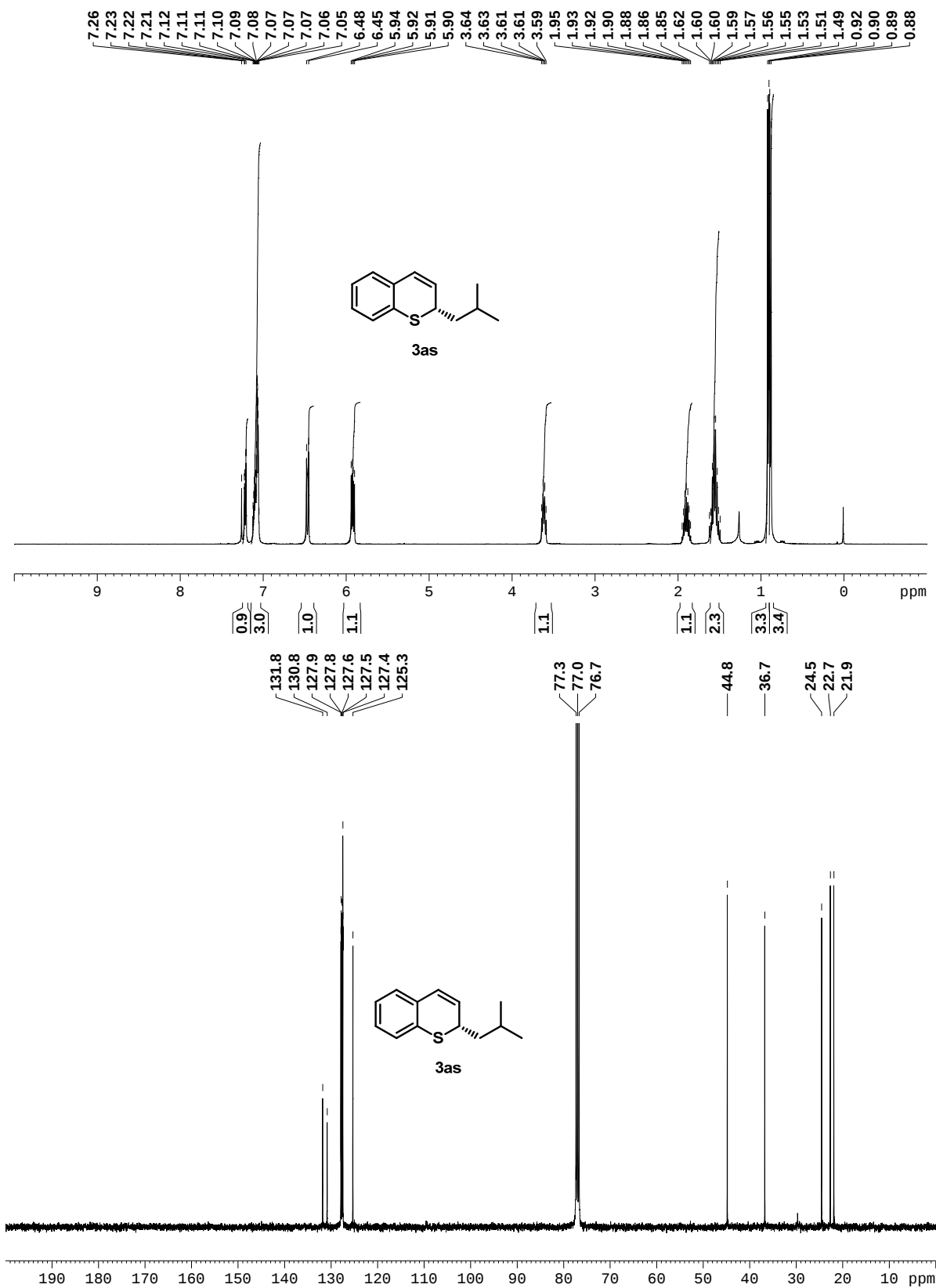


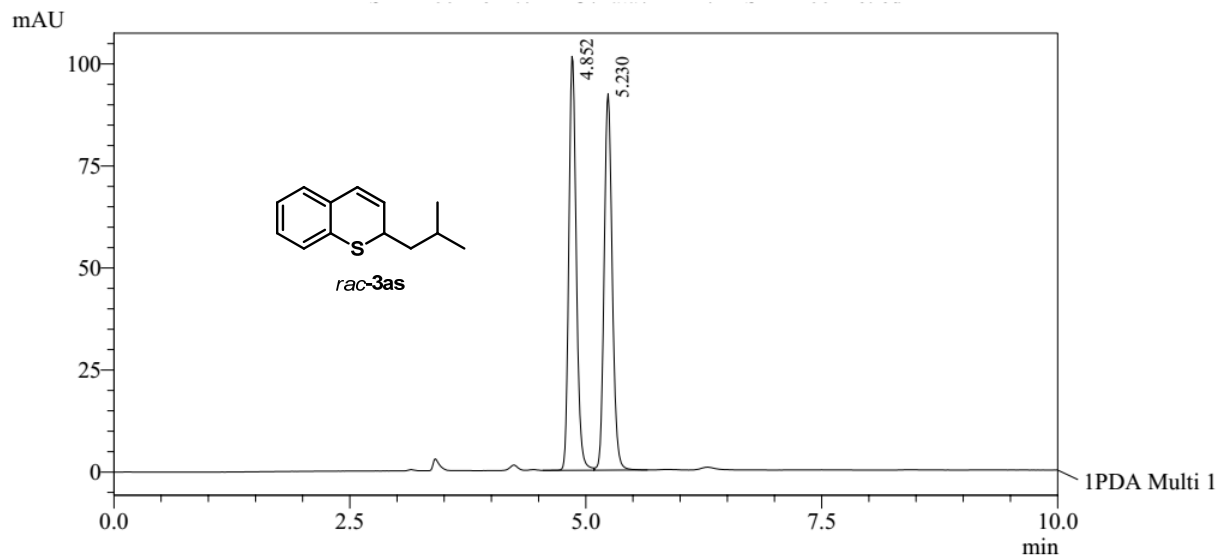
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.900	158577	5.097
2	5.225	2952876	94.903
Total		3111453	100.000





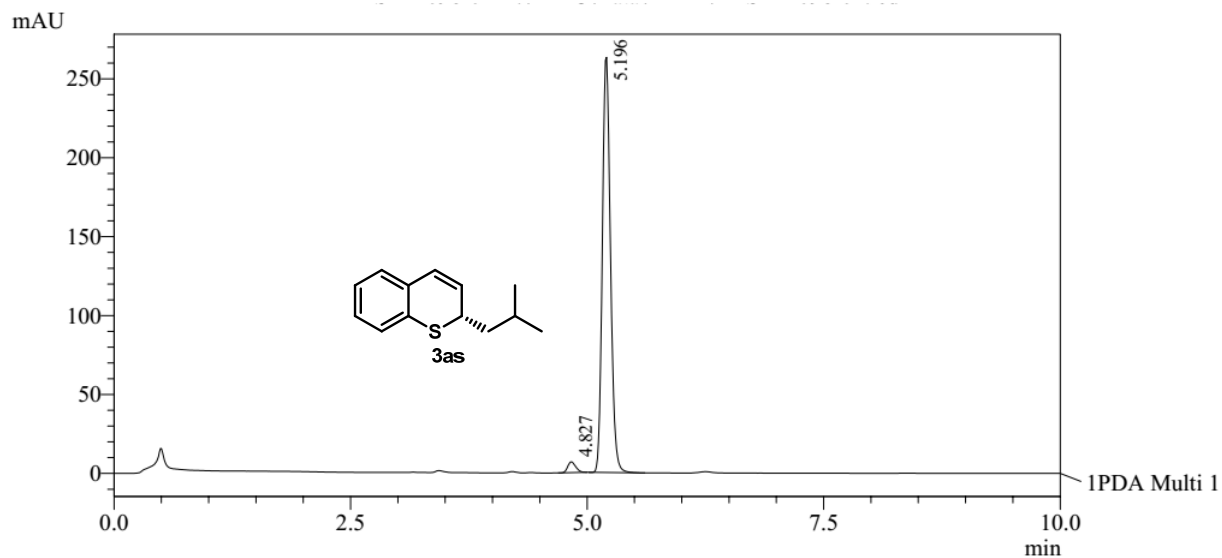
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.852	557905	50.349
2	5.230	550168	49.651
Total		1108073	100.000

Daicel Chiralpak IF column (95:5 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

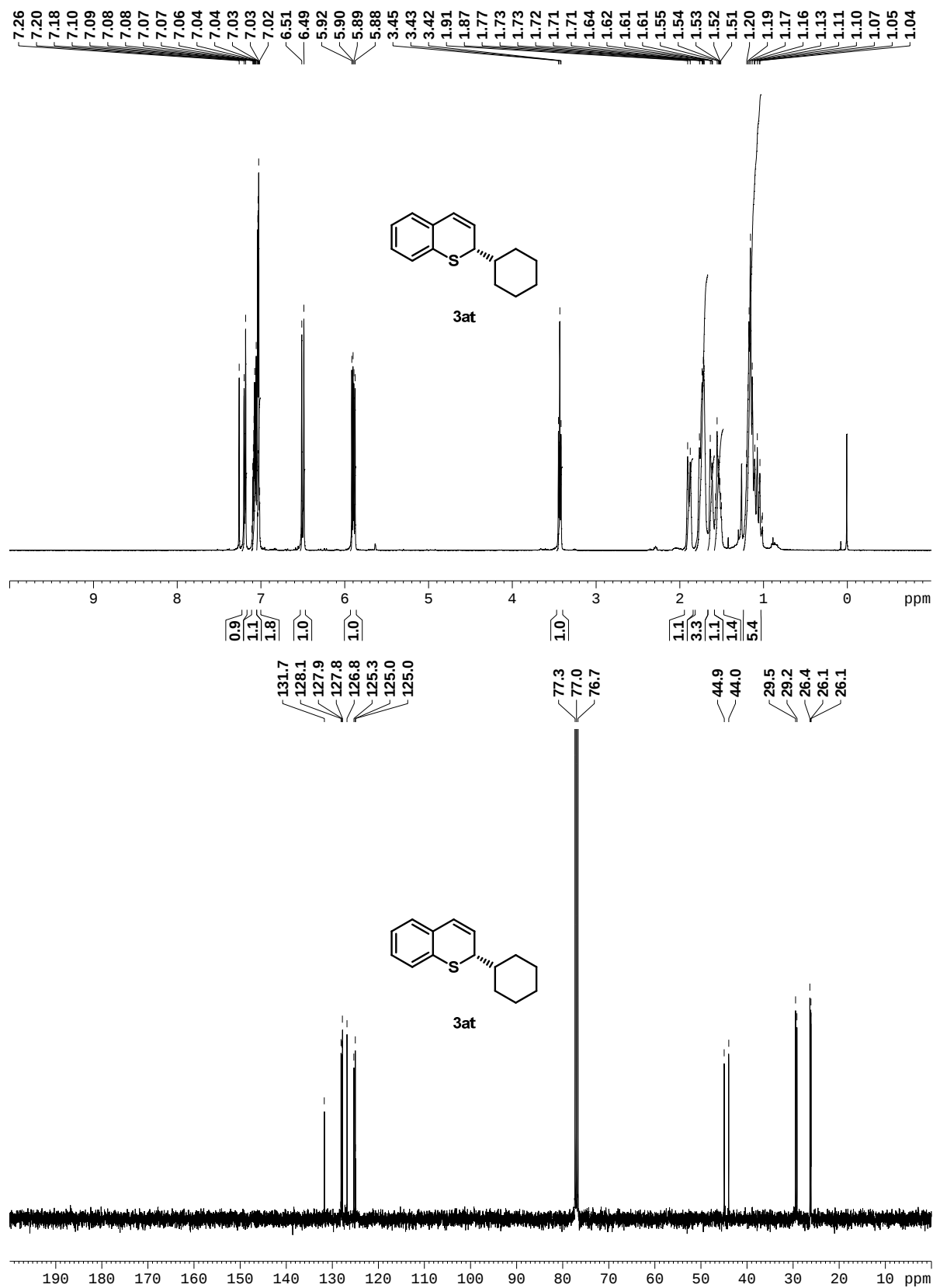


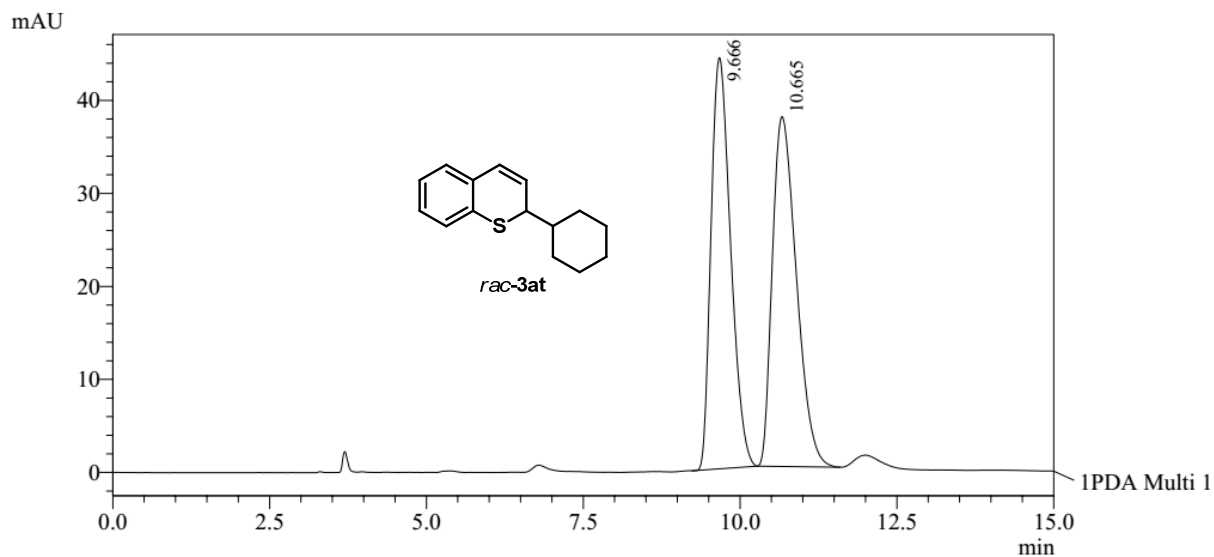
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.827	40408	2.506
2	5.196	1572248	97.494
Total		1612656	100.000





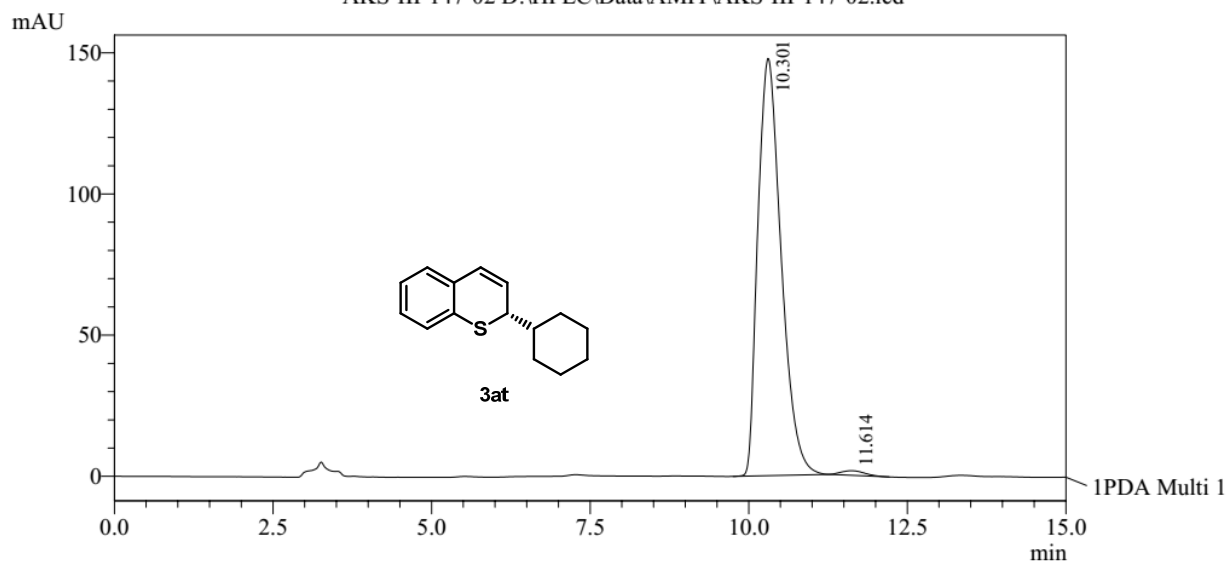
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	9.666	959531	48.921
2	10.665	1001863	51.079
Total		1961394	100.000

Daicel Chiralpak IE column (99:01 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

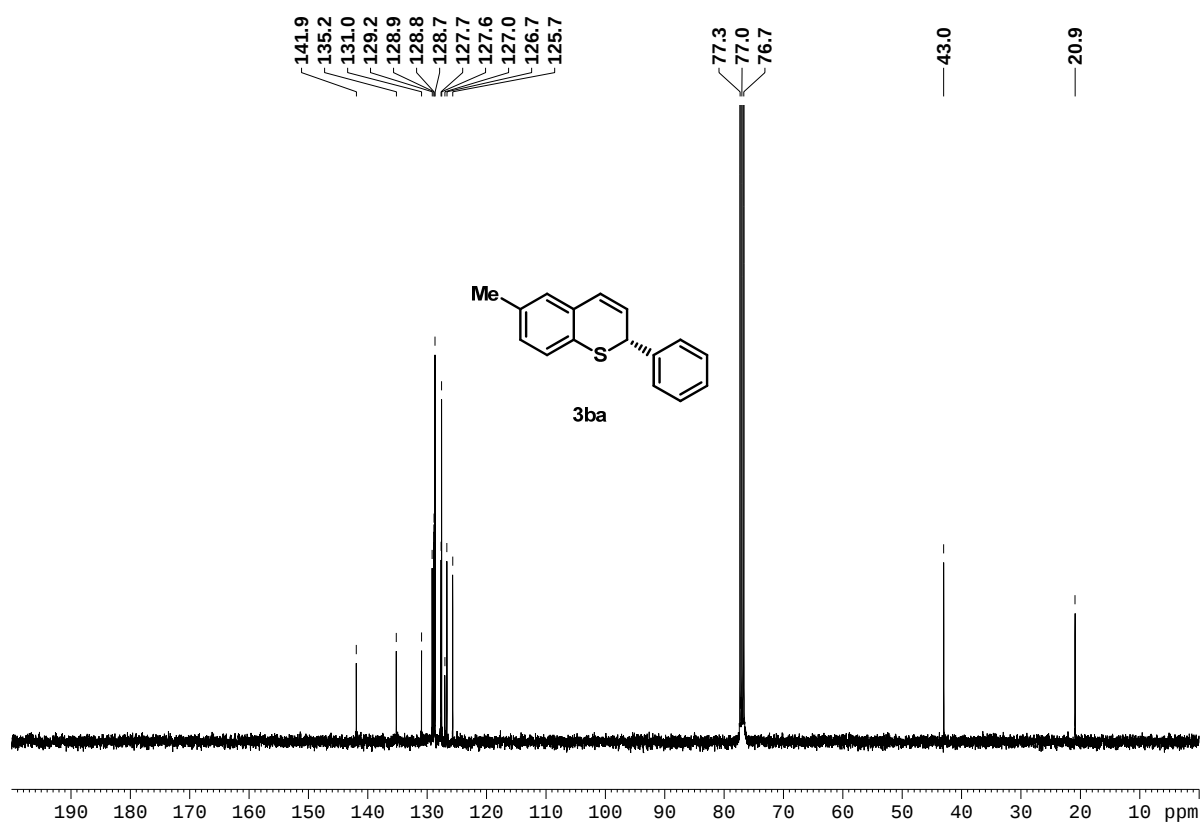
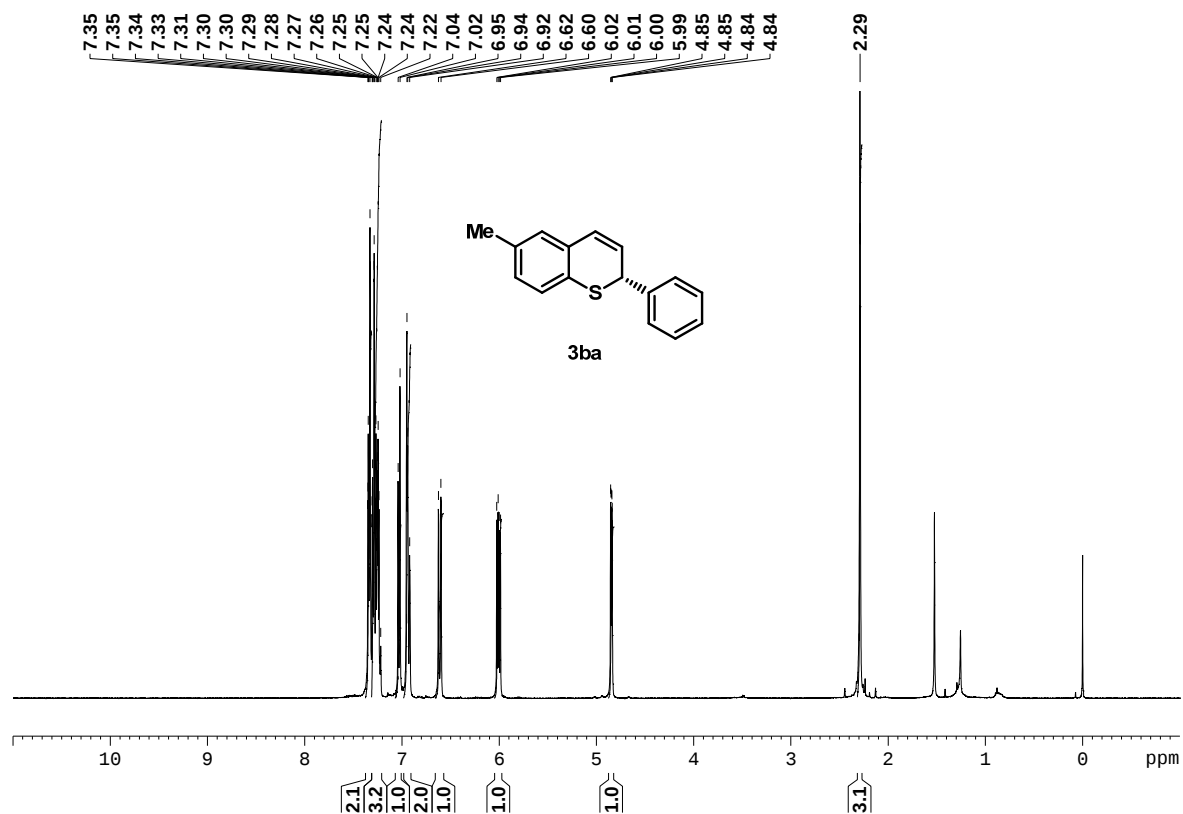


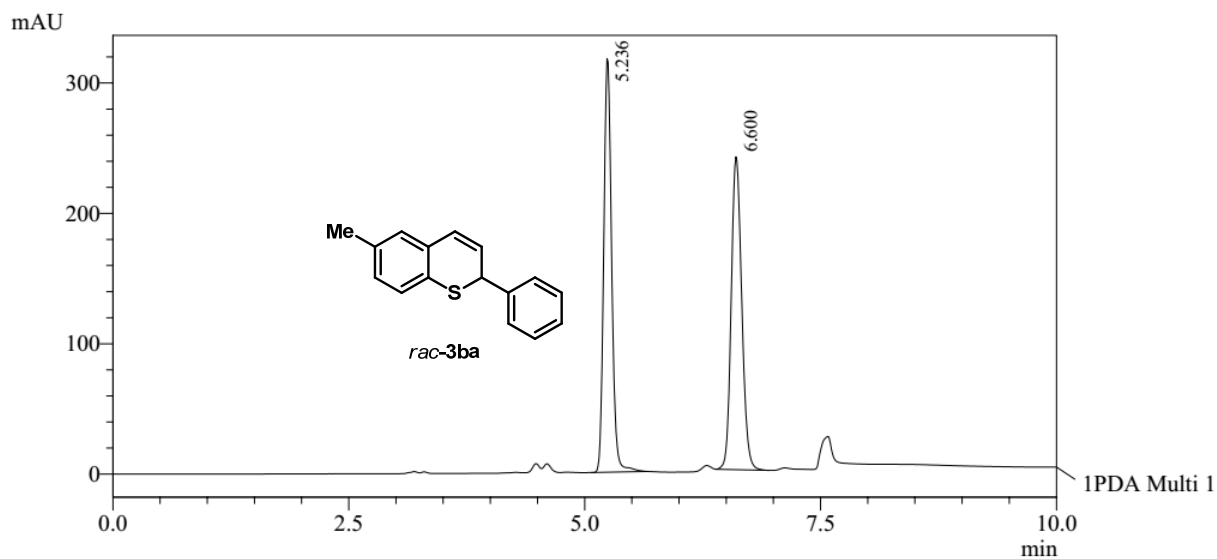
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	10.301	3775753	98.915
2	11.614	41435	1.085
Total		3817188	100.000





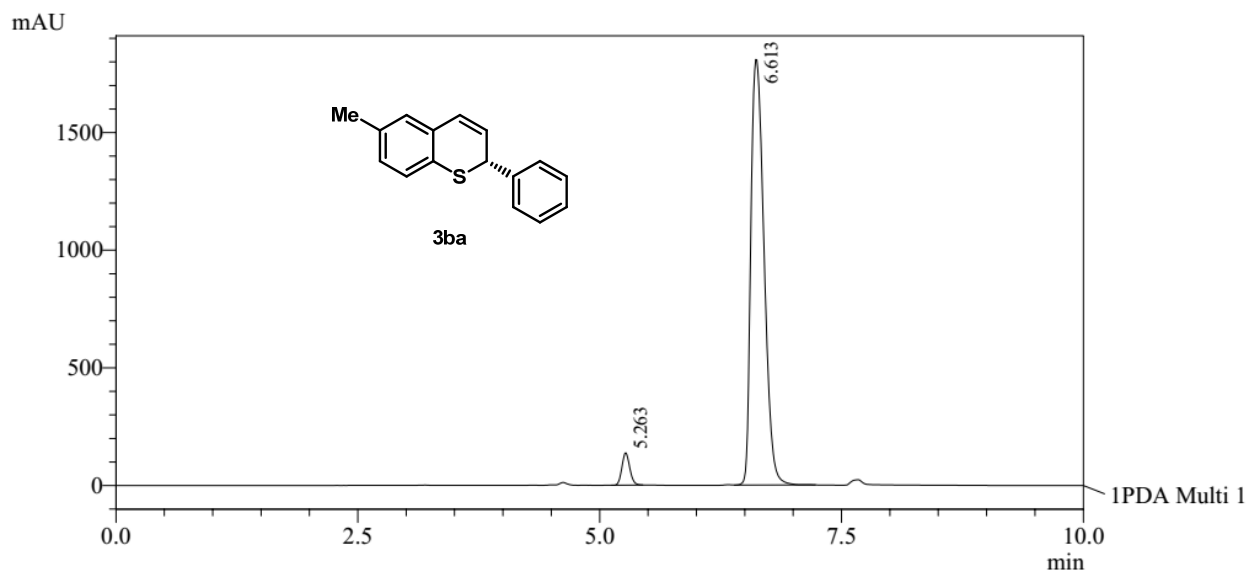
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.236	1854387	49.954
2	6.600	1857812	50.046
Total		3712198	100.000

Daicel Chiralpak IF column (80:20 *n*-Hexane/MTBE, 1.0 mL/min, 20 °C, 254 nm)

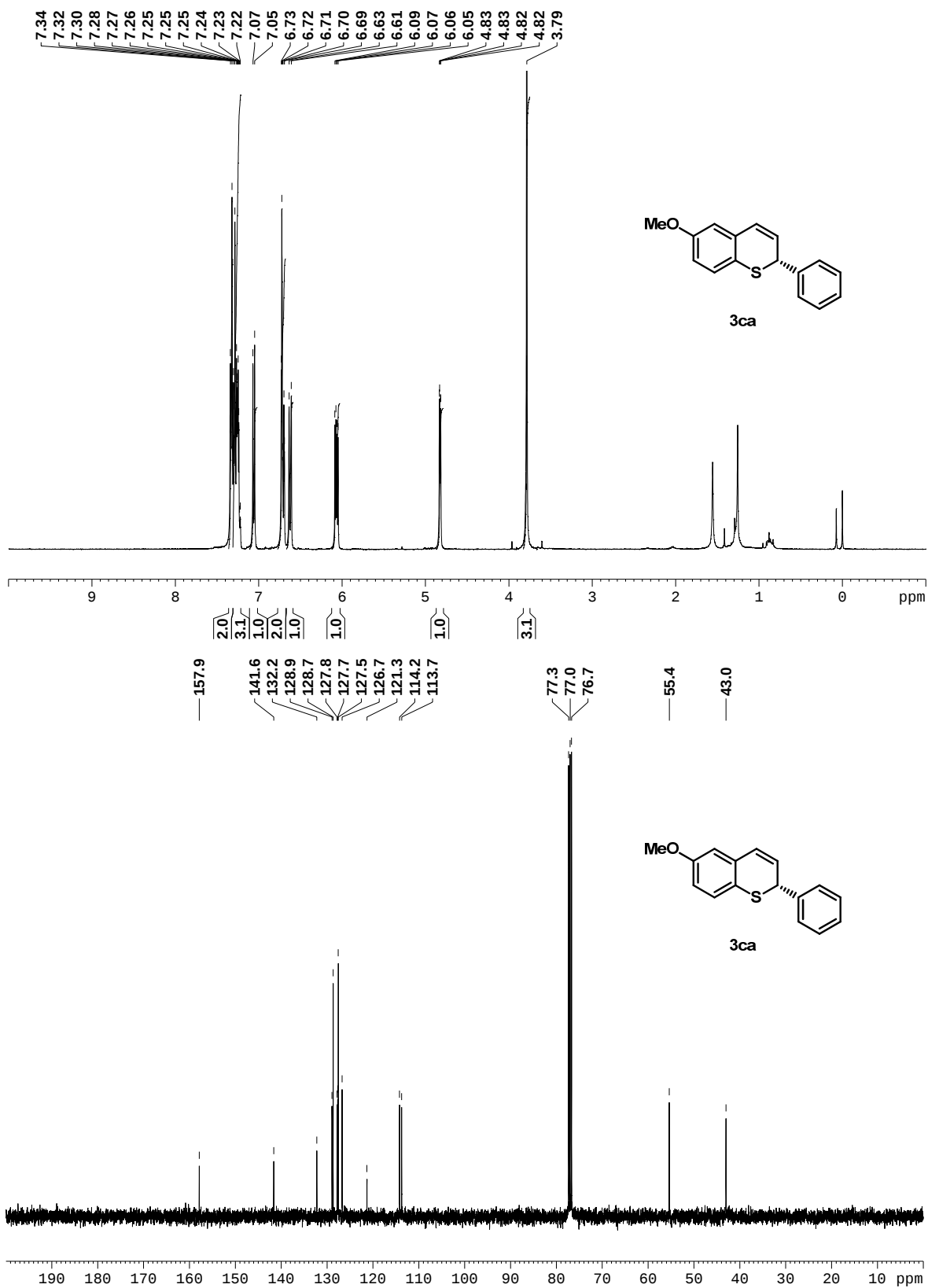


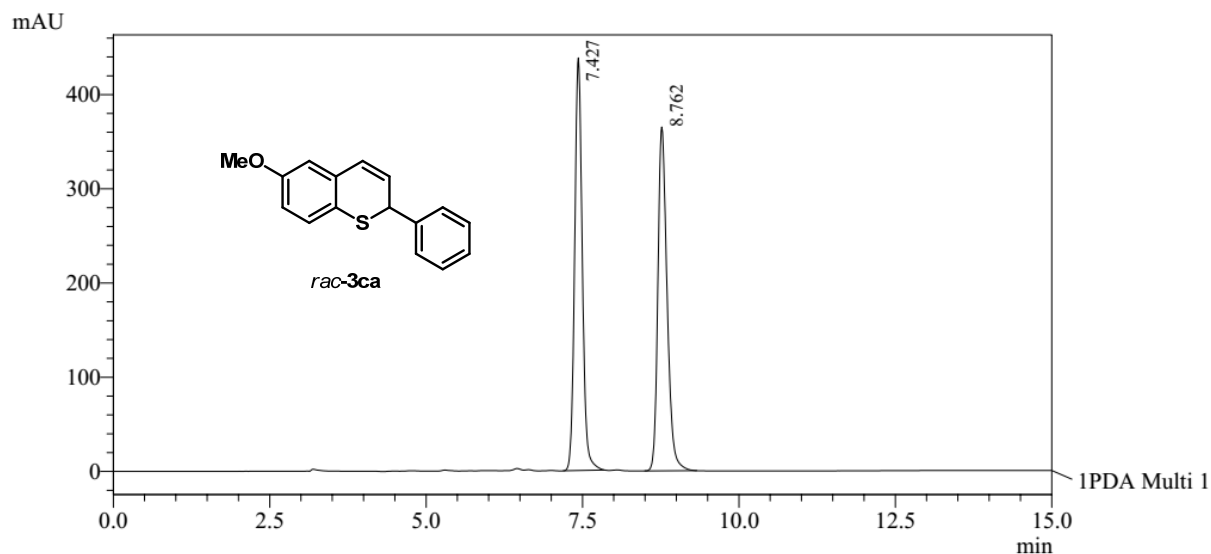
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.263	778269	4.339
2	6.613	17158836	95.661
Total		17937105	100.000





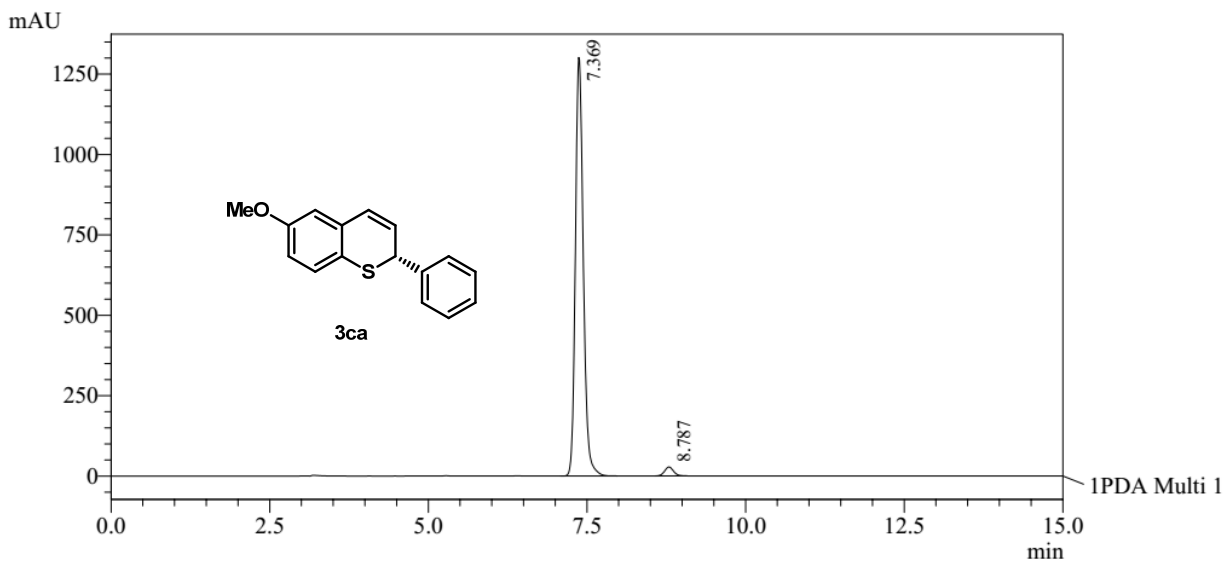
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	7.427	3719164	49.878
2	8.762	3737392	50.122
Total		7456556	100.000

Daicel Chiralpak IB column (97:03 *n*-Hexane/ *i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

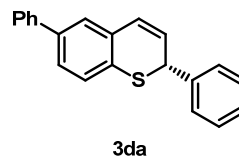
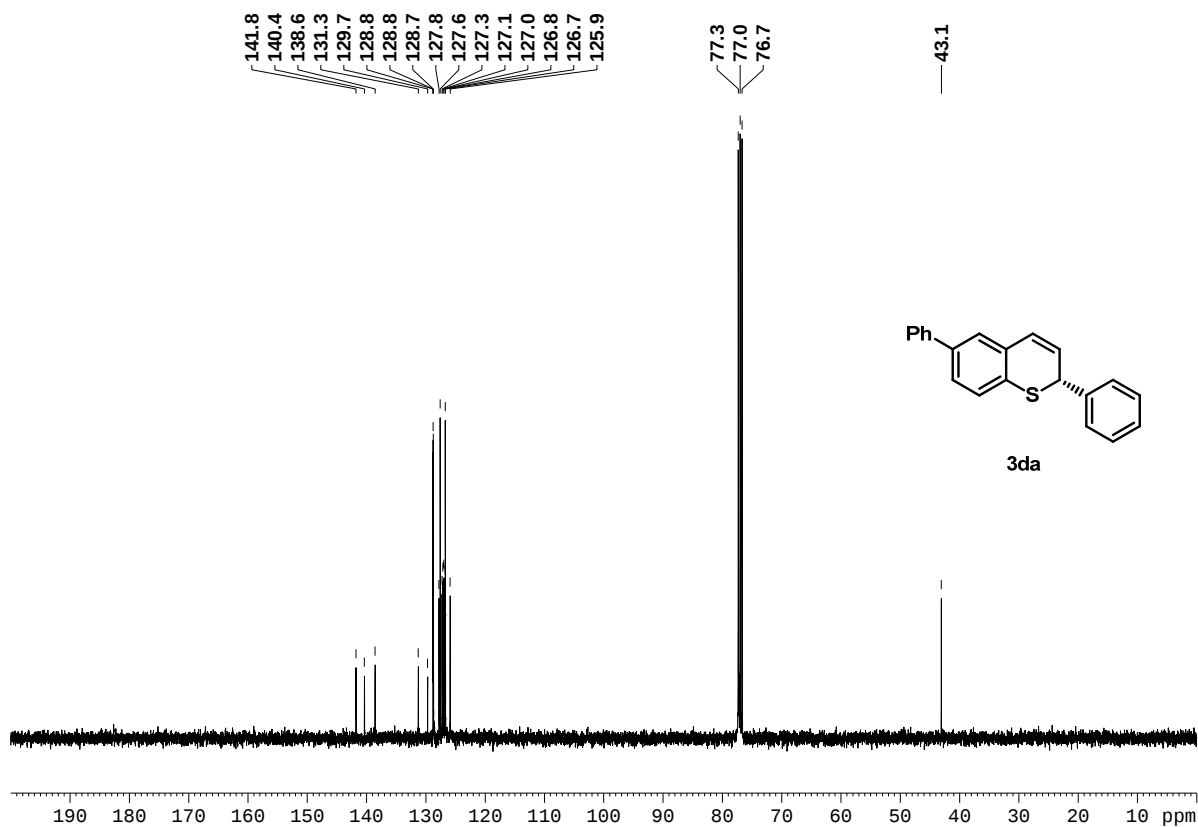
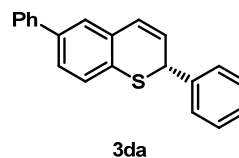
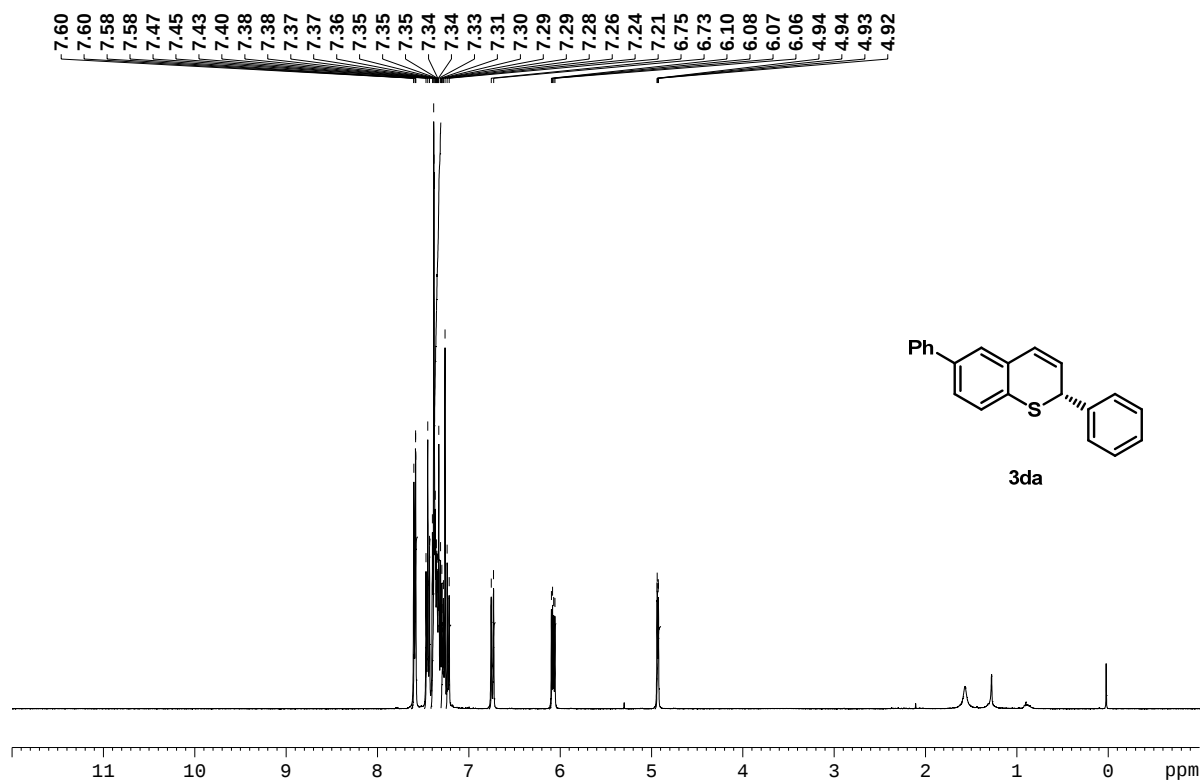


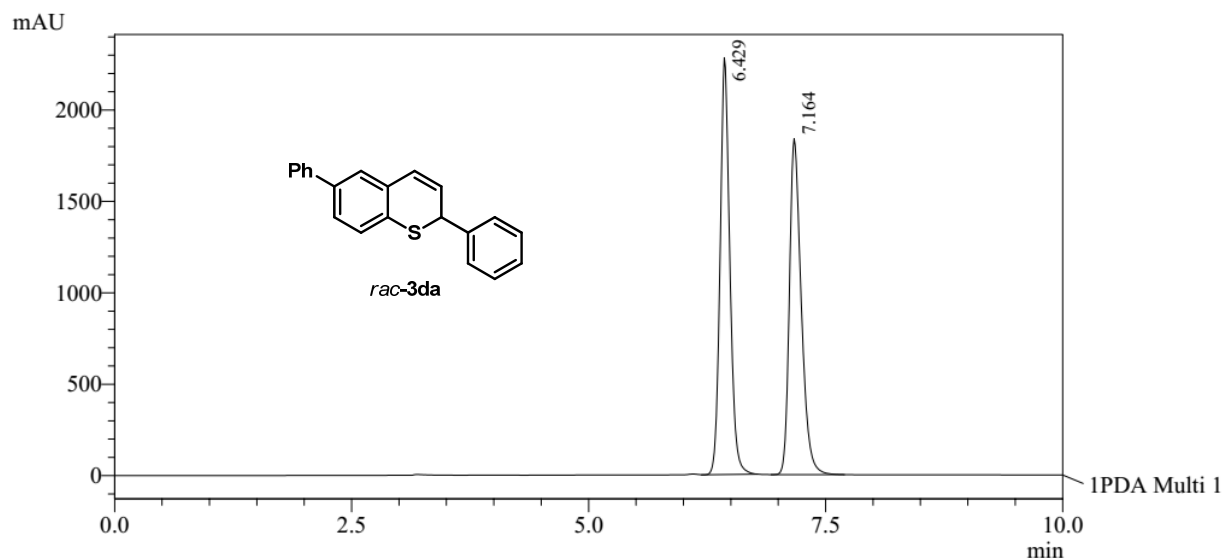
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	7.369	11305147	97.662
2	8.787	270657	2.338
Total		11575804	100.000





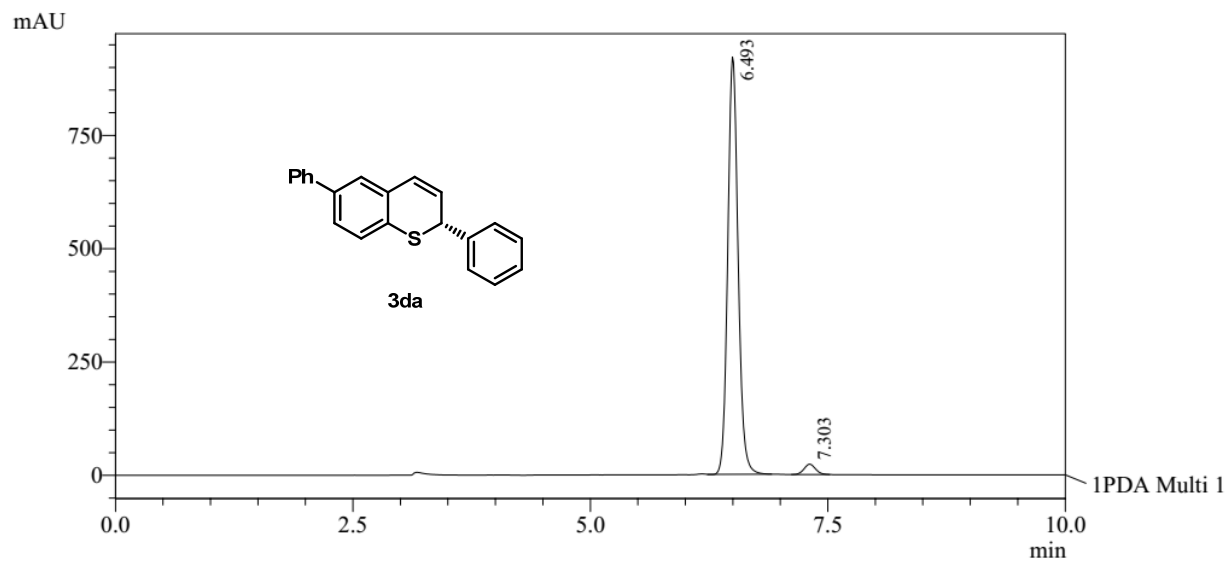
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.429	16278204	50.469
2	7.164	15975850	49.531
Total		32254054	100.000

Daicel Chiralpak IB column (97:03 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

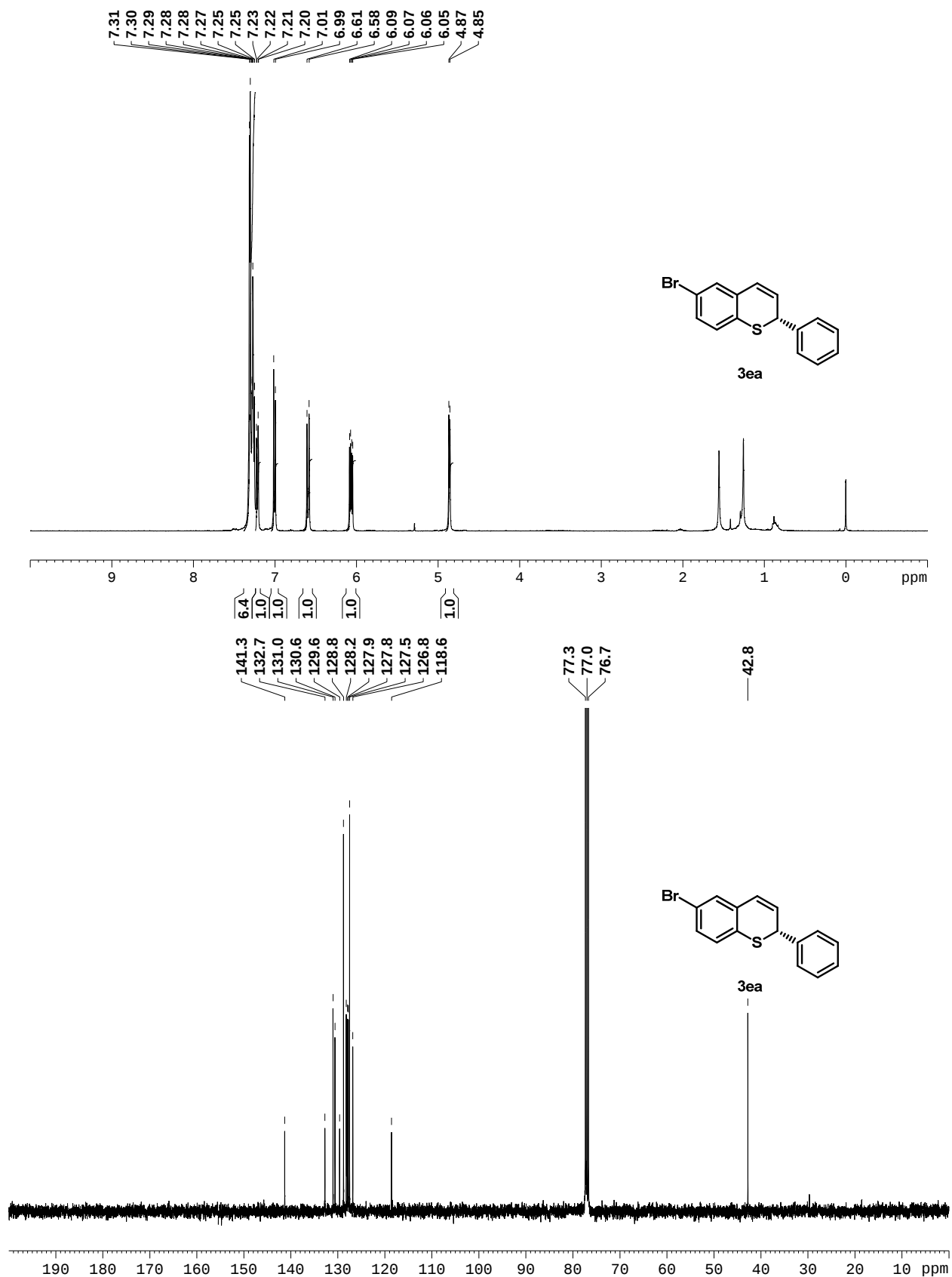


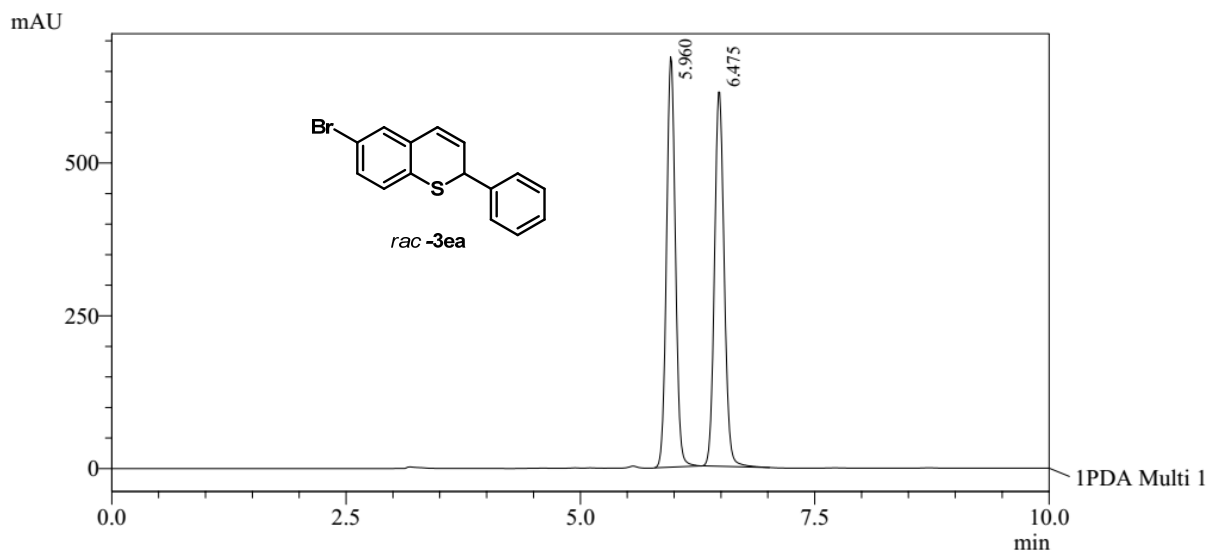
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.493	6977446	97.338
2	7.303	190808	2.662
Total		7168254	100.000



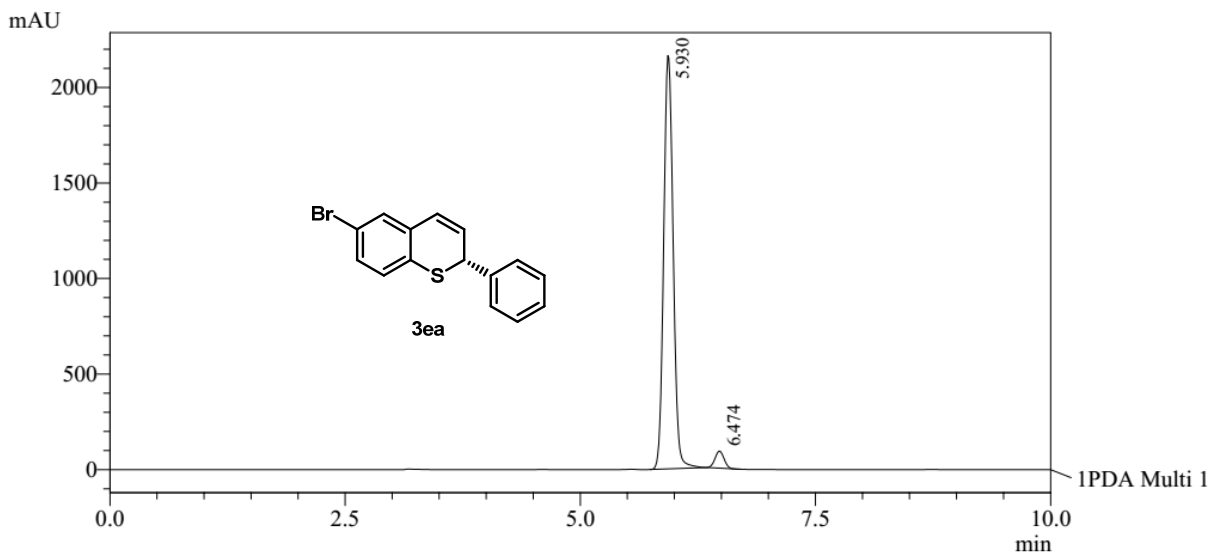


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.960	4419442	50.004
2	6.475	4418701	49.996
Total		8838143	100.000

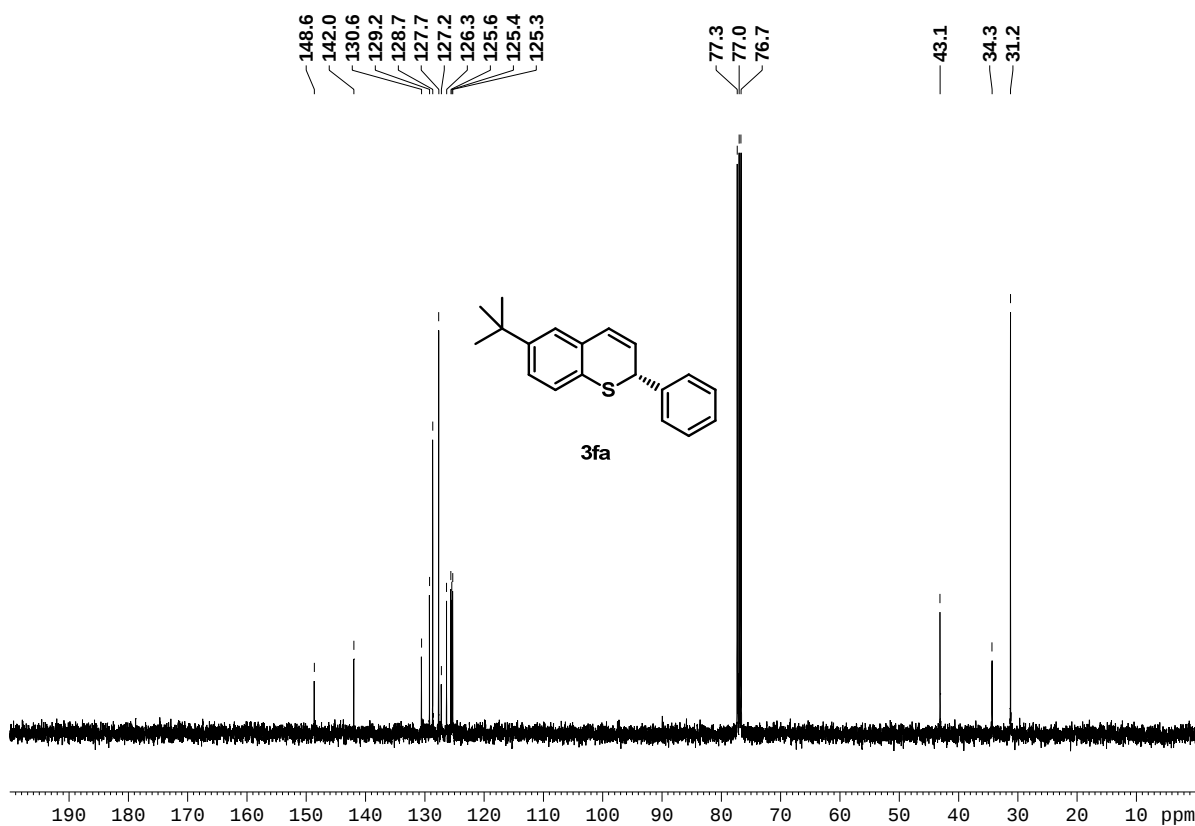
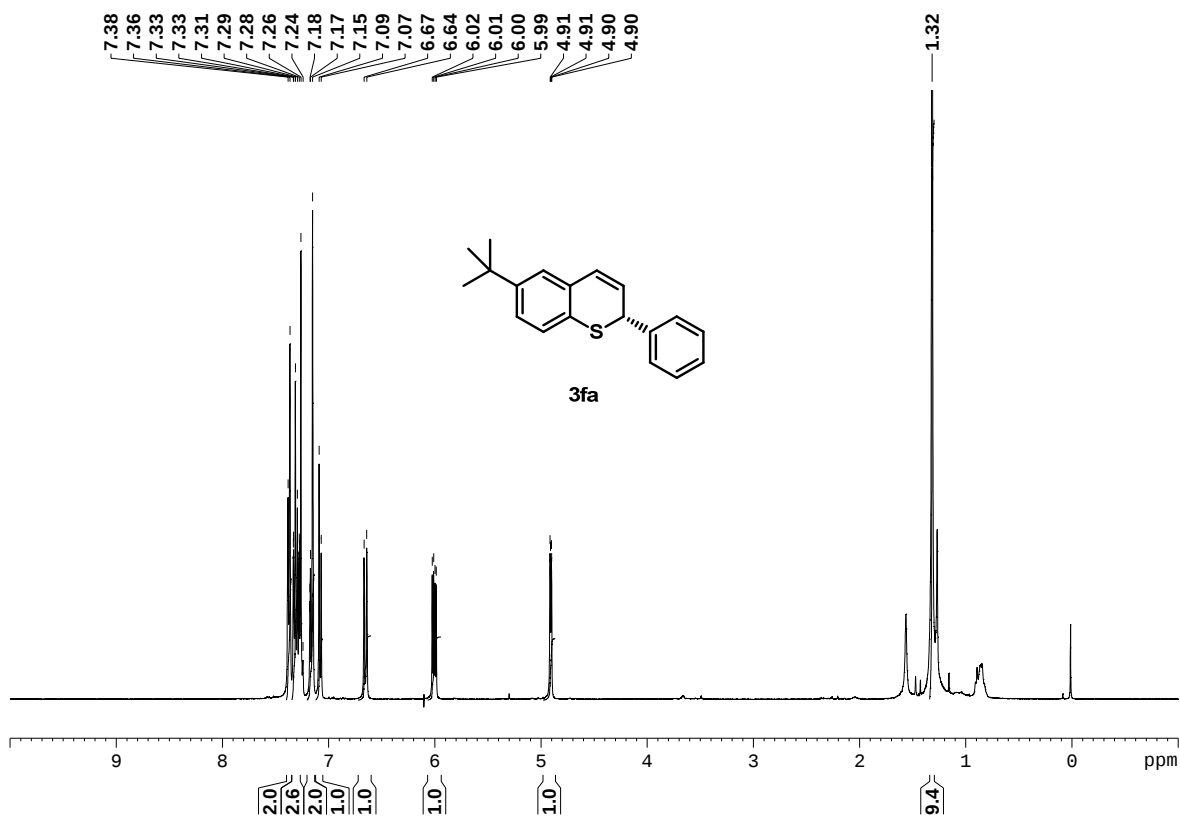
Daicel Chiralpak IB column (97:03 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

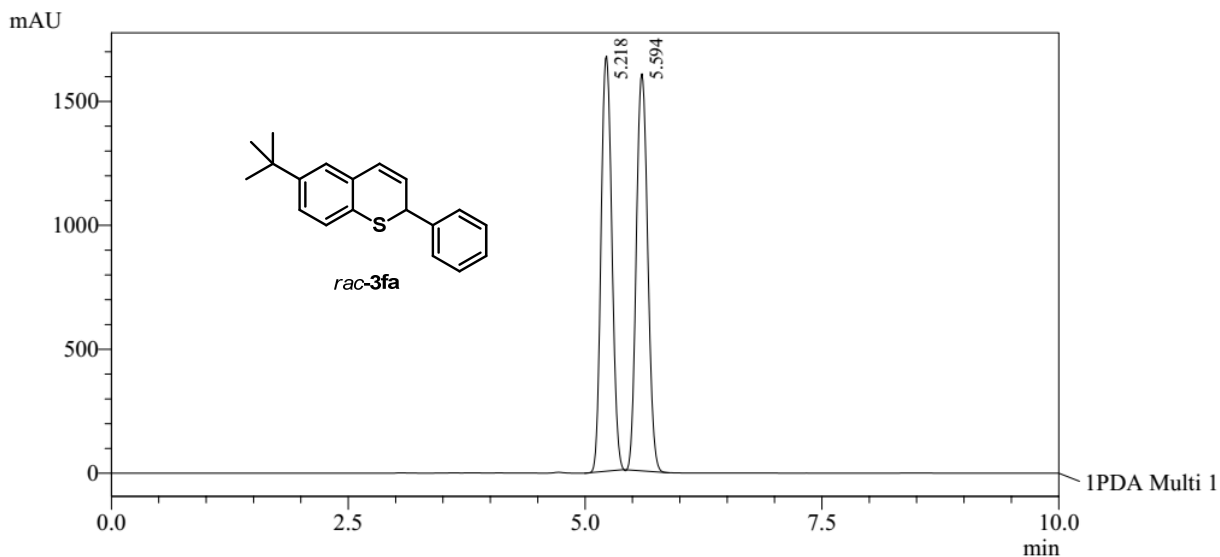


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.930	15174487	96.211
2	6.474	597673	3.789
Total		15772159	100.000





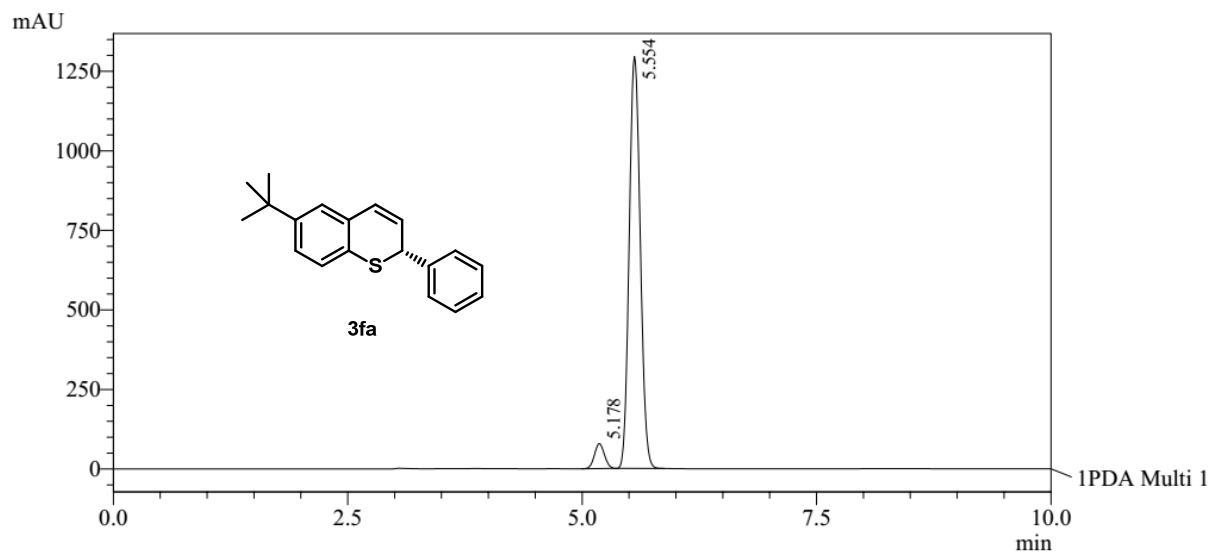
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.218	13254328	49.702
2	5.594	13413074	50.298
Total		26667402	100.000

Phenomenex Cellulose-1 column (97:03 *n*-Hexane/ *i*-PrOH, 1.0 mL/min, 20 °C, 254 nm

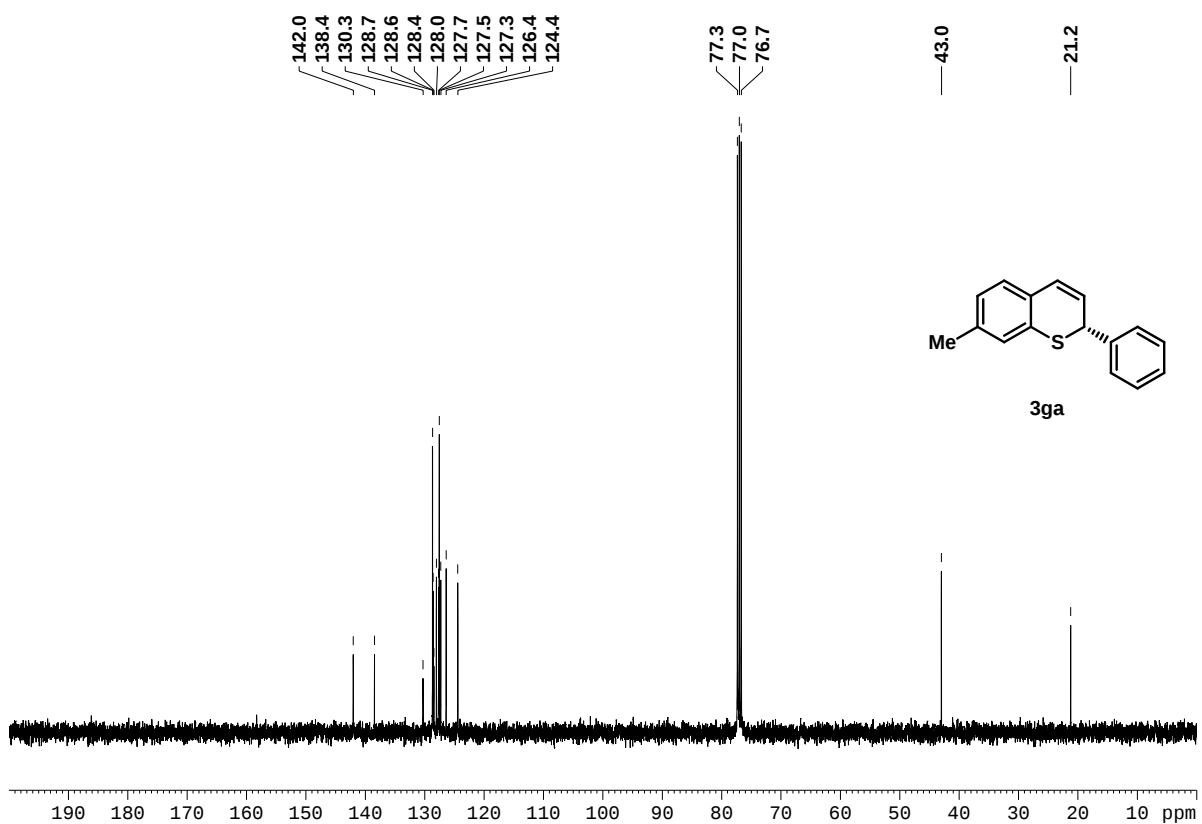
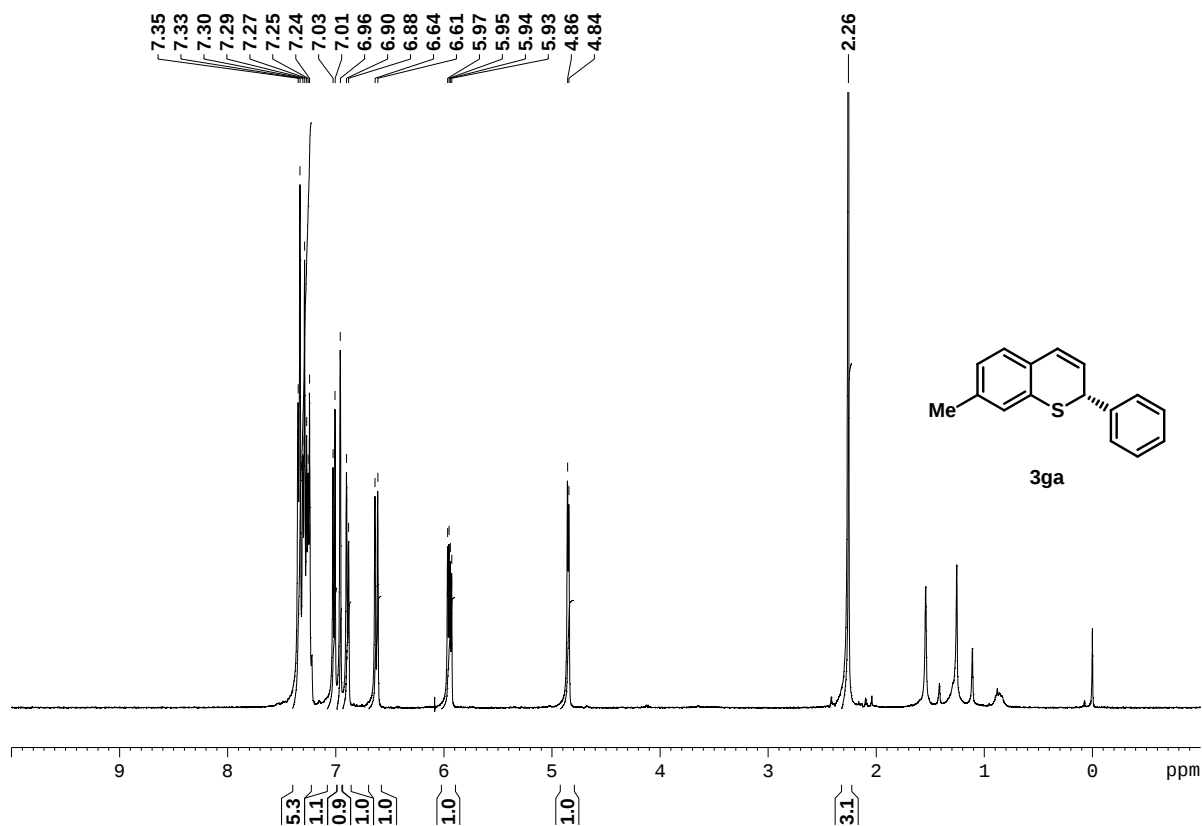


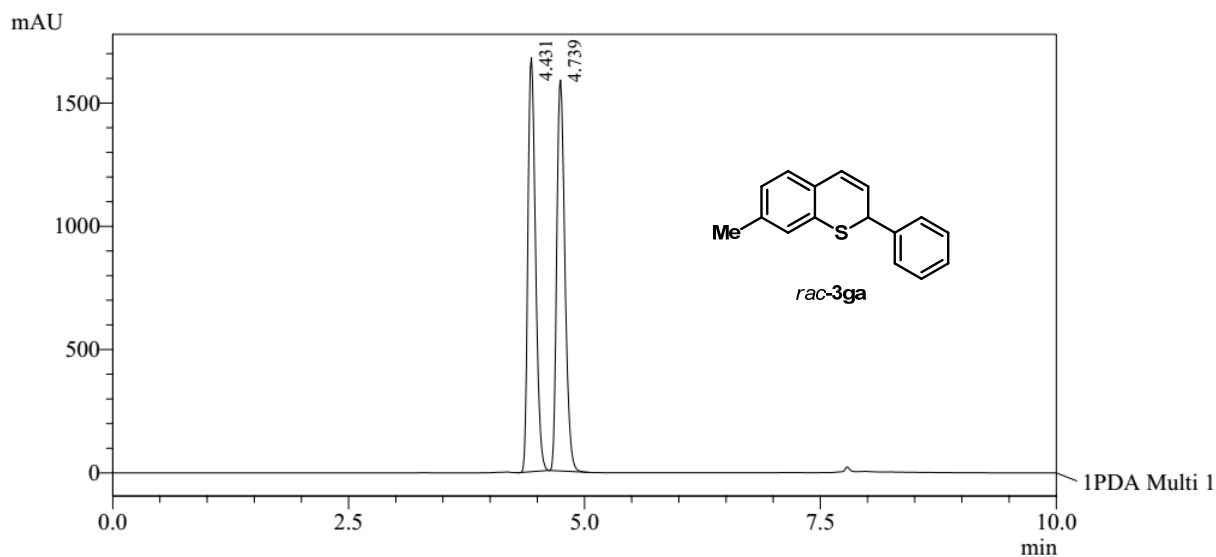
1 PDA Multi 1 / 254nm 4nm

PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.178	579865	5.170
2	5.554	10635523	94.830
Total		11215388	100.000



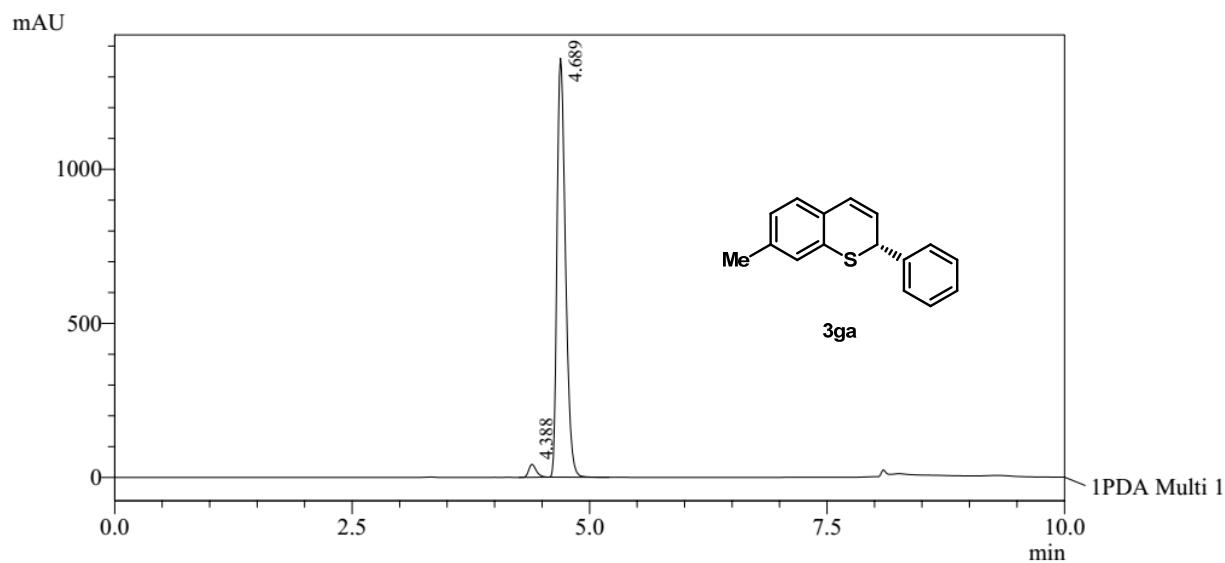


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.431	9456192	49.732
2	4.739	9558207	50.268
Total		19014399	100.000

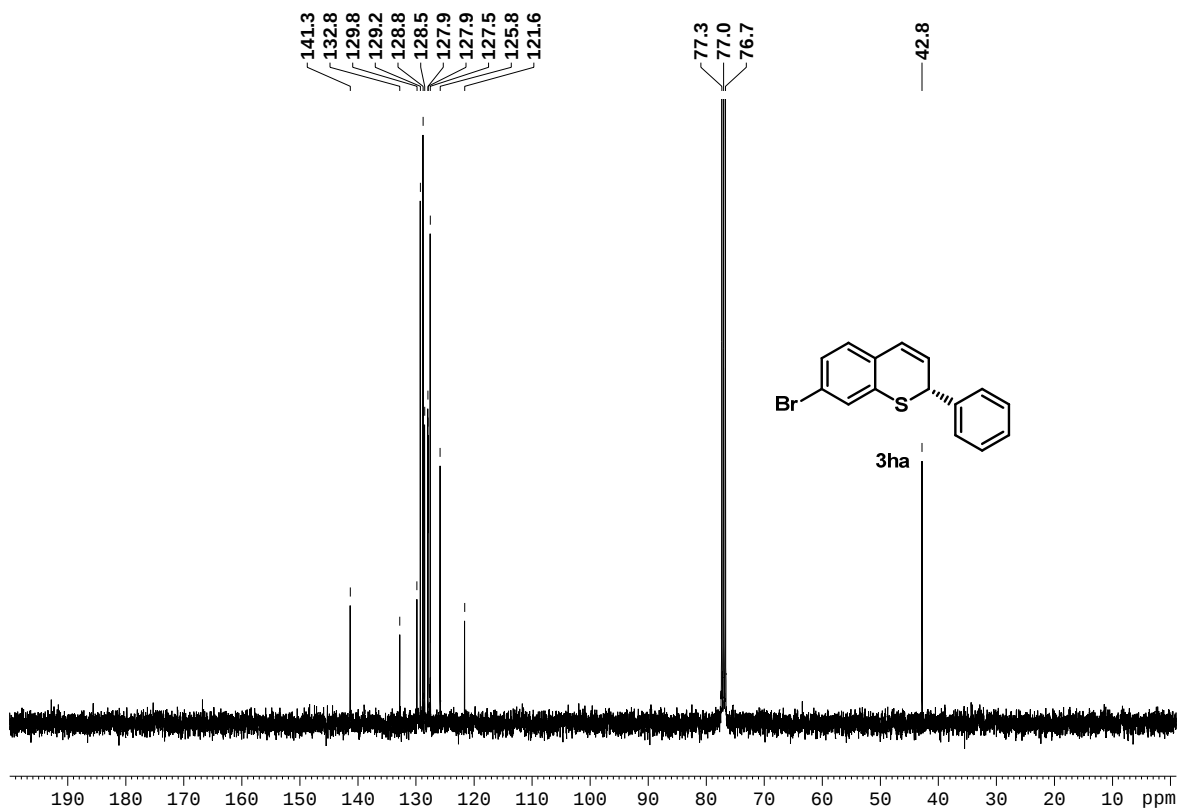
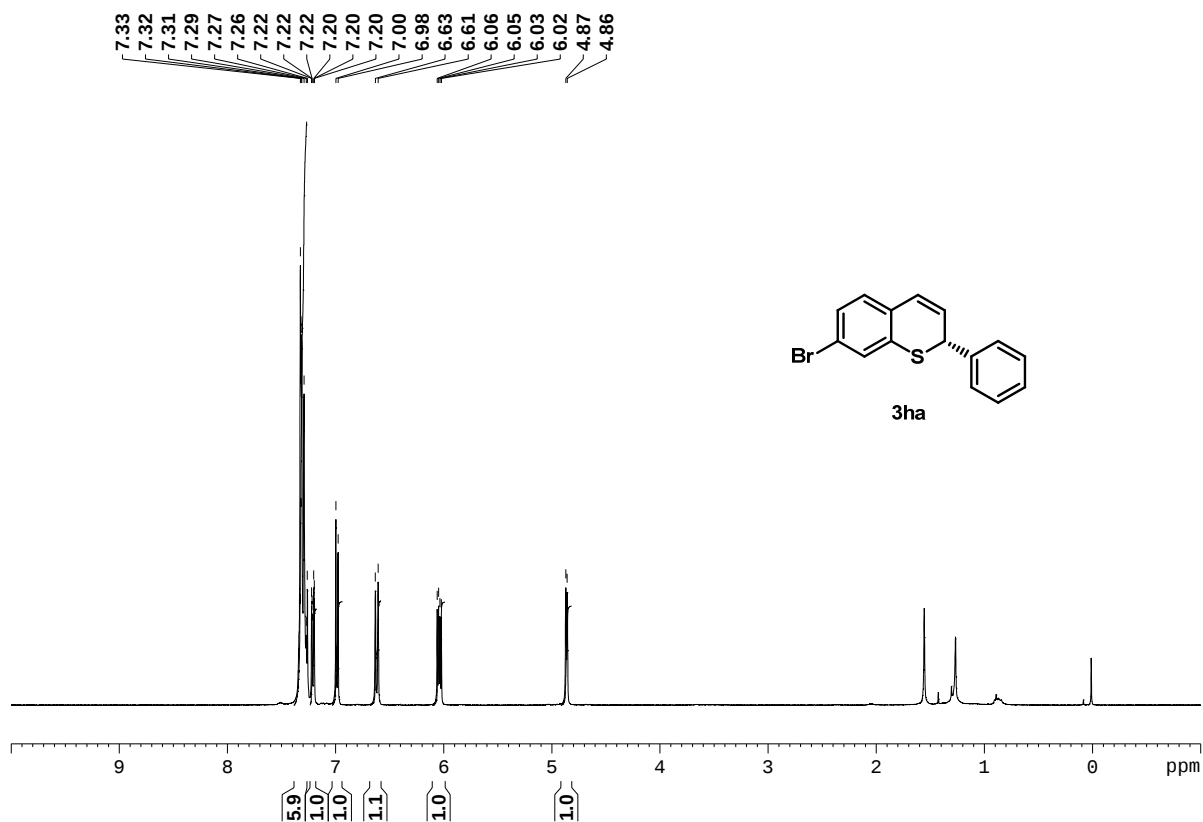
Daicel Chiralpak IC column (80:20 *n*-Hexane/ MTBE, 1.0 mL/min, 20 °C, 254 nm)

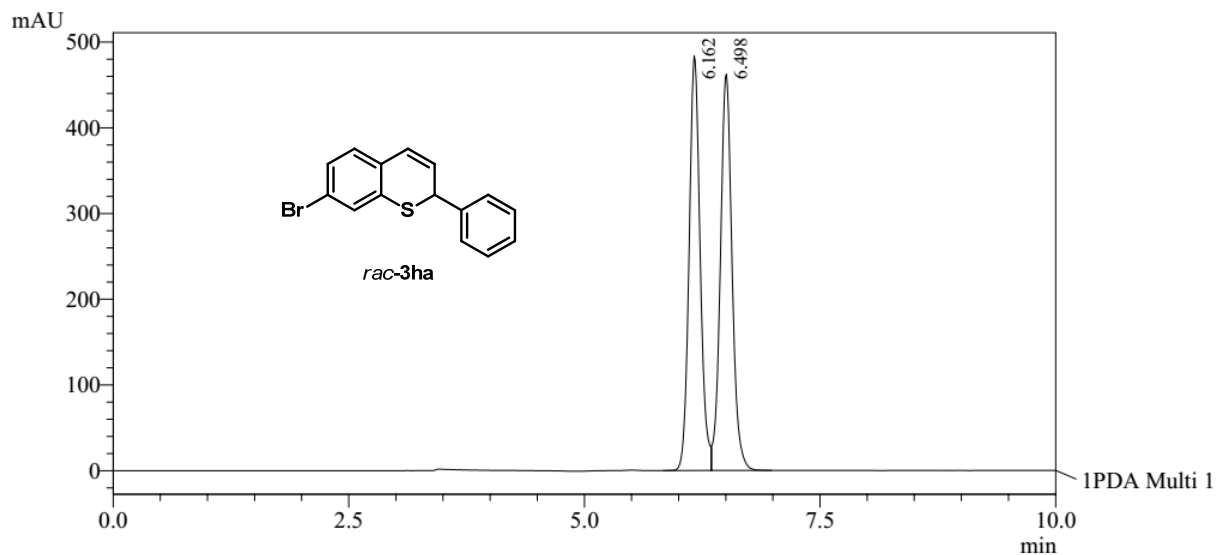


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	4.388	234268	2.707
2	4.689	8419625	97.293
Total		8653893	100.000



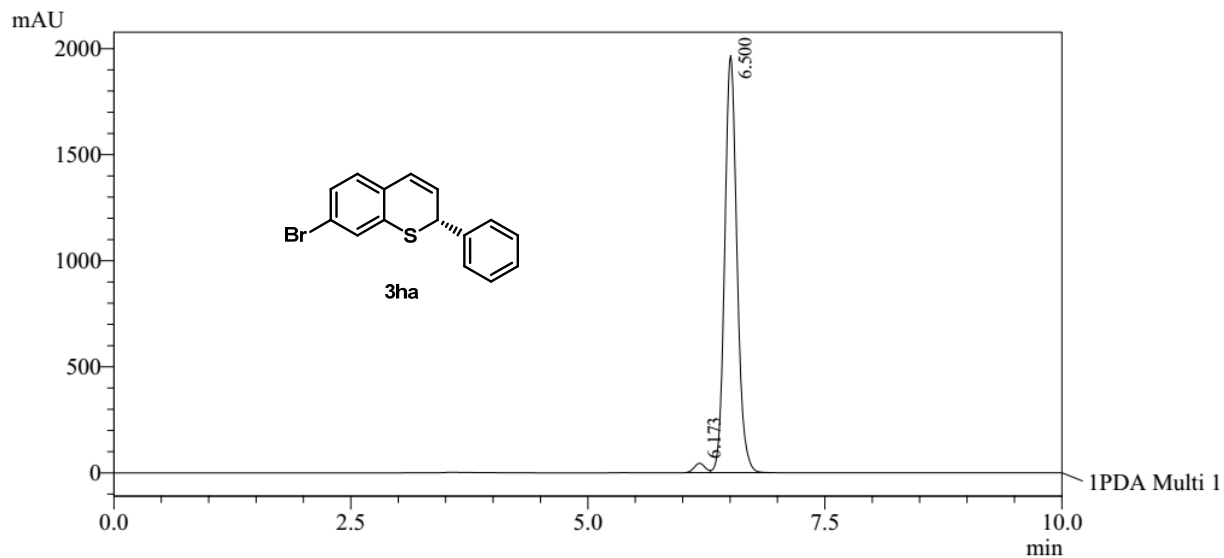


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.162	3950926	49.776
2	6.498	3986467	50.224
Total		7937393	100.000

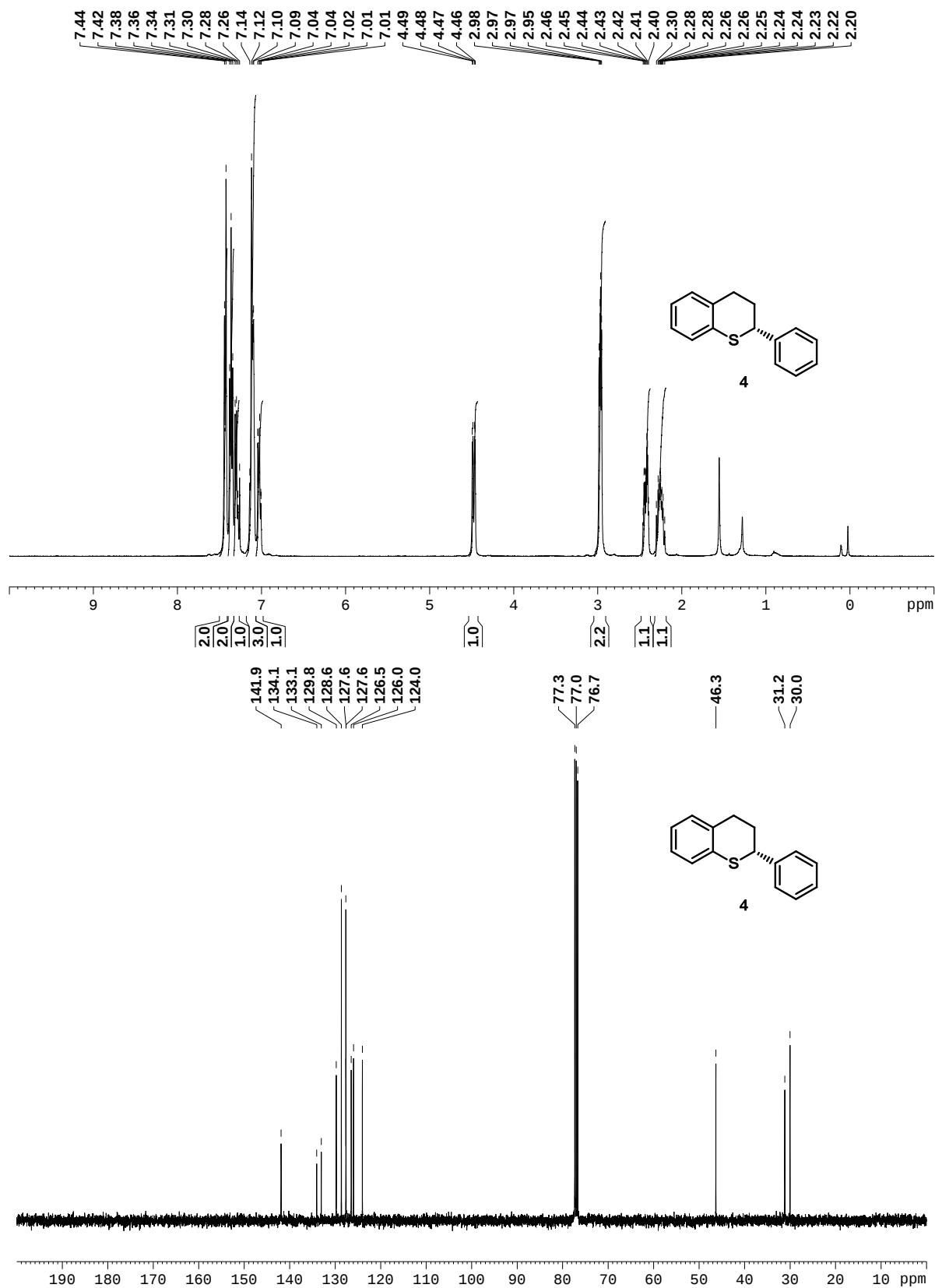
Daicel Chiralpak IB column (99:01 *n*-Hexane/ *i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)

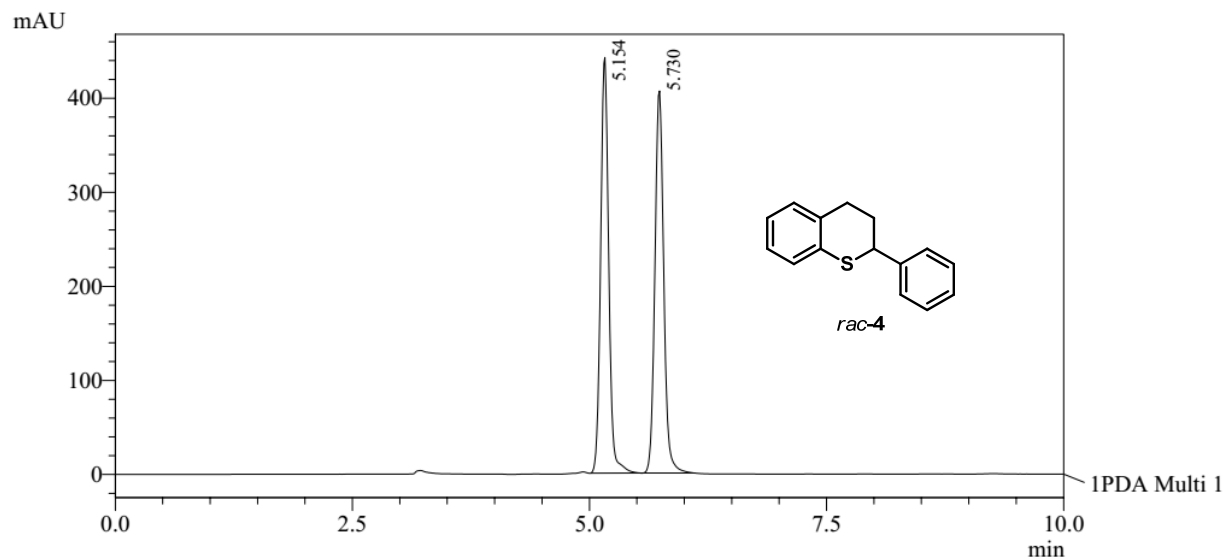


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	6.173	359708	1.992
2	6.500	17696988	98.008
Total		18056695	100.000



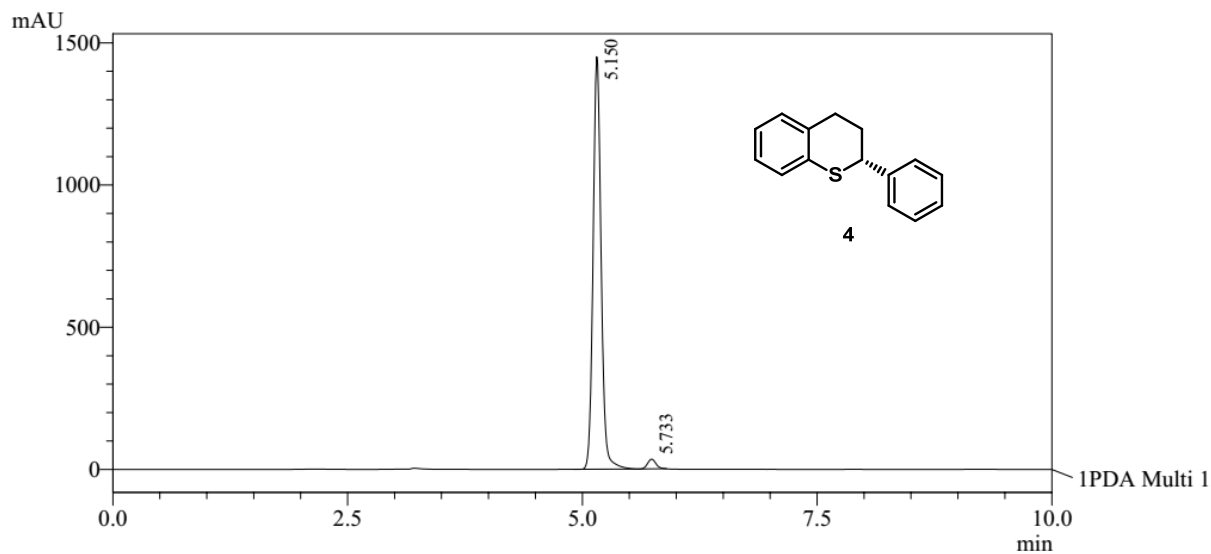


PeakTable

PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.154	2624016	49.947
2	5.730	2629538	50.053
Total		5253554	100.000

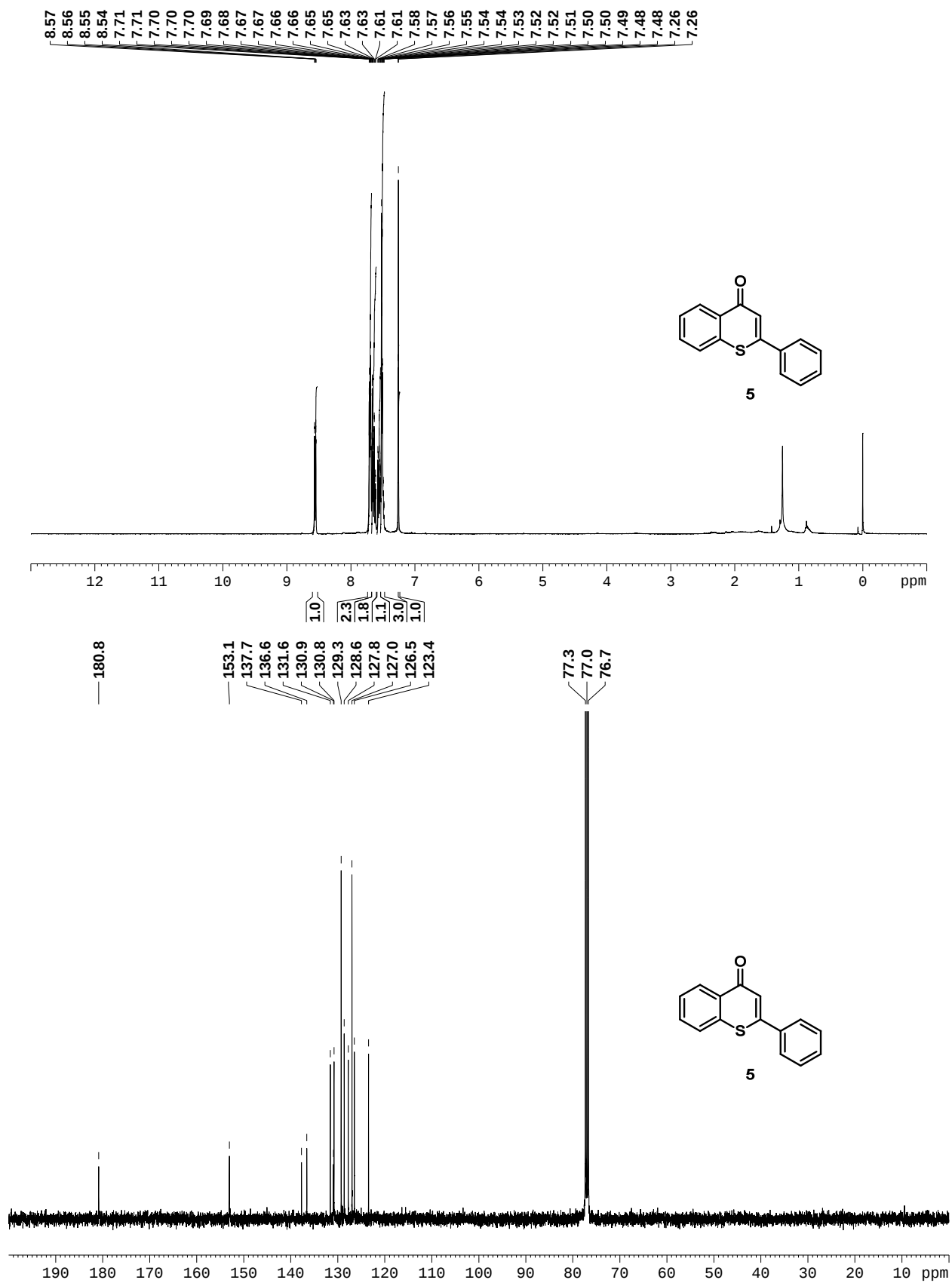
Daicel Chiralpak IB column (97:3 *n*-Hexane/*i*-PrOH, 1.0 mL/min, 20 °C, 254 nm)



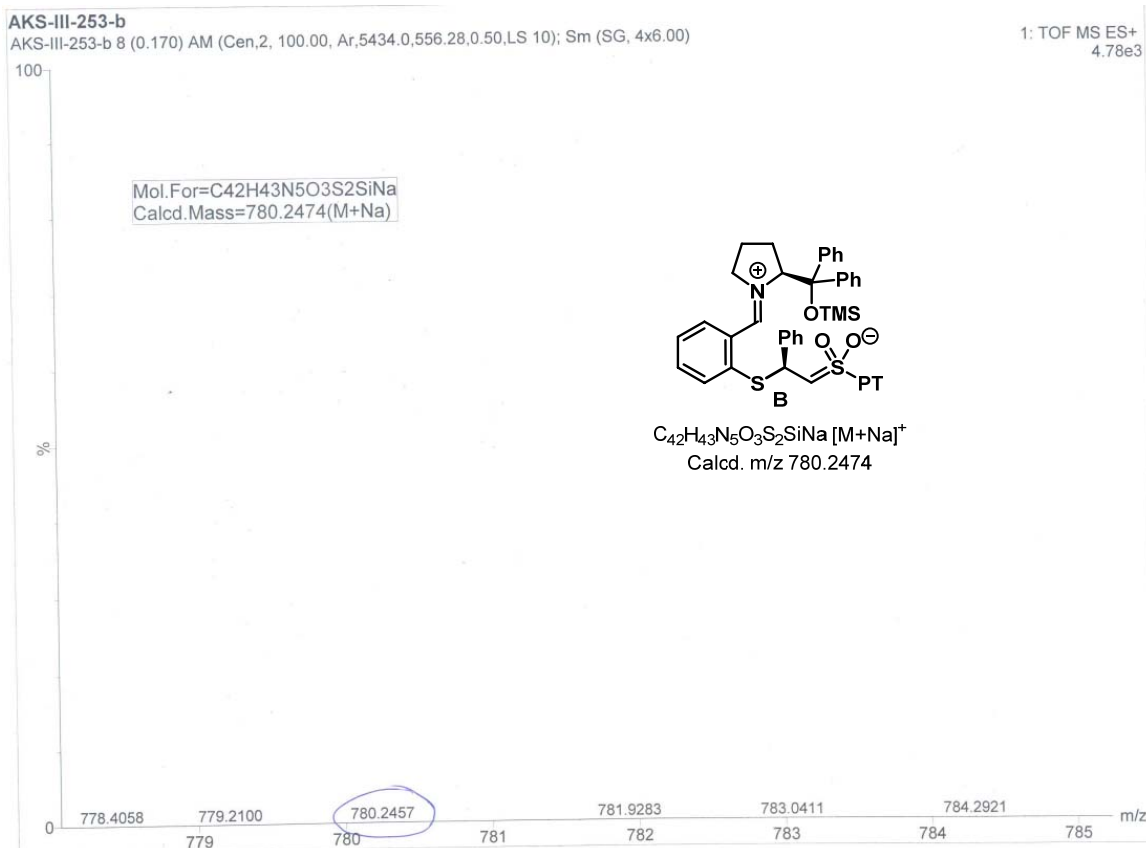
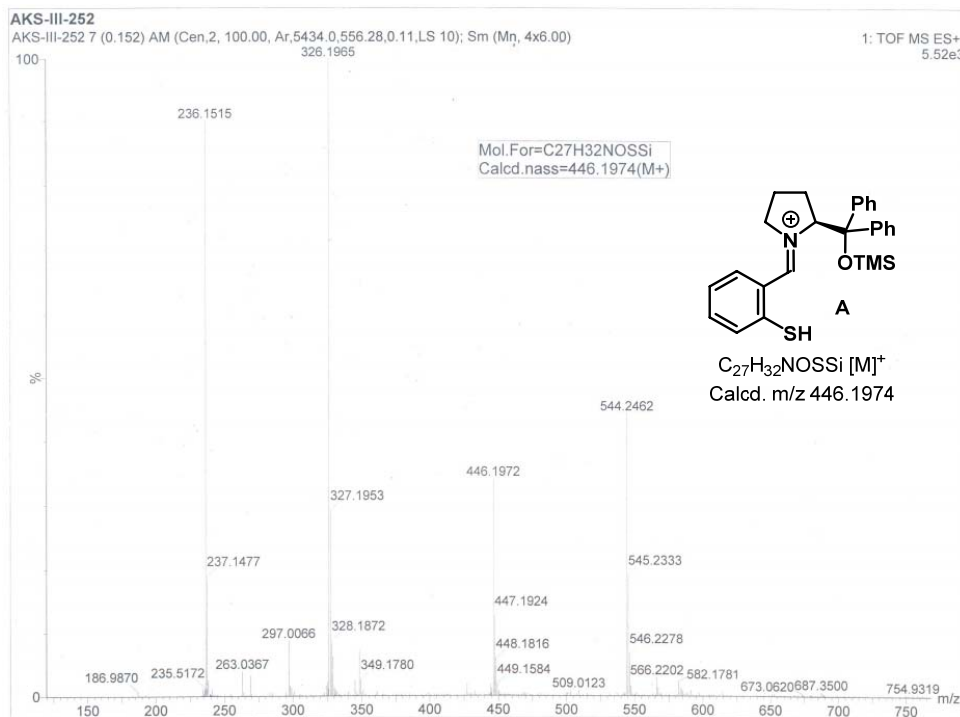
PeakTable

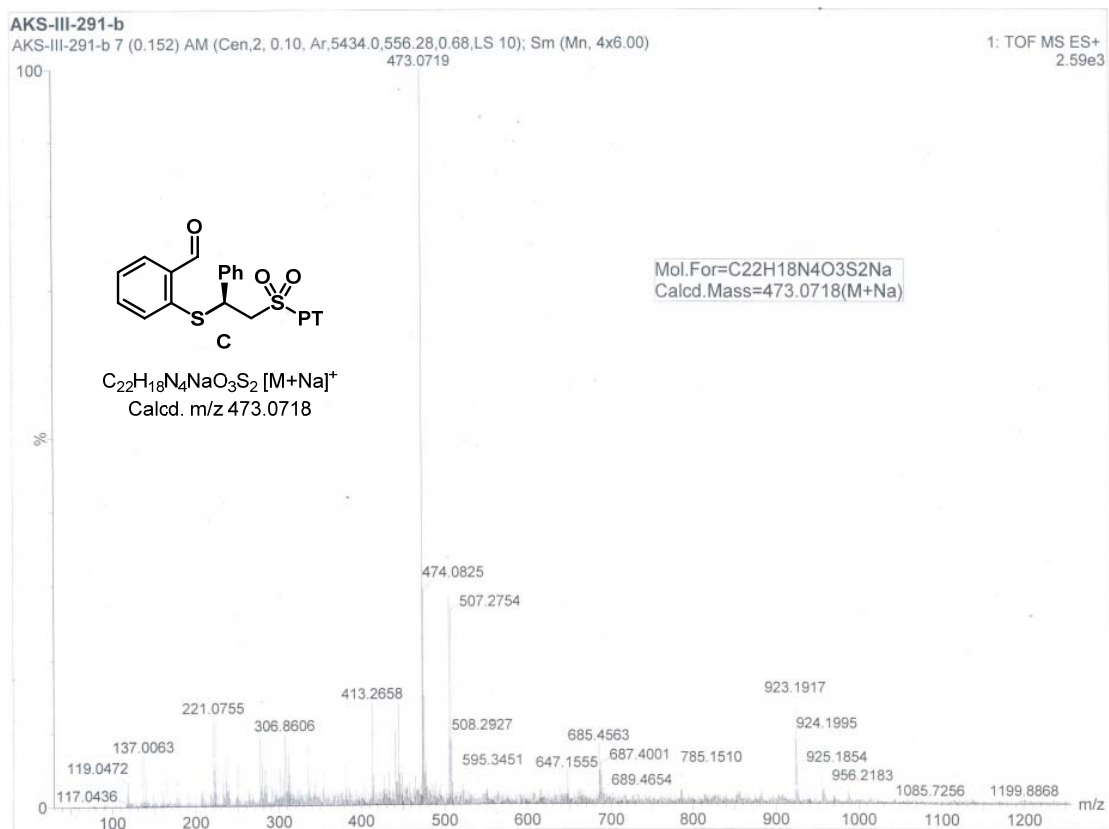
PDA Ch1 254nm 4nm

Peak#	Ret. Time	Area	Area %
1	5.150	8637611	97.626
2	5.733	210007	2.374
Total		8847618	100.000



Detection of reactive intermediates:





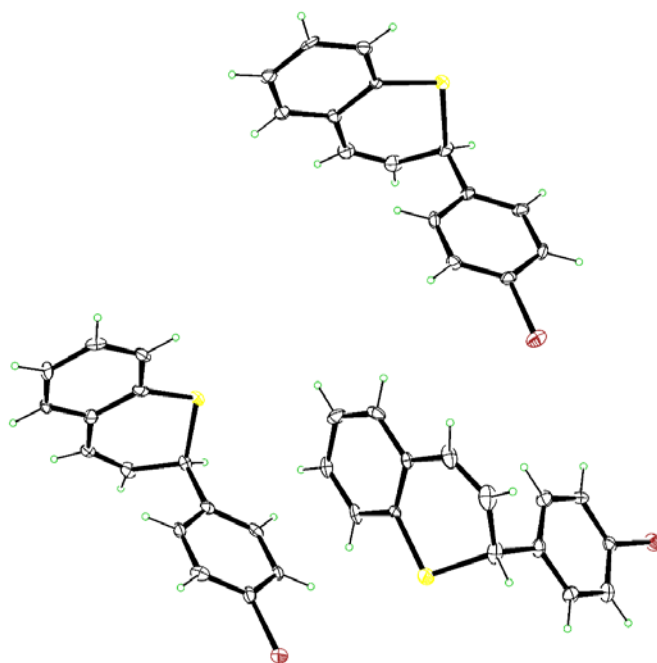
Single crystal X-ray diffraction analysis of 3ai and 3ak:**Single crystal X-ray diffraction analysis of 3ai:**

Single crystal of **3ai** (recrystallized from $\text{CHCl}_3/n\text{-Hexane} = 1:10$ at 25°C) was mounted and the diffraction data were collected at 273 K on a Bruker SMART APEX CCD diffractometer using SMART/SAINT software. Intensity data were collected using graphite-monochromatized Mo-K α radiation (71.073 pm). The structure was solved by direct method using the SHELX-97 and refined by full-matrix least-squares on F^2 . Empirical absorption corrections were applied with SADABS. All Non-hydrogen atoms were refined anisotropically and hydrogen atoms were included in geometric positions. Structure was drawn using Olex-2 and ORTEP-3. The crystallographic refinement parameters are given below:

Table S1. Crystal data and structure refinement for **3ai**

Identification code	3ai	
CCDC	1506906	
Empirical formula	$\text{C}_{15}\text{H}_{11}\text{BrS}$	
Formula weight	303.22	
Temperature	102(3) K	
Wavelength	0.71073 Å	
Crystal system	orthorhombic	
Space group	$P2_12_12_1$	
	$a = 10.9526(7)$	$\alpha = 90^\circ$
	$b = 11.6626(10)$ Å	$\beta = 90^\circ$
	$c = 29.560(2)$ Å	$\gamma = 90^\circ$
Volume	$3775.9(5)$ Å ³	
Z	12	
Density (calculated)	1.600 g/cm ³	
Absorption coefficient	3.404 mm ⁻¹	
F(000)	1824.0	
Crystal size	$0.171 \times 0.124 \times 0.047$ mm ³	
Theta range for data collection	5.102 to 50°	
Index ranges	$-12 \leq h \leq 13$, $-9 \leq k \leq 13$, $-24 \leq l \leq 35$	
Reflections collected	11268	
Independent reflections	6221 [$R_{\text{int}} = 0.0506$, $R\sigma = 0.1039$]	

Data/ restraints/ parameters	6221/ 0/ 460
Goodness-of-fit on F^2	0.998
Final R indexes [$I > 2\sigma(I)$]	$R_1 = 0.0557$, $\omega R_2 = 0.0797$
Final R indexes (all data)	$R_1 = 0.0842$, $\omega R_2 = 0.0892$
Largest diff. peak/hole	0.49/−0.46 e.Å ^{−3}
Flack parameter	0.002(11)



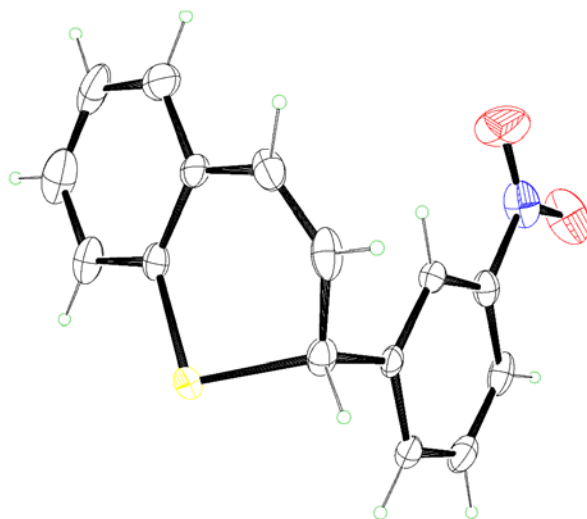
ORTEP representation of the X-ray structure of **3ai** (thermal ellipsoids at 30% probability)

Single crystal X-ray diffraction analysis of **3ak**:

Single crystal of **3ak** (recrystallized from 1:4 CHCl₃/*n*-Hexane at 25 °C) was mounted and the diffraction data was collected at 273 K on a Bruker SMART APEX CCD diffractometer using SMART/SAINT software. Intensity data were collected using graphite-monochromatized Mo-K α radiation (71.073 pm). The structure was solved by direct methods using the SHELX-97 and refined by full-matrix least-squares on F^2 . Empirical absorption corrections were applied with SADABS. All Non-hydrogen atoms were refined anisotropically and hydrogen atoms were included in geometric positions. Structure was drawn using Olex-2 and ORTEP-3. The crystallographic refinement parameters are given below:

Table S2. Crystal data and structure refinement for **3ak**

Identification code	3ak	
CCDC	1506907	
Empirical formula	C ₁₅ H ₁₁ NO ₂ S	
Formula weight	269.32	
Temperature	150(2) K	
Wavelength	0.71073 Å	
Crystal system	orthorhombic	
Space group	P2 ₁ 2 ₁ 2 ₁	
	a = 6.9573(7) Å	$\alpha = 90^\circ$
	b = 13.1585(13) Å	$\beta = 90^\circ$
	c = 13.8579(14) Å	$\gamma = 90^\circ$
Volume	1268.7(2) Å ³	
Z	4	
Density (calculated)	1.410 g/cm ³	
Absorption coefficient	0.251 mm ⁻¹	
F(000)	560.0	
Crystal size	0.26 × 0.25 × 0.24 mm ³	
Theta range for data collection	4.268 to 49.954 °	
Index ranges	-8 ≤ h ≤ 8, -15 ≤ k ≤ 15, -16 ≤ l ≤ 16	
Reflections collected	21761	
Independent reflections	21761 [<i>R</i> _{int} = 0.0649, <i>R</i> σ = 0.0757]	
Data/ restraints/ parameters	21761/ 0/ 173	
Goodness-of-fit on F ²	1.036	
Final R indexes [<i>I</i> > 2σ (<i>I</i>)]	<i>R</i> ₁ = 0.0488, ω <i>R</i> ₂ = 0.1274	
Final R indexes (all data)	<i>R</i> ₁ = 0.0554, ω <i>R</i> ₂ = 0.1326	
Largest diff. peak/hole	0.67/-0.35 e.Å ⁻³	
Flack parameter	0.09(3)	



ORTEP representation of the X-ray structure of **3ak** (thermal ellipsoids at 30% probability)