

Highly Enantioselective α -alkylation of 2-Oxindoles with (3-Indolyl)methanols by Cooperative Catalysis of Lewis Acid and Organocatalyst

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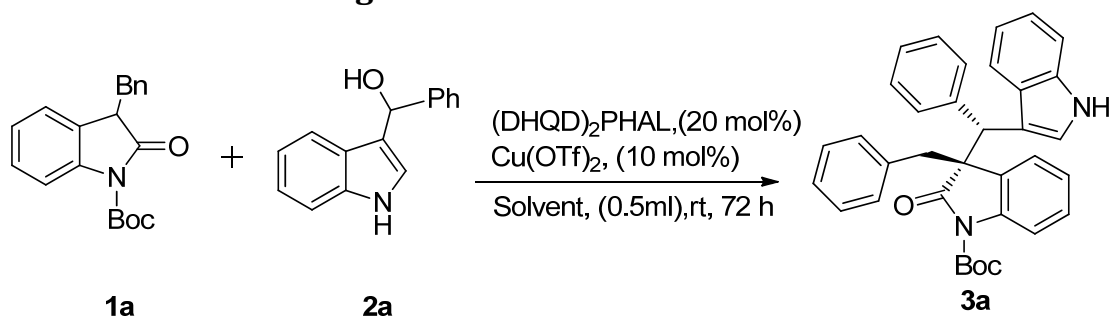
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1. Experimental section	S2
1.1 Solvent screening	S2
1.2 Effect of N-Substituent of oxindoles	S2
1.3 Screening catalyst	S3
2. Determination of absolute configuration of syn-3f.....	S3
2.1 Experimental.....	S4
2.2 Theoretical Calculation.....	S4
2.3 1D NOE of 3f.....	S6
3. NMR Spectrum and HPLC Chromatograms	S6

1. Experimental section

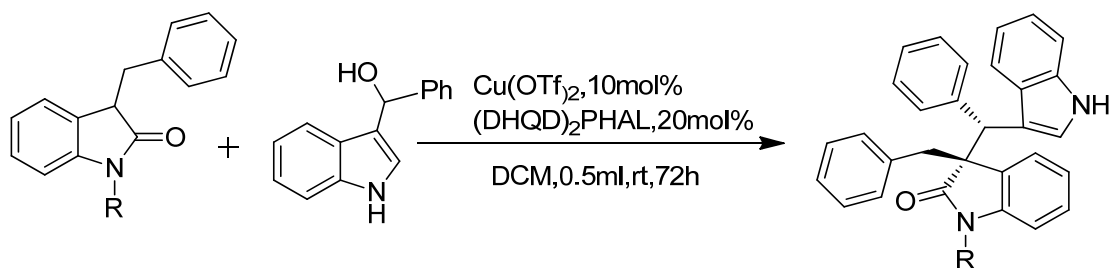
1.1 Solvent screening



Entry	Solvent	dr ^b	Yield(%) ^c	Ee ^d (%)
Syn-3a				
1	EtOAc	N.D.	57	62
2	DMSO	1:1	70	4
3	Toluene	2:1	72	57
4	CH_3NO_2	5:1	70	85
5	CHCl_3	3:1	80	82
6	1,1,2-trichloroethene	2:1	55	73
7	1,1,2,2-tetrachloroethane	3:1	80	75
8	1,1,2-trichloroethane	2:1	65	71
9	CH_2Cl_2	3:1	74	80

^a Reaction conditions: **1a** (0.2 mmol), **2a** (0.1 mmol). ^b Determined by ^1H NMR. ^c Isolated yield of the combined products. ^d Major diastereomer's ee determined by chiral HPLC.

1.2 Effect of N-Substituent of oxindoles



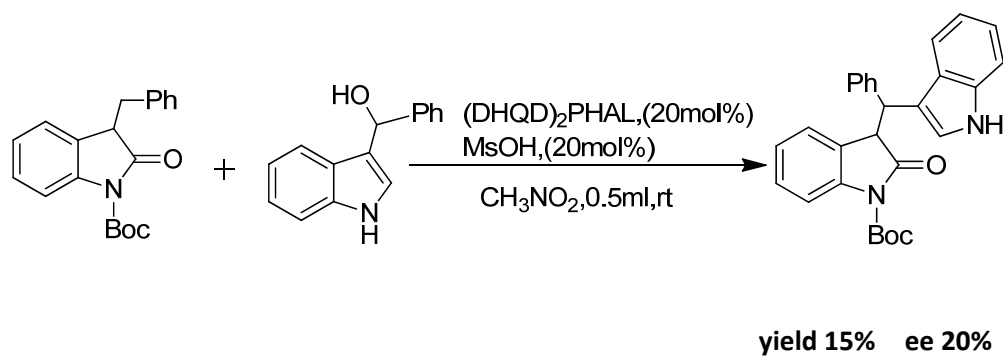
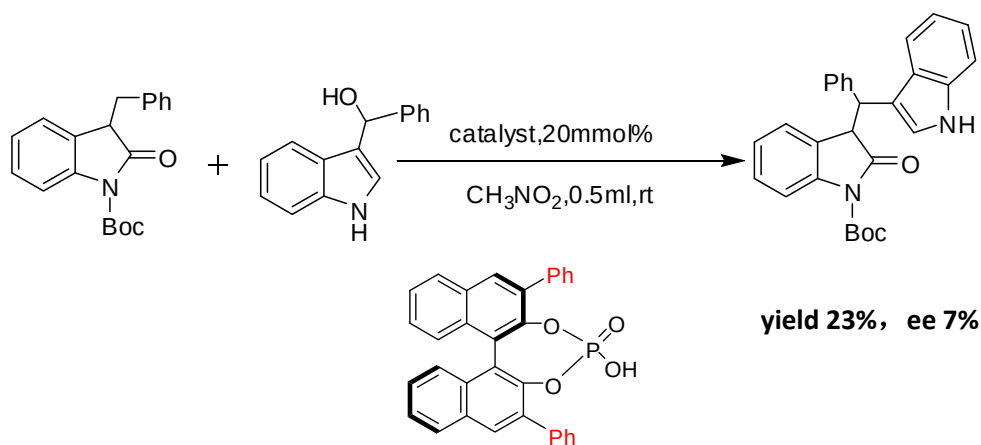
R=Boc, yield 74%, dr 3:1, ee 80%

R=Ac, yield 62%, dr 3:1, ee rac

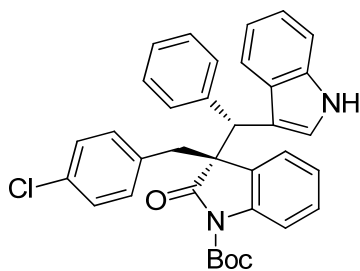
R=H, trace

R=Bn, trace

1.3 Screening catalyst



2. Determination of absolute configuration of syn-3f



2.1 Experimental

About 11 mg of sample was dissolved in 150 μ l CDCl_3 and placed in a BaF_2 cell with a path-length of 75 μ m. IR and VCD spectra were recorded on a Chiral IR-2X FT-VCD spectrometer (BioTools, Inc.) equipped with single photoelastic modulation, with 4 cm^{-1} resolution, 9h collection for both the chiral sample.

2.2 Theoretical Calculation

Since there are two chiral centers in 3g , we investigated the geometry of all the possible conformations, i.e., RR, RS, SR and SS. The IR and VCD spectra were calculated for all of them. We find that the calculated IR and VCD spectra of the RR configuration agree best with the experiment, as shown in Fig.2. Therefore the absolute configuration of 3g is assigned as (RR).

Geometry optimization, frequency, and IR and VCD intensity calculations were carried out at the B3LYP/6-31G(d,p) level of theory. Polarized continuum model (PCM) was adopted to consider solvent (chloroform) effects in the calculation of IR and VCD spectra. The IR and VCD intensities were converted to Lorentzian bands with 8 cm^{-1} half-width for comparison to experiment. All calculations were performed with the Gaussian 09 program suite.

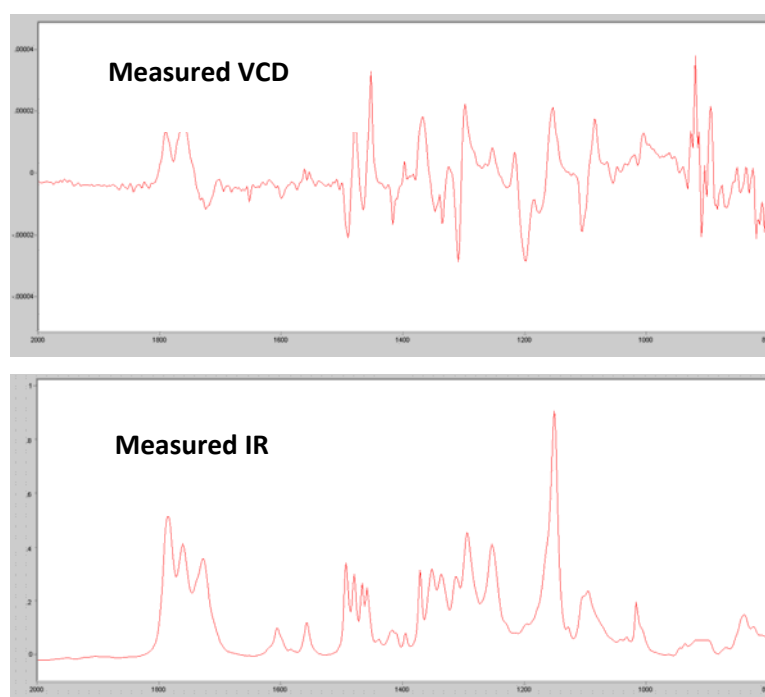


Fig.1 VCD and IR spectra of 3g in CDCl_3 (11mg/0.15ml)

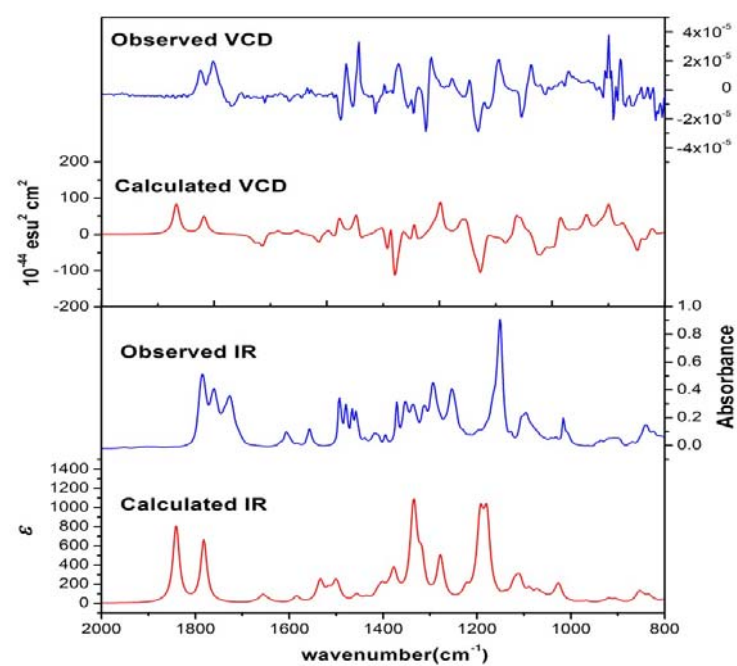
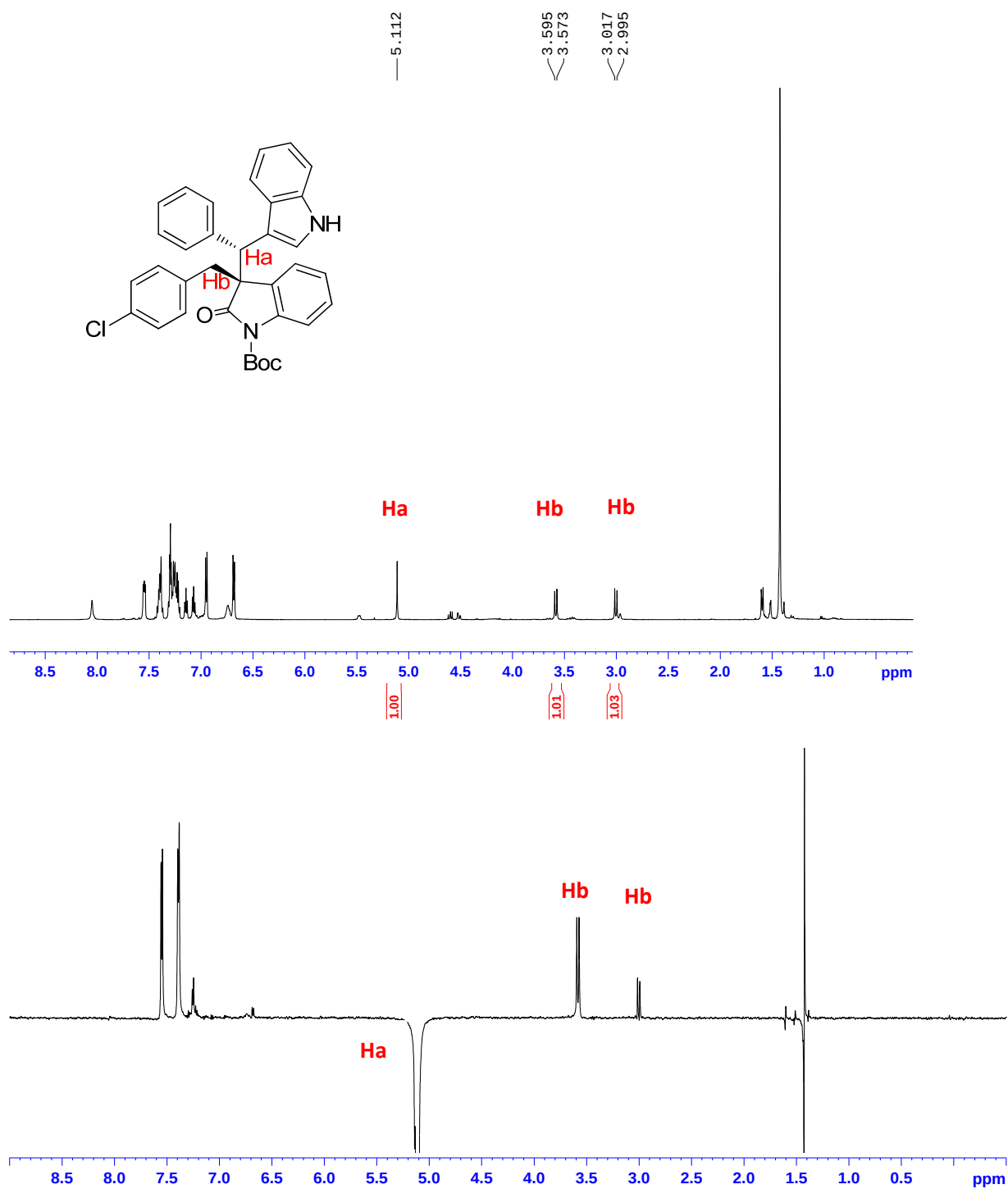
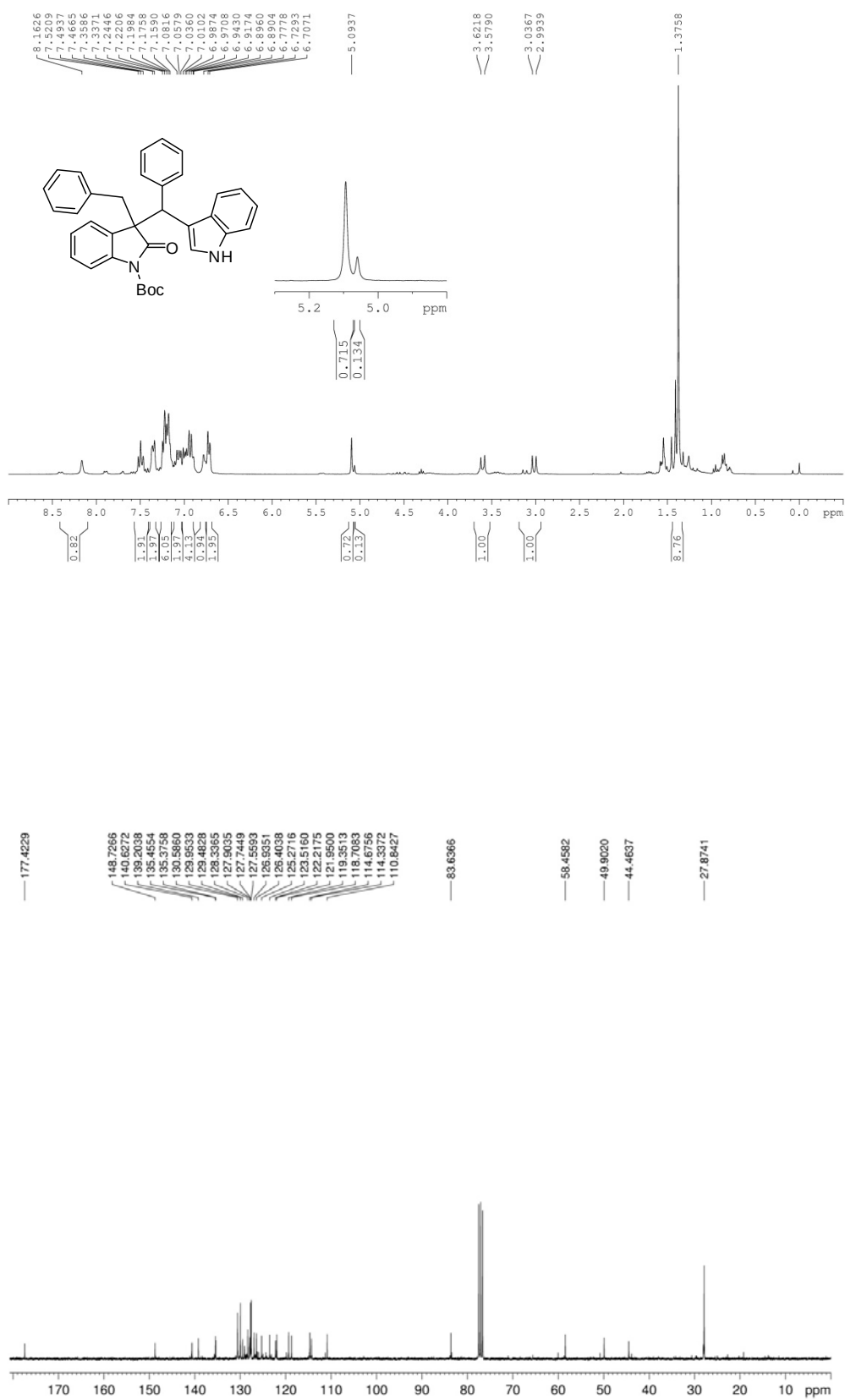


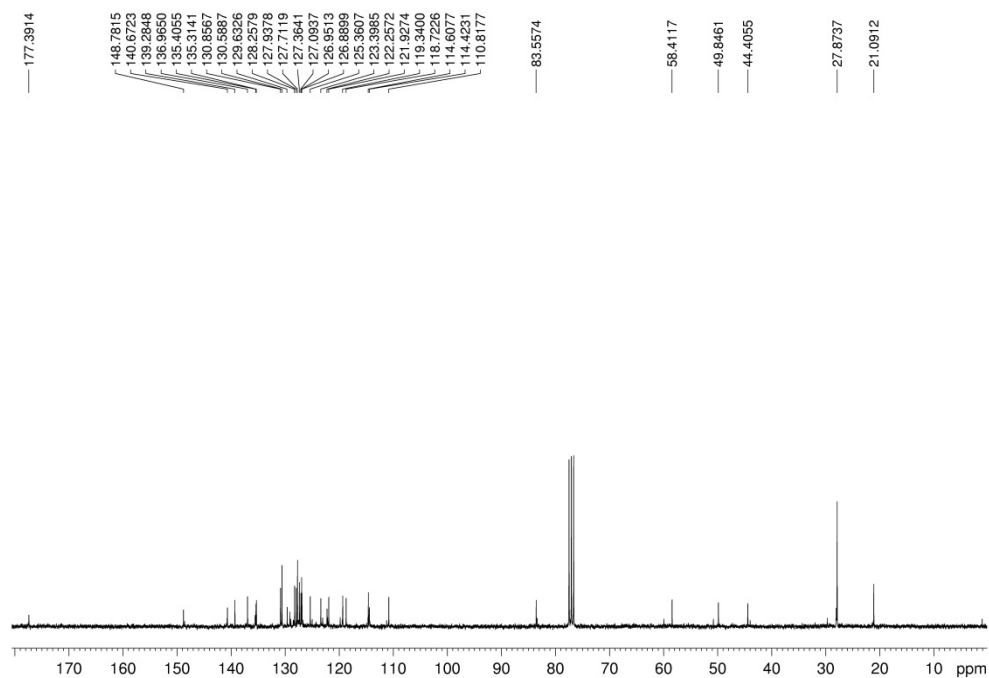
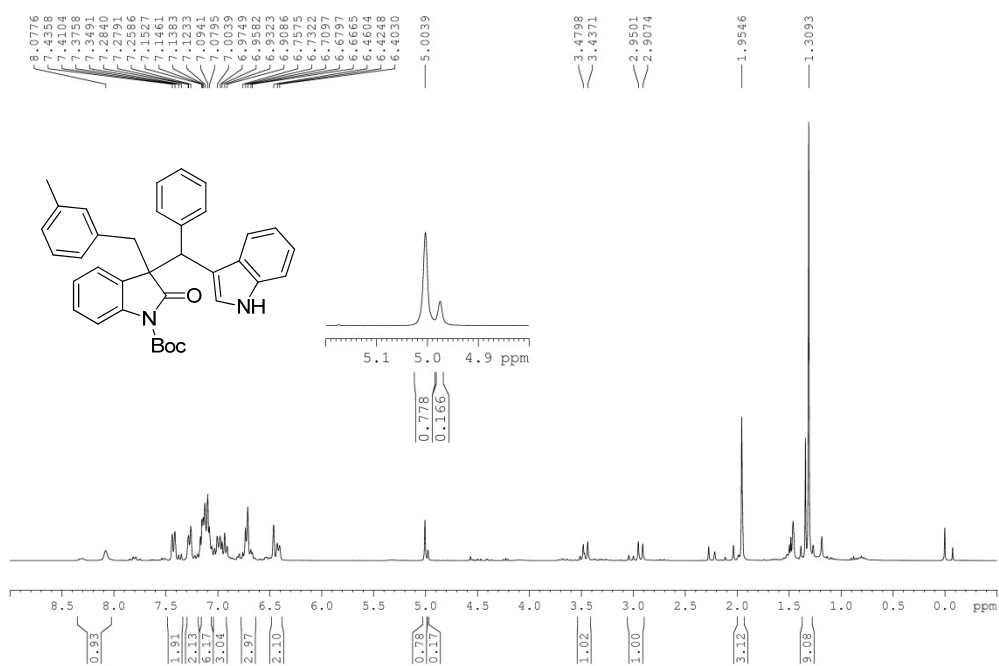
Fig.2 VCD and IR spectra observed for syn-3f compared with calculated spectra

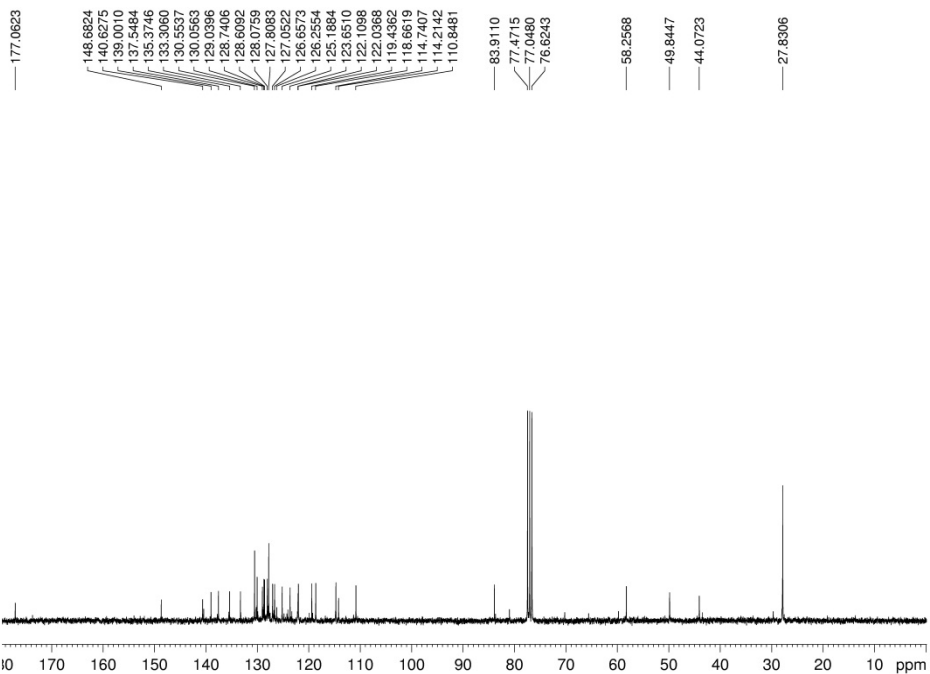
2.3 1D NOE of 3f

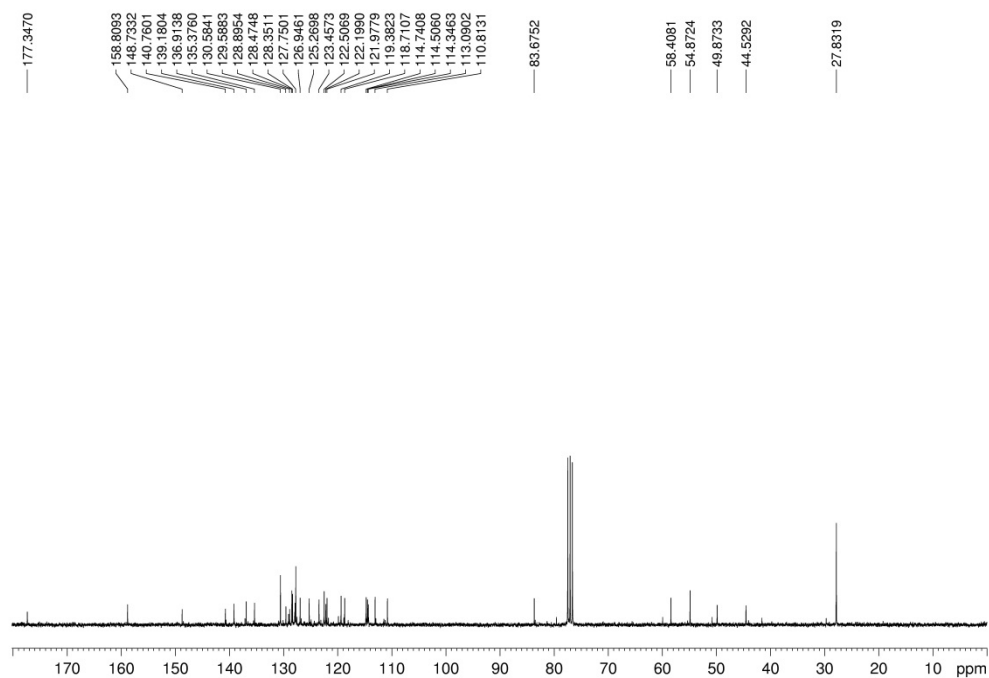
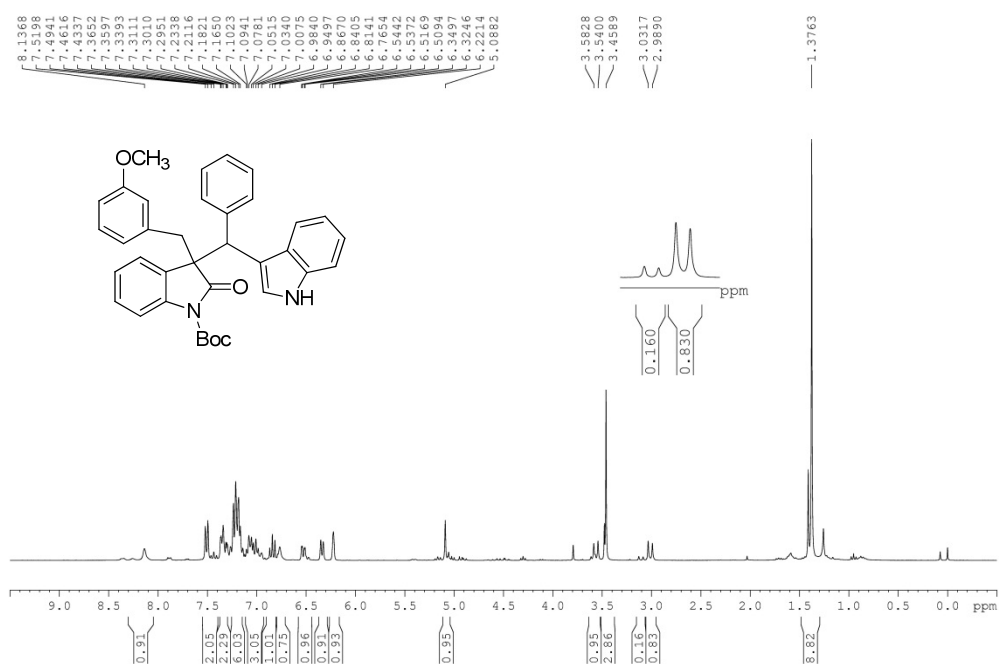


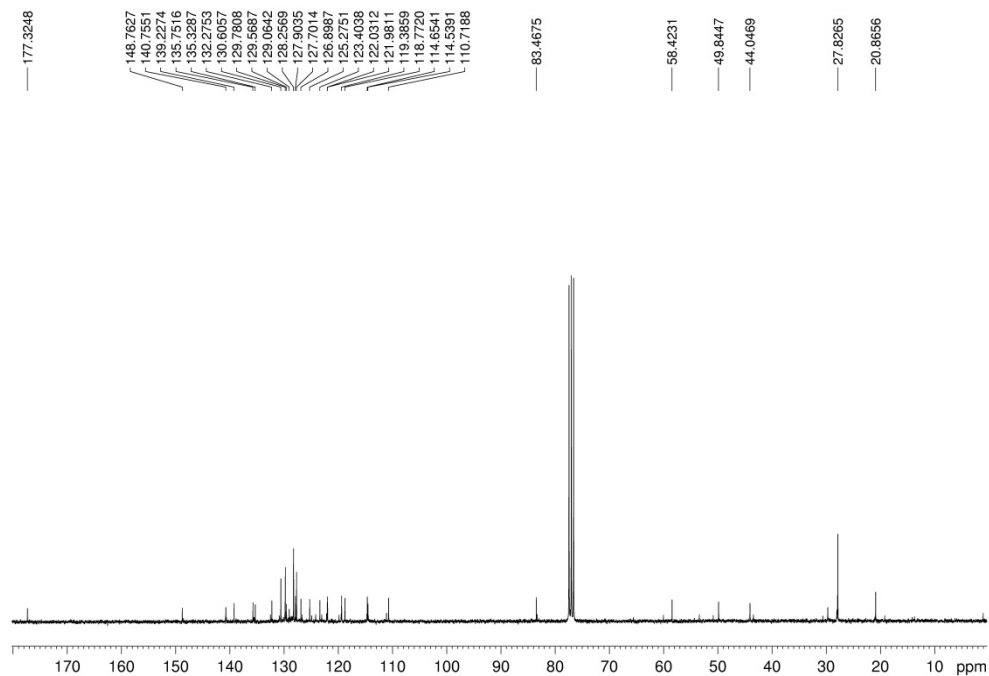
3. NMR Spectrum and HPLC Chromatograms

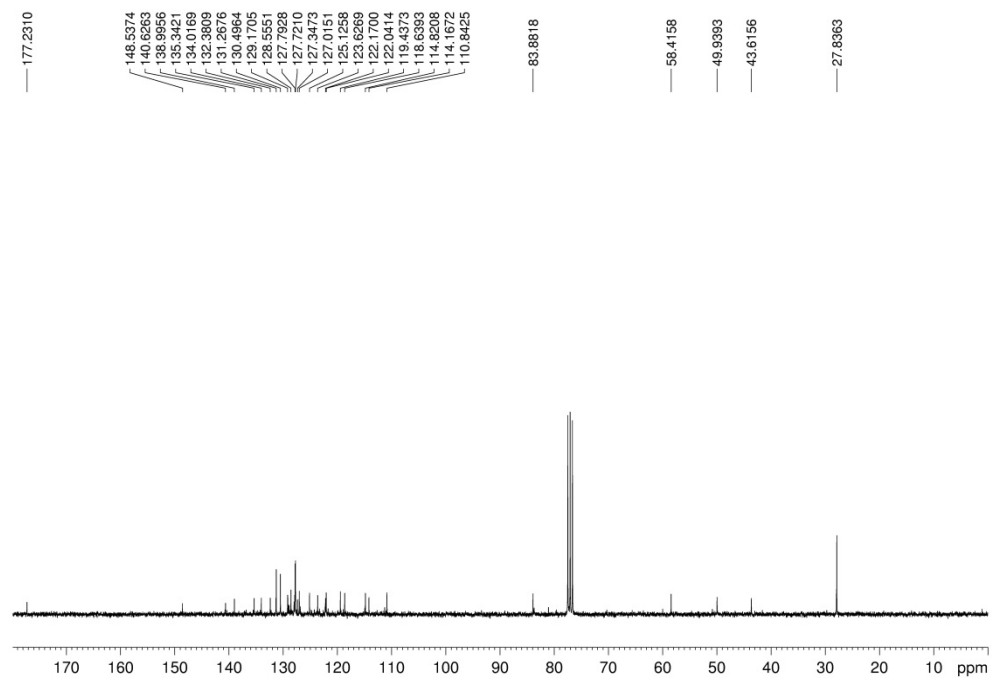


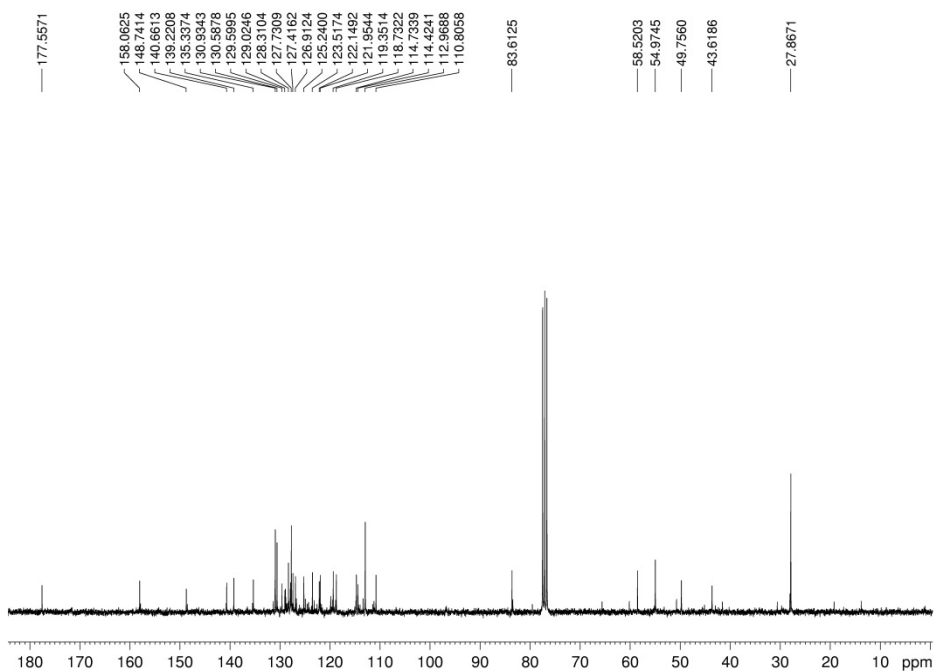
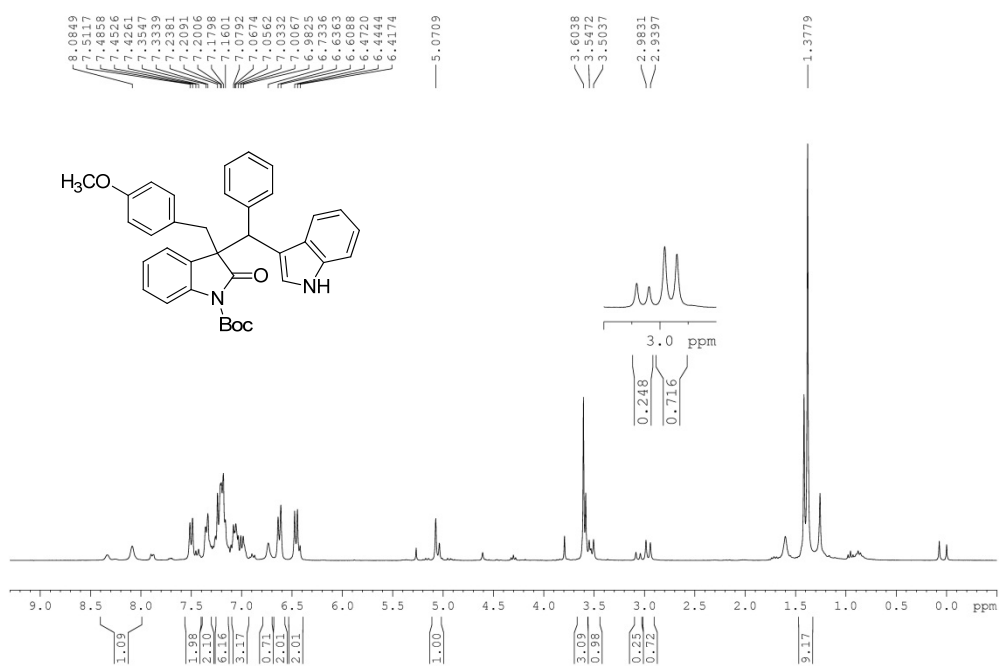


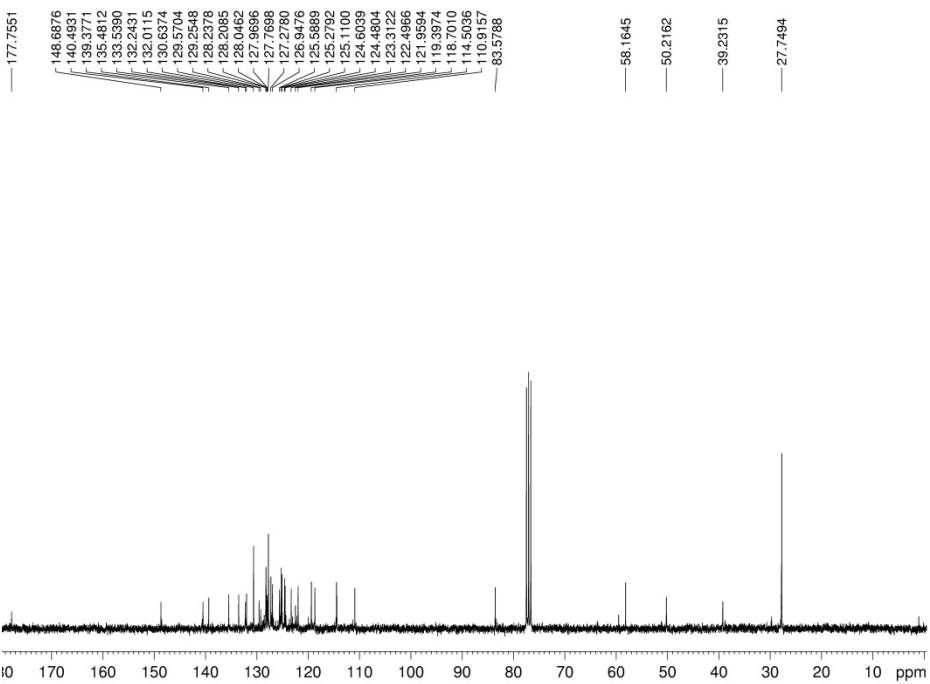


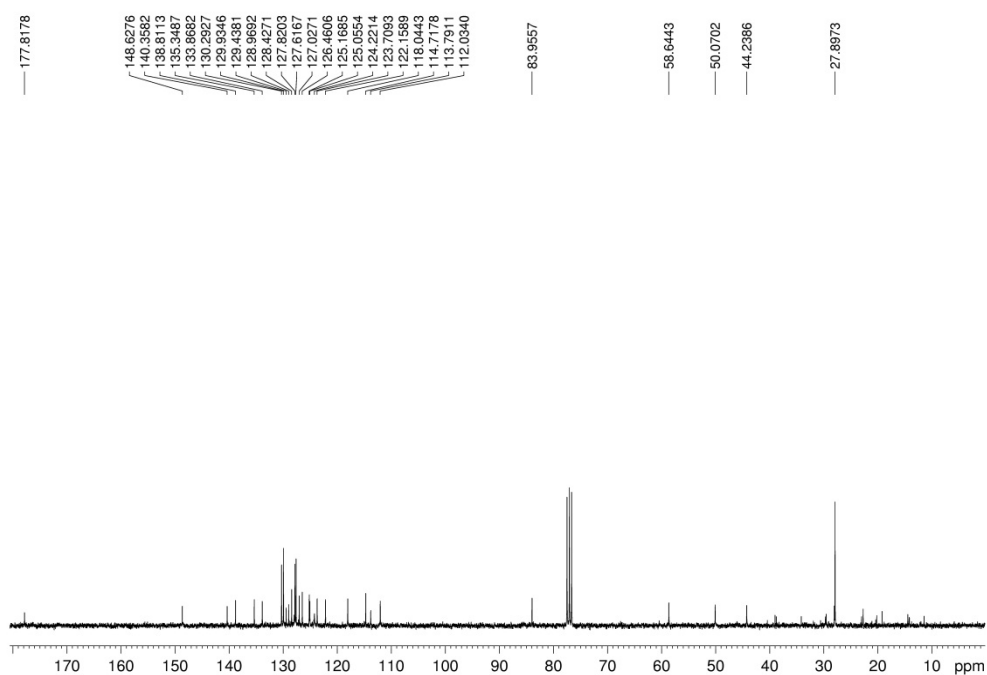
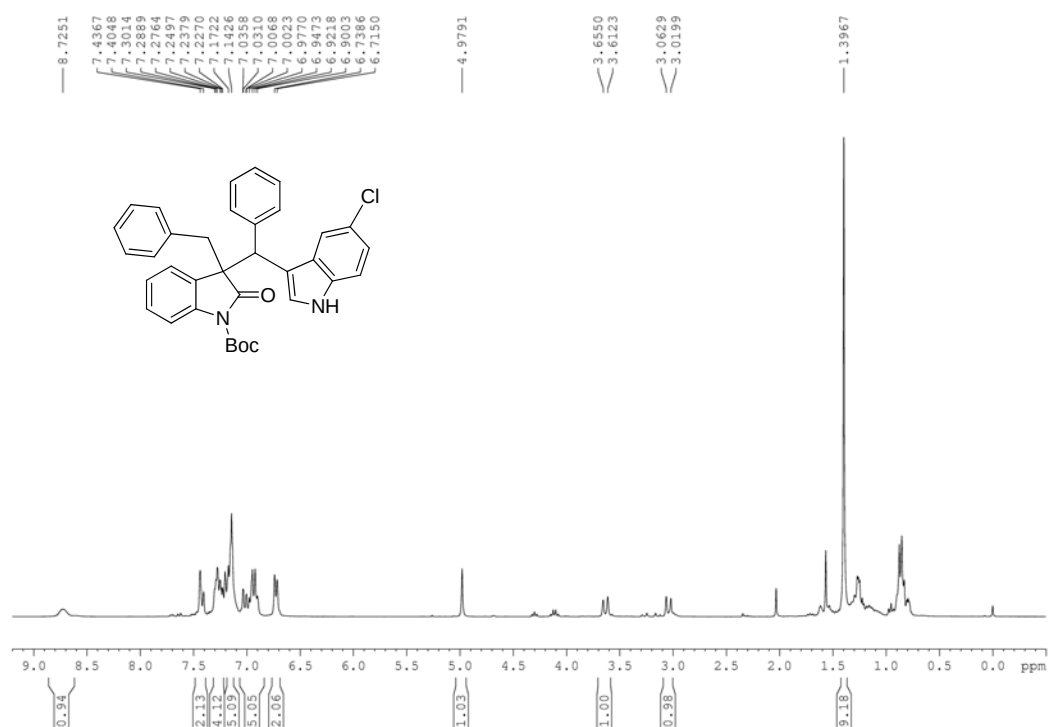


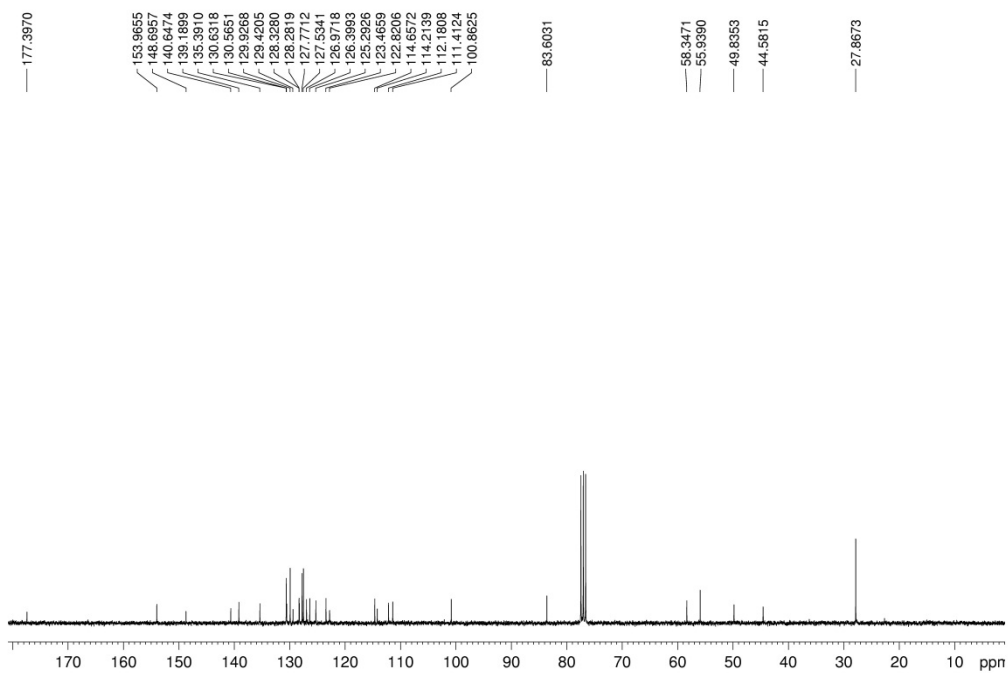
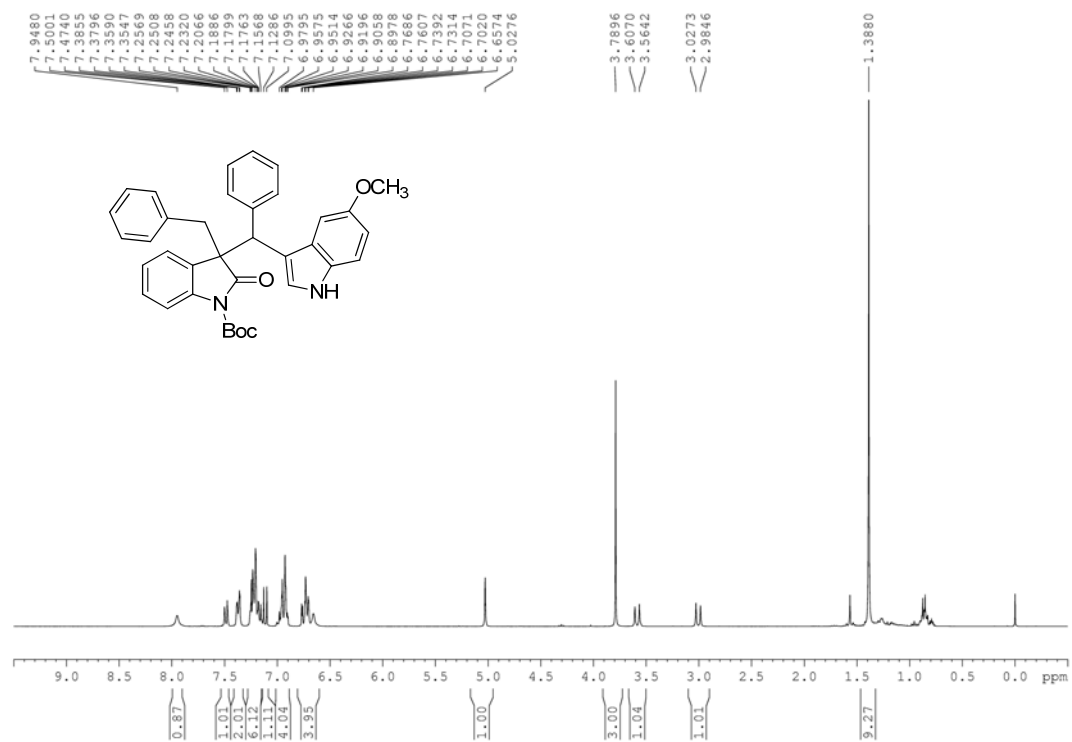


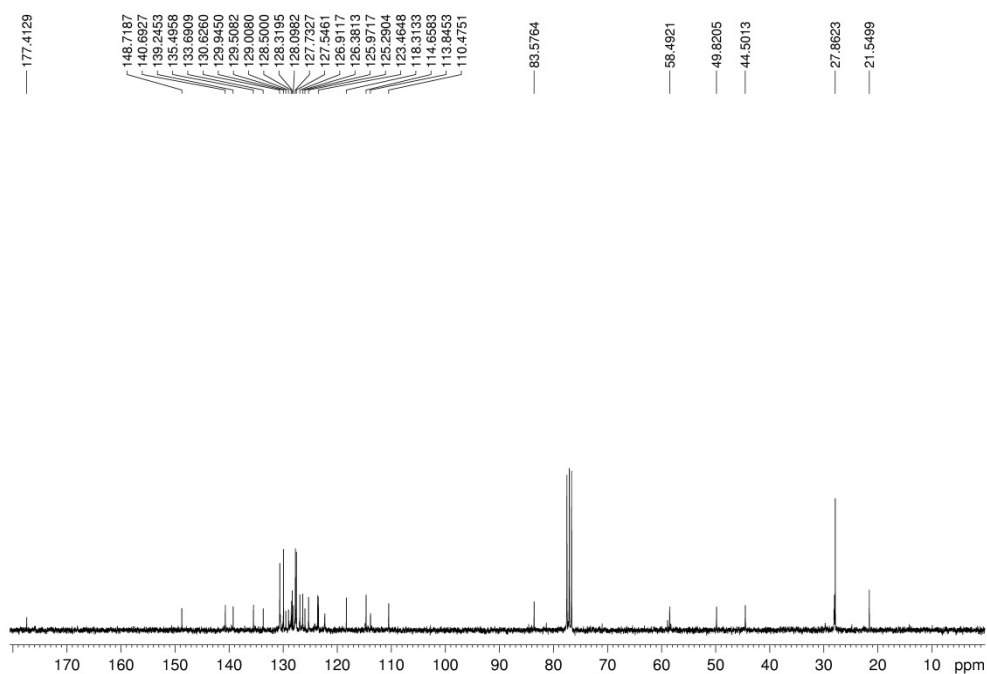
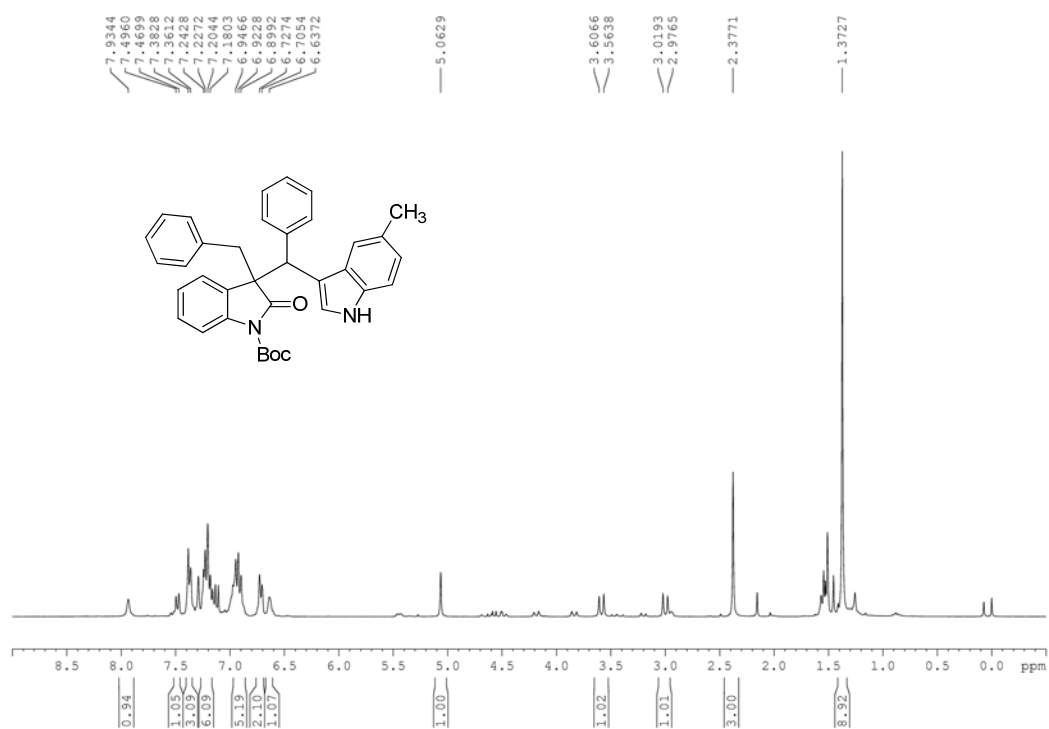


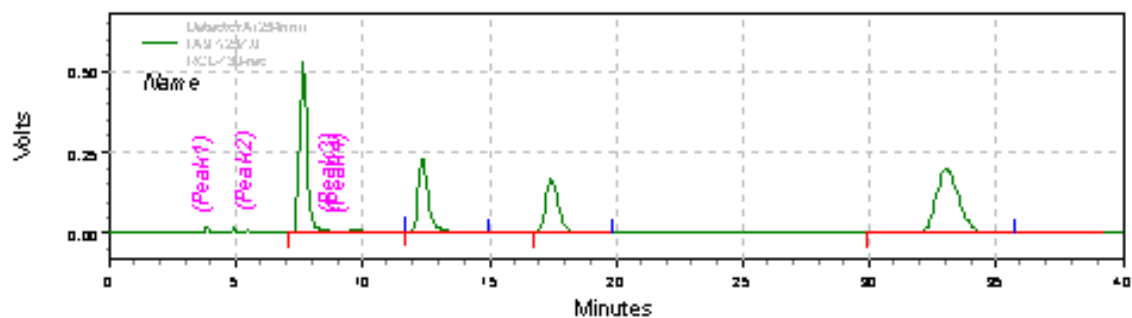
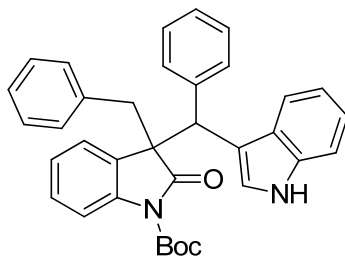






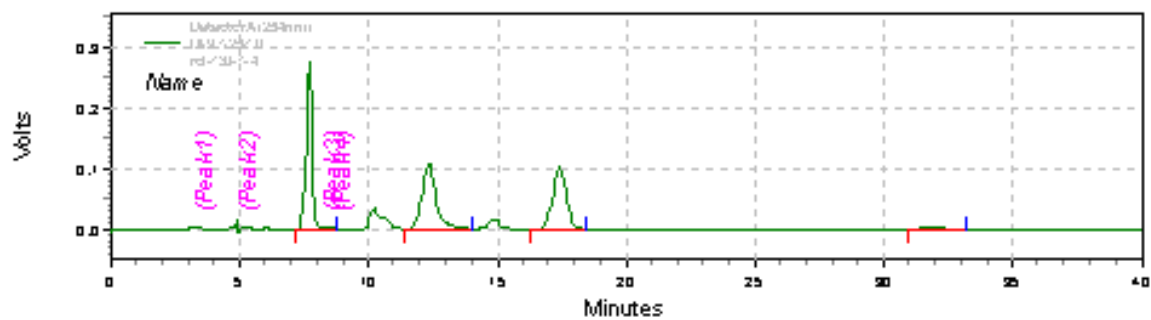






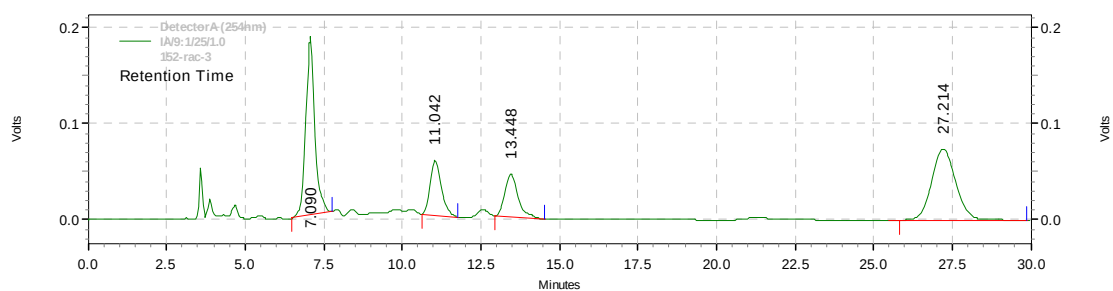
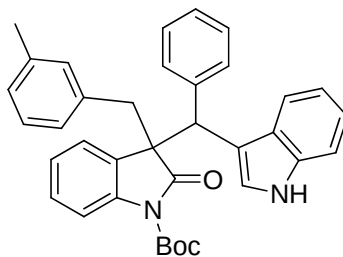
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.662	11820771	32.158
2	12.374	6494285	17.667
3	17.463	6043375	16.441
4	33.048	12400196	33.734



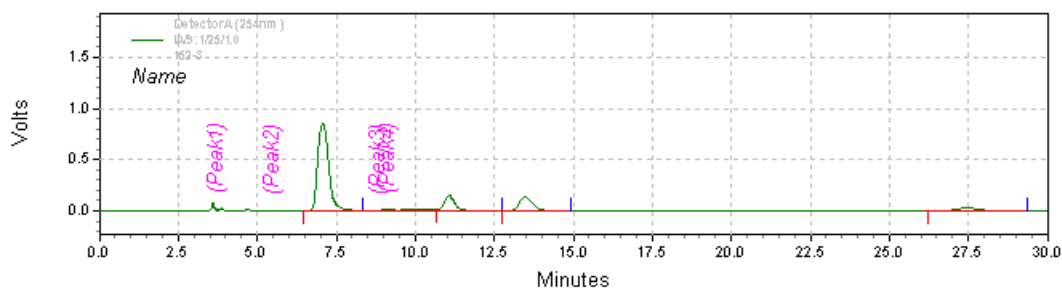
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.719	4532143	36.189
2	12.368	4040827	32.266
3	17.434	3751099	29.953
4	31.902	199401	1.592



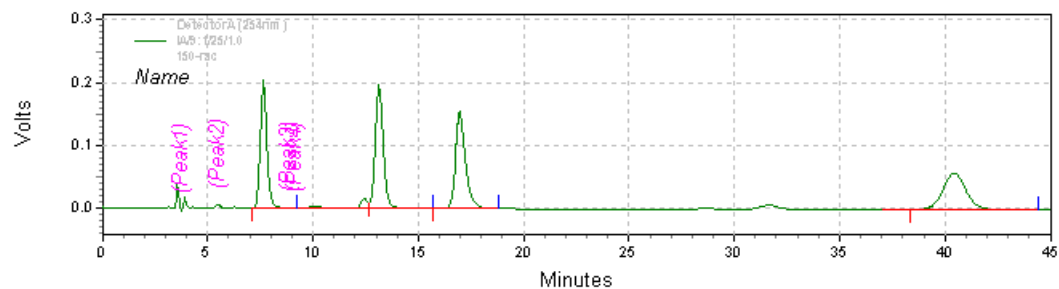
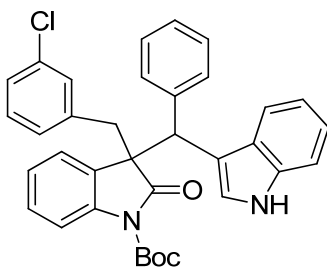
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.090	3854419	37.163
2	11.042	1383426	13.338
3	13.448	1288795	12.426
4	27.214	3845100	37.073



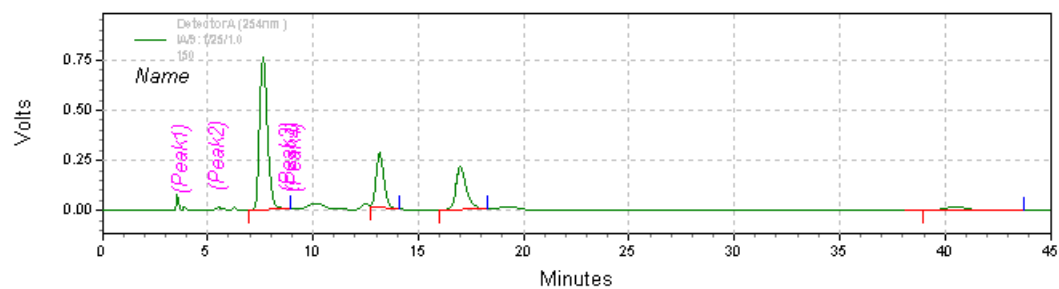
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.059	20825203	70.552
2	11.054	3754924	12.721
3	13.481	3907798	13.239
4	27.480	1029557	3.488



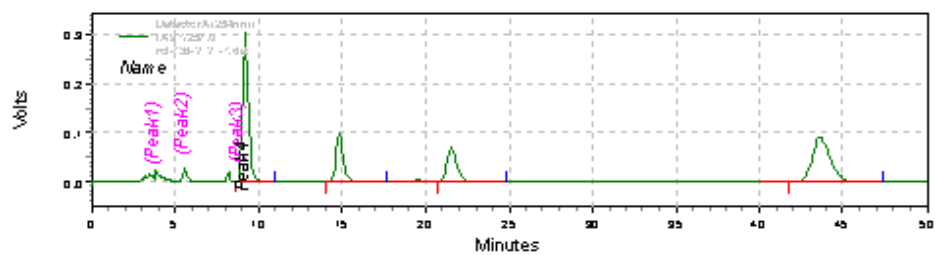
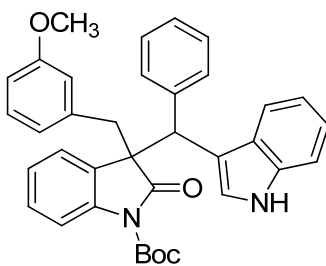
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.670	4392566	22.718
2	13.146	5173683	26.758
3	16.984	5466442	28.272
4	40.474	4302224	22.251



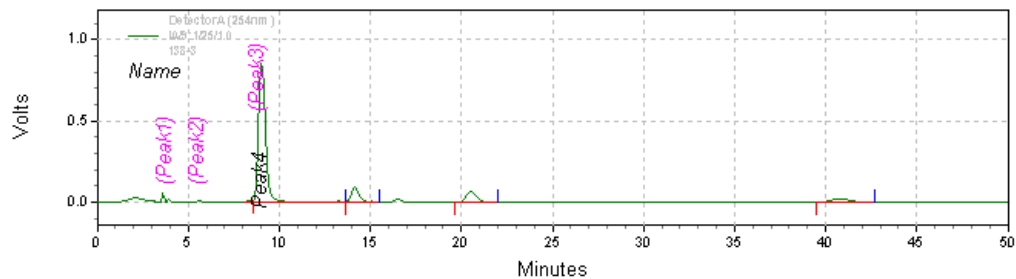
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.650	19783269	55.028
2	13.179	6853713	19.064
3	17.014	8011486	22.284
4	40.549	1302736	3.624



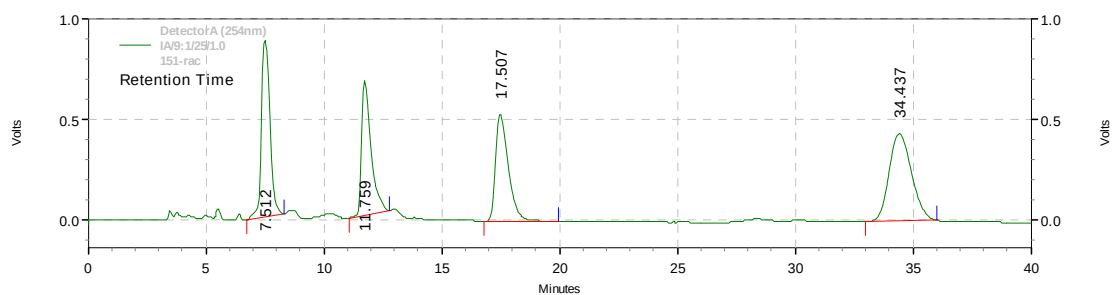
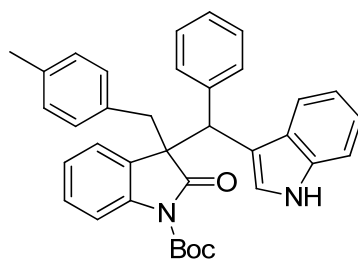
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	9.199	6971075	34.402
2	14.829	2917085	14.396
3	21.559	2950015	14.558
4	43.646	7425350	36.644



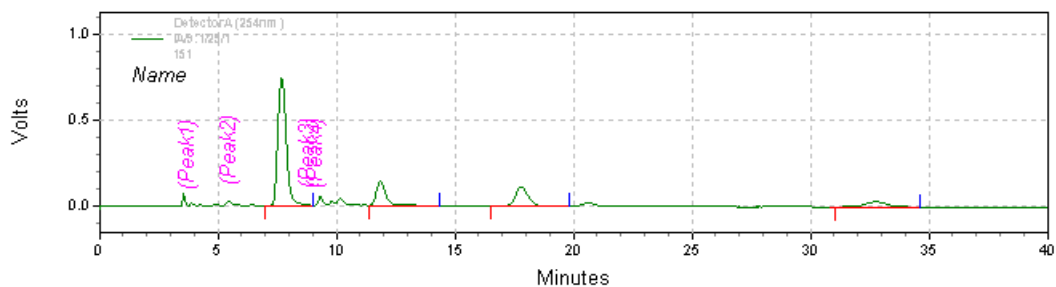
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	9.034	23285138	78.884
2	14.173	2642428	8.952
3	20.532	2459236	8.331
4	40.770	1131506	3.833



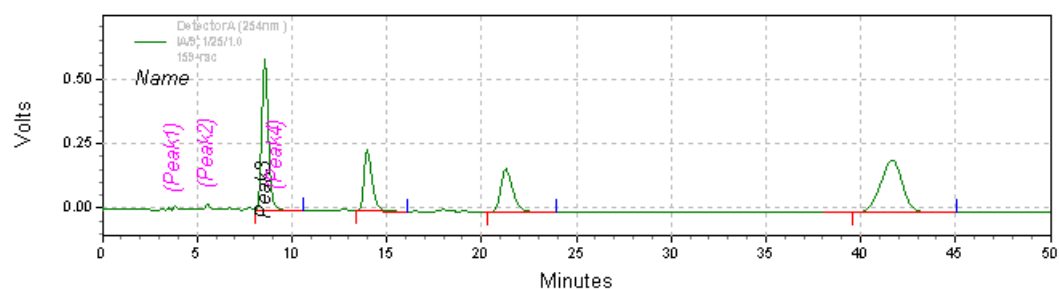
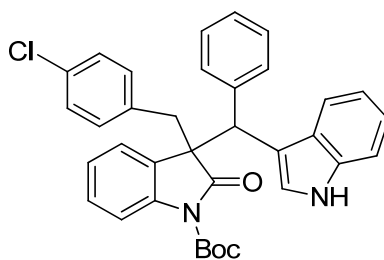
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.512	22234716	26.509
2	11.759	18421165	21.962
3	17.507	20050896	23.905
4	34.437	23170422	27.624



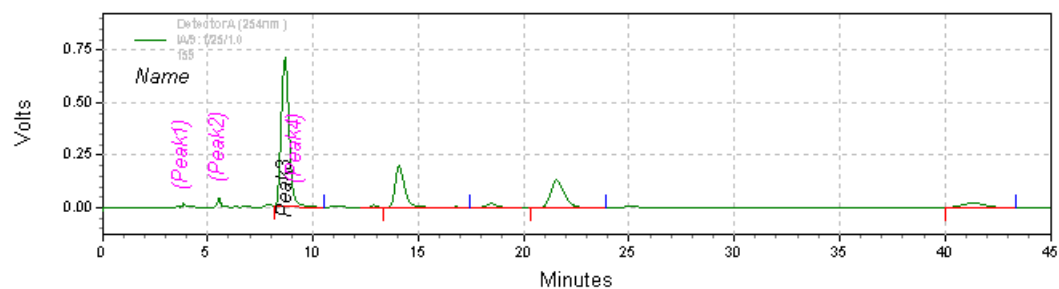
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.702	18365583	65.512
2	11.867	4042758	14.421
3	17.808	4376417	15.611
4	32.755	1249243	4.456



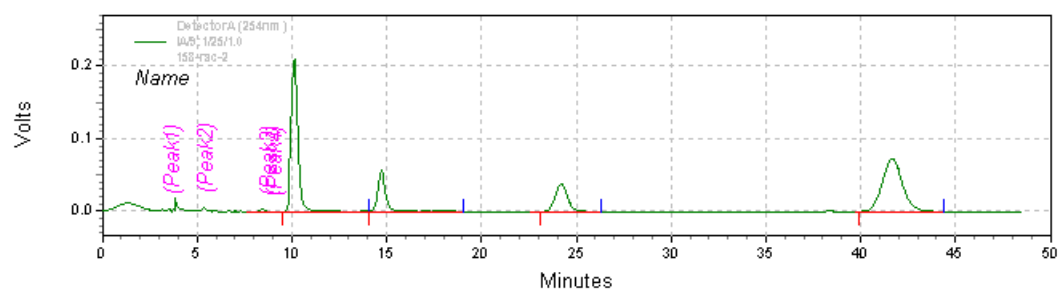
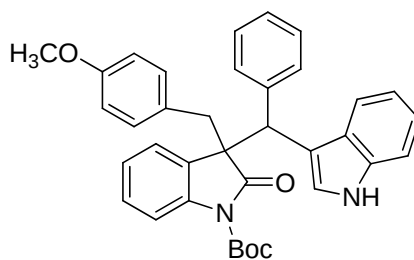
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	8.586	12739624	29.720
2	13.989	8334034	19.442
3	21.292	8546406	19.938
4	41.667	13245046	30.899



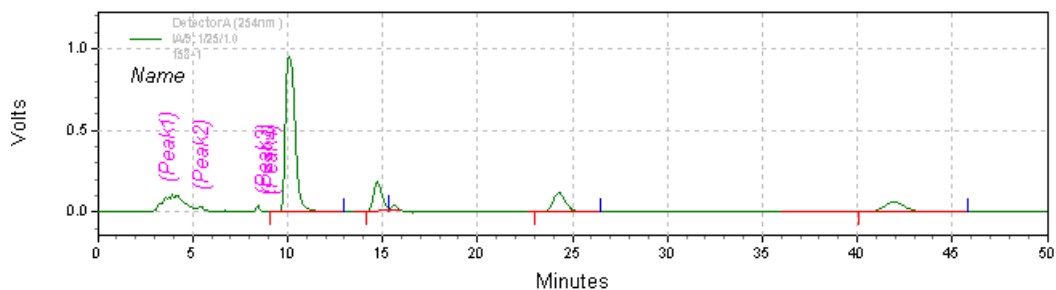
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	8.667	18355905	57.857
2	14.083	6185249	19.496
3	21.547	5795118	18.266
4	41.327	1389986	4.381



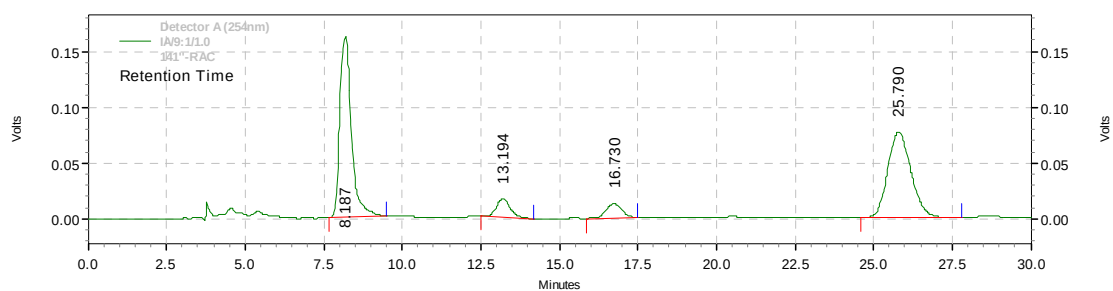
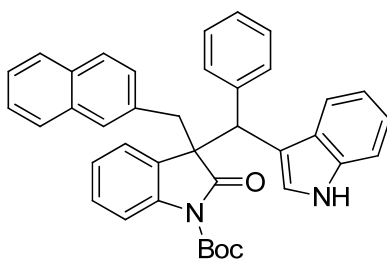
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	10.159	5327327	37.179
2	14.711	1776974	12.401
3	24.173	1731936	12.087
4	41.647	5492464	38.332



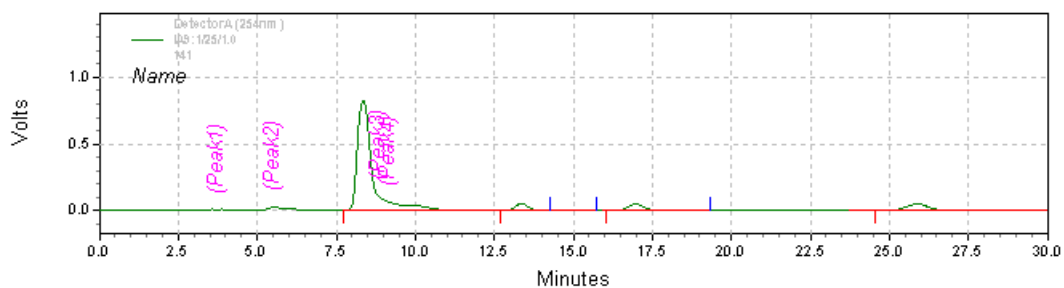
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	10.080	33374189	69.657
2	14.740	5470676	11.418
3	24.261	5462729	11.401
4	41.900	3604900	7.524



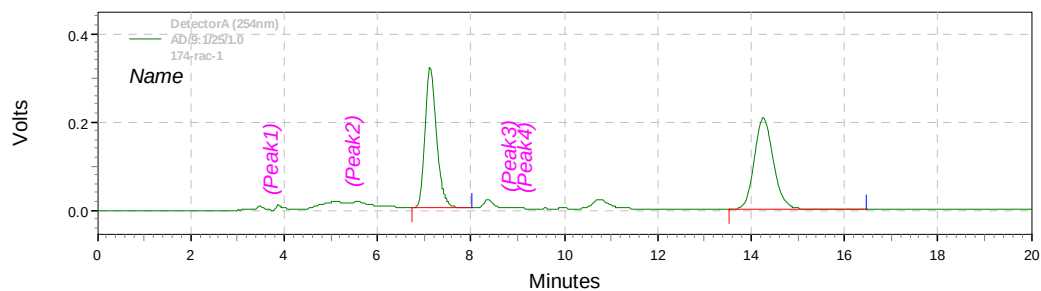
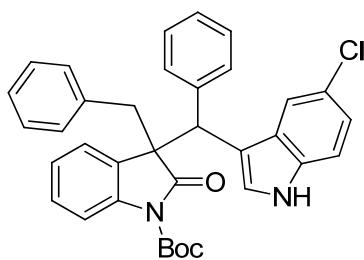
Detector A (254nm)

Pk #	Retention Time	Area	Area %
1	8.187	4058081	45.112
2	13.194	512538	5.698
3	16.730	449860	5.001
4	25.790	3975054	44.189



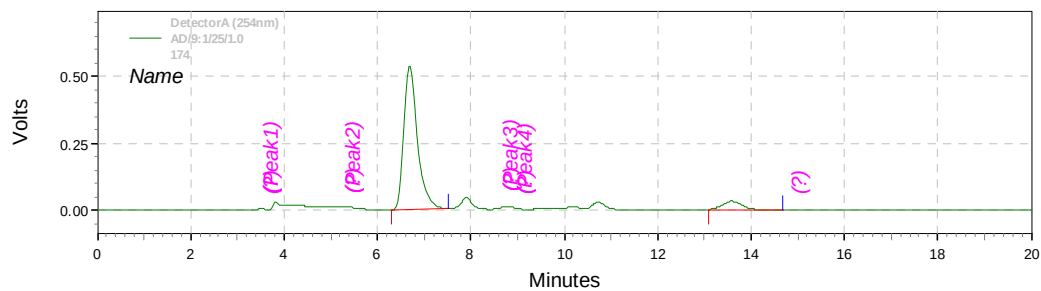
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	8.374	27277537	82.388
2	13.353	1477853	4.464
3	16.978	1749911	5.285
4	25.879	2603354	7.863



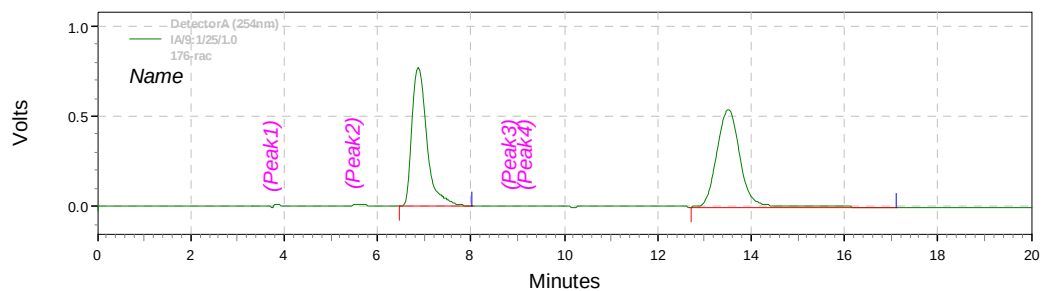
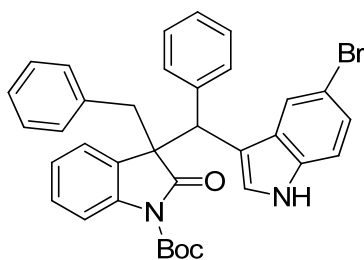
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	7.117	5503602	48.725
2	14.255	5791645	51.275



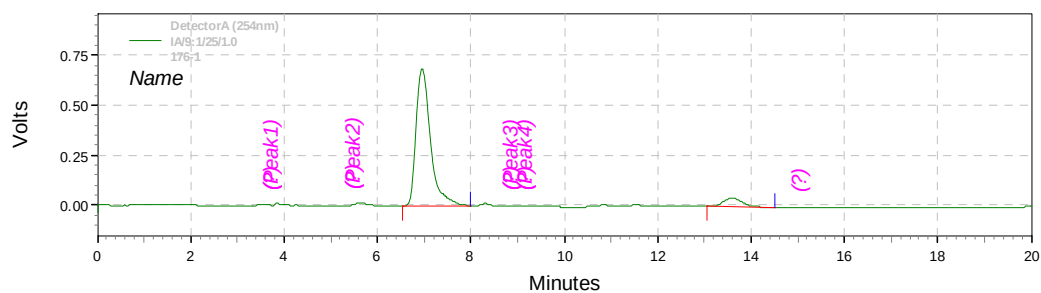
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	6.685	10682026	92.066
2	13.595	920605	7.934



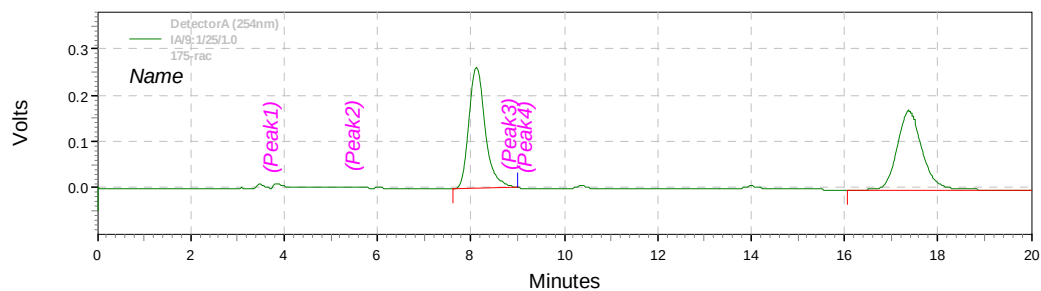
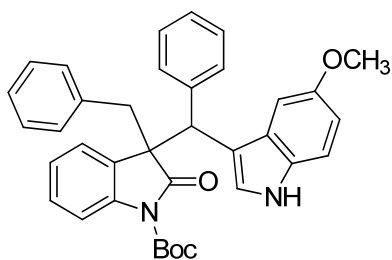
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	6.863	17157025	48.408
2	13.518	18285702	51.592



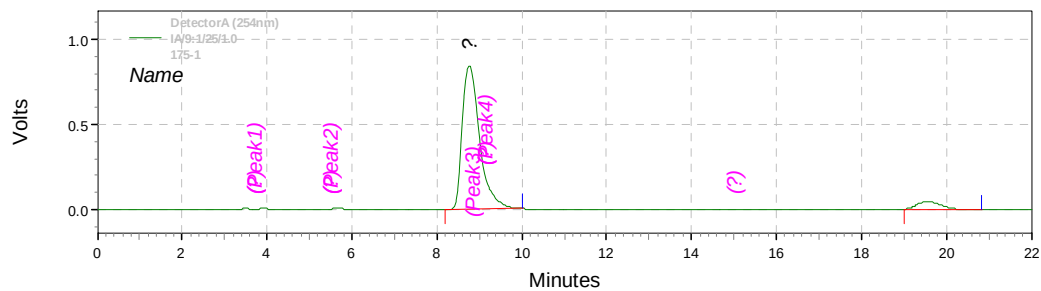
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	6.947	14683610	91.892
2	13.600	1295509	8.108



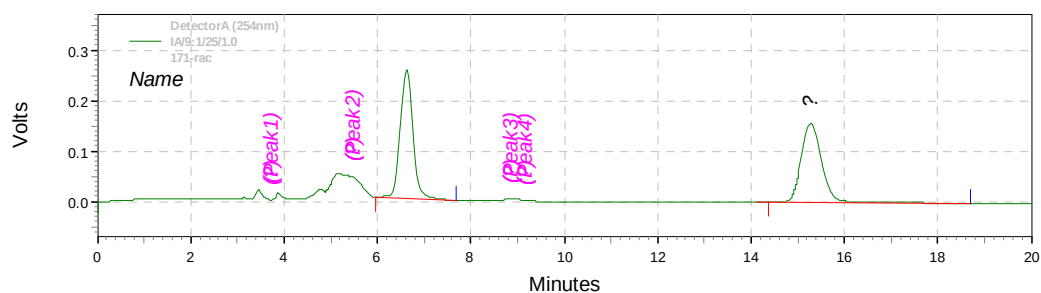
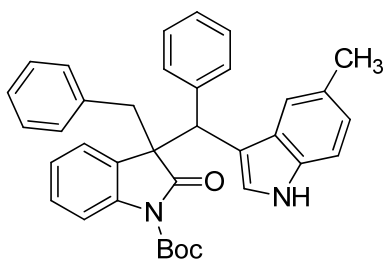
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	8.128	6304699	48.788
2	17.374	6618073	51.212



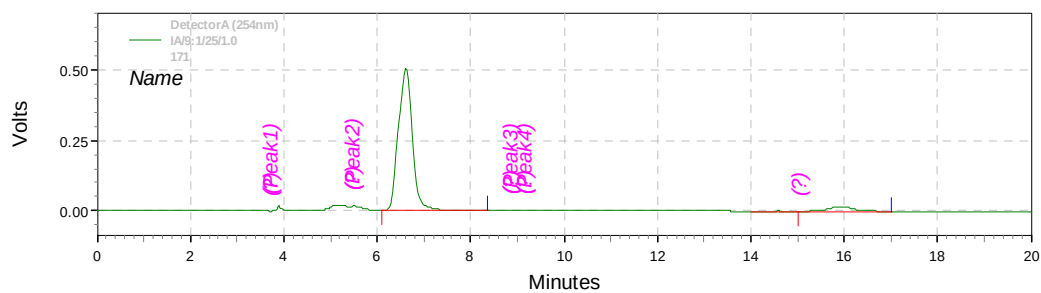
DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	8.774	23766794	93.268
2	19.578	1715526	6.732



DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	6.624	4796188	49.416
2	15.289	4909648	50.584



DetectorA (254nm)

Pk #	Retention Time	Area	Area %
1	6.611	10394244	94.883
2	15.927	560576	5.117