

Stereo- and Regioselective Photocycloaddition of Extended Alkenes Using γ -Cyclodextrin

Akshay Kashyap,^a Treyvon Bokosike,^a Nattamai Bhuvanesh,^b Mahesh Pattabiraman*^a

^a Department of Chemistry, University of Nebraska Kearney, NE – 68845, USA

^b Department of Chemistry, Texas A&M University, TX – 77845, USA

Table of Contents

Contents	Page
1. Synthesis of reactants	2
2. Complexation and irradiation	2
3. Isolation and purification of products	2
4. List of possible products	3
5. ¹ H and ¹³ C NMR spectra	3
6. Time-dependence experiment	8
7. NOESY spectra	9
8. Solution phase, and β -CD complex irradiations	10
9. Photoreactivity of 1d within γ -CD	10
10. X-Ray Crystallography	11
11. Job's plot	12
12. Computational chemistry	13
13. Isothermal titration calorimetry	28

1. Synthesis of reactants 1a, 1b, 1c, and 1d

Chalcones **1a** to **1d** were synthesized using aldol condensation reaction. Solid sodium hydroxide (2.5g) was placed in a 125 mL Erlenmeyer flask, and water (13 mL) added to it. The mixture was swirled until it is fully dissolved, and 13 mL of ethanol added to it. The mixture is brought to room temperature, and 1.8 mL acetone was added followed by 5.3 mL benzaldehyde. A yellow precipitate formed immediately, accumulated as the mixture was swirled for 15 minutes. The precipitate was filtered through a Büchner funnel, and transferred and washed with a small amount of cold ethanol. The crystals were air-dried. The crude, slightly wet product was spread out on a paper towel to dry for a short time, then weighed. The product was then recrystallized from ethyl acetate heated on a hot plate. Immiscible lower aqueous layer appeared in this step was decanted quickly. The resulting solution was allowed to cool during which recrystallization occurred. The final crystalline product was filtered and dried on a Büchner funnel. Final product was dried under vacuum overnight before recording NMR spectra. Typical product yield: 65-75%. Synthesis of other substrates followed same procedure involving the respective starting aldehydes.

2. Complexation and irradiation

Specific amounts of the guests (typically 20 mg) were weighed into a 20mL dram vial. Then half equivalent of γ -CD and water were added, and the mixture was sonicated for half hour followed by stirring for 12 hrs. Complexation is indicated by formation of chalky white or yellow precipitate, which forms a slurry of even consistency. The slurry was filtered using a filter funnel and dried to yield powdered complex. The precipitate was washed with cold water and methanol to remove uncomplexed host and guest.

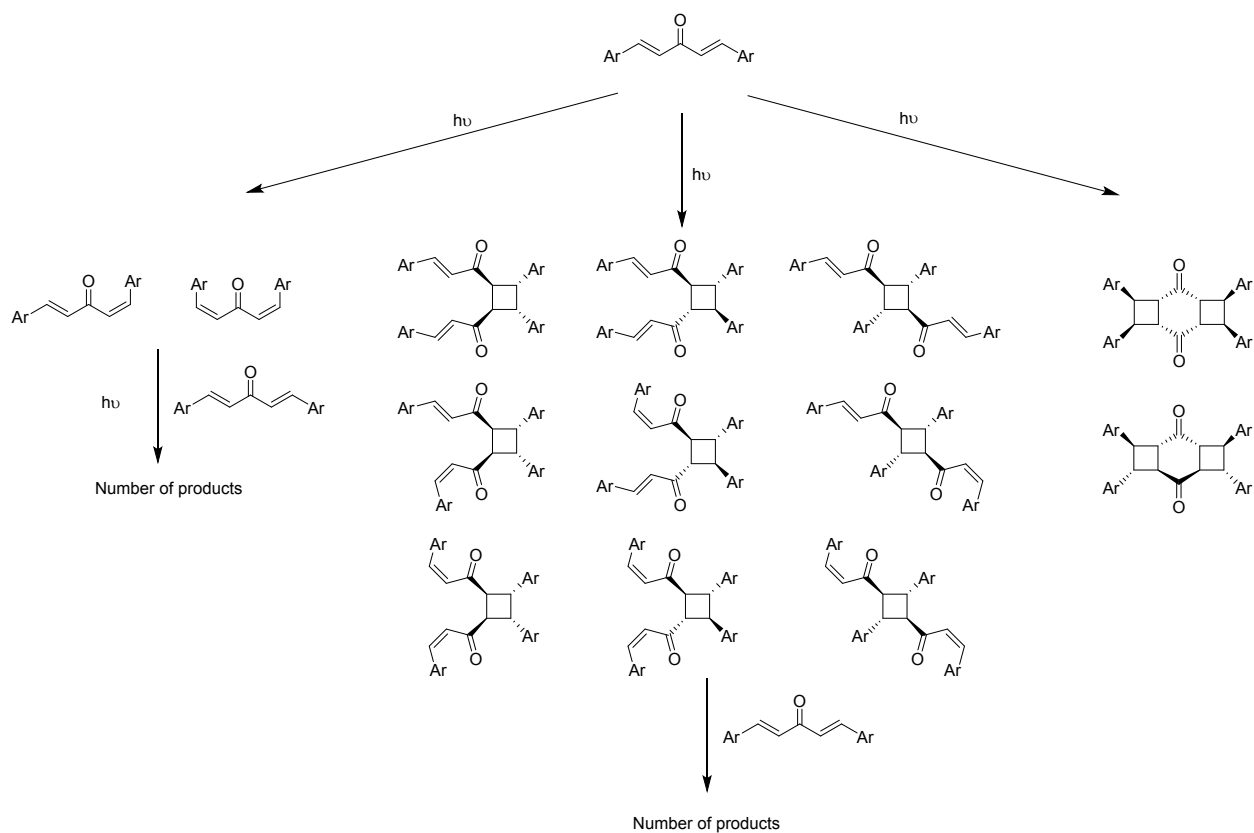
The resulting dried powder is now added to 10 mL of water and allowed to form a slurry again, and the complex is irradiated as a slurry for at least specific duration. A medium pressure photochemical lamp immersed in a water-cooled pyrex jacket was used for irradiation.

Irradiated complexes were decomplexed through biphasic extraction with 50 mL of water and 50 mL of ethyl acetate. The organic layer was separated from the aqueous layer, dried over anhydrous magnesium sulfate, and solvent removed *in vacuo*.

3. Isolation, purification of products, and characterization

Reaction mixtures with photoproducts were dissolved in 50% water/acetonitrile mixture. Compounds in the reaction mixtures were separated on a Gilson preparative HPLC. Preparative HPLC elution scheme involved initial solvent mixture of 20/80 acetonitrile/water mixture for 1 min followed by a 20 min ramp towards 80/20 acetonitrile/water mixture with the eluent flow rate maintained at 20 mL/min. Isolated products were dried in a lyophilizer to obtain pure product in solid form. Isolated compounds were characterizing primarily using ^1H and ^{13}C NMR spectra recorded using 400 MHz Bruker Avance spectrometer. Signals were analyzed using the *TopSpin 3.5pl7* software.

4. Scheme of possible products



Scheme S1. Possible products that could result from primary and secondary reaction of dibenzalacetones

5. ^1H and ^{13}C NMR spectra

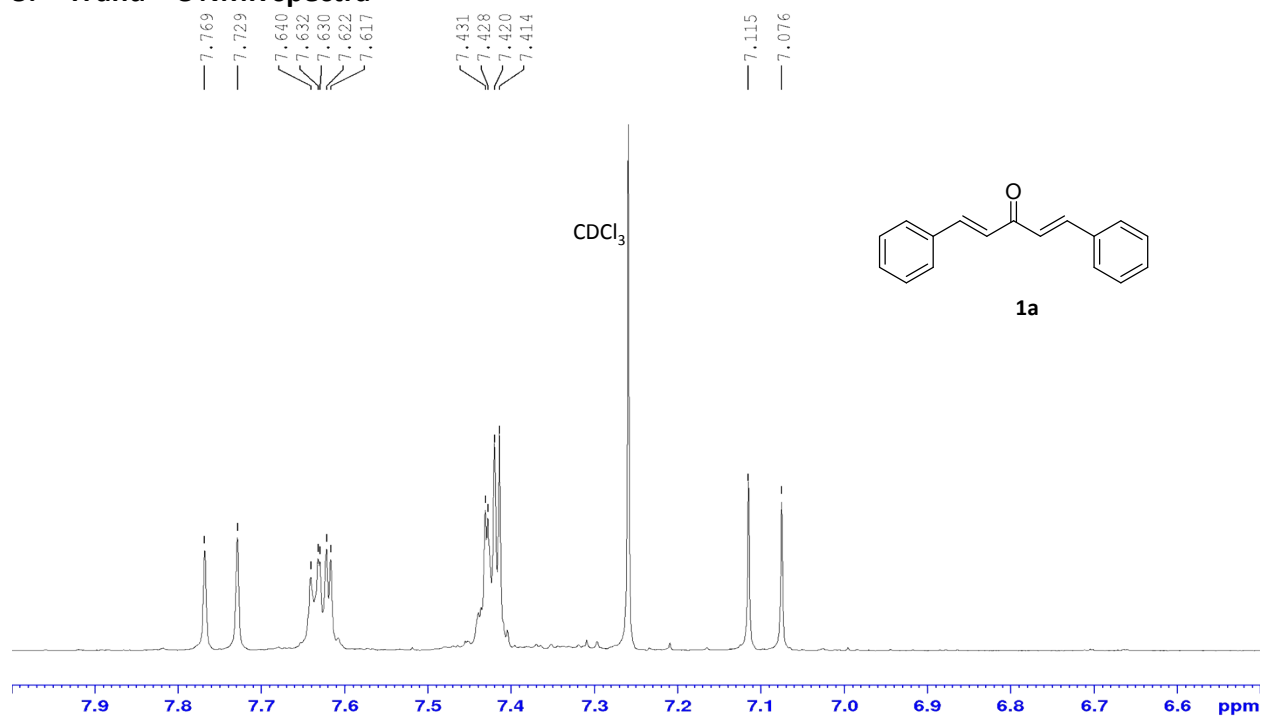


Figure S1. ^1H NMR spectrum of reactant **1a**

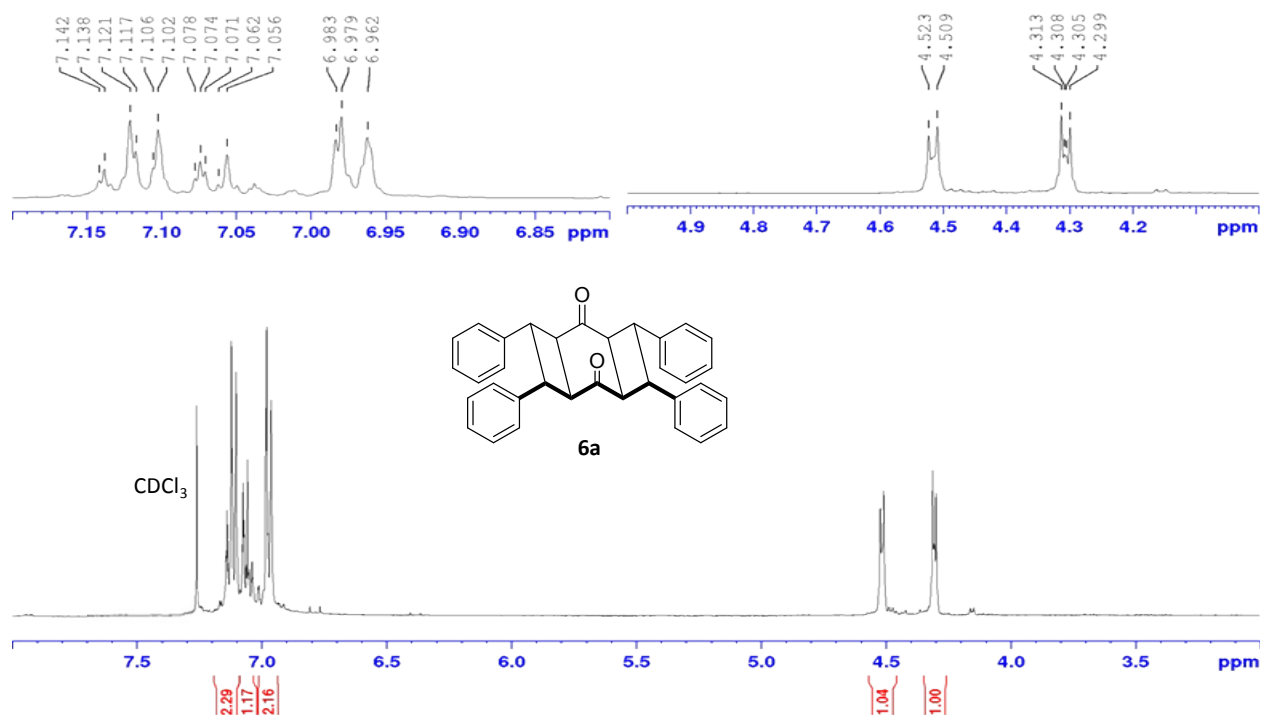


Figure S2 ^1H NMR spectrum of dimer **6a**

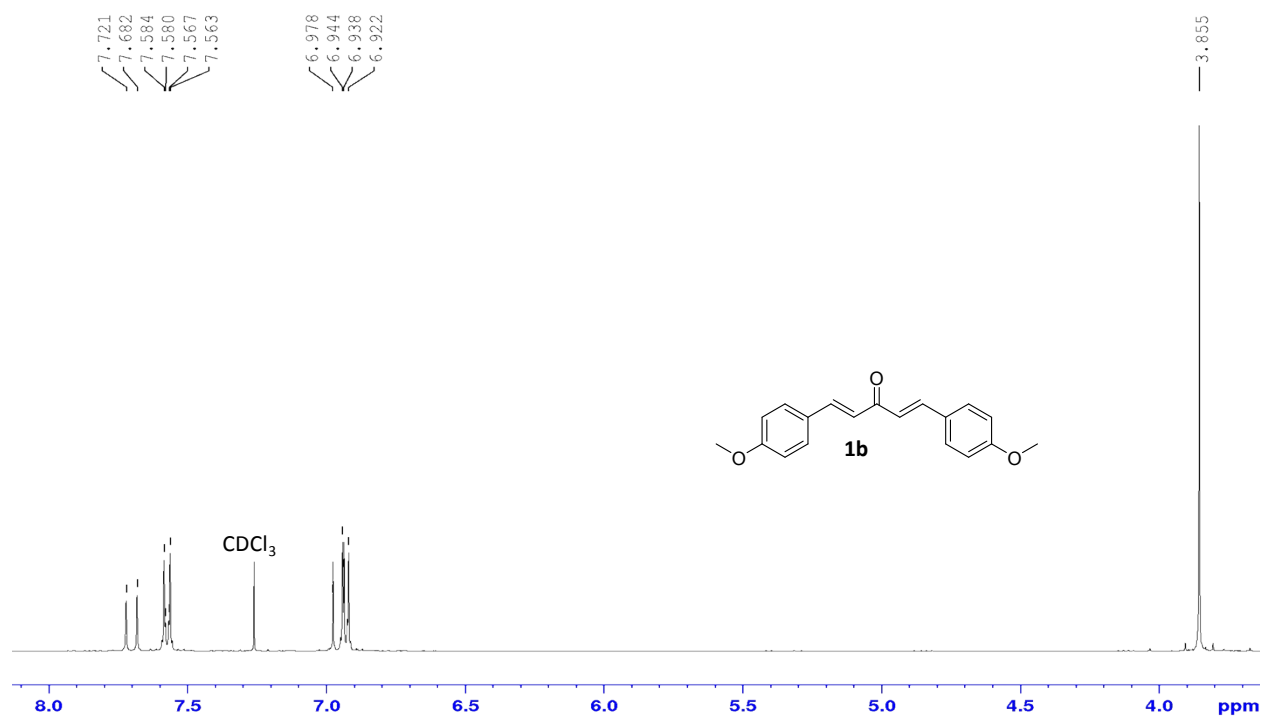


Figure S3 ^1H NMR spectrum of **1b**

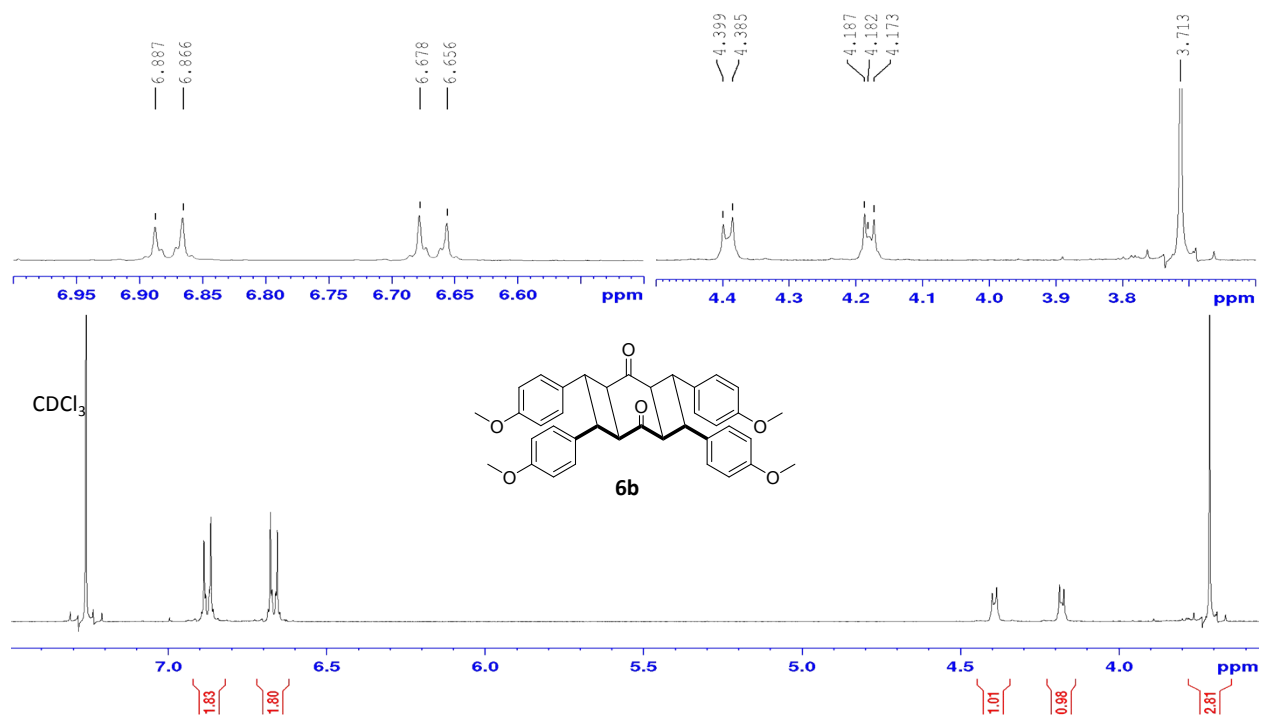


Figure S4 ¹H NMR spectrum of **6b**

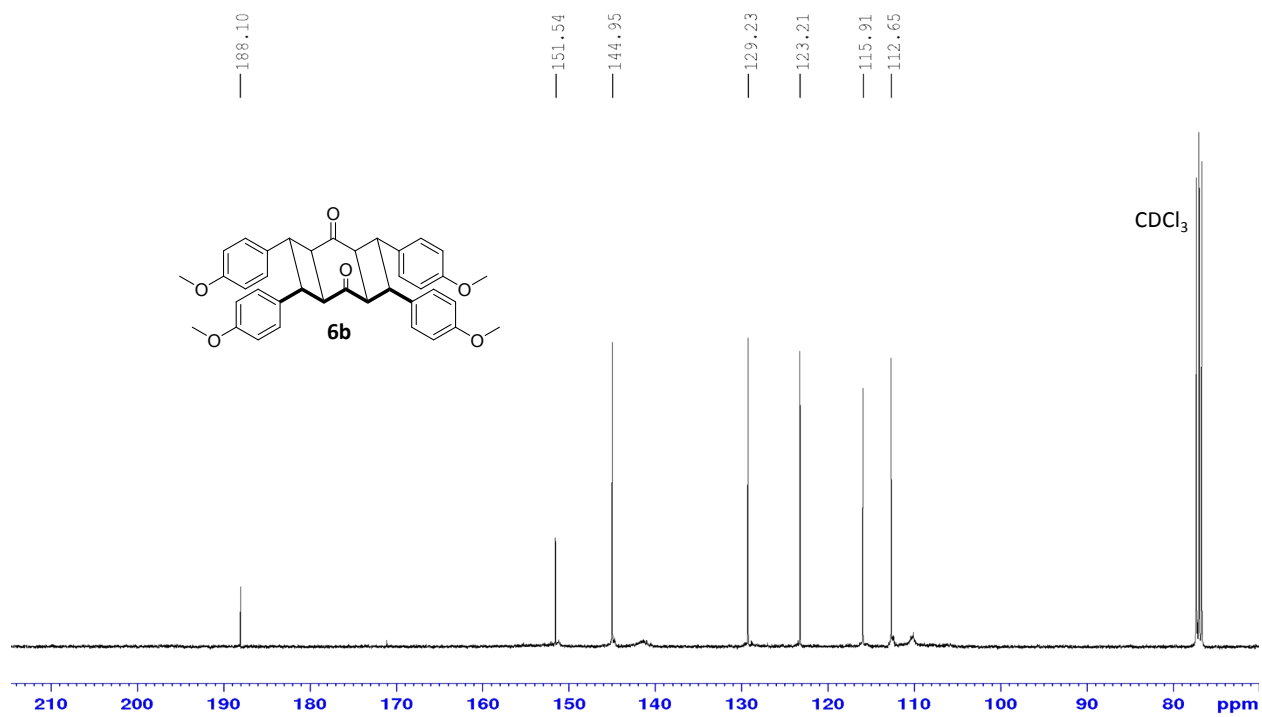


Figure S5 ¹³C NMR spectrum of **6b**

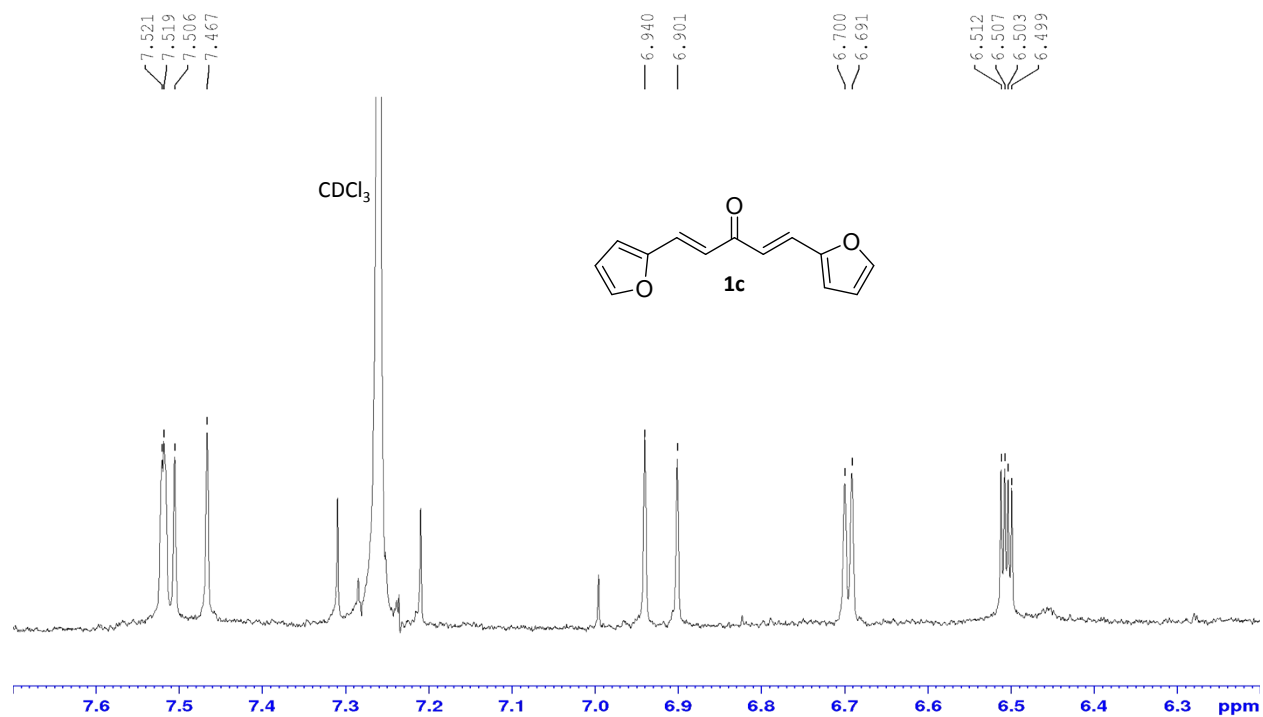


Figure S6 ¹H NMR spectrum of **1c**

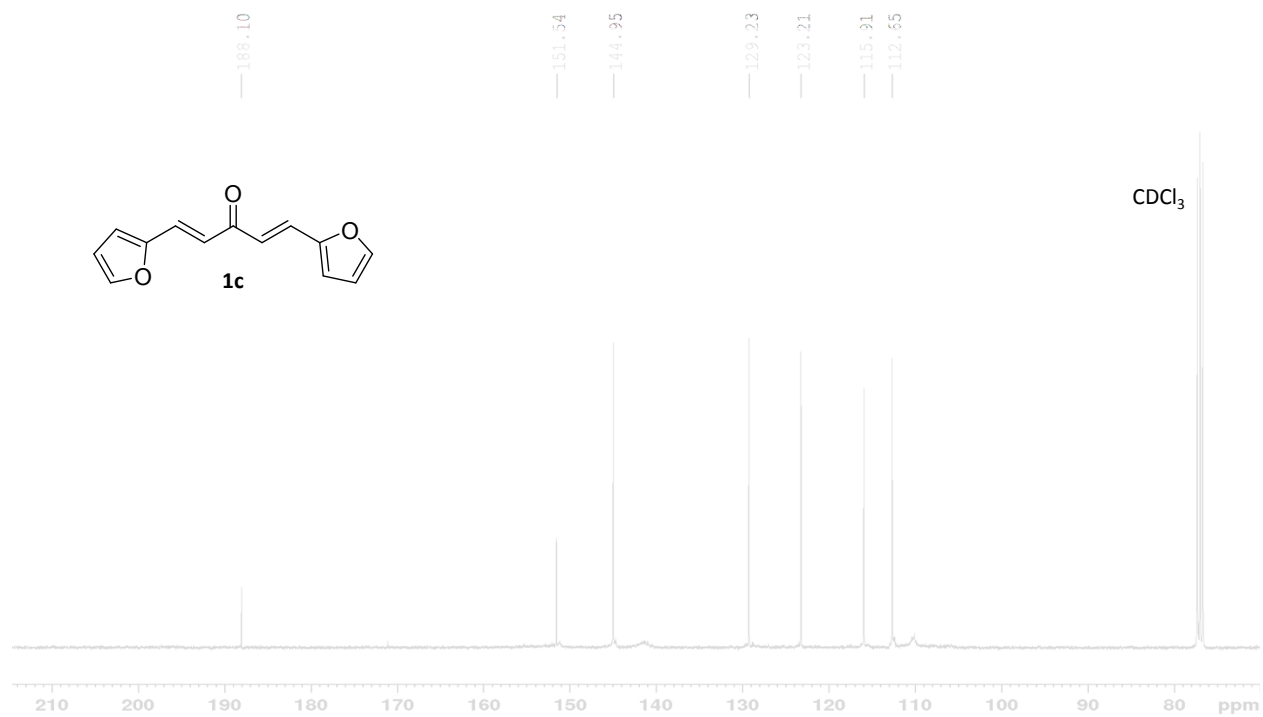


Figure S7 ¹³C NMR spectrum of **1c**

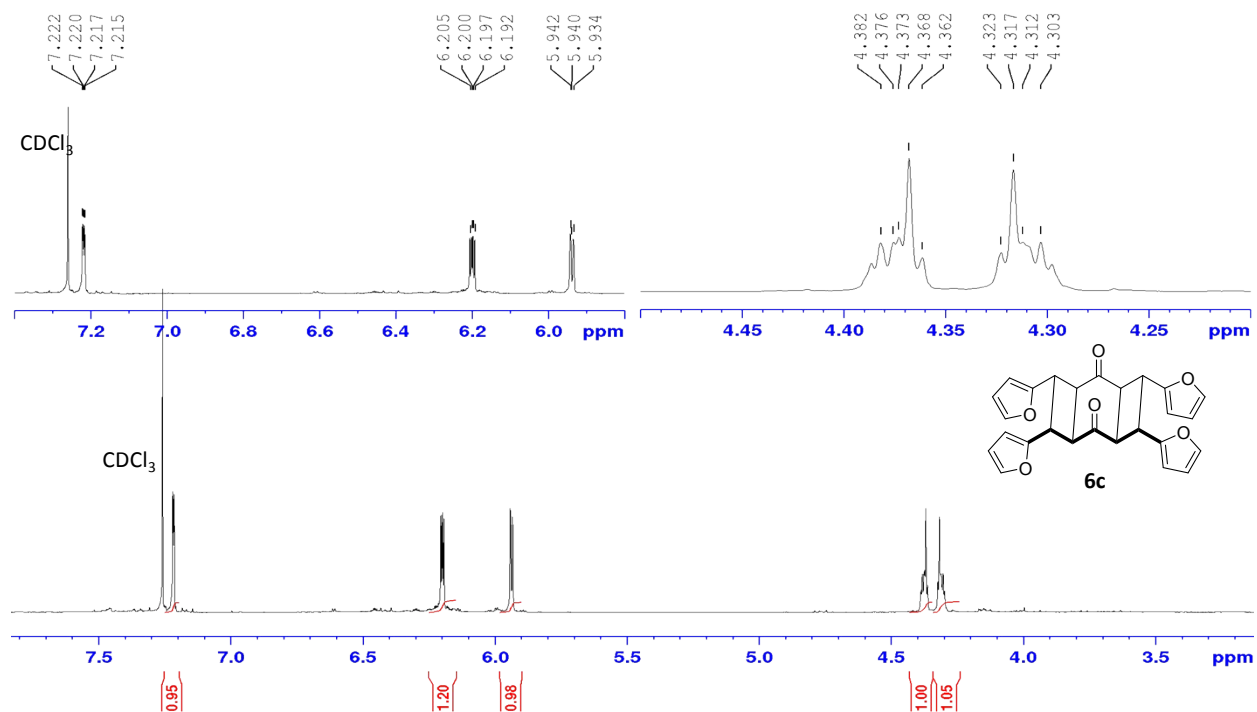


Figure S8 ^1H NMR spectrum of **6c**

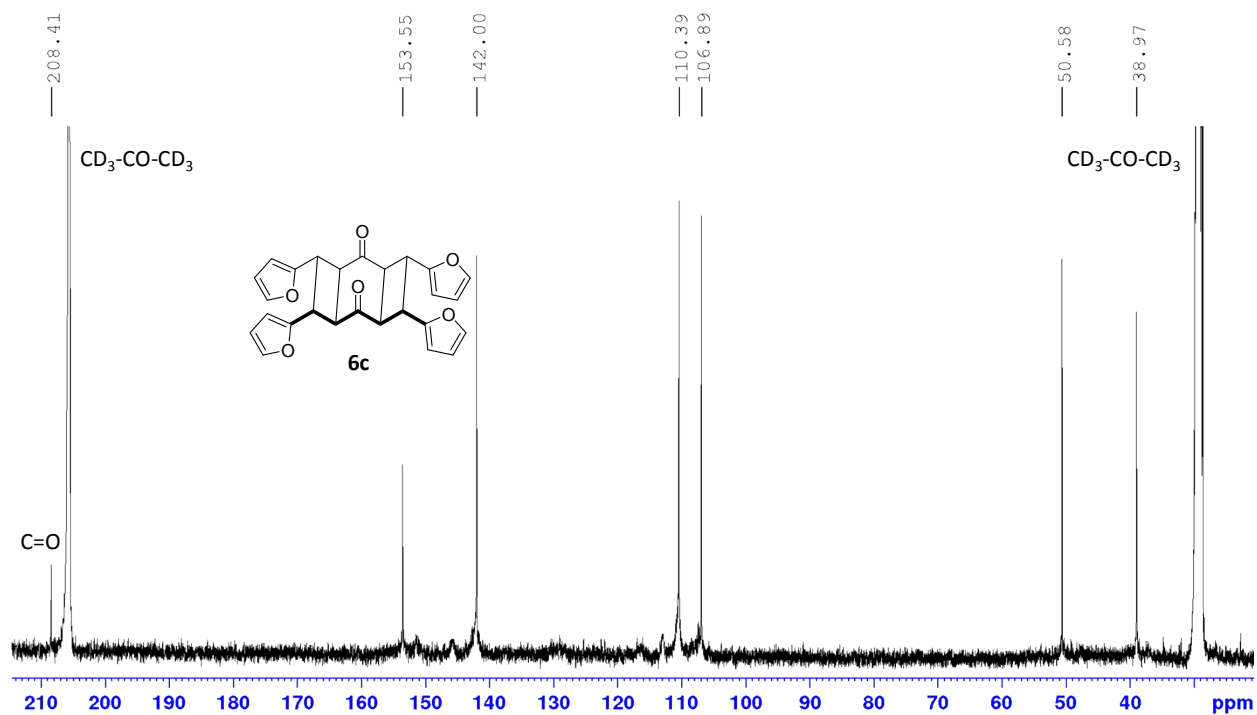


Figure S9 ^{13}C NMR spectrum of **6c**

6. Time-dependence experiment

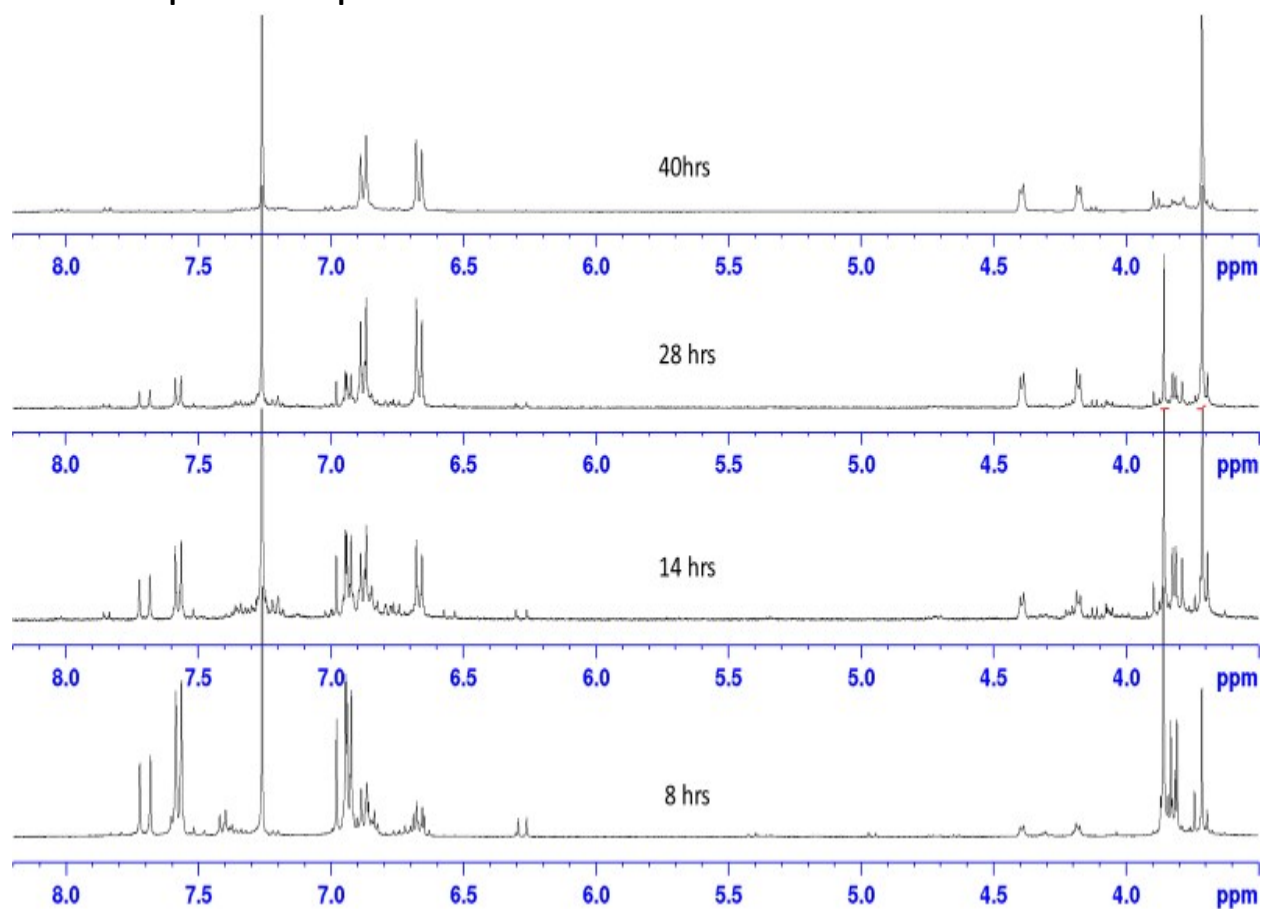


Figure S10 Time-dependent reaction of **1b**. NMR spectrum of reaction mixture of 1:2 complex irradiated for different durations as listed in the spectra.

7. NOESY spectra

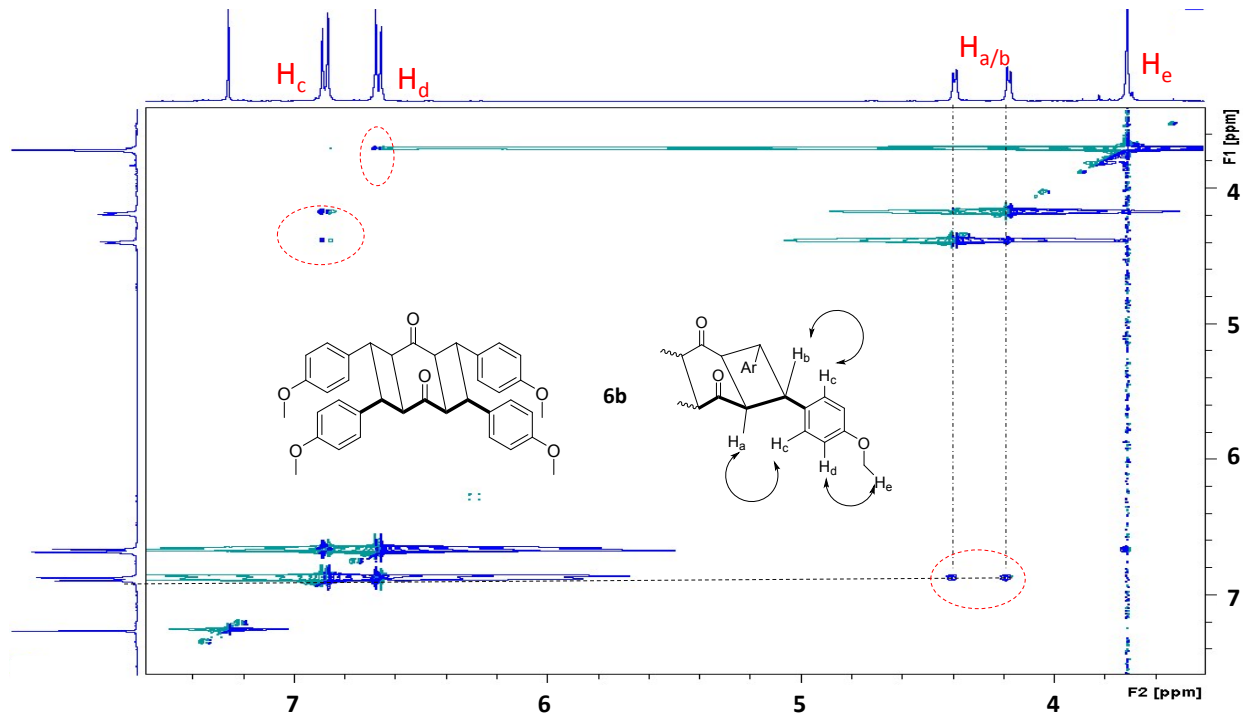


Figure S11 NOESY NMR spectrum of **6b**

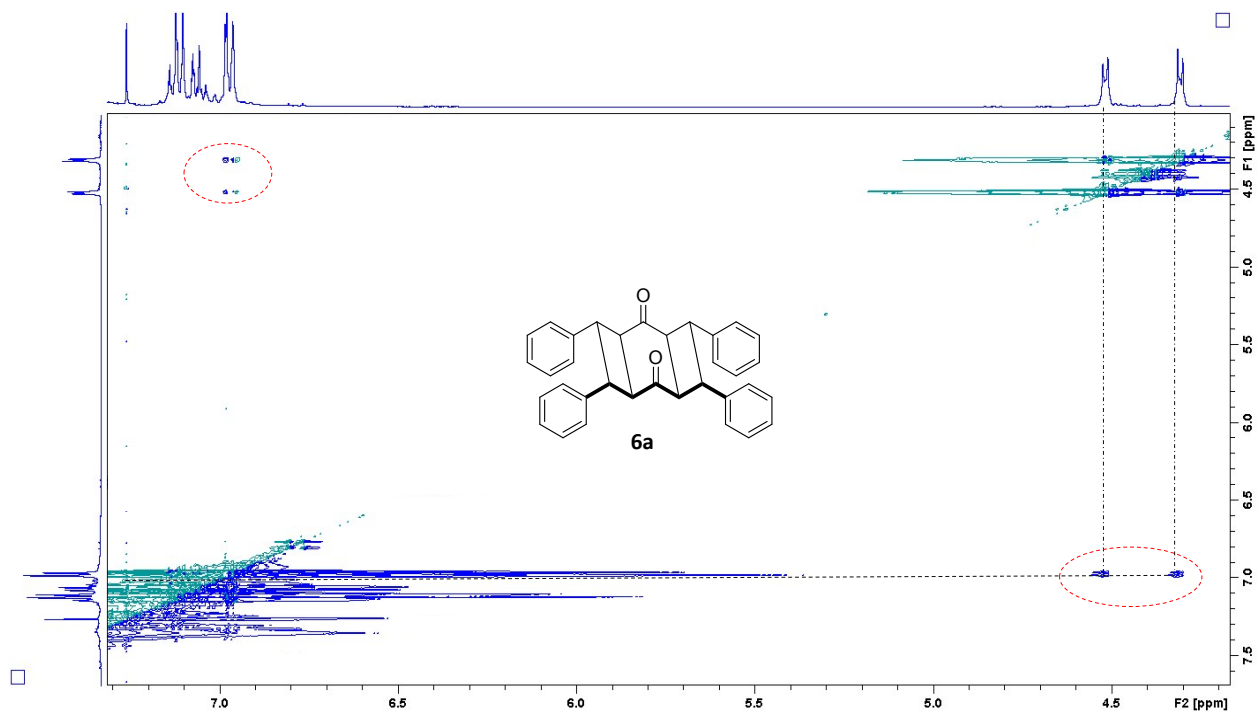


Figure S12 NOESY NMR spectrum of **6a**

8. Solution phase and irradiation of β -CD complex for compound **1b**

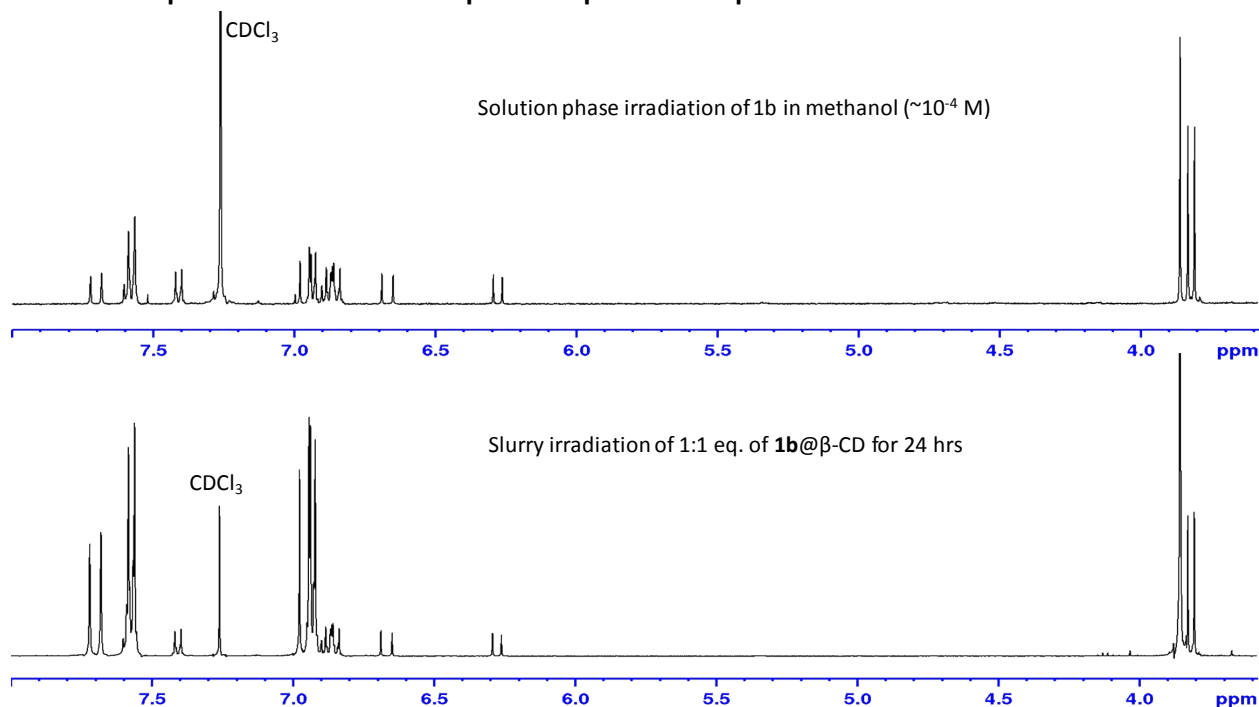


Figure S13 Product mixture obtained from irradiation of **1b** in methanol ($\sim 10^{-4}$ M, top NMR), and as a 1:1 complex with β -CD in slurry form.

9. Photoreactivity of **1d**@ γ -CD

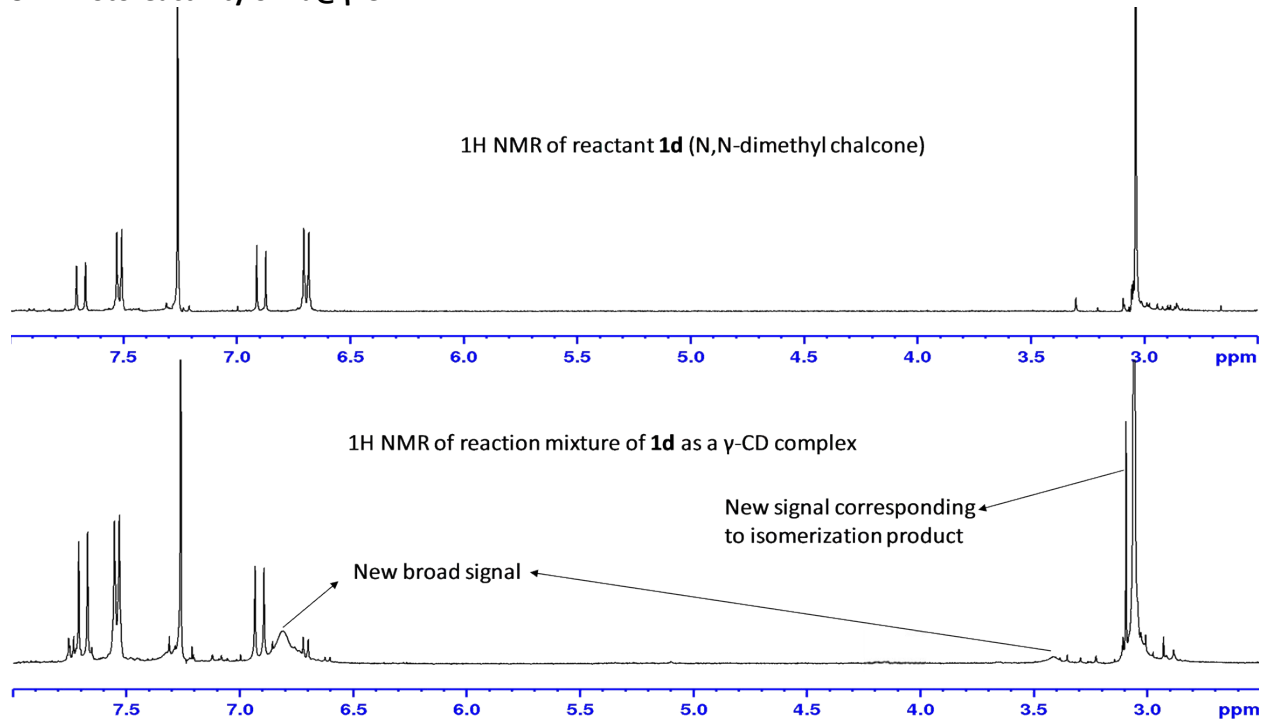


Figure S14 Product mixture obtained after irradiation of **1d**@ γ -CD after 36 hrs. Appearance of new signals in the NMR spectrum suggests photoactivity. Lack of dimers suggests supramolecular influence.

10. X-Ray crystallography

Data Collection

A Leica MZ 75 microscope was used to identify a suitable colorless plate with very well defined faces with dimensions (max, intermediate, and min) $0.243 \times 0.186 \times 0.016 \text{ mm}^3$ from a representative sample of crystals of the same habit. The crystal mounted on a nylon loop was then placed in a cold nitrogen stream (Oxford) maintained at 100 K.

A BRUKER Venture X-ray (kappa geometry) diffractometer was employed for crystal screening, unit cell determination, and data collection. The goniometer was controlled using the APEX3 software suite.¹ The sample was optically centered with the aid of a video camera such that no translations were observed as the crystal was rotated through all positions. The X-ray radiation employed was generated from a Cu- $\text{I}\mu\text{s}$ X-ray tube ($K_{\alpha} = 1.5418\text{\AA}$ with a potential of 50 kV and a current of 1.0mA). 45 data frames were taken at widths of 1° . These reflections were used to determine the unit cell. The unit cell was verified by examination of the $h k l$ overlays on several frames of data. No super-cell or erroneous reflections were observed. After careful examination of the unit cell, an extended data collection procedure (26 sets) was initiated using omega and phi scans.

Data Reduction, Structure Solution, and Refinement

Integrated intensity information for each reflection was obtained by reduction of the data frames with the program APEX3.¹ The integration method employed a three-dimensional profiling algorithm and all data were corrected for Lorentz and polarization factors, as well as for crystal decay effects. Finally the data was merged and scaled to produce a suitable data set. The absorption correction program SADABS² was employed to correct the data for absorption effects.

Systematic reflection conditions and statistical tests of the data suggested the space group $P-1$. A solution was obtained readily using XT/XS in APEX3.^{1,3} Hydrogen atoms were placed in idealized positions and were set riding on the respective parent atoms. All non-hydrogen atoms were refined with anisotropic thermal parameters. Absence of additional symmetry or void were confirmed using PLATON (ADDSYM).[#] The structure was refined (weighted least squares refinement on F^2) to convergence.^{3,4}

Olex2 was employed for the final data presentation and structure plots.⁴

¹ APEX3 “Program for Data Collection on Area Detectors” BRUKER AXS Inc., 5465 East Cheryl Parkway, Madison, WI 53711-5373 USA

² SADABS, Sheldrick, G.M. “Program for Absorption Correction of Area Detector Frames”, BRUKER AXS Inc., 5465 East Cheryl Parkway, Madison, WI 53711-5373 USA.

³ Sheldrick, G.M. (2008). Acta Cryst. A64, 112-122. Sheldrick, G. M. (2015), Acta Cryst. A71, 3-8. Sheldrick, G. M. (2015). Acta Cryst. C71, 3-8. XT, XS, BRUKER AXS Inc., 5465 East Cheryl Parkway, Madison, WI 53711-5373 USA.

⁴ Dolomanov, O. V, Bourhis, L. J., Gildea, R. J., Howard, J. A. K., and Puschmann, H. “OLEX2: A Complete Structure Solution, Refinement and Analysis Program”, *J. Appl. Cryst.* **2009**, *42*, 339-341.

[#] Spek, A. L., "PLATON - A Multipurpose Crystallographic Tool" *J. Appl. Cryst.* **2003**, *36*, 7-13.; Spek, A. L., Utrecht University, Utrecht, The Netherlands **2008**.

* Sheldrick, G. M. “Cell_Now (version **2008/1**): Program for Obtaining Unit Cell Constants from Single Crystal Data”: University of Göttingen, Germany

§ Spek, A. L. (2015) Acta Cryst. C71, 9-18.

11. Job's continuous variation method

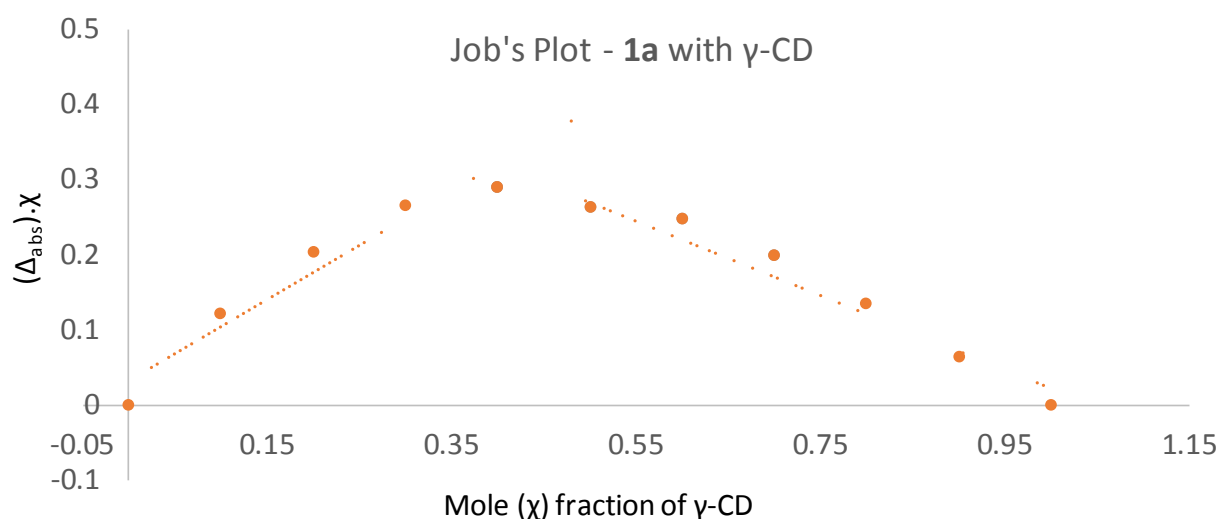


Figure S15 Job's plot for **1a** (dibenzalacetone) with γ -CD

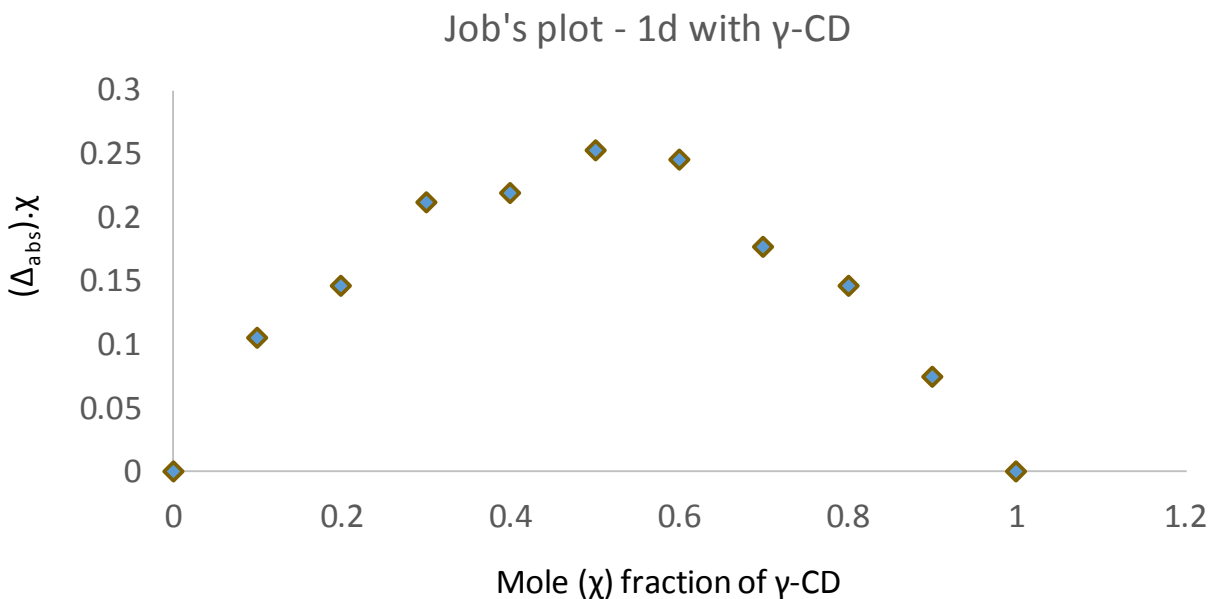


Figure S16 Job's plot for **1d** (N,N-dimethyl chalcone) with γ -CD

12. Computational chemistry

Computational chemistry calculations were performed using Gaussian '09. Windows 10 desktop with eight dual core processors and 16 GB RAM were used for the purposes. Calculations were performed at different levels of theory to obtain complex structures for gaining insight into relative dimensions and arrangement of the components. Theory performed at molecular mechanics (MM, UFF) and semi empirical (SE PM6) using solvent effects based on IPCM were first obtained. However, the geometry-optimized structures showed the reactant chalcones in which the aromatic rings were orthogonal to the pentadienone skeleton. This was realized to be an unrealistic due to the common knowledge that the molecule is fully conjugated and should remain planar. Hence *ab initio* calculations were performed using ONIOM wB97XD/631++G(2d,2p)/Auto:PM6.

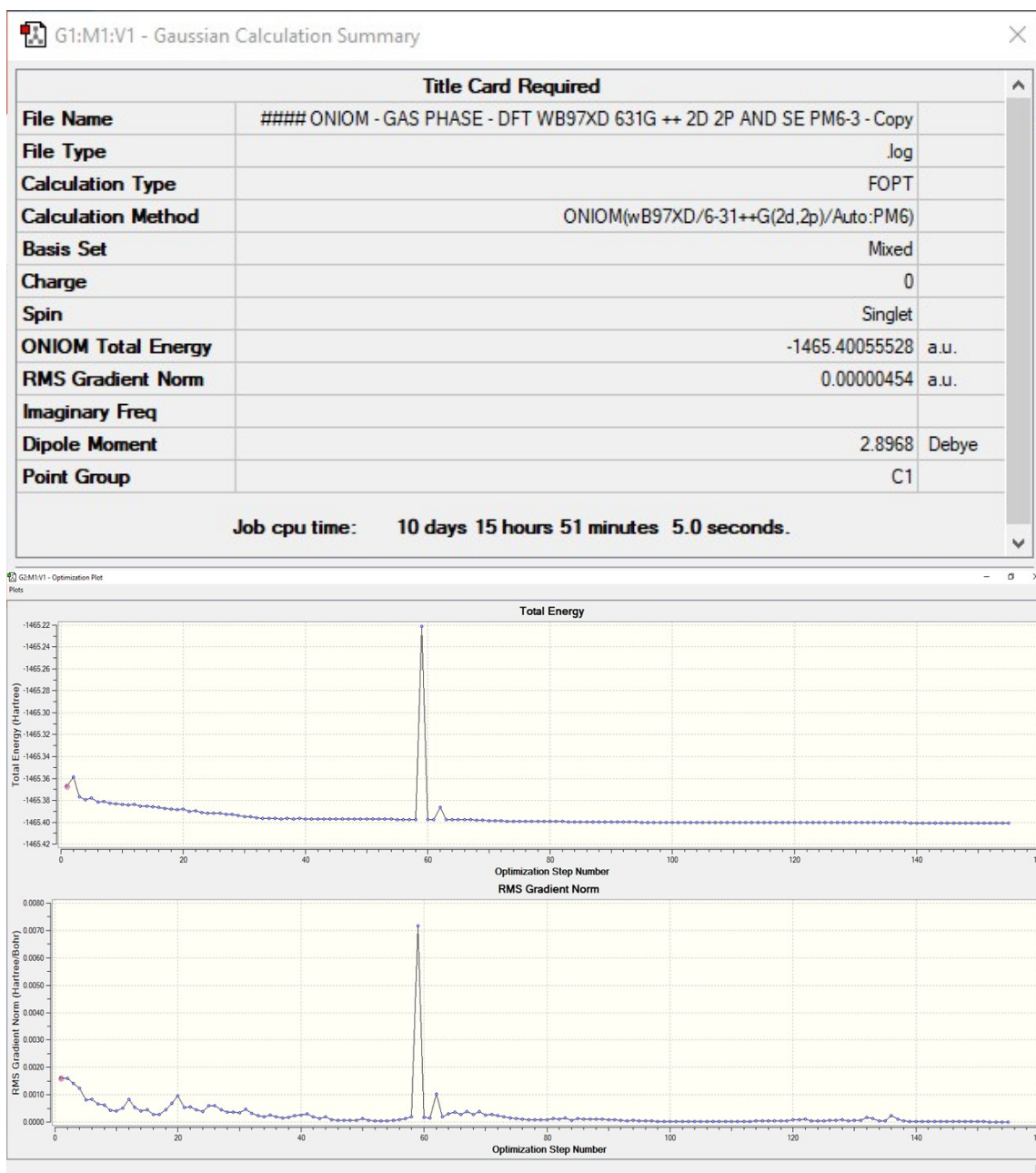


Figure S17. Output screen from calculation performed on $1a_2@Y\text{-Cd}$
Coordinates of final structure obtained after geometry optimization (.pdb file format)

```

TITLE          Required
REMARK        1 File created by GaussView 5.0.9
HETATM        1  C              0      -2.272    5.956    1.448
C
HETATM        2  H              0      -1.973    7.011    1.647
H
HETATM        3  C              0      -3.132    5.300    2.558
C
HETATM        4  H              0      -3.951    5.988    2.878
H

```

HETATM	5	C	0	-3.689	3.946	2.061
C						
HETATM	6	H	0	-2.883	3.176	1.952
H						
HETATM	7	C	0	-4.450	4.151	0.726
C						
HETATM	8	H	0	-5.390	4.727	0.869
H						
HETATM	9	C	0	-3.538	4.819	-0.327
C						
HETATM	10	H	0	-2.705	4.139	-0.614
H						
HETATM	11	C	0	-4.281	5.408	-1.547
C						
HETATM	12	H	0	-4.600	6.452	-1.338
H						
HETATM	13	H	0	-3.636	5.394	-2.444
H						
HETATM	14	O	0	-4.754	2.803	0.273
O						
HETATM	15	O	0	-2.953	6.036	0.216
O						
HETATM	16	C	0	3.014	5.470	1.140
C						
HETATM	17	H	0	3.960	6.026	1.355
H						
HETATM	18	C	0	2.420	4.729	2.370
C						
HETATM	19	H	0	2.739	5.239	3.311
H						
HETATM	20	C	0	0.885	4.548	2.342
C						
HETATM	21	H	0	0.621	3.537	1.921
H						
HETATM	22	C	0	0.142	5.666	1.589
C						
HETATM	23	H	0	0.008	6.572	2.227
H						
HETATM	24	C	0	0.825	6.039	0.262
C						
HETATM	25	H	0	0.872	5.168	-0.440
H						
HETATM	26	C	0	0.179	7.265	-0.399
C						
HETATM	27	H	0	0.795	7.642	-1.234
H						
HETATM	28	H	0	-0.851	7.028	-0.748
H						
HETATM	29	O	0	-1.141	5.084	1.263
O						
HETATM	30	O	0	2.176	6.462	0.574
O						

HETATM	31	C	0	6.478	2.799	-1.719
C						
HETATM	32	H	0	7.252	2.608	-2.497
H						
HETATM	33	C	0	6.845	3.871	-0.667
C						
HETATM	34	H	0	7.287	4.774	-1.155
H						
HETATM	35	C	0	5.624	4.263	0.195
C						
HETATM	36	H	0	5.408	3.502	0.983
H						
HETATM	37	C	0	4.381	4.546	-0.668
C						
HETATM	38	H	0	4.430	5.569	-1.104
H						
HETATM	39	C	0	4.131	3.439	-1.718
C						
HETATM	40	H	0	3.802	2.497	-1.217
H						
HETATM	41	C	0	3.137	3.891	-2.802
C						
HETATM	42	H	0	3.388	4.886	-3.212
H						
HETATM	43	H	0	3.081	3.157	-3.629
H						
HETATM	44	O	0	3.187	4.441	0.153
O						
HETATM	45	C	0	6.798	-2.472	-1.173
C						
HETATM	46	H	0	7.269	-3.427	-1.484
H						
HETATM	47	C	0	7.558	-1.705	-0.063
C						
HETATM	48	H	0	8.655	-1.680	-0.284
H						
HETATM	49	C	0	7.015	-0.267	0.114
C						
HETATM	50	H	0	6.051	-0.248	0.672
H						
HETATM	51	C	0	6.903	0.442	-1.248
C						
HETATM	52	H	0	7.911	0.721	-1.639
H						
HETATM	53	C	0	6.107	-0.405	-2.262
C						
HETATM	54	H	0	5.057	-0.559	-1.917
H						
HETATM	55	C	0	6.168	0.176	-3.680
C						
HETATM	56	H	0	5.843	-0.564	-4.433
H						

HETATM	57	H	0	5.550	1.100	-3.751
H						
HETATM	58	O	0	6.127	1.627	-0.965
O						
HETATM	59	O	0	6.741	-1.712	-2.383
O						
HETATM	60	C	0	-6.118	-2.477	-1.796
C						
HETATM	61	H	0	-6.888	-2.036	-2.472
H						
HETATM	62	C	0	-6.603	-3.676	-0.950
C						
HETATM	63	H	0	-7.112	-4.434	-1.598
H						
HETATM	64	C	0	-5.427	-4.313	-0.180
C						
HETATM	65	H	0	-5.129	-3.691	0.694
H						
HETATM	66	C	0	-4.238	-4.604	-1.108
C						
HETATM	67	H	0	-4.384	-5.590	-1.618
H						
HETATM	68	C	0	-3.900	-3.480	-2.102
C						
HETATM	69	H	0	-3.321	-2.676	-1.577
H						
HETATM	70	C	0	-3.167	-3.991	-3.363
C						
HETATM	71	H	0	-3.741	-3.713	-4.276
H						
HETATM	72	H	0	-2.146	-3.572	-3.424
H						
HETATM	73	O	0	-3.040	-4.634	-0.277
O						
HETATM	74	O	0	-5.071	-2.836	-2.670
O						
HETATM	75	C	0	-6.121	2.595	-0.134
C						
HETATM	76	H	0	-6.672	3.560	-0.246
H						
HETATM	77	C	0	-6.784	1.598	0.845
C						
HETATM	78	H	0	-7.896	1.611	0.706
H						
HETATM	79	C	0	-6.250	0.155	0.682
C						
HETATM	80	H	0	-5.238	0.025	1.123
H						
HETATM	81	C	0	-6.298	-0.259	-0.799
C						
HETATM	82	H	0	-7.353	-0.406	-1.135
H						

HETATM	83	C	0	-5.554	0.765	-1.681
C						
HETATM	84	H	0	-4.464	0.786	-1.444
H						
HETATM	85	C	0	-5.799	0.519	-3.179
C						
HETATM	86	H	0	-5.570	1.425	-3.776
H						
HETATM	87	H	0	-5.181	-0.343	-3.535
H						
HETATM	88	O	0	-5.570	-1.502	-0.876
O						
HETATM	89	O	0	-6.080	2.102	-1.466
O						
HETATM	90	C	0	2.624	-5.717	-0.285
C						
HETATM	91	H	0	2.408	-6.803	-0.436
H						
HETATM	92	C	0	3.572	-5.422	0.908
C						
HETATM	93	H	0	4.252	-6.290	1.086
H						
HETATM	94	C	0	4.389	-4.122	0.737
C						
HETATM	95	H	0	3.803	-3.212	1.055
H						
HETATM	96	C	0	4.890	-3.949	-0.718
C						
HETATM	97	H	0	5.651	-4.700	-1.017
H						
HETATM	98	C	0	3.676	-3.977	-1.667
C						
HETATM	99	H	0	2.919	-3.215	-1.363
H						
HETATM	100	C	0	3.979	-3.934	-3.177
C						
HETATM	101	H	0	3.621	-4.877	-3.654
H						
HETATM	102	H	0	3.487	-3.069	-3.653
H						
HETATM	103	O	0	5.449	-2.614	-0.725
O						
HETATM	104	O	0	3.080	-5.305	-1.548
O						
HETATM	105	C	0	-2.650	-5.930	0.192
C						
HETATM	106	H	0	-3.512	-6.631	0.271
H						
HETATM	107	C	0	-1.924	-5.646	1.532
C						
HETATM	108	H	0	-1.851	-6.573	2.149
H						

HETATM	109	C	0	-0.532	-5.003	1.318
C						
HETATM	110	H	0	-0.617	-3.917	1.066
H						
HETATM	111	C	0	0.250	-5.772	0.234
C						
HETATM	112	H	0	0.551	-6.787	0.586
H						
HETATM	113	C	0	-0.555	-5.863	-1.076
C						
HETATM	114	H	0	-0.762	-4.851	-1.490
H						
HETATM	115	C	0	0.119	-6.777	-2.111
C						
HETATM	116	H	0	-0.567	-7.025	-2.942
H						
HETATM	117	H	0	1.048	-6.305	-2.504
H						
HETATM	118	O	0	1.415	-4.962	-0.021
O						
HETATM	119	O	0	-1.827	-6.518	-0.807
O						
HETATM	120	O	0	-7.620	-3.279	-0.049
O						
HETATM	121	H	0	-7.322	-2.497	0.509
H						
HETATM	122	O	0	-5.811	-5.615	0.293
O						
HETATM	123	H	0	-6.665	-5.552	0.784
H						
HETATM	124	O	0	-2.749	-4.800	2.319
O						
HETATM	125	H	0	-3.040	-4.014	1.787
H						
HETATM	126	O	0	0.208	-4.972	2.534
O						
HETATM	127	H	0	0.447	-5.876	2.851
H						
HETATM	128	O	0	-6.616	2.022	2.179
O						
HETATM	129	H	0	-5.627	2.188	2.382
H						
HETATM	130	O	0	-7.154	-0.740	1.348
O						
HETATM	131	H	0	-7.316	-0.428	2.272
H						
HETATM	132	O	0	-4.545	3.356	3.039
O						
HETATM	133	H	0	-5.137	4.020	3.462
H						
HETATM	134	O	0	-2.353	5.161	3.726
O						

HETATM	135	H	0	-1.552	4.563	3.560
H						
HETATM	136	O	0	0.362	4.457	3.671
O						
HETATM	137	H	0	0.594	5.243	4.212
H						
HETATM	138	O	0	3.056	3.464	2.463
O						
HETATM	139	H	0	2.840	2.911	1.656
H						
HETATM	140	O	0	5.909	5.520	0.836
O						
HETATM	141	H	0	6.751	5.444	1.348
H						
HETATM	142	O	0	7.889	3.402	0.166
O						
HETATM	143	H	0	7.697	2.468	0.497
H						
HETATM	144	O	0	7.989	0.507	0.836
O						
HETATM	145	H	0	8.177	0.077	1.704
H						
HETATM	146	O	0	7.482	-2.414	1.155
O						
HETATM	147	H	0	6.516	-2.654	1.376
H						
HETATM	148	O	0	5.490	-4.099	1.645
O						
HETATM	149	H	0	6.076	-4.880	1.540
H						
HETATM	150	O	0	2.786	-5.389	2.086
O						
HETATM	151	H	0	2.076	-4.668	2.026
H						
HETATM	152	O	0	-3.102	-5.412	-3.451
O						
HETATM	153	H	0	-2.600	-5.795	-2.686
H						
HETATM	154	O	0	-7.139	0.094	-3.433
O						
HETATM	155	H	0	-7.762	0.844	-3.291
H						
HETATM	156	O	0	-5.505	4.732	-1.814
O						
HETATM	157	H	0	-5.325	3.830	-2.193
H						
HETATM	158	O	0	-0.029	8.307	0.560
O						
HETATM	159	H	0	0.842	8.657	0.859
H						
HETATM	160	O	0	1.807	3.933	-2.277
O						

HETATM	161	H	0	1.814	4.310	-1.341
H						
HETATM	162	O	0	5.333	3.164	-2.476
O						
HETATM	163	O	0	0.594	-7.979	-1.496
O						
HETATM	164	H	0	-0.171	-8.544	-1.240
H						
HETATM	165	O	0	5.370	-3.931	-3.476
O						
HETATM	166	H	0	5.748	-3.017	-3.344
H						
HETATM	167	O	0	7.483	0.643	-3.998
O						
HETATM	168	H	0	8.101	-0.122	-4.061
H						
HETATM	169	C	0	2.511	-1.827	5.392
C						
HETATM	170	C	0	1.615	-2.295	4.423
C						
HETATM	171	C	0	0.511	-3.051	4.836
C						
HETATM	172	C	0	0.303	-3.315	6.182
C						
HETATM	173	C	0	1.191	-2.829	7.138
C						
HETATM	174	C	0	2.296	-2.083	6.740
C						
HETATM	175	H	0	3.379	-1.257	5.081
H						
HETATM	176	H	0	-0.171	-3.462	4.095
H						
HETATM	177	H	0	-0.554	-3.904	6.486
H						
HETATM	178	H	0	1.027	-3.038	8.189
H						
HETATM	179	H	0	2.997	-1.711	7.478
H						
HETATM	180	C	0	1.875	-1.998	3.012
C						
HETATM	181	H	0	2.900	-1.746	2.750
H						
HETATM	182	C	0	0.967	-1.981	2.025
C						
HETATM	183	H	0	-0.084	-2.154	2.236
H						
HETATM	184	C	0	1.360	-1.691	0.632
C						
HETATM	185	C	0	0.266	-1.461	-0.349
C						
HETATM	186	H	0	-0.755	-1.597	-0.005
H						

HETATM	187	C	0	0.557	-1.058	-1.592
C						
HETATM	188	H	0	1.612	-0.911	-1.807
H						
HETATM	189	C	0	-0.366	-0.773	-2.692
C						
HETATM	190	C	0	0.154	-0.258	-3.884
C						
HETATM	191	C	0	-1.747	-0.989	-2.611
C						
HETATM	192	C	0	-0.674	0.033	-4.963
C						
HETATM	193	H	0	1.220	-0.076	-3.961
H						
HETATM	194	C	0	-2.573	-0.711	-3.689
C						
HETATM	195	H	0	-2.185	-1.388	-1.699
H						
HETATM	196	C	0	-2.041	-0.196	-4.870
C						
HETATM	197	H	0	-0.253	0.447	-5.872
H						
HETATM	198	H	0	-3.644	-0.900	-3.608
H						
HETATM	199	H	0	-2.691	0.034	-5.706
H						
HETATM	200	O	0	2.535	-1.651	0.291
O						
HETATM	201	C	0	-2.786	3.571	-5.309
C						
HETATM	202	C	0	-1.419	3.771	-5.145
C						
HETATM	203	C	0	-0.804	3.413	-3.951
C						
HETATM	204	C	0	-1.540	2.850	-2.903
C						
HETATM	205	C	0	-2.916	2.665	-3.079
C						
HETATM	206	C	0	-3.532	3.021	-4.269
C						
HETATM	207	H	0	-3.269	3.846	-6.239
H						
HETATM	208	H	0	-0.832	4.205	-5.946
H						
HETATM	209	H	0	0.265	3.570	-3.814
H						
HETATM	210	H	0	-3.507	2.230	-2.279
H						
HETATM	211	H	0	-4.599	2.858	-4.394
H						
HETATM	212	C	0	-0.834	2.474	-1.677
C						

HETATM	213	H	0	0.246	2.634	-1.680
H						
HETATM	214	C	0	-1.360	1.955	-0.561
C						
HETATM	215	H	0	-2.423	1.745	-0.467
H						
HETATM	216	C	0	-0.510	1.656	0.611
C						
HETATM	217	C	0	-1.187	1.009	1.751
C						
HETATM	218	H	0	-2.229	0.731	1.622
H						
HETATM	219	C	0	-0.551	0.804	2.915
C						
HETATM	220	H	0	0.477	1.154	2.984
H						
HETATM	221	C	0	-1.111	0.186	4.117
C						
HETATM	222	C	0	-0.464	0.387	5.342
C						
HETATM	223	C	0	-2.274	-0.595	4.095
C						
HETATM	224	C	0	-0.985	-0.139	6.517
C						
HETATM	225	H	0	0.451	0.969	5.369
H						
HETATM	226	C	0	-2.789	-1.130	5.267
C						
HETATM	227	H	0	-2.774	-0.793	3.154
H						
HETATM	228	C	0	-2.151	-0.895	6.483
C						
HETATM	229	H	0	-0.474	0.030	7.456
H						
HETATM	230	H	0	-3.687	-1.737	5.233
H						
HETATM	231	H	0	-2.555	-1.315	7.397
H						
HETATM	232	O	0	0.679	1.949	0.644
O						
END						
CONECT	1	2	3	15	29	
CONECT	2	1				
CONECT	3	1	4	5	134	
CONECT	4	3				
CONECT	5	3	6	7	132	
CONECT	6	5				
CONECT	7	5	8	9	14	
CONECT	8	7				
CONECT	9	7	10	11	15	
CONECT	10	9				
CONECT	11	9	12	13	156	

CONNECT	12	11			
CONNECT	13	11			
CONNECT	14	7	75		
CONNECT	15	1	9		
CONNECT	16	17	18	30	44
CONNECT	17	16			
CONNECT	18	16	19	20	138
CONNECT	19	18			
CONNECT	20	18	21	22	136
CONNECT	21	20			
CONNECT	22	20	23	24	29
CONNECT	23	22			
CONNECT	24	22	25	26	30
CONNECT	25	24			
CONNECT	26	24	27	28	158
CONNECT	27	26			
CONNECT	28	26			
CONNECT	29	1	22		
CONNECT	30	16	24		
CONNECT	31	32	33	58	162
CONNECT	32	31			
CONNECT	33	31	34	35	142
CONNECT	34	33			
CONNECT	35	33	36	37	140
CONNECT	36	35			
CONNECT	37	35	38	39	44
CONNECT	38	37			
CONNECT	39	37	40	41	162
CONNECT	40	39			
CONNECT	41	39	42	43	160
CONNECT	42	41			
CONNECT	43	41			
CONNECT	44	16	37		
CONNECT	45	46	47	59	103
CONNECT	46	45			
CONNECT	47	45	48	49	146
CONNECT	48	47			
CONNECT	49	47	50	51	144
CONNECT	50	49			
CONNECT	51	49	52	53	58
CONNECT	52	51			
CONNECT	53	51	54	55	59
CONNECT	54	53			
CONNECT	55	53	56	57	167
CONNECT	56	55			
CONNECT	57	55			
CONNECT	58	31	51		
CONNECT	59	45	53		
CONNECT	60	61	62	74	88
CONNECT	61	60			
CONNECT	62	60	63	64	120
CONNECT	63	62			

CONNECT	64	62	65	66	122
CONNECT	65	64			
CONNECT	66	64	67	68	73
CONNECT	67	66			
CONNECT	68	66	69	70	74
CONNECT	69	68			
CONNECT	70	68	71	72	152
CONNECT	71	70			
CONNECT	72	70			
CONNECT	73	66	105		
CONNECT	74	60	68		
CONNECT	75	14	76	77	89
CONNECT	76	75			
CONNECT	77	75	78	79	128
CONNECT	78	77			
CONNECT	79	77	80	81	130
CONNECT	80	79			
CONNECT	81	79	82	83	88
CONNECT	82	81			
CONNECT	83	81	84	85	89
CONNECT	84	83			
CONNECT	85	83	86	87	154
CONNECT	86	85			
CONNECT	87	85			
CONNECT	88	60	81		
CONNECT	89	75	83		
CONNECT	90	91	92	104	118
CONNECT	91	90			
CONNECT	92	90	93	94	150
CONNECT	93	92			
CONNECT	94	92	95	96	148
CONNECT	95	94			
CONNECT	96	94	97	98	103
CONNECT	97	96			
CONNECT	98	96	99	100	104
CONNECT	99	98			
CONNECT	100	98	101	102	165
CONNECT	101	100			
CONNECT	102	100			
CONNECT	103	45	96		
CONNECT	104	90	98		
CONNECT	105	73	106	107	119
CONNECT	106	105			
CONNECT	107	105	108	109	124
CONNECT	108	107			
CONNECT	109	107	110	111	126
CONNECT	110	109			
CONNECT	111	109	112	113	118
CONNECT	112	111			
CONNECT	113	111	114	115	119
CONNECT	114	113			
CONNECT	115	113	116	117	163

CONNECT	116	115	
CONNECT	117	115	
CONNECT	118	90	111
CONNECT	119	105	113
CONNECT	120	62	121
CONNECT	121	120	
CONNECT	122	64	123
CONNECT	123	122	
CONNECT	124	107	125
CONNECT	125	124	
CONNECT	126	109	127
CONNECT	127	126	
CONNECT	128	77	129
CONNECT	129	128	
CONNECT	130	79	131
CONNECT	131	130	
CONNECT	132	5	133
CONNECT	133	132	
CONNECT	134	3	135
CONNECT	135	134	
CONNECT	136	20	137
CONNECT	137	136	
CONNECT	138	18	139
CONNECT	139	138	
CONNECT	140	35	141
CONNECT	141	140	
CONNECT	142	33	143
CONNECT	143	142	
CONNECT	144	49	145
CONNECT	145	144	
CONNECT	146	47	147
CONNECT	147	146	
CONNECT	148	94	149
CONNECT	149	148	
CONNECT	150	92	151
CONNECT	151	150	
CONNECT	152	70	153
CONNECT	153	152	
CONNECT	154	85	155
CONNECT	155	154	
CONNECT	156	11	157
CONNECT	157	156	
CONNECT	158	26	159
CONNECT	159	158	
CONNECT	160	41	161
CONNECT	161	160	
CONNECT	162	31	39
CONNECT	163	115	164
CONNECT	164	163	
CONNECT	165	100	166
CONNECT	166	165	
CONNECT	167	55	168

CONNECT	168	167		
CONNECT	169	170	174	175
CONNECT	170	169	171	180
CONNECT	171	170	172	176
CONNECT	172	171	173	177
CONNECT	173	172	174	178
CONNECT	174	169	173	179
CONNECT	175	169		
CONNECT	176	171		
CONNECT	177	172		
CONNECT	178	173		
CONNECT	179	174		
CONNECT	180	170	181	182
CONNECT	181	180		
CONNECT	182	180	183	184
CONNECT	183	182		
CONNECT	184	182	185	200
CONNECT	185	184	186	187
CONNECT	186	185		
CONNECT	187	185	188	189
CONNECT	188	187		
CONNECT	189	187	190	191
CONNECT	190	189	192	193
CONNECT	191	189	194	195
CONNECT	192	190	196	197
CONNECT	193	190		
CONNECT	194	191	196	198
CONNECT	195	191		
CONNECT	196	192	194	199
CONNECT	197	192		
CONNECT	198	194		
CONNECT	199	196		
CONNECT	200	184		
CONNECT	201	202	206	207
CONNECT	202	201	203	208
CONNECT	203	202	204	209
CONNECT	204	203	205	212
CONNECT	205	204	206	210
CONNECT	206	201	205	211
CONNECT	207	201		
CONNECT	208	202		
CONNECT	209	203		
CONNECT	210	205		
CONNECT	211	206		
CONNECT	212	204	213	214
CONNECT	213	212		
CONNECT	214	212	215	216
CONNECT	215	214		
CONNECT	216	214	217	232
CONNECT	217	216	218	219
CONNECT	218	217		
CONNECT	219	217	220	221

CONNECT	220	219		
CONNECT	221	219	222	223
CONNECT	222	221	224	225
CONNECT	223	221	226	227
CONNECT	224	222	228	229
CONNECT	225	222		
CONNECT	226	223	228	230
CONNECT	227	223		
CONNECT	228	224	226	231
CONNECT	229	224		
CONNECT	230	226		
CONNECT	231	228		
CONNECT	232	216		

Frequency calculation of optimized structure showed that there were no negative frequencies

13. Isothermal titration calorimetry

Isothermal titration experiments were performed on a TA nano ITC instrument with titration cell and reference cell volumes of 1000 μL , and a 100 μL syringe used to inject the titrant. Stock solutions (10^{-2}M) of γ -cyclodextrin in water and 4-methoxy chalcone **1b** in methanol were prepared, which were further diluted with water. The host concentration was diluted to $5 \times 10^{-4}\text{ M}$, and that of guest was diluted to $5 \times 10^{-5}\text{ M}$. The solutions were degassed for 15 mins before titration ITC experiments were performed with the cell temperature set to be maintained at $25.0\text{ }^{\circ}\text{C}$ with each injection volume set at $4.81\text{ }\mu\text{L}$ for nineteen injections. The stirring rate during injection rate was set at 350 rpm. Blank heat from titration of $5 \times 10^{-4}\text{ M}$ host was subtracted from the host-guest complex titration before isotherm fitting algorithm. The independent binding model available in nanoAnalyze software issued along with the VP nano ITC instrument package was used to use for curve fitting.

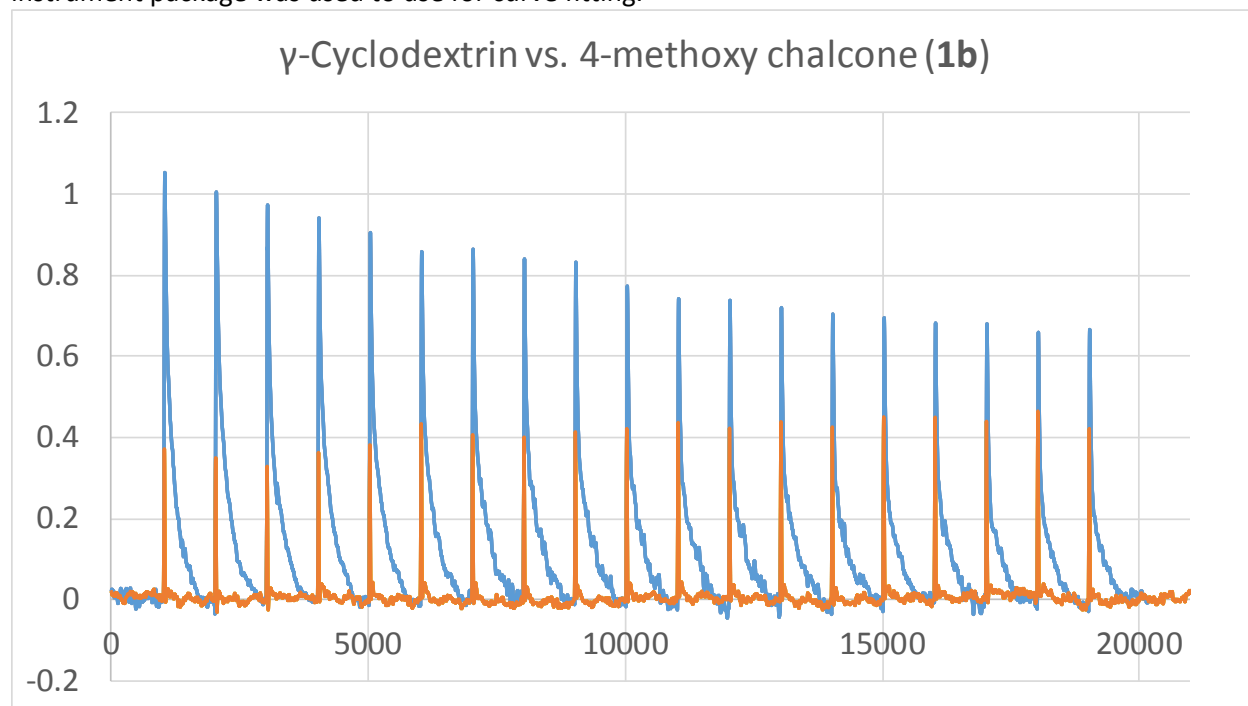


Figure S18 ITC data for titration of $5 \times 10^{-4}\text{ M}$ γ -CD to $5 \times 10^{-5}\text{ M}$ **1b** (blue), and the background titration of $5 \times 10^{-4}\text{ M}$ γ -CD to water (orange).

