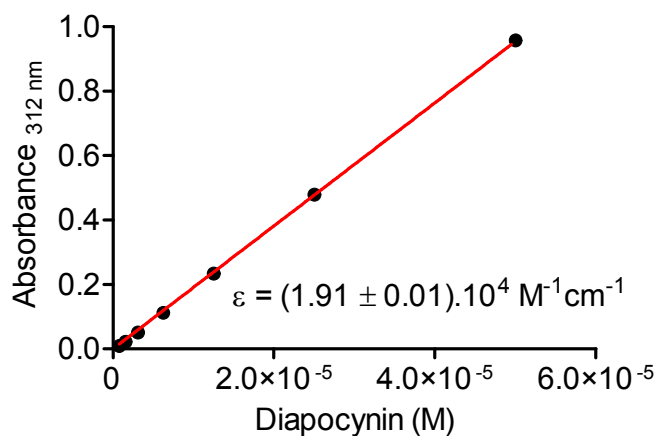
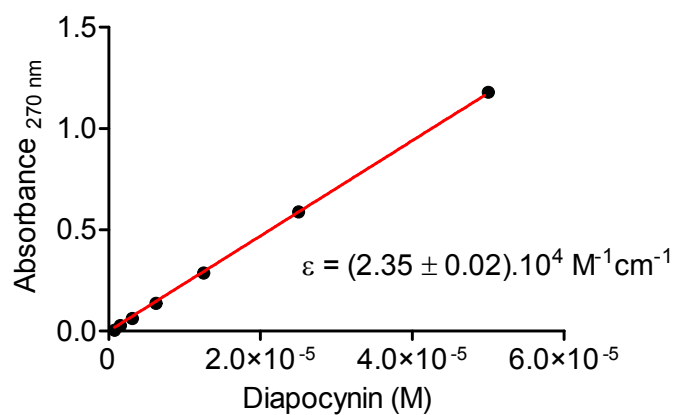
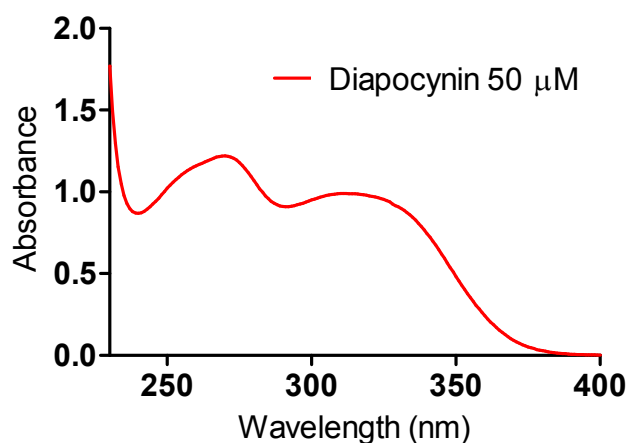


## Supporting Information

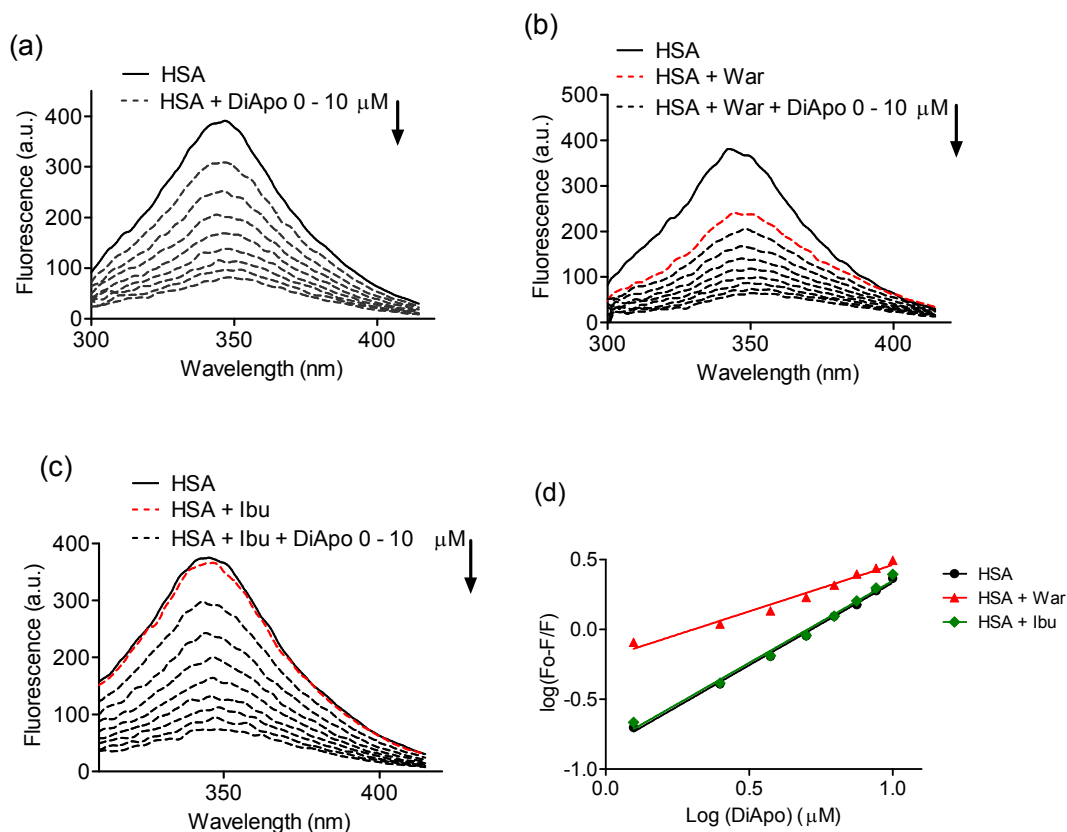
### Molar Absorption Coefficient of Diapocynin

Absorption spectrum and determination of molar absorption coefficient of diapocynin  
in 0.05 mM phosphate buffer, pH 7.0



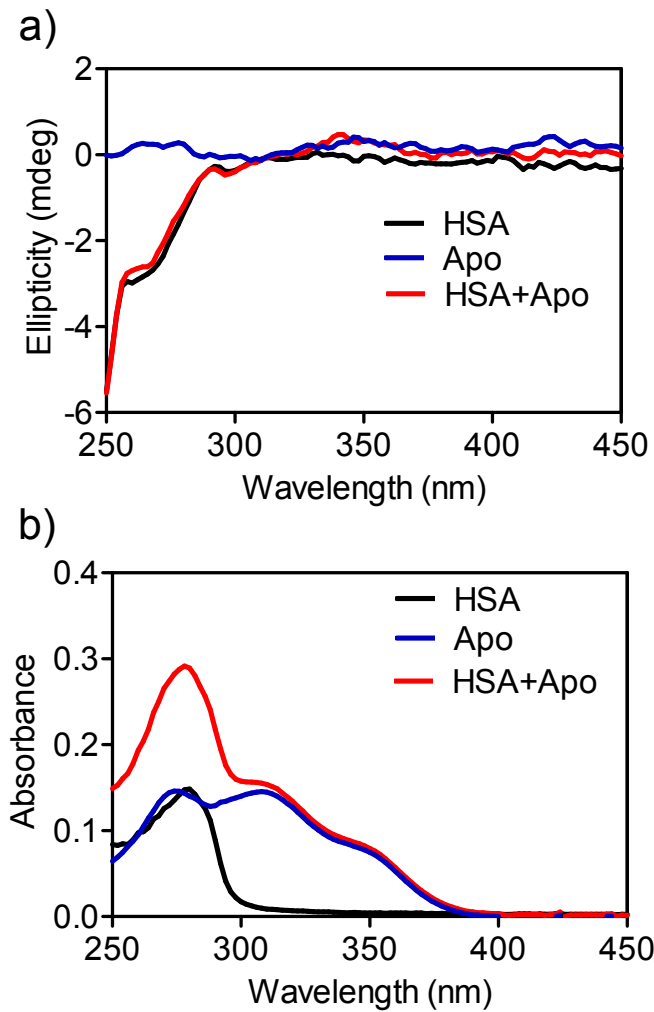
## Number of Binding Sites

Quenching of the intrinsic fluorescence of HSA ( $5 \mu\text{M}$ ;  $\lambda_{\text{ex}} = 280 \text{ nm}$ ) by diapocynin in the absence **(a)** or presence of  $5 \mu\text{M}$  warfarin **(b)** or  $5 \mu\text{M}$  ibuprofen **(c)**. The experiments were performed in  $0.05 \text{ M}$  phosphate buffer,  $\text{pH } 7.0$  at  $25^\circ\text{C}$ . The determinations of the binding constants **(d)** were performed by fitting the data in the double logarithm equation:  $\log [(F_0-F)/F] = \log K_a + n \cdot \log [Q]$ . The results are the average and SD of experiments performed in triplicate. The binding constants ( $K_a$ ) and stoichiometry of the binding ( $n$ ) were: warfarin + diapocynin ( $6.27 \times 10^5 \text{ mol}^{-1} \cdot \text{L}$ ,  $n = 0.66$ ); ibuprofen + diapocynin ( $1.49 \times 10^5 \text{ mol}^{-1} \cdot \text{L}$ ,  $n = 1.18$ ); only diapocynin ( $1.42 \times 10^5 \text{ mol}^{-1} \cdot \text{L}$ ,  $n = 1.17$ ).



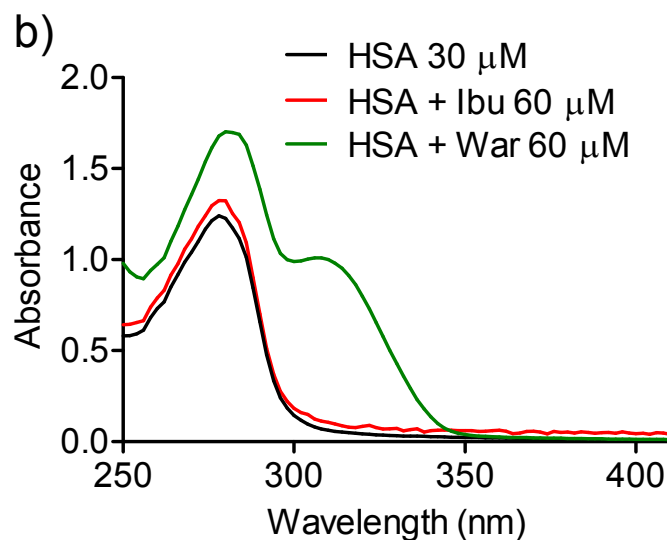
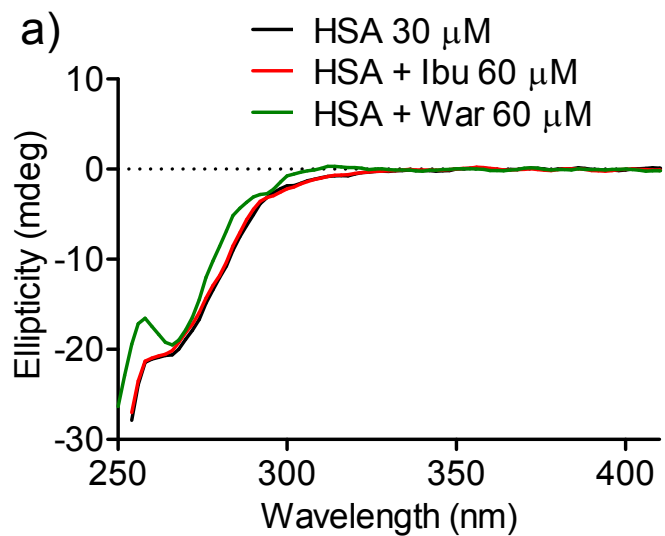
### Absence of Induced Ellipticity for Apocynin

The binding of apocynin with HSA did not provoke the generation of an induced circular dichroism signal. **(a)** Near-UV-CD and **(b)** UV-Vis spectra of HSA in the presence or absence of apocynin (Apo). The mixtures consisted of 30  $\mu\text{M}$  HSA and 30  $\mu\text{M}$  apocynin in 0.05 M phosphate buffer, pH 7.0 at 25°C.



### Absence of Induced Ellipticity for Warfarin and Ibuprofen

The binding of warfarin and ibuprofen with HSA did not provoke the generation of an induced circular dichroism signal. **(a)** Near-UV-CD and **(b)** UV-Vis spectra of HSA in the presence or absence of warfarin (War) or ibuprofen (Ibu). The mixtures consisted of 30  $\mu\text{M}$  HSA and 30  $\mu\text{M}$  ligands in 0.05 M phosphate buffer, pH 7.0 at 25°C.



**Cartesian coordinates of the *syn-aS* conformer.**

<b>Atom</b>	<b>X</b>	<b>Y</b>	<b>Z</b>
C	-1.462416	-1.002983	0.519662
C	-0.737174	0.057761	-0.019818
C	-1.432232	1.105077	-0.623767
C	-2.837561	1.093448	-0.655056
C	-3.546329	0.041448	-0.105855
C	-2.851739	-1.024949	0.483748
C	0.745291	0.064166	0.018713
C	1.432256	1.116597	0.615099
C	2.840900	1.111860	0.650715
C	3.553205	0.063916	0.110211
C	2.866501	-1.008393	-0.477664
C	1.478385	-0.998798	-0.516610
O	0.757635	2.128366	1.191998
O	3.371506	2.207492	1.267085
C	4.782402	2.274405	1.370918
C	3.678943	-2.124831	-1.048875
O	4.888752	-2.094151	-1.004063
O	-0.761982	2.114530	-1.208638
O	-3.377269	2.183854	-1.276817
C	-4.786541	2.260883	-1.360287
C	-3.562312	-2.192616	1.089107
C	-5.076985	-2.196106	1.086398
C	2.955273	-3.292262	-1.682752
O	-2.944876	-3.115064	1.570283
H	5.005094	3.204507	1.888210
H	5.167678	1.428751	1.947300
H	5.243527	2.283319	0.379387
H	1.402589	2.755780	1.542421
H	3.693904	-4.007293	-2.037852
H	2.294161	-3.773522	-0.958382
H	2.340404	-2.952891	-2.519652
H	0.931016	-1.812180	-0.978514
H	4.634798	0.030866	0.124752
H	-4.627458	0.040345	-0.136773
H	-0.942738	-1.825541	0.997449
H	-1.406618	2.744806	-1.553860
H	-5.011056	3.189163	-1.880171
H	-5.189376	1.416558	-1.927662
H	-5.234246	2.282644	-0.362229
H	-5.459126	-2.164914	0.063078
H	-5.421972	-3.104280	1.575279
H	-5.465065	-1.323227	1.616744

Cartesian coordinates of the *anti-aS* conformer.

Atom	X	Y	Z
C	1.560067	-0.947025	0.447803
C	0.732243	0.068503	-0.035084
C	1.328594	1.238376	-0.505713
C	2.727449	1.379246	-0.493663
C	3.533650	0.366109	-0.013276
C	2.941487	-0.811553	0.466928
C	-0.744753	-0.077505	-0.037755
C	-1.338717	-1.237507	-0.527474
C	-2.741318	-1.382357	-0.510266
C	-3.544481	-0.381769	-0.012065
C	-2.954085	0.791887	0.480698
C	-1.574547	0.932605	0.462861
O	-0.583709	-2.232067	-1.038024
O	-3.169096	-2.570128	-1.028460
C	-4.566517	-2.798315	-1.059395
C	-3.859726	1.856911	1.007502
O	-5.060718	1.701131	1.011818
O	0.573160	2.249252	-0.991460
O	3.158233	2.575944	-0.995529
C	4.554114	2.789719	-1.070798
C	3.760945	-1.939831	1.009350
C	5.269493	-1.800630	1.006295
C	-3.240830	3.133950	1.531787
O	3.235303	-2.938470	1.443132
H	-4.703861	-3.781743	-1.502529
H	-4.982625	-2.786482	-0.048251
H	-5.067818	-2.043256	-1.671332
H	-1.175943	-2.936666	-1.330607
H	-4.038155	3.795492	1.862820
H	-2.570417	2.918763	2.367060
H	-2.653260	3.622675	0.751152
H	-1.109046	1.834591	0.837757
H	-4.623399	-0.462362	0.013571
H	4.608902	0.482685	-0.007523
H	1.127121	-1.867239	0.818548
H	1.169781	2.941024	-1.304947
H	4.688457	3.777010	-1.506403
H	5.005340	2.764664	-0.074464
H	5.027680	2.038338	-1.709666
H	5.575561	-0.946658	1.615886
H	5.702067	-2.711506	1.413755
H	5.641023	-1.639795	-0.008523

Cartesian coordinates of the *anti-aR* conformer.

Atom	X	Y	Z
C	1.560083	-0.947000	-0.447786
C	0.732241	0.068524	0.035080
C	1.328571	1.238417	0.505683
C	2.727424	1.379311	0.493630
C	3.533643	0.366179	0.013265
C	2.941500	-0.811504	-0.466915
C	-0.744752	-0.077511	0.037755
C	-1.338694	-1.237514	0.527499
C	-2.741292	-1.382390	0.510295
C	-3.544474	-0.381827	0.012074
C	-2.954100	0.791829	-0.480714
C	-1.574565	0.932574	-0.462882
O	-0.583666	-2.232049	1.038067
O	-3.169047	-2.570158	1.028515
C	-4.566464	-2.798371	1.059456
C	-3.859762	1.856826	-1.007539
O	-5.060750	1.701023	-1.011849
O	0.573120	2.249290	0.991410
O	3.158186	2.576028	0.995471
C	4.554063	2.789833	1.070730
C	3.760977	-1.939778	-1.009313
C	5.269523	-1.800553	-1.006258
C	-3.240890	3.133864	-1.531853
O	3.235352	-2.938434	-1.443079
H	-4.703789	-3.781792	1.502610
H	-5.067778	-2.043308	1.671377
H	-4.982573	-2.786567	0.048312
H	-1.175888	-2.936653	1.330666
H	-4.038227	3.795383	-1.862901
H	-2.653331	3.622618	-0.751226
H	-2.570471	2.918672	-2.367118
H	-1.109081	1.834560	-0.837796
H	-4.623390	-0.462441	-0.013559
H	4.608893	0.482774	0.007508
H	1.127152	-1.867229	-0.818511
H	1.169730	2.941079	1.304883
H	4.688388	3.777135	1.506312
H	5.027647	2.038475	1.709613
H	5.005286	2.764764	0.074395
H	5.575578	-0.946591	-1.615868
H	5.641047	-1.639688	0.008558
H	5.702113	-2.711432	-1.413696

Cartesian coordinates of the *syn-aR* conformer.

Atom	X	Y	Z
C	-1.462401	-1.002909	-0.519896
C	-0.737153	0.057842	0.019551
C	-1.432215	1.105225	0.623365
C	-2.837542	1.093646	0.654551
C	-3.546312	0.041618	0.105428
C	-2.851729	-1.024840	-0.484024
C	0.745306	0.064201	-0.018813
C	1.432340	1.116614	-0.615151
C	2.840966	1.111871	-0.650610
C	3.553184	0.063917	-0.110001
C	2.866409	-1.008383	0.477803
C	1.478299	-0.998767	0.516605
O	0.757783	2.128410	-1.192145
O	3.371609	2.207494	-1.266963
C	4.782495	2.274286	-1.370994
C	3.678816	-2.124774	1.049132
O	4.888632	-2.093988	1.004651
O	-0.761987	2.114758	1.208187
O	-3.377253	2.184145	1.276135
C	-4.786492	2.260499	1.360721
C	-3.562268	-2.192539	-1.089331
C	-5.076924	-2.196164	-1.086170
C	2.955022	-3.292254	1.682746
O	-2.944851	-3.114837	-1.570824
H	5.005173	3.204308	-1.888430
H	5.243766	2.283280	-0.379529
H	5.167620	1.428520	-1.947319
H	1.402814	2.755809	-1.542500
H	3.693533	-4.007463	2.037751
H	2.340216	-2.952920	2.519714
H	2.293840	-3.773271	0.958281
H	0.930846	-1.812120	0.978469
H	4.634776	0.030861	-0.124426
H	-4.627449	0.040579	0.136264
H	-0.942713	-1.825506	-0.997601
H	-1.406688	2.745004	1.553388
H	-5.011025	3.188587	1.880937
H	-5.234999	2.282224	0.363019
H	-5.188455	1.415855	1.928247
H	-5.458835	-2.164166	-0.062794
H	-5.465095	-1.323680	-1.617121
H	-5.422037	-3.104684	-1.574324



### Cartesian coordinates of the Transition State (TS).

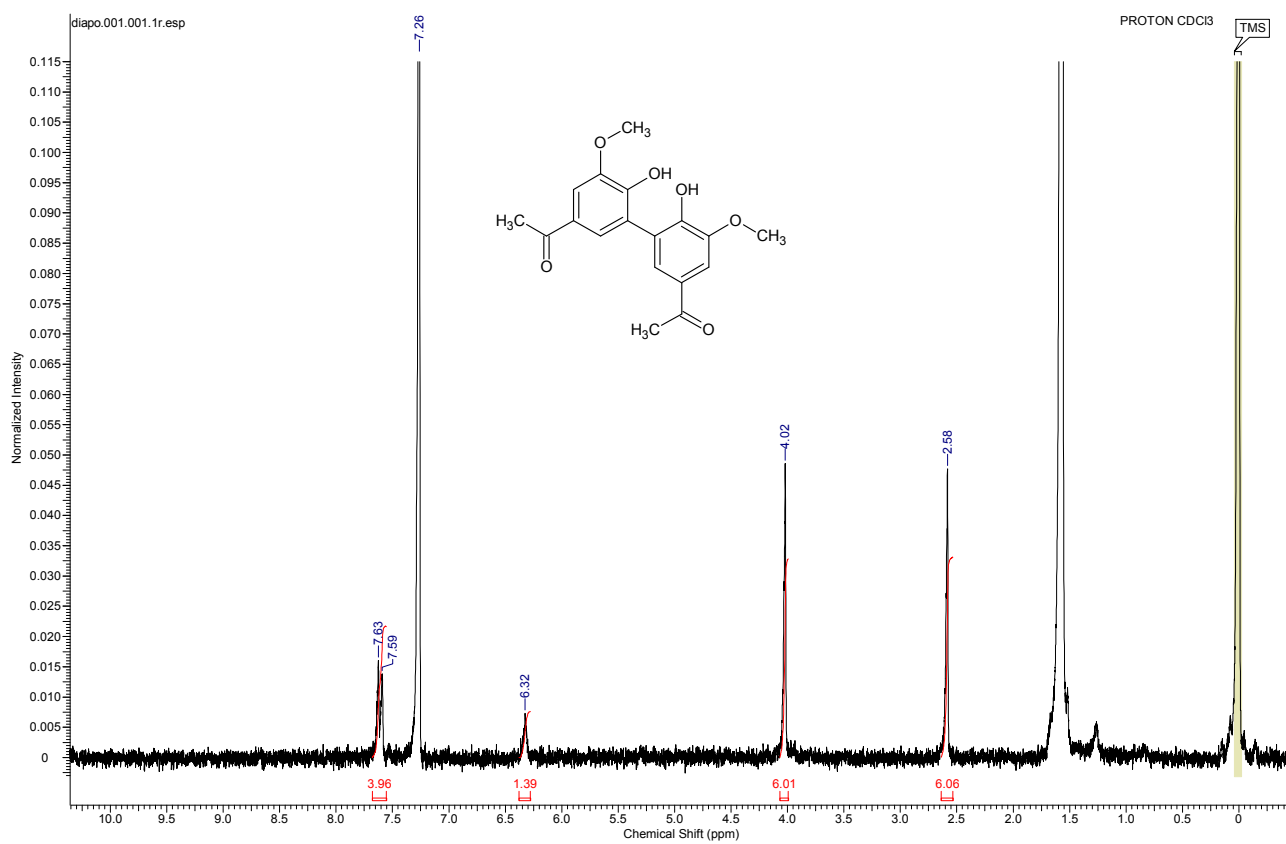
Atom	X	Y	Z
C	-1.529733	-0.747877	-0.789299
C	-0.742159	0.051382	0.042711
C	-1.381117	0.970742	0.883960
C	-2.789262	1.080729	0.879538
C	-3.557279	0.281449	0.047178
C	-2.925474	-0.647889	-0.803380
C	0.747754	-0.056814	0.041144
C	1.386915	-1.007095	0.843054
C	2.797490	-1.119343	0.829375
C	3.560203	-0.289631	0.030163
C	2.929000	0.676313	-0.778861
C	1.534746	0.779049	-0.761358
O	0.660400	-1.830062	1.638471
O	3.268489	-2.101773	1.666702
C	4.674753	-2.300872	1.733126
C	3.799018	1.547128	-1.620161
O	5.017072	1.421094	-1.602990
O	-0.651925	1.764849	1.706460
O	-3.260452	2.028656	1.758525
C	-4.664597	2.211205	1.860598
C	-3.690428	-1.535996	-1.726853
C	-5.211160	-1.449594	-1.738939
C	3.150380	2.601517	-2.504776
O	-3.115719	-2.323239	-2.465984
H	4.830955	-3.115357	2.442322
H	5.080424	-2.582574	0.753732
H	5.184979	-1.398129	2.090845
H	1.295910	-2.406183	2.101138
H	3.935982	3.136703	-3.040898
H	2.464980	2.144057	-3.228076
H	2.568204	3.313935	-1.908329
H	1.029881	1.515601	-1.377330
H	4.641630	-0.349136	-0.003343
H	-4.636651	0.373544	0.051735
H	-1.061478	-1.471169	-1.448034
H	-1.285210	2.311052	2.206460
H	-4.815076	2.990470	2.609528
H	-5.093443	2.536757	0.904170
H	-5.161980	1.289093	2.187313
H	-5.549689	-0.442967	-2.013065
H	-5.593387	-2.166838	-2.467177
H	-5.629669	-1.680028	-0.751683

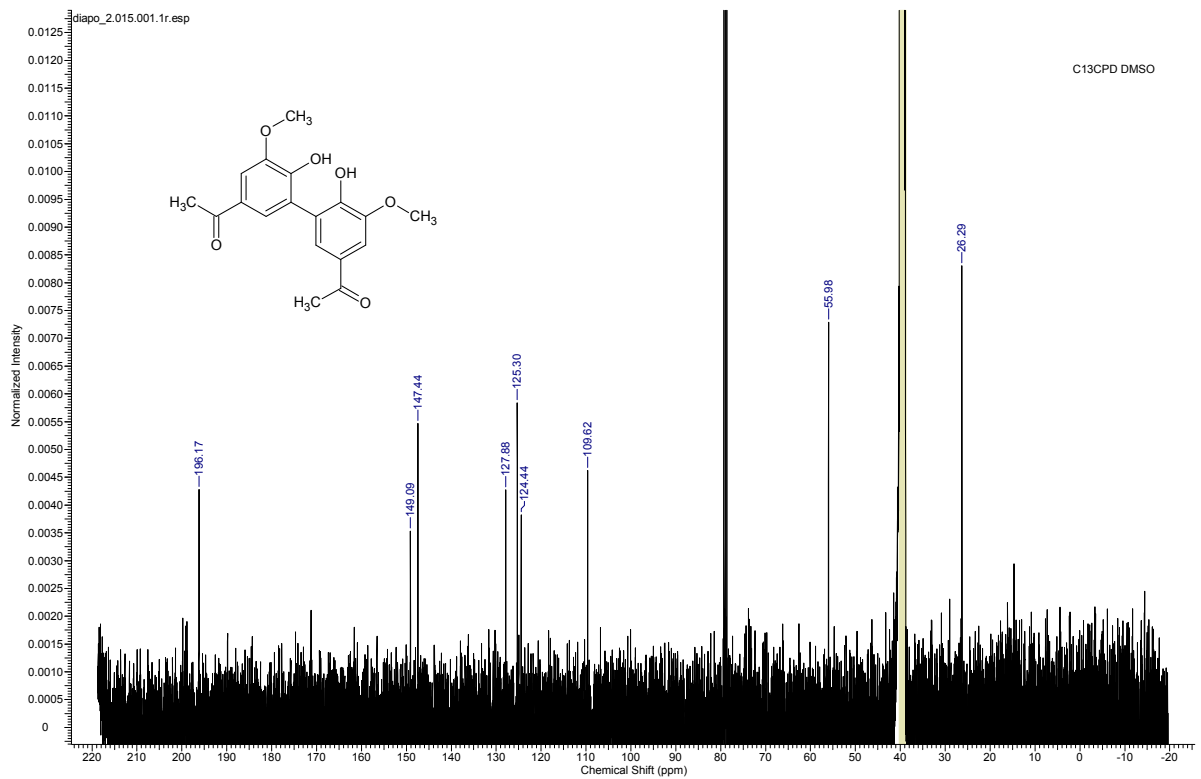
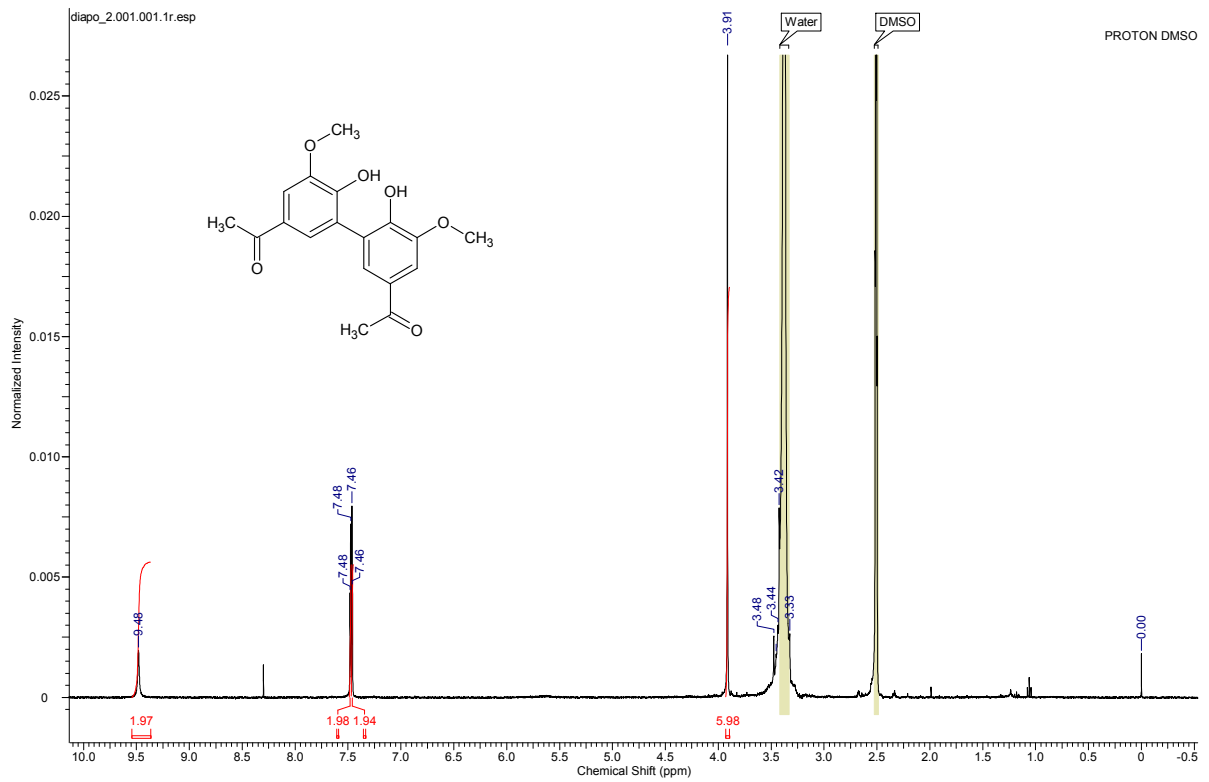
Thermodynamic parameters (atomic unit) to the four conformers (*syn-aS*, *anti-aS*, *anti-aR* and *syn-aR*) and to the transition state (TS).

Conformer	ZPE	Thermal Correction to		
		Energy	Enthalpy	Gibbs Free Energy
<i>syn-aS</i>	0.3329	0.3564	0.3573	0.2791
<i>anti-aS</i>	0.3335	0.3568	0.3578	0.2804
<i>anti-aR</i>	0.3335	0.3568	0.3578	0.2804
<i>syn-aR</i>	0.3329	0.3564	0.3573	0.2791
TS <sup>(a)</sup>	0.3309	0.3538	0.3547	0.2786

<sup>(a)</sup>  $\nu_1 = -18.51 \text{ cm}^{-1}$ . The normal mode of vibration associated with this frequency corresponds to the torsional movement of the dihedral angle. This angle appears between the phenyl rings.

# NMR spectra of diapocynin (CDCl<sub>3</sub> and DMSO-D<sub>6</sub> solutions)





# MS and MS/MS spectra of diapocynin

