

Supplementary Information

Computational Analysis and Biological Activities of Oxyresveratrol Analogues, the Putative Cyclooxygenase-2 Inhibitors

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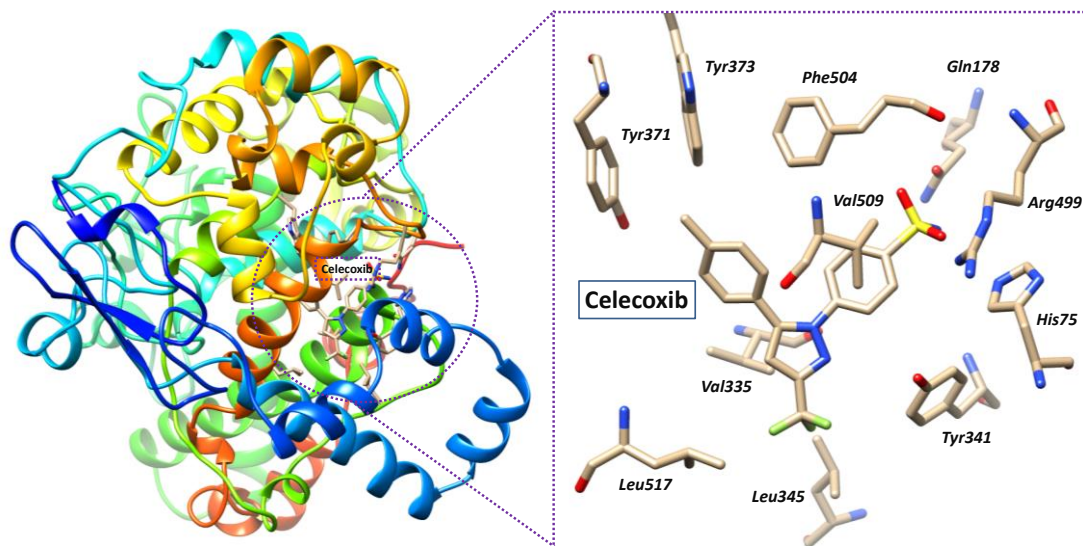
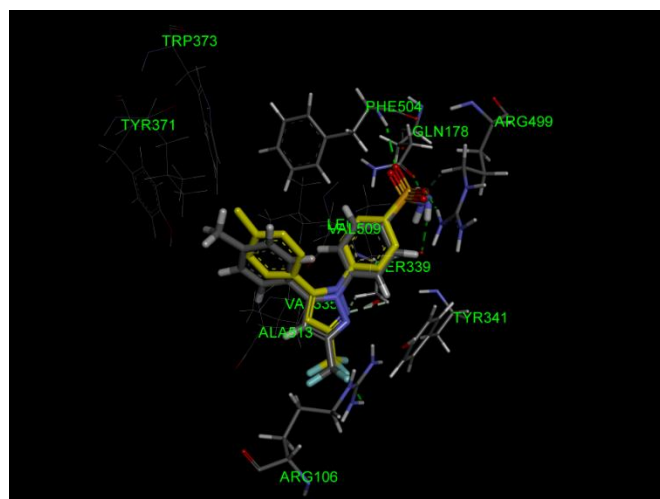
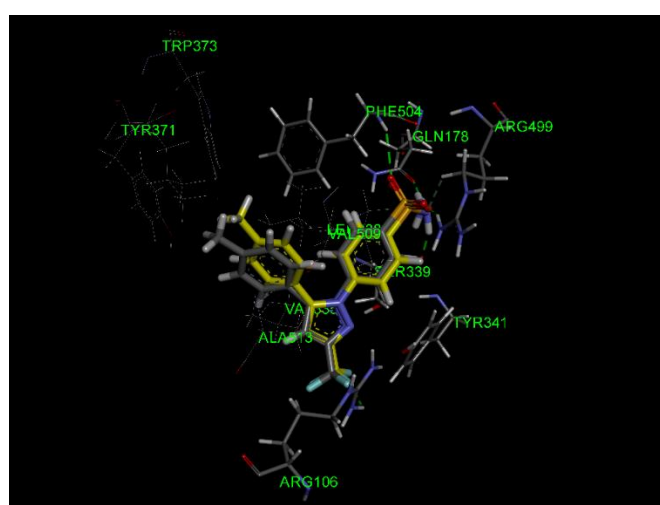


Figure S1. X-ray crystallographic pose of celecoxib bound to COX-2 receptor (PDB entry: 3LN1).

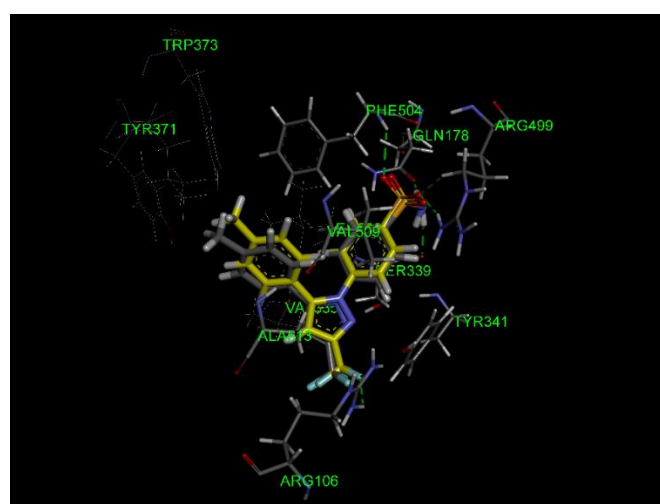
Right panel shows the co-crystallized celecoxib and key residues of amino acids at the catalytic site of COX-2.



(a)



(b)



(c)

Figure S2. The superposition of the one ns MD structure (grey) and the best-docked poses (yellow) of celecoxib from (a) AutoDock with the RMSD value of 0.68 Å, (b) ChemPLP with the RMSD value of 0.55 Å and (c) GoldScore with the RMSD value of 0.56 Å was within 6 Å of COX-2 binding pocket. Only the amino acid residues in proximity to the binding pocket are highlighted for clarity.

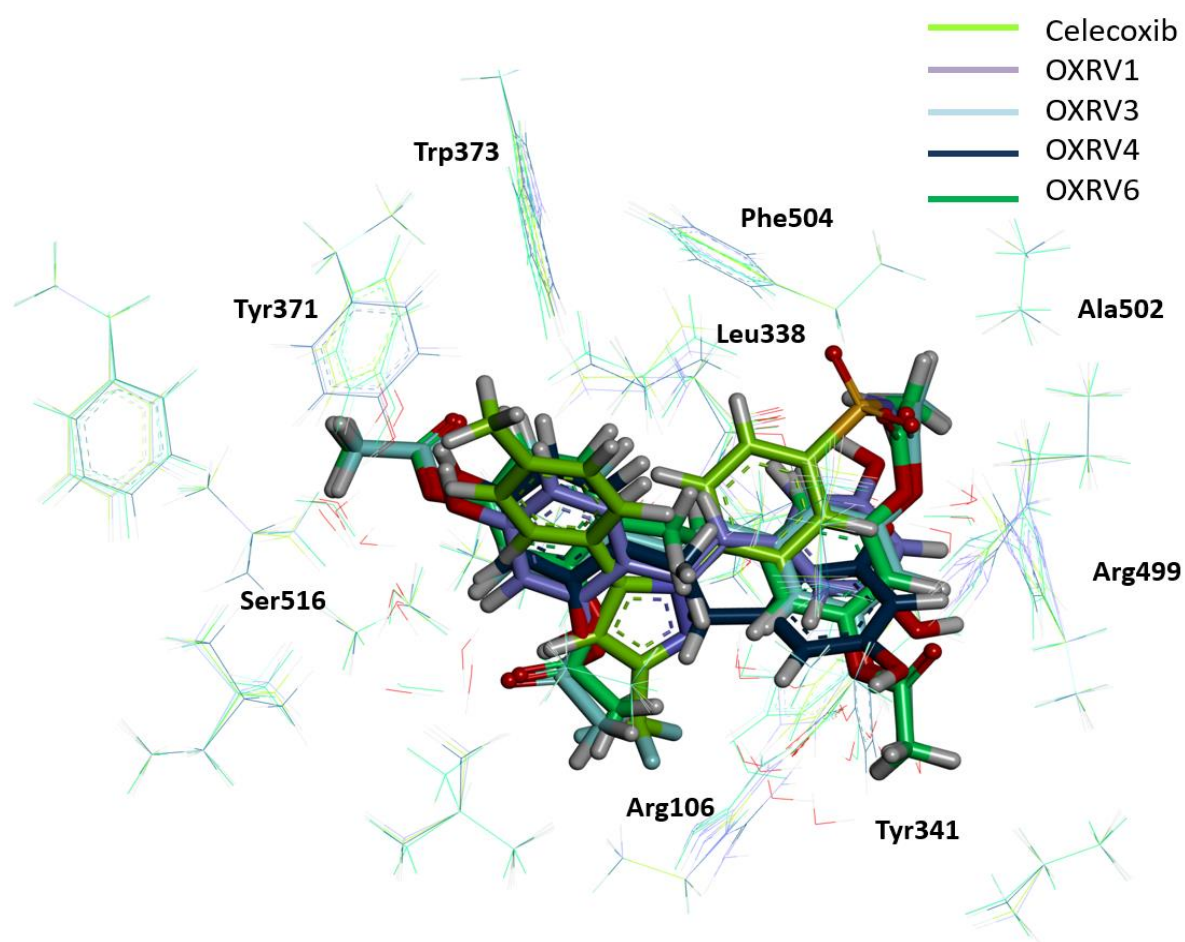
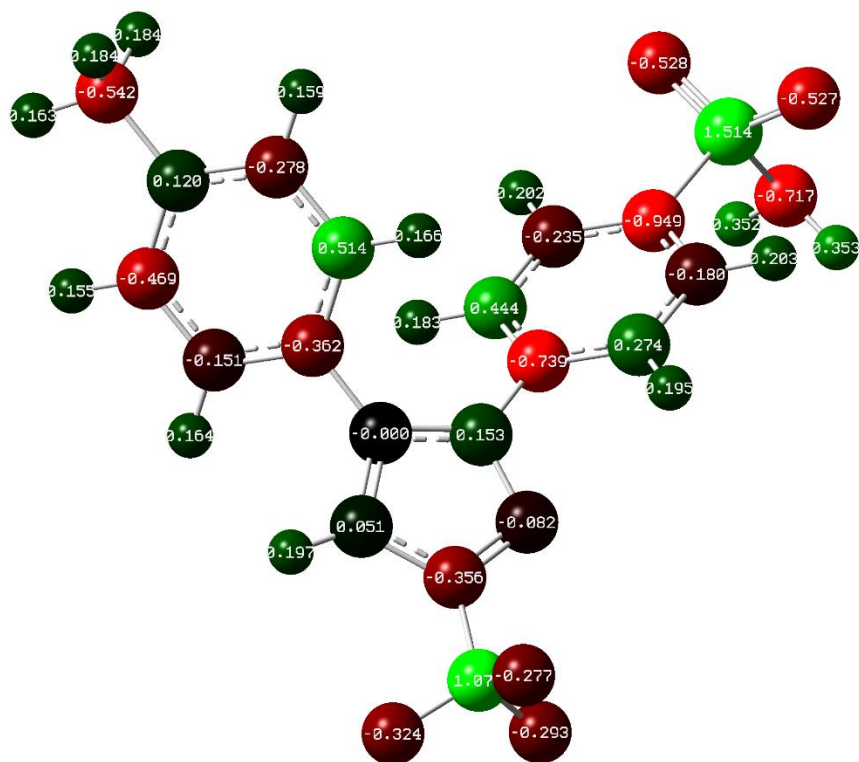
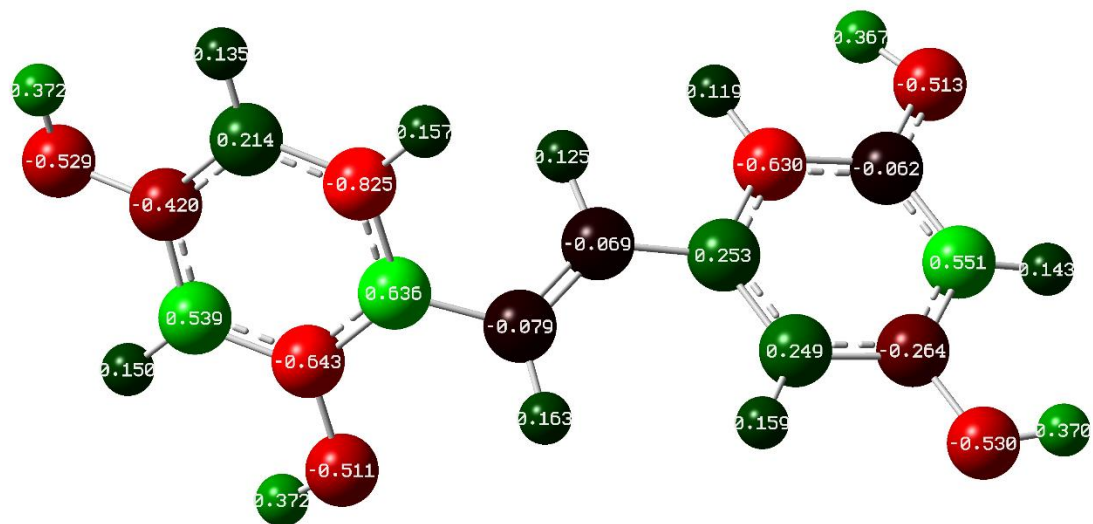


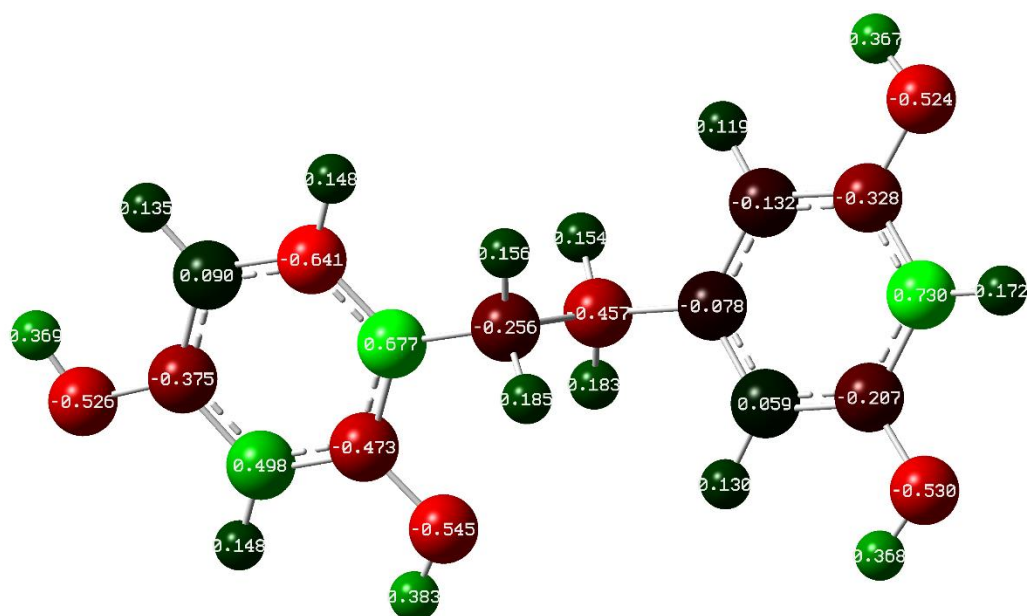
Figure S3. Overlay of the lowest-QM-energy poses (stick representation) of celecoxib and oxyresveratrol derivatives in catalytic pocket of COX-2.



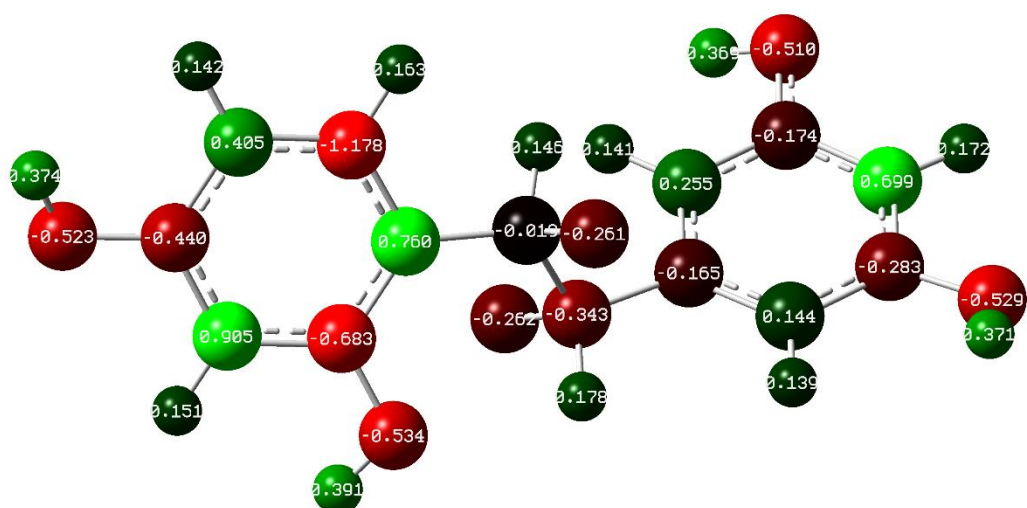
(a)

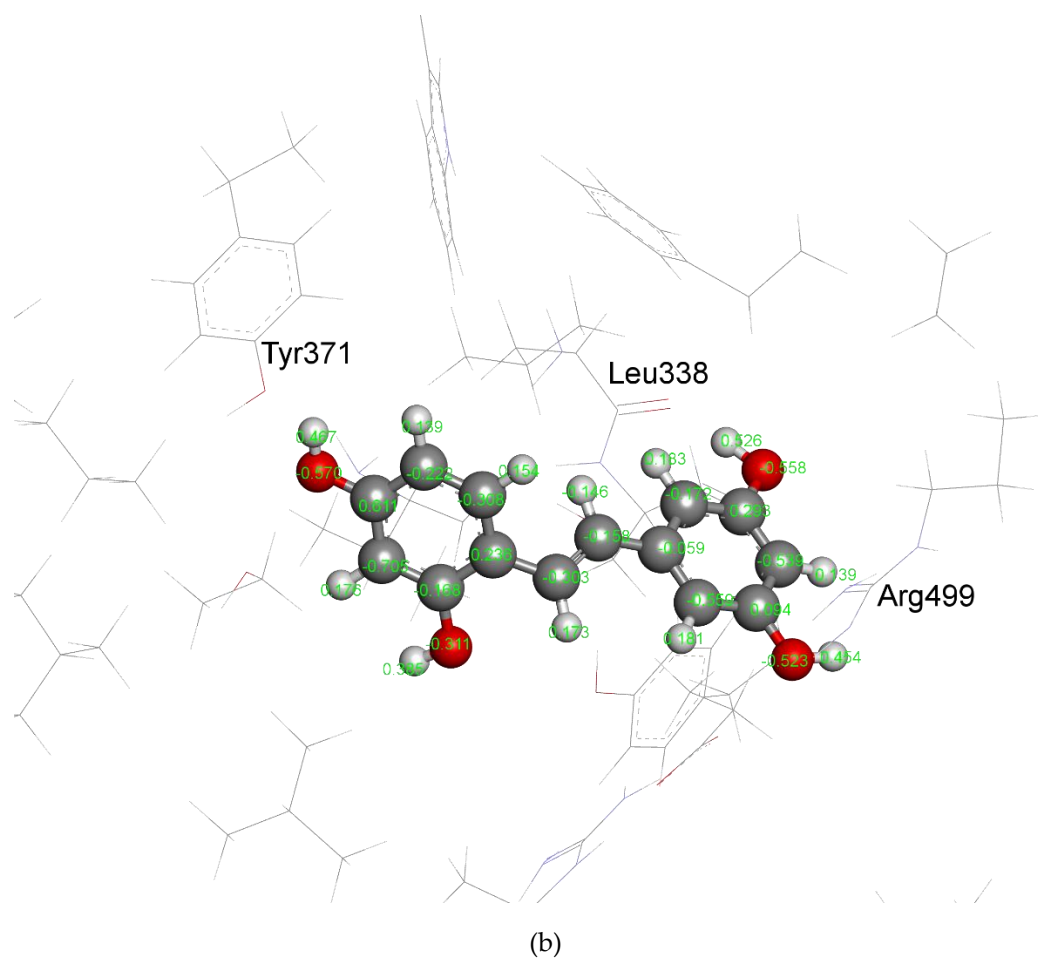
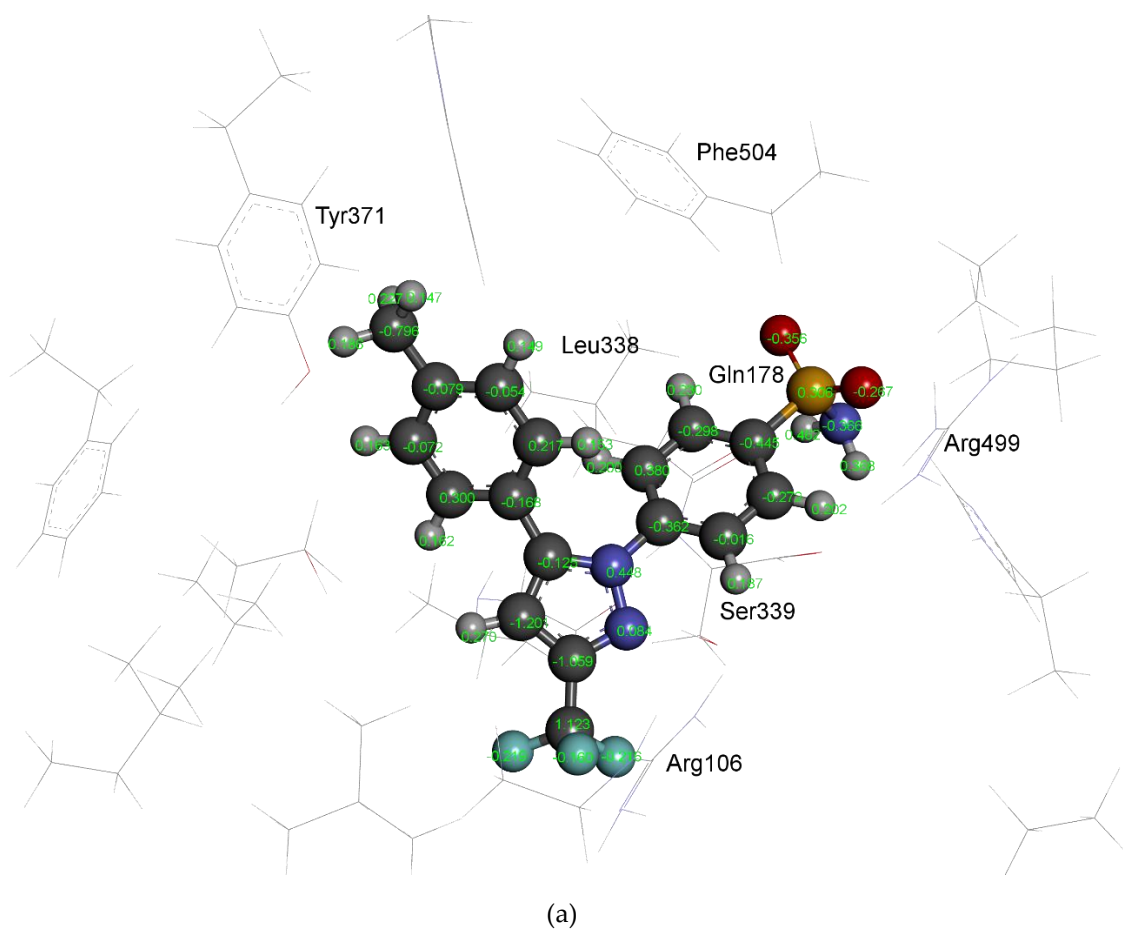


(b)



(c)





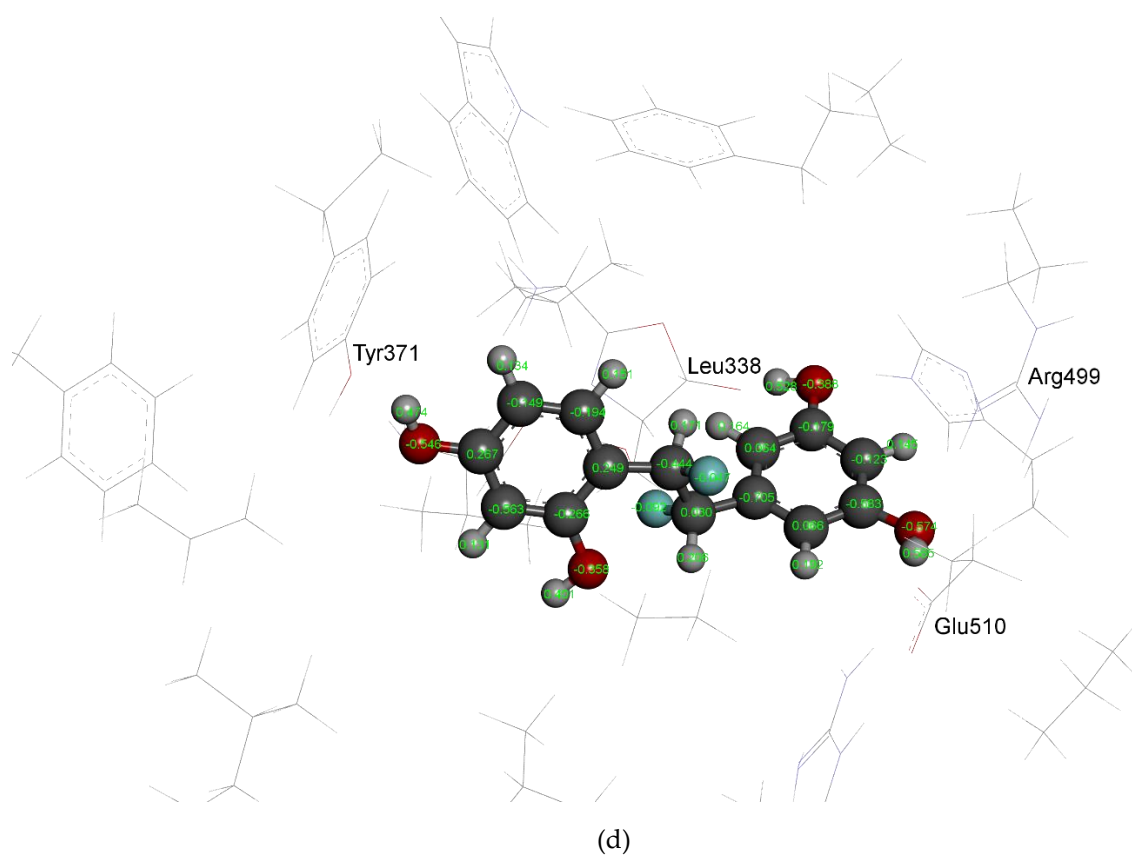
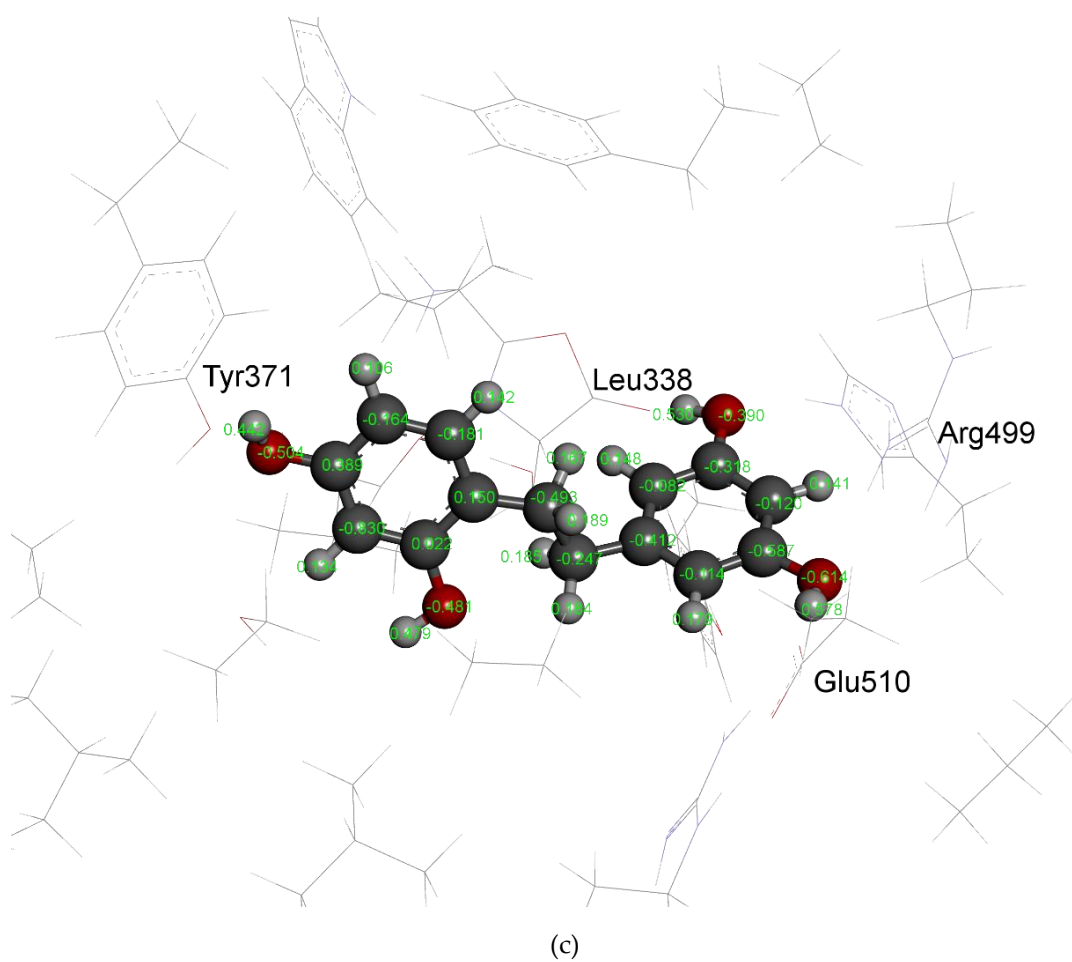


Table S1. Hydrophobic interactions of relaxed pose of celecoxib obtained from MD simulation in the catalytic pocket of COX-2.

Residues	Distance (Å)	Interaction type
Val335	4.47, 5.34	Pi-Alkyl
Val335	5.27	Alkyl-Alkyl
Leu338	5.38, 5.50	Pi-Alkyl
Ser339	2.26	Pi-Sigma
Tyr341	5.18	Pi-Alkyl
Leu370	5.15	Alkyl-Alkyl
Tyr371	4.57	Pi-Alkyl
Trp373	4.88	Pi-Alkyl
Val509	3.87	Pi-Alkyl
Ala513	3.99, 4.42	Pi-Alkyl

Table S2. Dunnett t-test of docking scores.

Compounds	Mean Difference	Significance	95% Confidence Interval
			Upper Bound
1	-18.03	0.031	-1.571
2	-20.65	0.013	-4.191
3	-11.46	0.195	4.999
4	-16.08	0.056	0.376
5	-17.51	0.036	-1.047
6	-4.85	0.633	11.613
7	-17.83	0.033	-1.374
8	-17.72	0.034	-1.261
9	-12.56	0.149	3.899

Mean square error = 59.086; Standard error = 6.276

Table S3 Dunnett t-test of QM interaction energy.

Compounds	Mean Difference	Significance	95% Confidence Interval
			Upper Bound
1	-5.502	0.151	1.652
2	9.332	1.000	16.487
3	-6.468	0.081	0.687
4	-15.998	0.000	-8.843
5	6.308	1.000	13.462
6	-8.040	0.025	-0.886
7	-17.248	0.000	-10.093
8	11.472	1.000	18.627
9	-7.138	0.051	0.017

Mean square error = 15.730; Standard error = 2.804

Characteristics of oxyresveratrol (1)

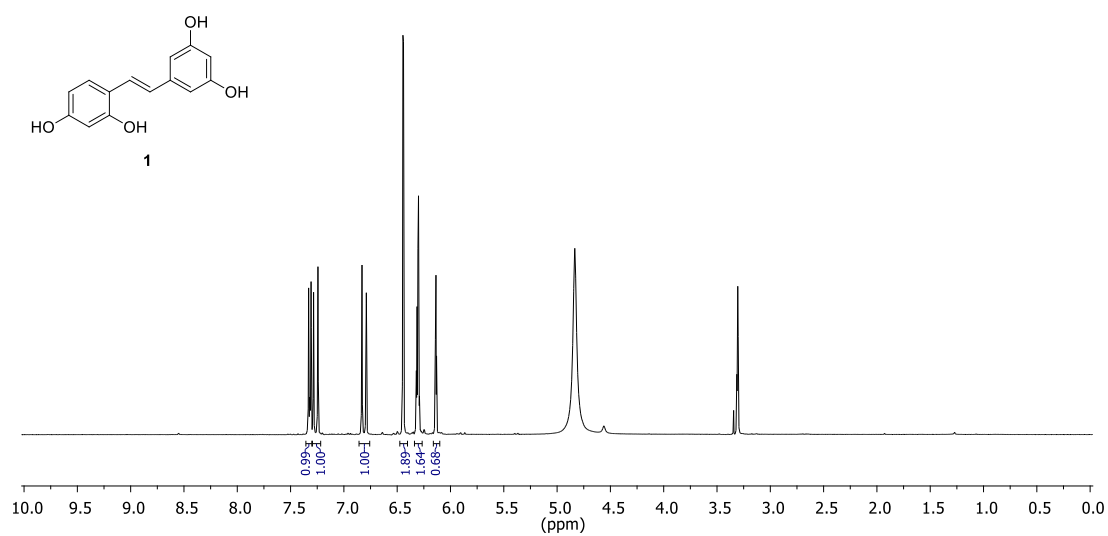


Figure S6 ¹H NMR (Methanol-*d*₄, 400 MHz) of oxyresveratrol (1).

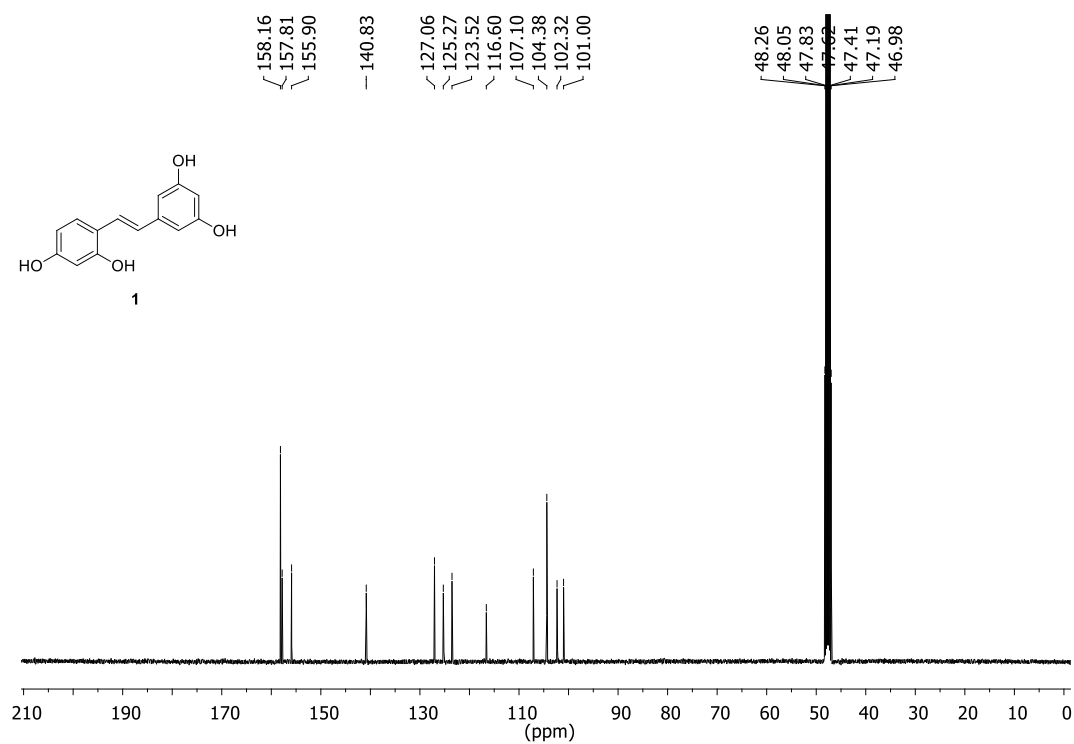


Figure S7 ¹³C NMR (Methanol-*d*₄, 100 MHz) of oxyresveratrol (1).

Characteristics of 2,3',4,5'-tetramethoxystilbene (2)

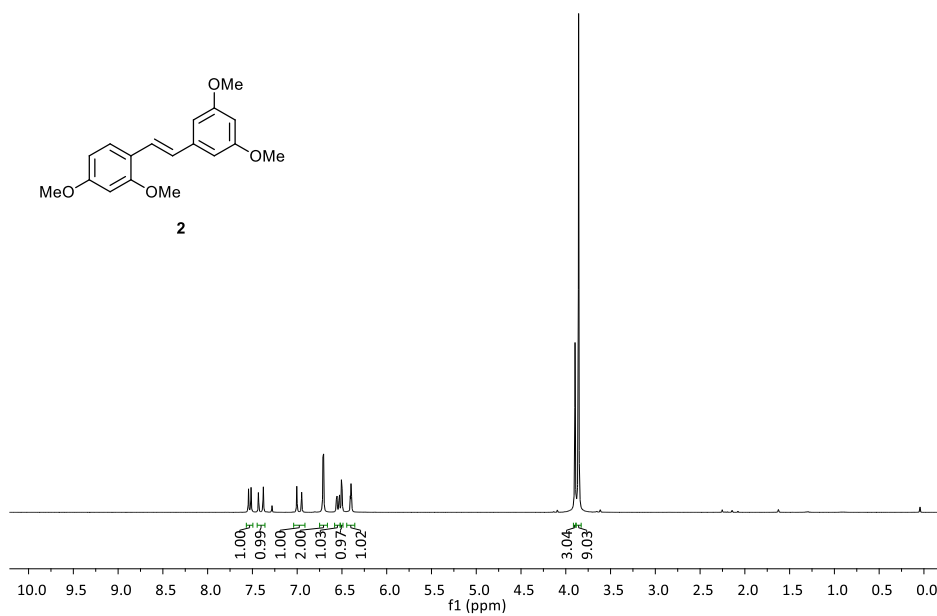


Figure S8 ¹H NMR (CDCl₃, 300 MHz) of 2,3',4,5'-tetramethoxystilbene (2).

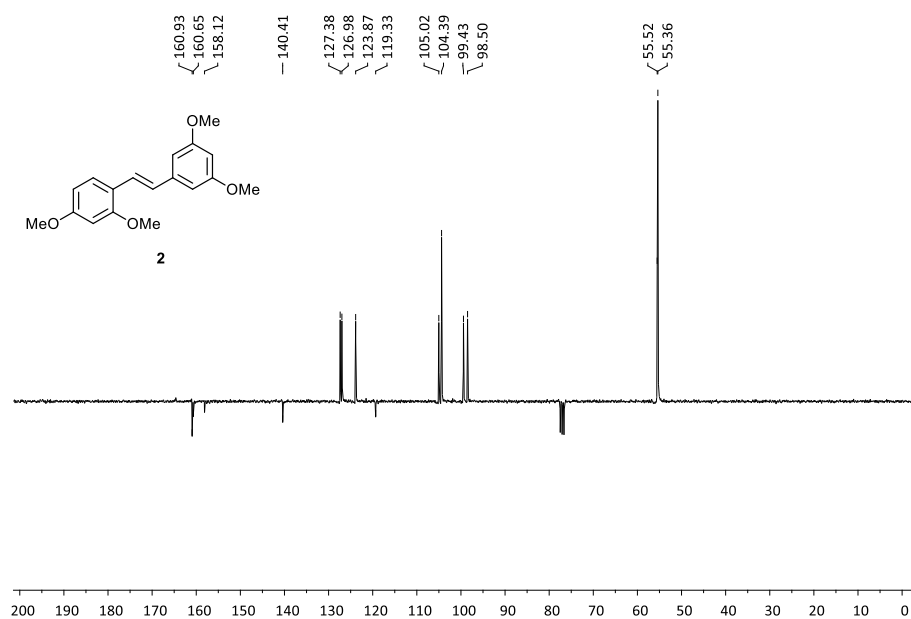


Figure S9 ¹³C NMR (CDCl₃, 75 MHz) of 2,3',4,5'-tetramethoxystilbene (2).

Characteristics of 2,3',4,5'-tetrahydroxybibenzyl (4)

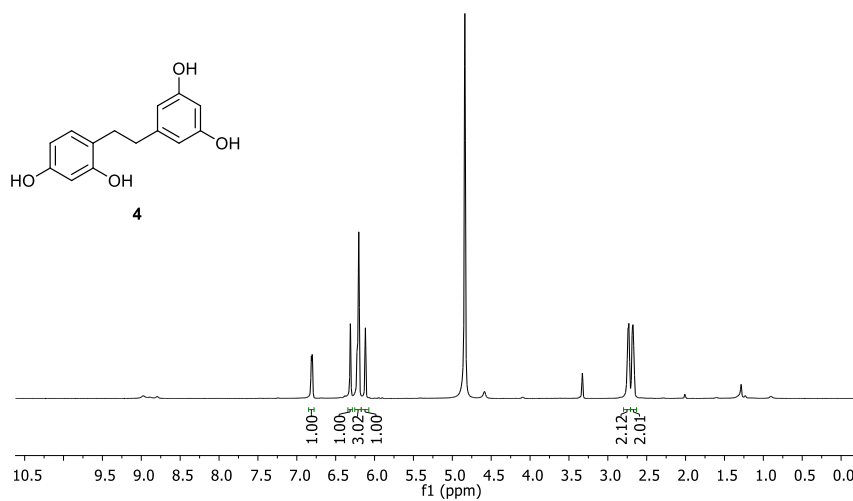


Figure S10 ¹H NMR (Methanol-*d*₄, 600 MHz) of 2,3',4,5'-tetrahydroxybibenzyl (4).

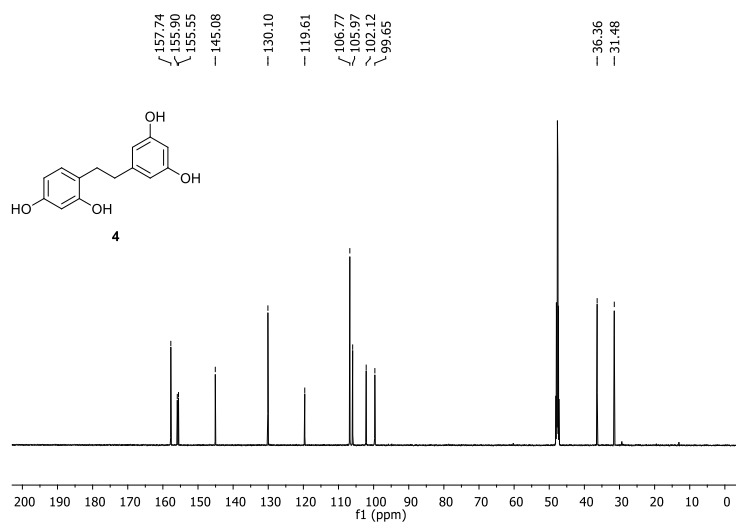


Figure S11 ¹³C NMR (Methanol-*d*₄, 150 MHz) of 2,3',4,5'-tetrahydroxybibenzyl (4).

Characteristics of 2,3',4,5'-tetramethoxybibenzyl (5)

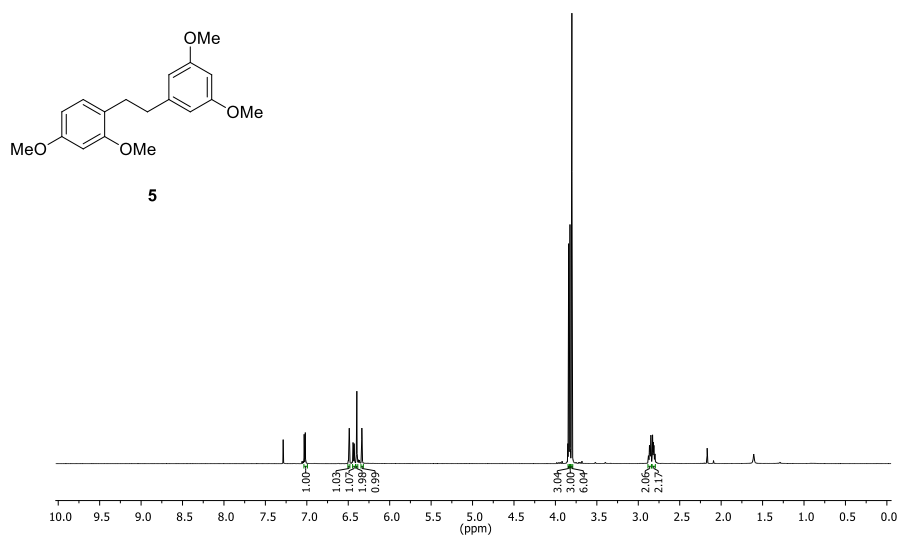


Figure S12 ¹H NMR (CDCl₃, 600 MHz) of 2,3',4,5'-tetramethoxybibenzyl (5).

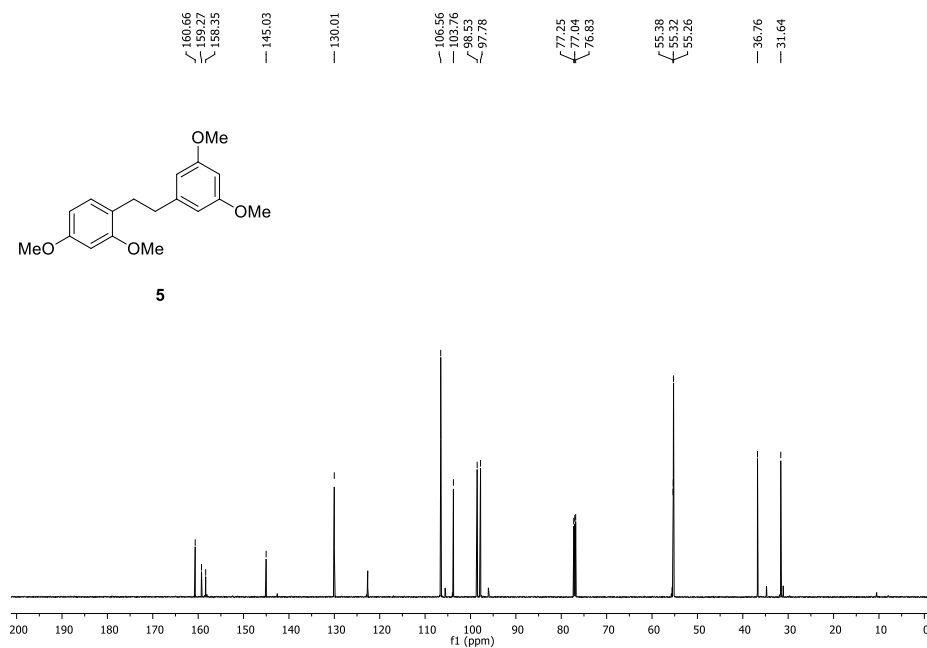


Figure S13 ¹³C NMR (CDCl₃, 150 MHz) of 2,3',4,5'-tetramethoxybibenzyl (5).

Characteristics of 2,3',4,5'-tetraacetoxystilbene (3)

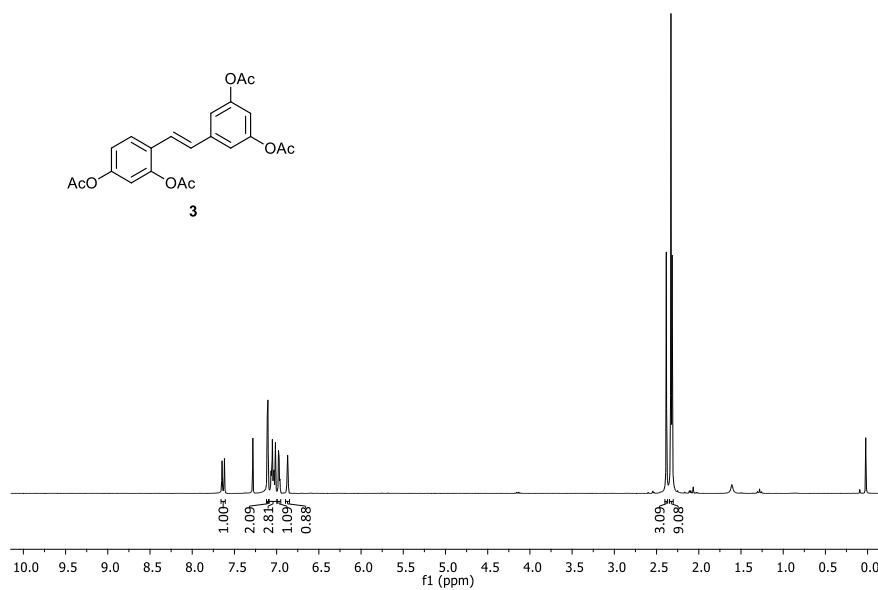


Figure S14 ¹H NMR (CDCl₃, 300 MHz) of 2,3',4,5'-tetraacetoxystilbene (3).

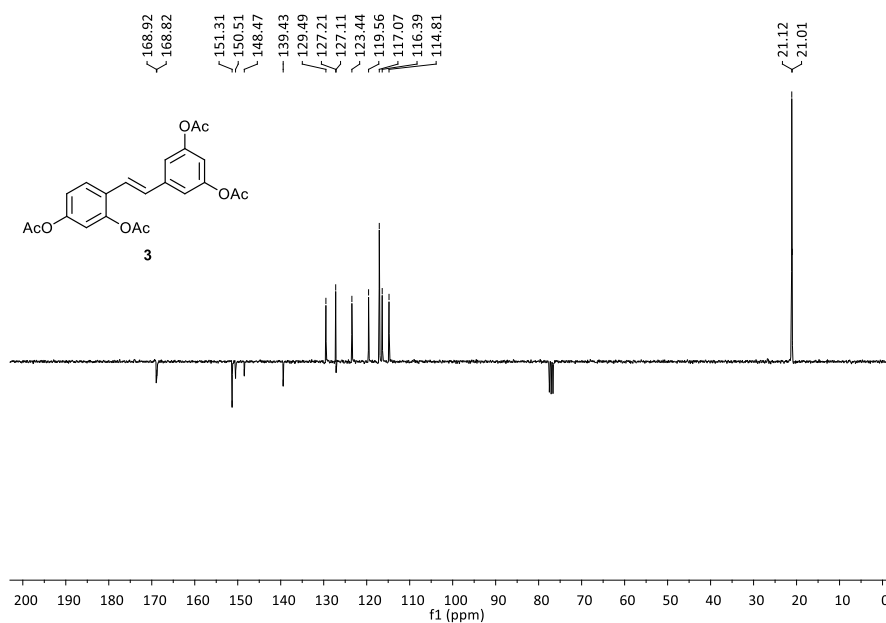


Figure S15 ¹³C NMR (CDCl₃, 75 MHz) of 2,3',4,5'-tetraacetoxystilbene (3)

Characteristics of 2,3',4,5'-tetraacetoxybibenzyl (6)

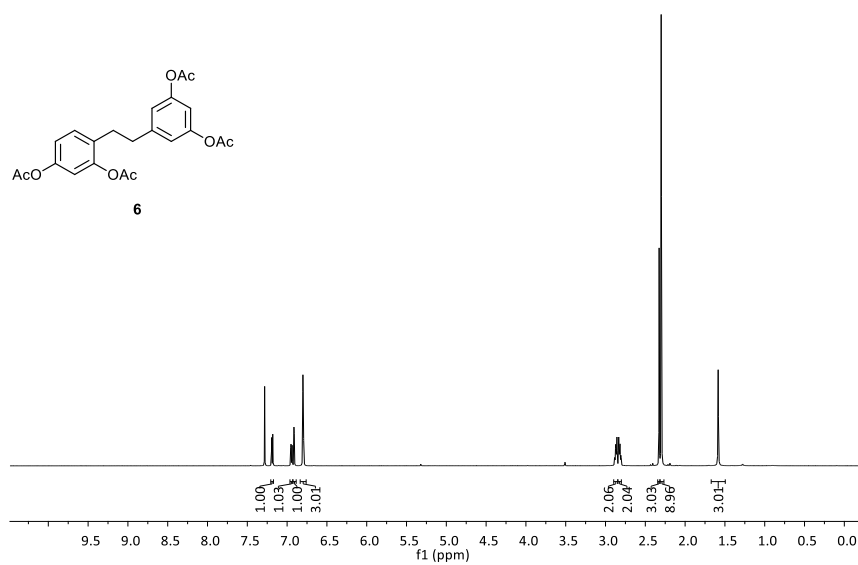


Figure S16 ¹H NMR (CDCl₃, 600 MHz) of 2,3',4,5'-tetraacetoxybibenzyl (6).

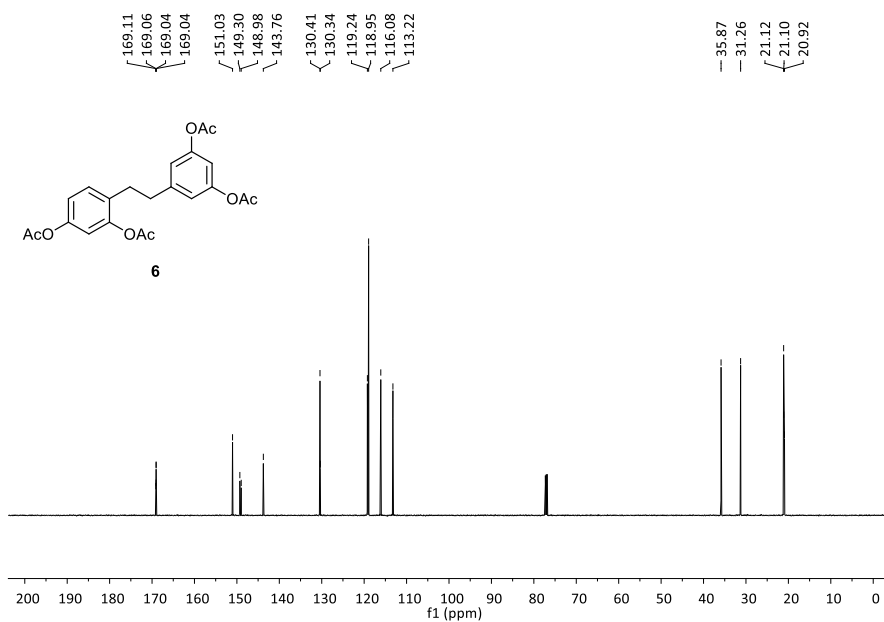


Figure S17 ¹³C NMR (CDCl₃, 150 MHz) of 2,3',4,5'-tetraacetoxybibenzyl (6).