

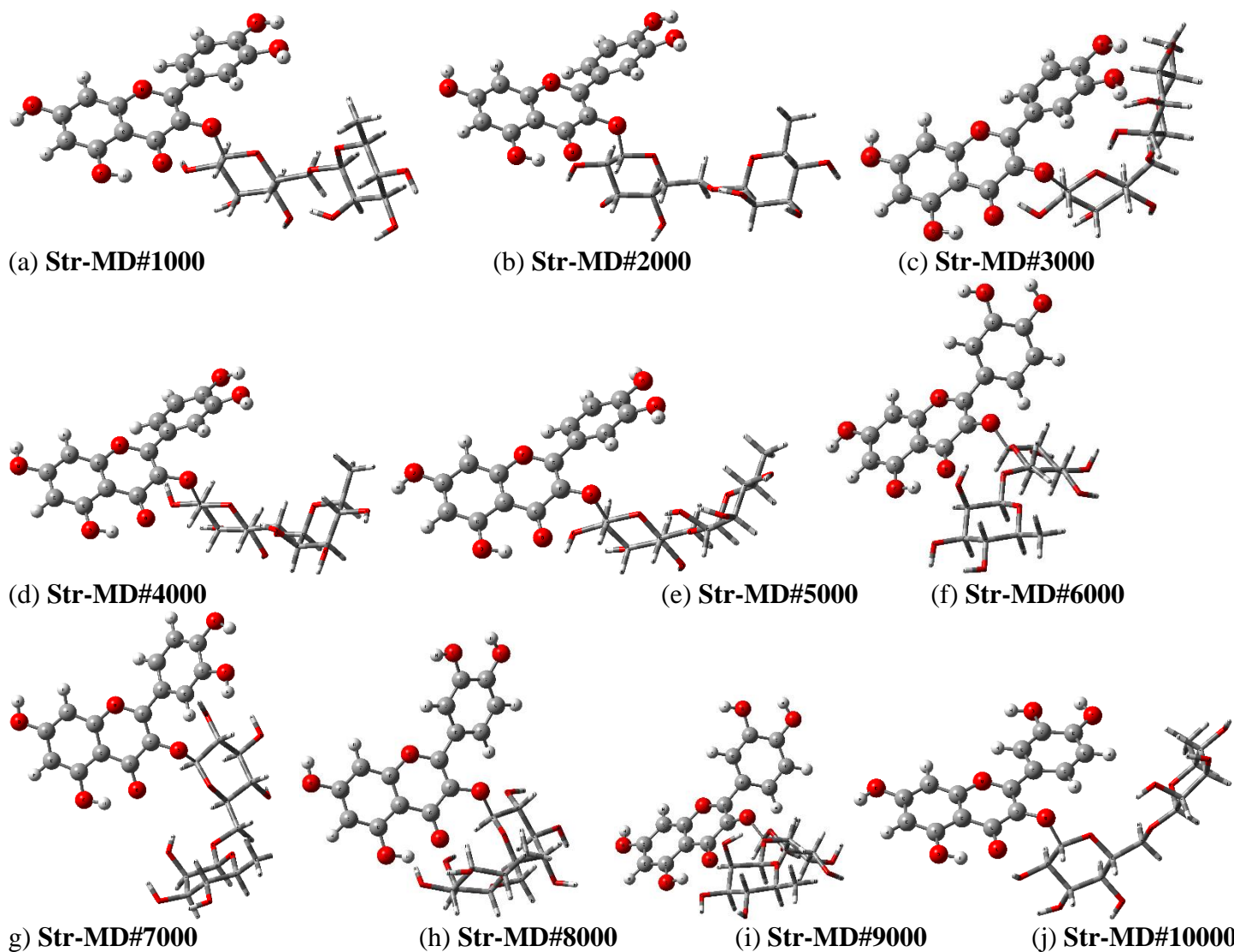
## SUPPLEMENTARY MATERIAL

### ON THE USE OF $^1\text{H}$ NMR CHEMICAL SHIFTS AND THERMODYNAMIC DATA FOR THE PREDICTION OF THE PREDOMINANT CONFORMATION OF ORGANIC MOLECULES IN SOLUTION: The example of the flavonoid Rutin

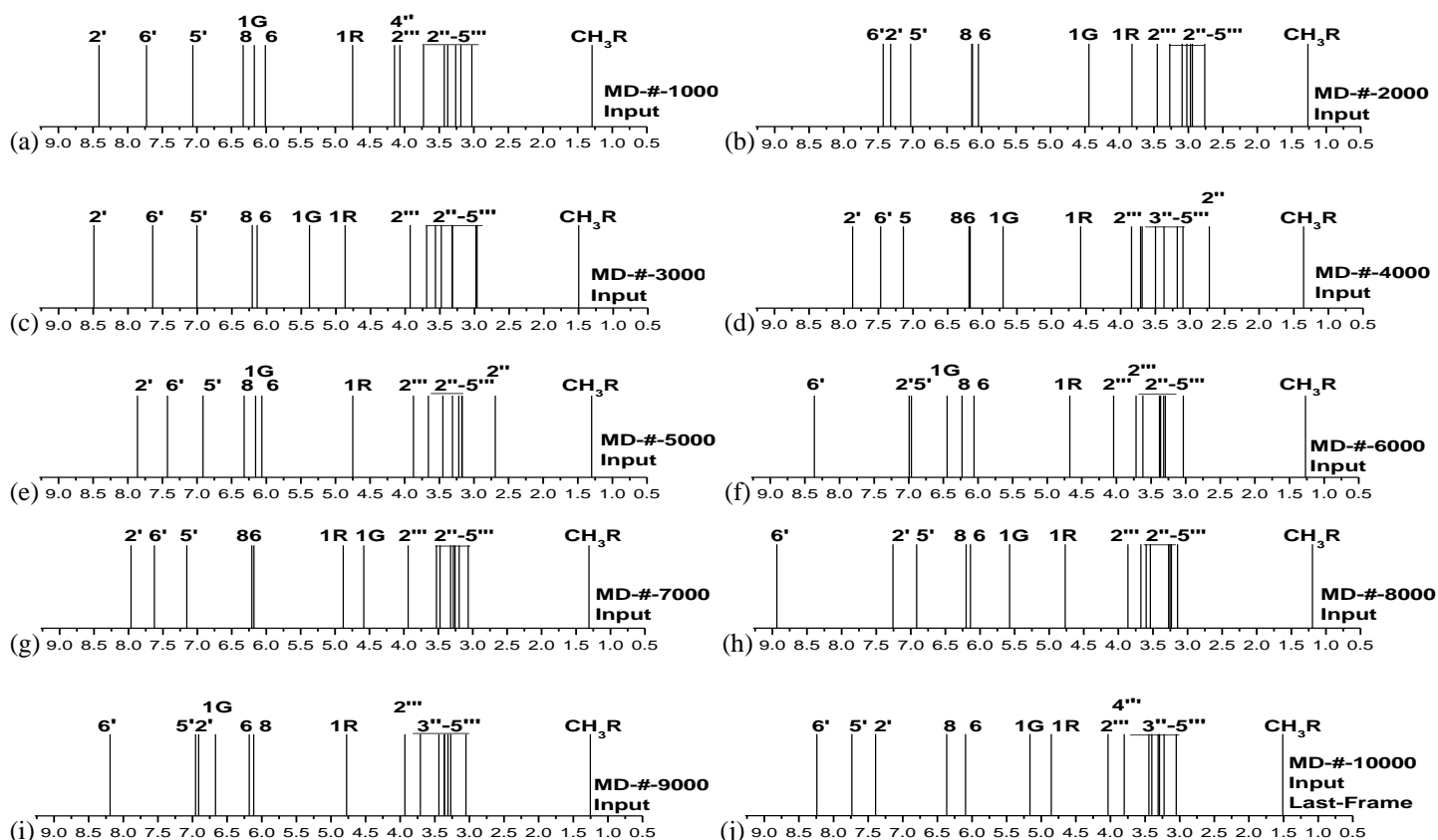
Haroldo C. Da Silva<sup>a,b</sup>, Wagner B. De Almeida<sup>a,\*</sup>

<sup>a</sup> Laboratório de Química Computacional e Modelagem Molecular (LQC-MM), Departamento de Química Inorgânica, Instituto de Química, Universidade Federal Fluminense (UFF), Outeiro de São João Batista s/n, Campus do Valonguinho, 24020-141, Centro, Niterói, RJ, Brazil.

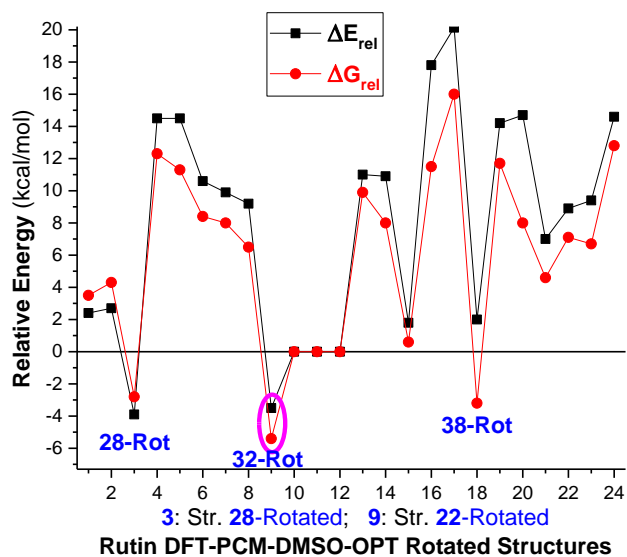
<sup>b</sup> Laboratório de Química Teórica e Simulação Molecular (LQTSM), Departamento de Físico-Química, Instituto de Química, Pavilhão Haroldo Lisboa da Cunha, Universidade do Estado do Rio de Janeiro (UERJ), Rua São Francisco Xavier, 524, 20550-013, Maracanã, Rio de Janeiro, RJ, Brazil.



**Figure S1.**  $\omega$ B97x-D /6-31G(d,p) new fully optimized structures of rutin obtained from ten frames (from a total of 10000) of MD simulation in DMSO. The optimized torsion angle values are given in Table S1.



**Figure S2.** B3LYP/6-31G(d,p)-PCM-DMSO  $^1\text{H}$  NMR spectra for DFT optimized rutin structures (MD inputs)

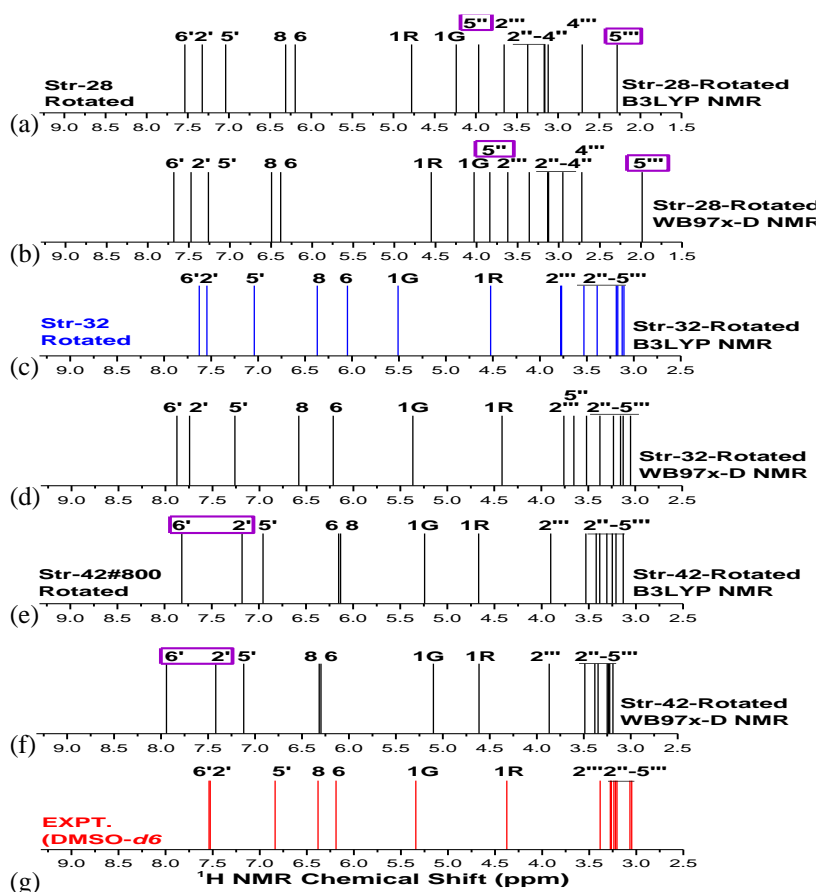


**Figure S3.**  $\omega$ B97x-D/6-31G(d,p)-PCM-DMSO relative energies ( $\Delta E_{\text{rel}}$  and  $\Delta G_{\text{rel}}$  in kcal mol<sup>-1</sup>) for relevant fully optimized and  $\phi_1$ ,  $\phi_s$  rotated structures located on the PES for rutin.  $\Delta G_{\text{rel}}$  was estimated using thermal corrections for DFT fully optimized structures (the  $\phi_1$ ,  $\phi_s$  rotated structures are not true minimum on the PES, so harmonic frequency calculations, needed for evaluation of thermal correction, are meaningless).

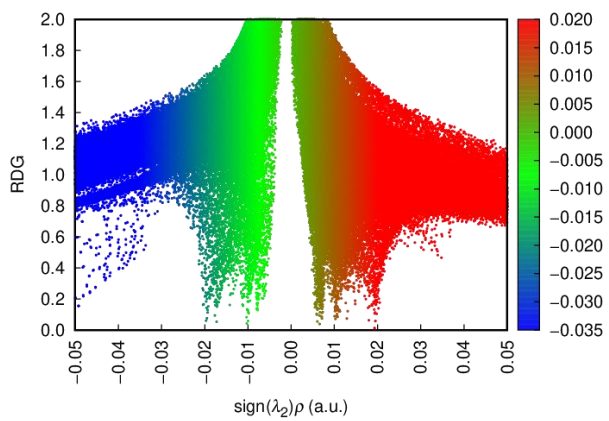
**1:** Str-28-PCM-OPT; **2:** Str-28 Vacuum-OPT; **3: Str-28-PCM-OPT-Rotated;** **4:** Str-30-PCM-OPT; **5:** Str-30 Vacuum-OPT; **6: Str-30-PCM-OPT-Rotated;** **7:** Str-32-PCM-OPT; **8:** Str-32 Vacuum-OPT; **9: Str-32-PCM-OPT-Rotated;** **10:** Str-34-PCM-OPT; **11:** Str-34 Vacuum-OPT; **12: Str-34-PCM-OPT-Rotated;** **13:** Str-37-PCM-OPT; **14:** Str-37 Vacuum-OPT; **15: Str-37-PCM-OPT-Rotated;** **16:** Str-38-PCM-OPT; **17:** Str-38 Vacuum-OPT; **18: Str-38-PCM-OPT-Rotated;** **19:** Str-42-PCM-OPT; **20:** Str-42 Vacuum-OPT; **21: Str-42-PCM-OPT-Rotated;** **22:** Str-44-PCM-OPT; **23:** Str-44 Vacuum-OPT; **24: Str-44-PCM-OPT-Rotated.**

**Table S1.** Deviation between  $\omega$ B97x-D/6-31G(d,p)-PCM-DMSO and B3LYP/6-31G(d,p)-PCM-DMSO  $^1\text{H}$  NMR chemical shifts (in ppm) for rotated structures of rutin. MAE (in ppm) values are also given.

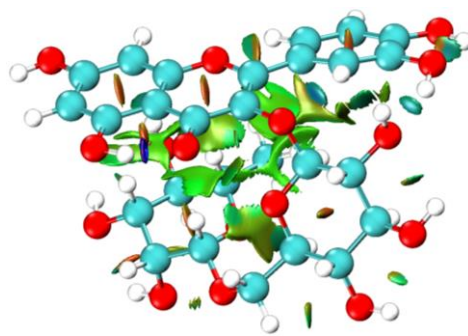
CH <sub>n</sub> Protons	Str.-28-Rot	Str.-30-Rot	Str.-32-Rot	Str.-34-Rot	Str.-37-Rot	Str.-42-Rot	Str.-44-Rot
H-C6	0.19	0.19	0.17	0.24	0.29	0.16	0.25
H-C8	0.18	0.22	0.2	0.18	0.13	0.16	0.19
H-C2'	0.15	0.18	0.20	0.16	0.08	0.24	0.14
H-C5'	0.22	0.21	0.22	0.20	0.21	0.16	0.16
H-C6'	0.15	0.18	0.25	0.21	0.19	0.13	0.18
H-C1-R	-0.23	-0.12	-0.11	-0.10	-0.07	-0.05	-0.09
H-C1-G	-0.21	-0.05	-0.14	-0.09	-0.09	-0.14	-0.07
H-CH3	-0.05	0.003	-0.006	0.03	0.02	0.04	0.03
H-2''	-0.01	-0.11	0.05	-0.03	-0.24	-0.07	-0.09
H-3''	-0.03	-0.01	-0.01	0.013	0.01	-0.002	0.03
H-4''	0.004	-0.10	-0.01	-0.13	-0.14	-0.02	-0.10
H-5''	-0.13	-0.11	-0.12	-0.04	-0.06	-0.04	-0.06
H-2'''	-0.04	-0.07	-0.02	-0.07	-0.03	-0.04	-0.05
H-3'''	-0.21	0.04	0.03	0.07	0.14	0.12	0.09
H-4'''	0.01	-0.02	-0.04	-0.05	-0.14	-0.12	-0.11
H-5'''	-0.30	-0.05	-0.06	0.04	0.03	0.01	-0.001
<b>MAE</b>	<b>0.13</b>	<b>0.10</b>	<b>0.10</b>	<b>0.10</b>	<b>0.12</b>	<b>0.09</b>	<b>0.10</b>



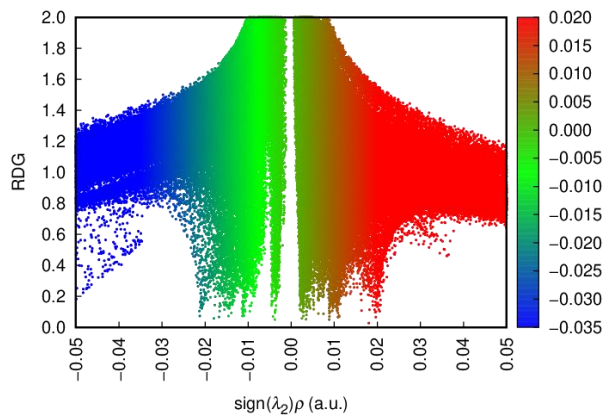
**Figure S4.** B3LYP/6-31G(d,p)-PCM-DMSO and  $\omega$ B97x-D /6-31G(d,p)-PCM-DMSO  $^1\text{H}$  NMR spectra for selected structures of rutin ( $\omega$ B97x-D /6-31G(d,p)-PCM-DMSO optimized geometries) (a-f) and experimental spectra (in DMSO-*d*6) (g). The protons having large deviation from experimental data are highlighted in pink rectangle.



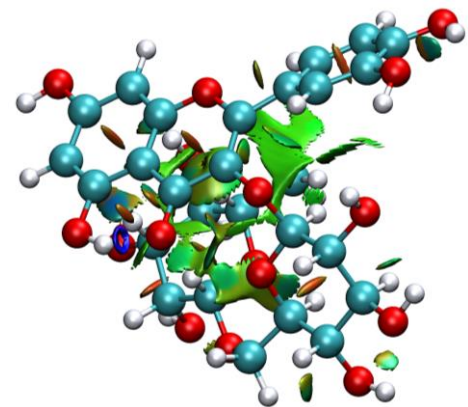
(a) 28 full opt



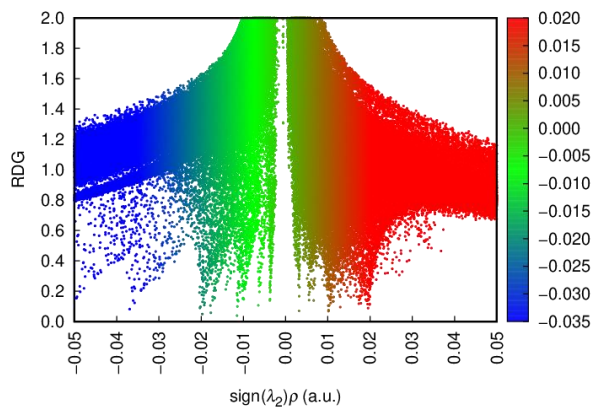
(b) 28 full opt



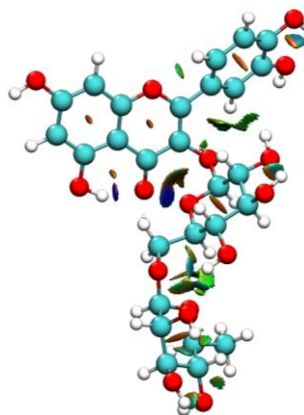
(c) 28 rot



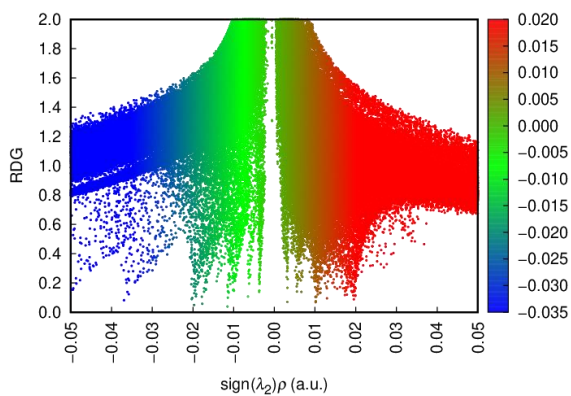
(d) 28 rot



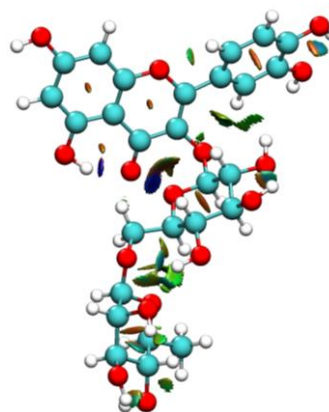
**(e) 30 FO**



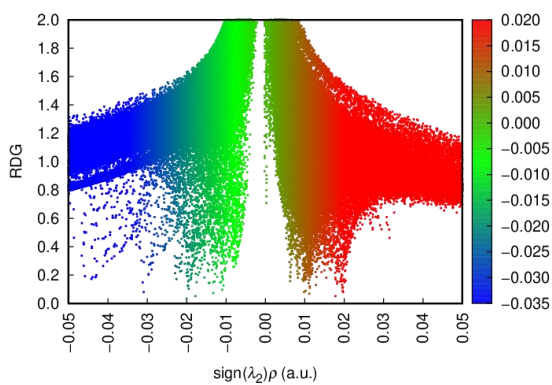
**(f) 30 FO**



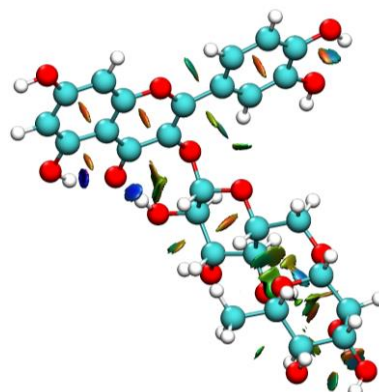
**(g) 30 ROT**



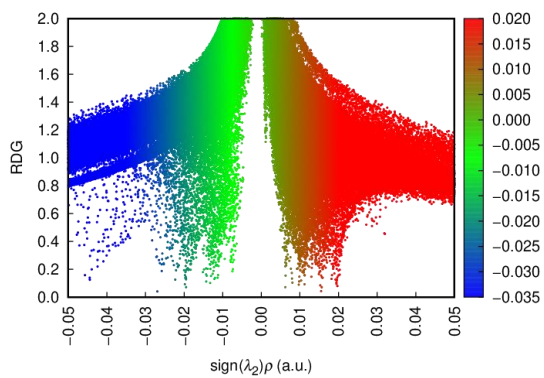
**(h) 30 ROT**



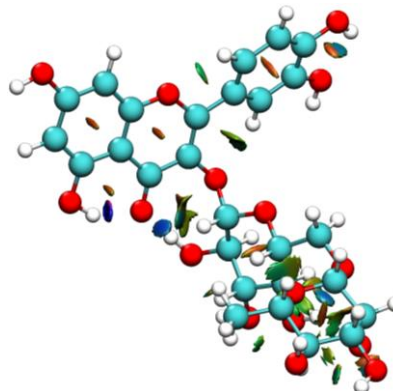
**(i) 32FO**



**(j) 32FO**

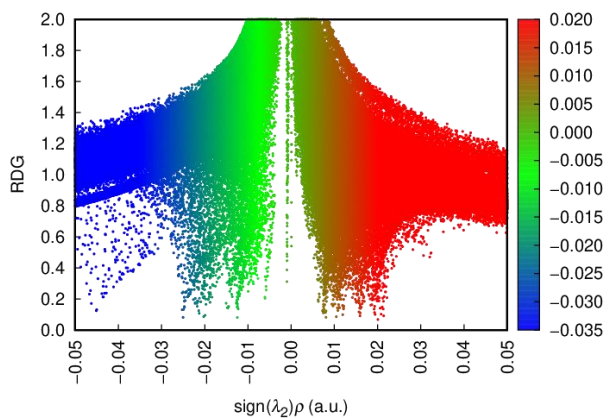


**(k) 32ROT**

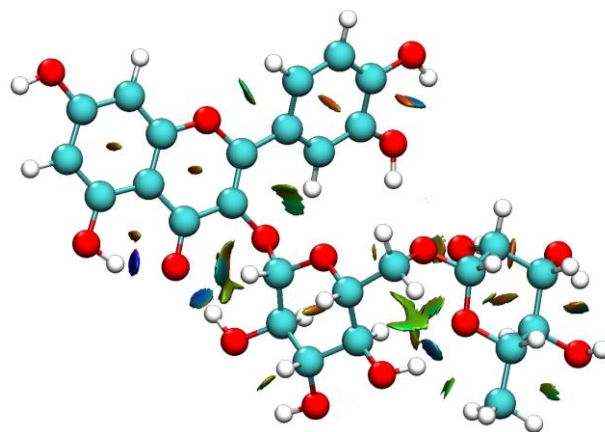


**(l) 32 ROT**

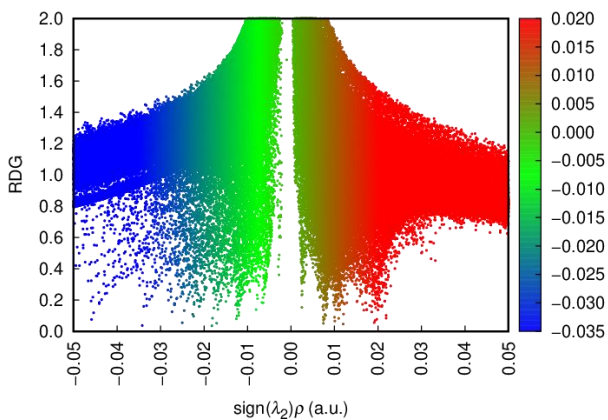




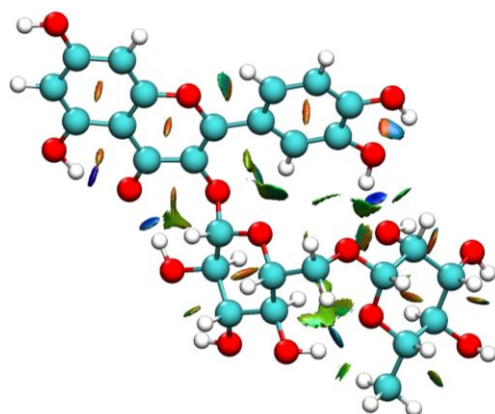
**(m) 34 FO**



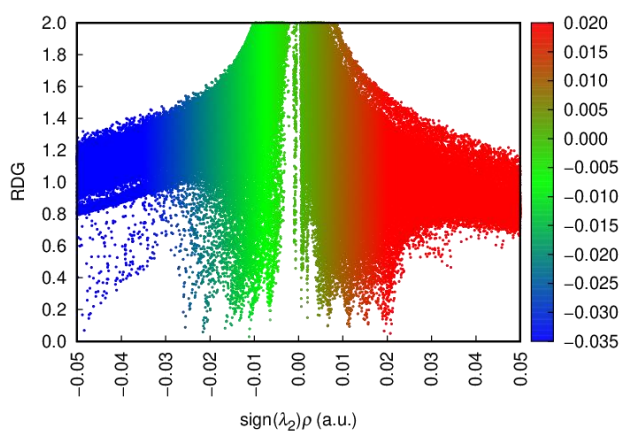
**(n) 34 FO**



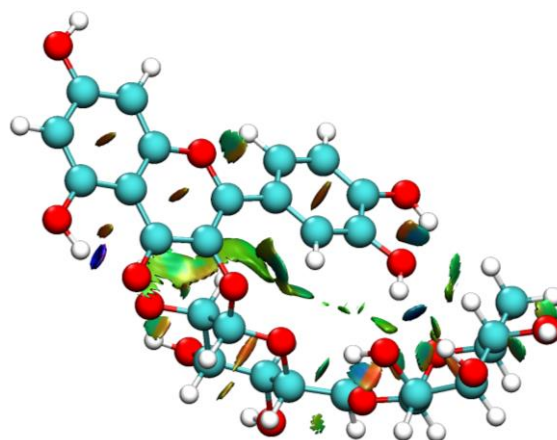
**(o) 34 ROT**



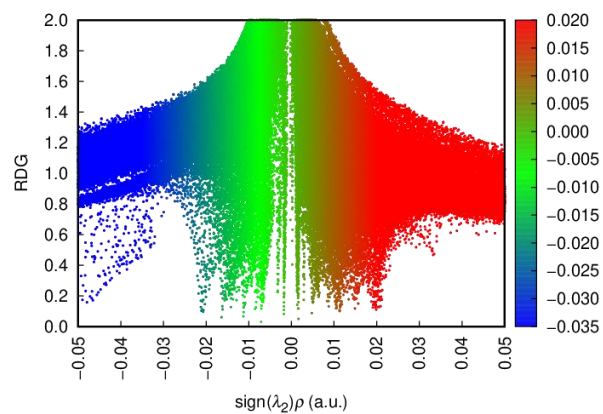
**(p) 34 ROT**



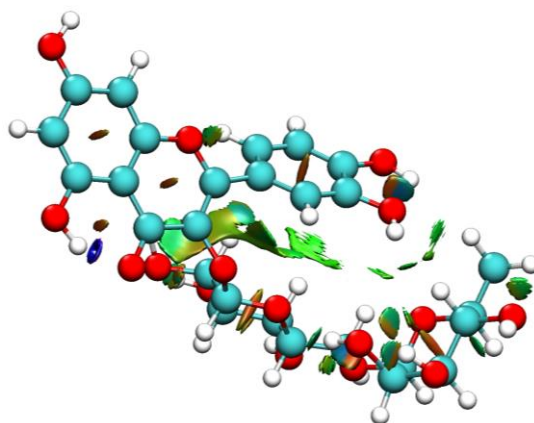
**(q) 37 FO**



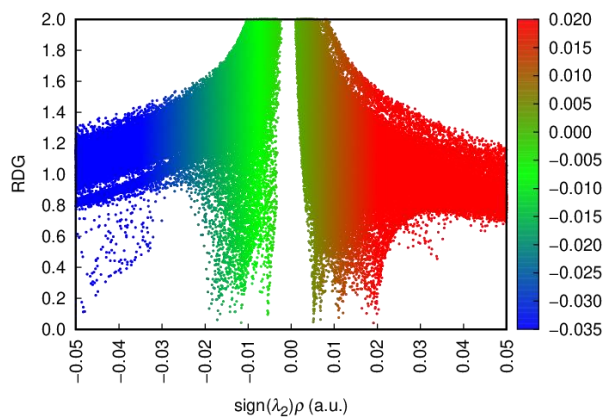
**(r) 37 FO**



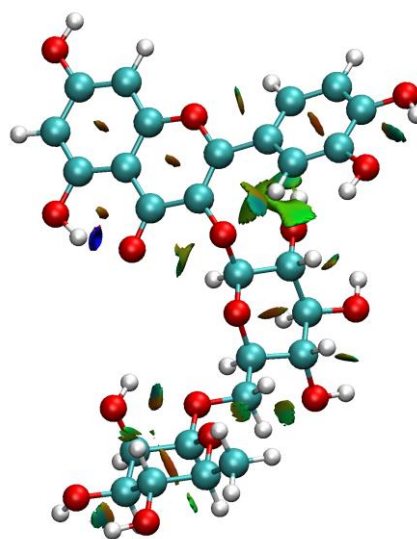
**(s) 37 ROT**



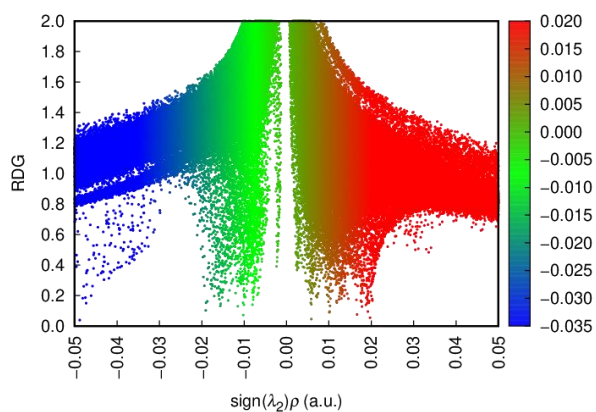
**(t) 37 ROT**



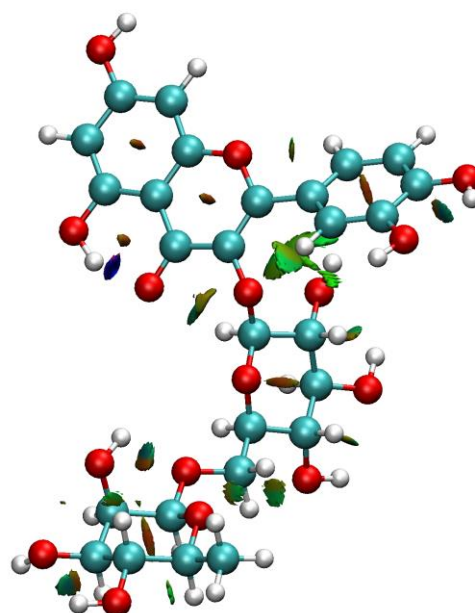
**(u) 38 FO**



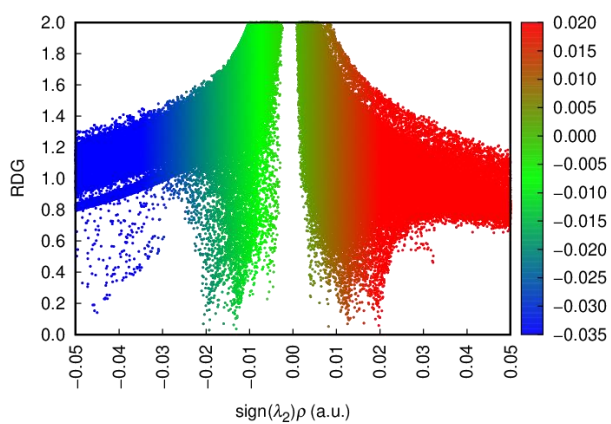
**(v) 38 FO**



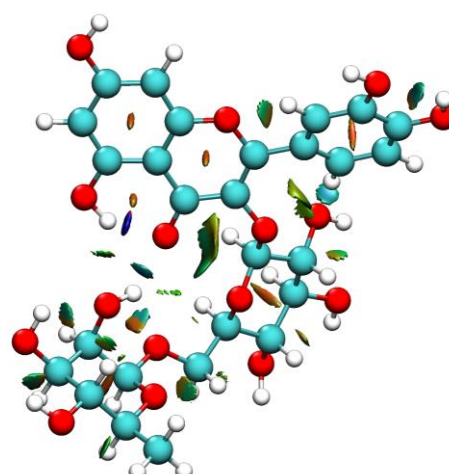
**(w) 38 ROT**



**(x) 38 ROT**

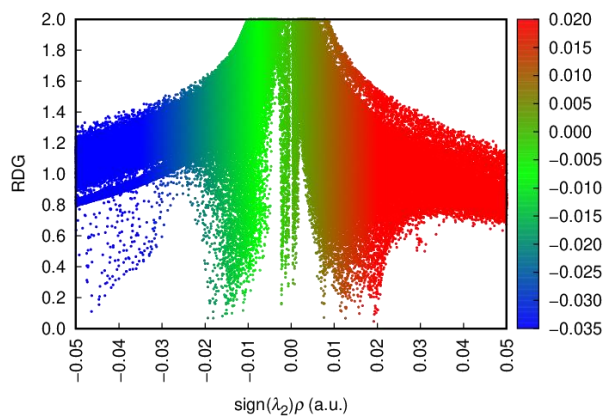


**(y) 42 OPT**

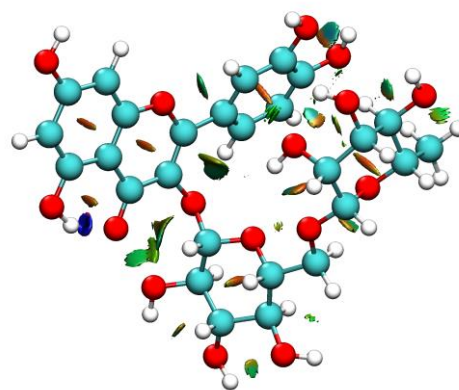


**(z) 42 OPT**

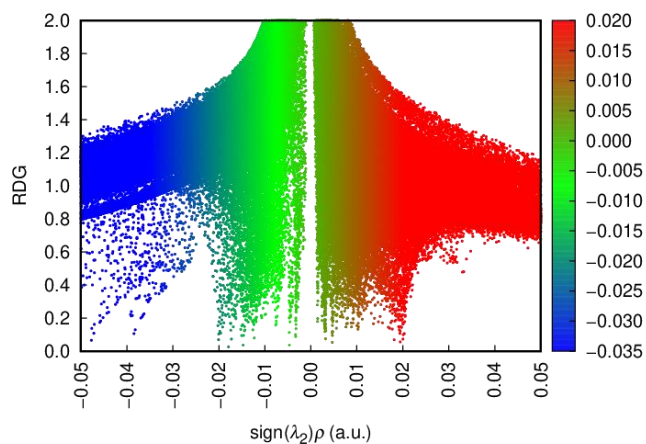




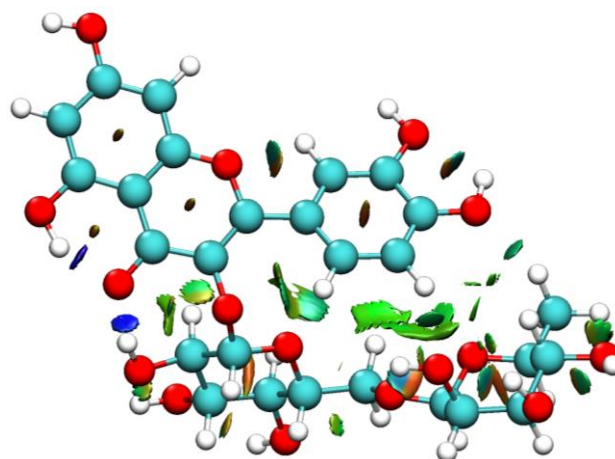
(aa) 42 ROT



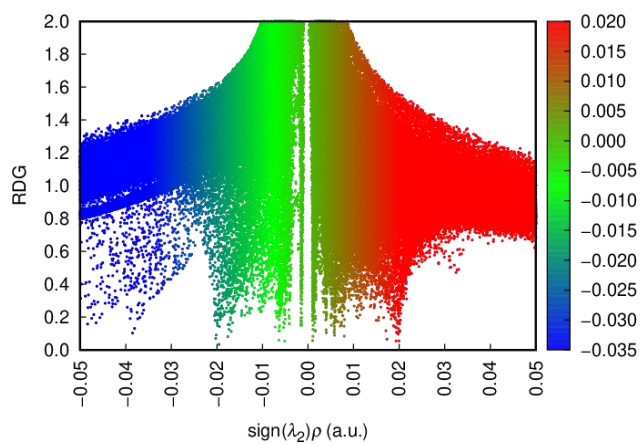
(ab) 42 ROT



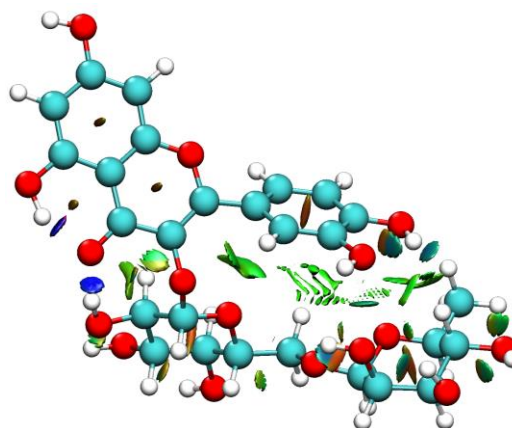
(ac) 44 FO



(ad) 44 FO



(ae) 44 ROT



(af) 44 ROT

**Figure S5.** Bi and tridimensional NCI plots calculated at  $\omega$ B97x-D/6-31G(d,p) – PCM DMSO level of theory (FO = Fully Optimized and ROT = rotated).

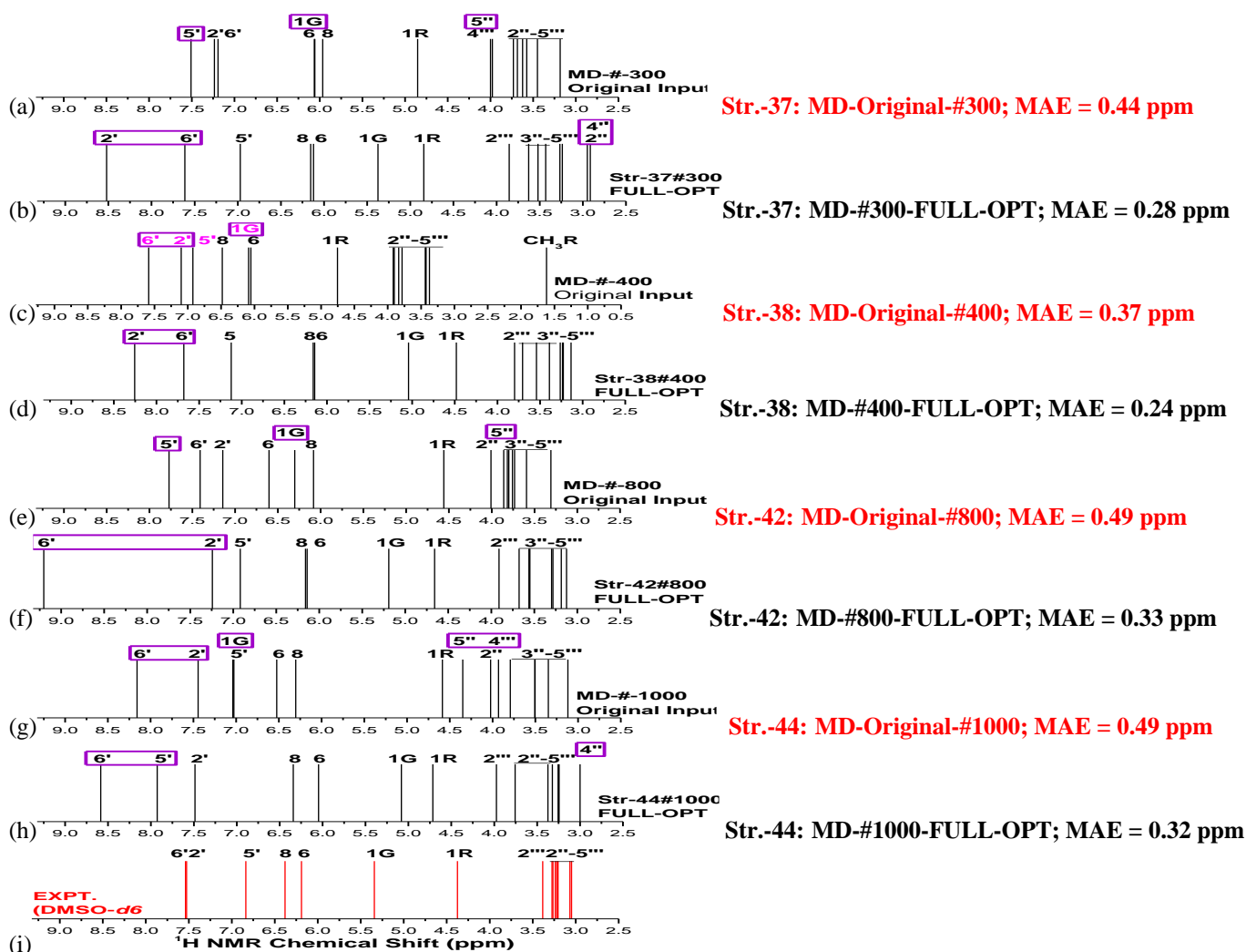
**Table S2.** Selected  $\omega$ B97x-D /6-31G(d,p)-PCM-DMSO optimized torsion angles ( $^{\circ}$ ) for fully optimized structures located on the PES for rutin using relevant MD frames as input and MD original frames (#300, #400, #800 and #1000) not optimized structures. Relative energy values ( $\Delta E_{rel}$  in kcal mol $^{-1}$ ) for DFT-PCM optimized geometries and original MD frames inputs (DFT-PCM single point calculations) are also given.

Structures	$\Phi_1$ [O-C2-C1'-C2']	$\Phi_2$ [C1''-O-C3-C2]	$\Phi_3$ [C2''-C1''-O-C3]	$\Phi_4$ [O-C6''-C5''-O]	$\Phi_5$ [C1'''-O-C6'''-C5''']	$\Phi_6$ [C1''''-O-C6''''-C5''']	$\Delta E_{rel}$
<b>37 PCM-DMSO-FULL-OPT (DM-#300)</b> <sup>a</sup>	158.3	-92.5	14.3	57.9	-166.2	168.1	<b>2.2</b>
<b>37 DM-#300 Original Frame (non-opt)</b> <sup>b</sup>	<b>106.0</b>	<b>-110.0</b>	<b>57.7</b>	<b>39.5</b>	<b>-167.6</b>	<b>174.1</b>	<b>-13.4</b> <sup>c</sup> (197) <sup>d</sup>
<b>38 PCM-DMSO-FULL-OPT (DM-#400)</b> <sup>a</sup>	153.2	-113.8	93.0	69.4	170.8	-175.2	<b>8.9</b>
<b>38 DM-#400 Original Frame (non-opt)</b> <sup>b</sup>	<b>114.4</b>	<b>-117.6</b>	<b>70.8</b>	<b>102.3</b>	<b>-165.0</b>	<b>167.4</b>	<b>8.6</b> <sup>c</sup> (194) <sup>d</sup>
<b>42 PCM-DMSO-FULL-OPT (DM-#800)</b> <sup>a</sup>	29.3	-141.2	134.3	67.5	-169.8	175.3	<b>5.3</b>
<b>42 DM-#800 Original Frame (non-opt)</b> <sup>b</sup>	<b>72.6</b>	<b>-141.3</b>	<b>77.7</b>	<b>84.9</b>	<b>-157.7</b>	<b>174.8</b>	<b>-8.7</b> <sup>c</sup> (133) <sup>d</sup>
<b>44 PCM-DMSO-FULL-OPT (DM (#1000))</b> <sup>a</sup>	-20.9	-92.0	-37.7	68.8	165.6	-179.8	<b>0.0</b>
<b>44 DM-#1000 Original Frame (non-opt)</b> <sup>b</sup>	<b>34.8</b>	<b>-141.9</b>	<b>65.3</b>	<b>72.9</b>	<b>-170.1</b>	<b>178.3</b>	<b>0.0</b> <sup>c</sup> (0.0) <sup>d</sup>

<sup>a</sup>  $\omega$ B97x-D /6-31G(d,p)-PCM-DMSO optimized torsion angles ( $^{\circ}$ ); <sup>b</sup> DM original frame (non-optimized) torsion angle.

<sup>c</sup>  $\omega$ B97x-D /6-31G(d,p)-PCM-DMSO relative energy single-point using DM original frame (non-optimized) structure

<sup>d</sup> Classical DM relative energy (using Force Field) using DM original frame (non-optimized) structure



**Figure S6.** B3LYP/6-31G(d,p)-PCM-DMSO  $^1\text{H}$  NMR spectra for selected structures **37**, **42** and **44** of rutin obtained from DM frames. Three distinct structures were used in the DFT-PCM NMR calculations: (i) **Original DM frames (not optimized)** (a,c,e,g) (ii) DFT-PCM-DMSO fully optimized DM inputs (b,d,f,h) and (iii)  $\phi_1$ ,  $\phi_s$  torsion angles rotated structures (c,f,i). The experimental  $^1\text{H}$  NMR spectrum (in DMSO-*d*6) is also shown (i).

The protons having large deviation from experimental data are highlighted in pink rectangle.