

# **Structural Analysis and Multipole Modelling of Quercetin**

## **Monohydrate – A Quantitative and Comparative Study**

*Sławomir Domagała, Parthapratim Munshi, Maqsood Ahmed, Benoît Guillot,*

*Christian Jelsch\**

Laboratoire de Cristallographie, Résonance Magnétique et Modélisations (CRM2),  
CNRS, UMR 7036, Institut Jean Barriol, Faculté des Sciences et Techniques, Nancy  
Université, BP 70239, 54506 Vandoeuvre-lès-Nancy CEDEX, France

## **SUPPORTING INFORMATION**

### **Supplementary Tables S1-S7**

### **Supplementary Figures S1-S9**

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\*Corresponding author. E-mail: christian.jelsch@crm2.uhp-nancy.fr. Tel.: +33 (0)383

684 899. Fax: +33 (0)383 406 492.

**Table S1.** Atom-types assignment in quercetin monohydrate.

Atom label	Atom-type
O1	Occ
O3, O5, O7, O13, O14	Och
O4	O=c[cc]
C2, C3, C5, C7, C9, C13, C14,	Co[x]cc
C4	C=occ
C6, C8, C12, C15, C16	Ccch
C10	C1.5c1.5c1.5c
C11	C1c1.5c1.5c
H3, H5, H7, H13, H14	Ho[cc]
H6, H8, H12, H15, H16	Hc[cc]
O1W	Ohh
H1W, H2W	Ho[h]

The numbers in atom-type name denote bond orders *i.e.* 1 – single, 1.5 – intermediate, 2 – double bonds. The atomic symbols in between the brackets code the neighbours of the neighbours of the first atom.

**Table S2.** Comparison of the X-H distances for the IAM\_UR and TAAM\_UR models.

Bond	$d_{\text{IAM\_UR}}(\text{\AA})$	$d_{\text{TAAM\_UR}}(\text{\AA})$
O3-H3	0.909(27)	0.967(14)
O5-H5	0.921(25)	1.000(13)
O7-H7	0.822(26)	0.908(13)
O13-H13	0.841(26)	0.972(13)
O14-H14	0.880(25)	0.965(13)
O1W-H1W	0.900(28)	1.007(14)
O1W-H2W	0.923(28)	1.014(14)
Average (rmsd)	0.885(37)	0.976(33)
Average (neutron)	0.967(10)	
$(d_{\text{model}} - d_{\text{neutron}})/\sigma_{\text{neutron}}$	-8.21	0.91
C6-H6	0.904(22)	1.046(10)
C8-H8	0.955(22)	1.040(10)
C12-H12	0.976(22)	1.045(09)
C15-H15	1.010(23)	1.083(10)
C16-H16	0.943(23)	1.059(10)
Average (rmsd)	0.957(35)	1.055(16)
Average (neutron)	1.083(11)	
$(d_{\text{model}} - d_{\text{neutron}})/\sigma_{\text{neutron}}$	-11.42	-2.57

**Table S3.** Hirshfeld test values with the corresponding distances for the quercetin molecule for the IAM\_R, TAAM\_R and TAAM\_THEO\_R models. DMSDA ( $\text{\AA}^2 * 10^4$ )

Bond	IAM_R DMSD A	IAM_R <i>d</i> (Å)	TAAM _R DMSD A	TAAM_R <i>d</i> (Å)	TAAM_ THEO_R DMSDA	TAAM_ THEO_R <i>d</i> (Å)
O1-C2	-8	1.3708(16)	-2	1.3666(9)	-6	1.3654(8)
O1-C9	-5	1.3645(16)	1	1.3582(9)	-4	1.3560(8)
O3-C3	2	1.3576(16)	-3	1.3534(9)	-2	1.3515(8)
O4-C4	-16	1.2752(16)	5	1.2752(9)	0	1.2688(8)
O5-C5	-7	1.3603(16)	-7	1.3524(9)	-5	1.3512(8)
O7-C7	-20	1.3525(16)	-6	1.3466(9)	-3	1.3448(8)
O13-C13	1	1.3707(15)	5	1.3656(9)	5	1.3633(8)
C14-C14	-7	1.3584(16)	-4	1.3529(9)	-2	1.3511(8)
C2-C3	-4	1.3744(18)	3	1.3766(10)	1	1.3760(9)
C2-C11	17	1.4635(18)	3	1.4640(10)	-4	1.4633(9)
C3-C4	-12	1.4330(18)	-15	1.4341(10)	-3	1.4357(9)
C6-C5	-10	1.3718(19)	-6	1.3800(10)	3	1.3800(9)
C6-C7	-3	1.4051(18)	-4	1.4078(10)	2	1.4070(9)
C8-C7	-2	1.3896(19)	-6	1.3922(10)	1	1.3923(9)
C8-C9	-12	1.3836(19)	-8	1.3900(10)	2	1.3894(9)
C10-C4	-20	1.4278(18)	-8	1.4267(10)	-1	1.4284(9)
C10-C5	-14	1.4192(18)	-5	1.4214(10)	-7	1.4204(9)
C10-C9	-2	1.4009(18)	1	1.3999(9)	-2	1.4014(9)
C11-C12	-5	1.4087(18)	4	1.4088(10)	2	1.4083(9)
C11-C16	-4	1.4027(18)	-4	1.4044(10)	-5	1.4045(9)
C12-C13	-5	1.3830(19)	-3	1.3865(10)	1	1.3877(9)
C13-C14	2	1.3987(18)	1	1.4028(10)	2	1.4009(9)
C14-C15	8	1.3881(19)	6	1.3937(10)	0	1.3937(9)
C16-C15	2	1.3914(19)	1	1.3912(10)	0	1.3916(9)
rmsd	7.7(60)		4.7(30)		2.6(19)	

**Table S4.** Topological properties of the electron density for the covalent bonds of quercetin monohydrate for the TAAM\_OPT (first entry) and THEO\_OPT (second entry in italics) models.

Bond	$d_{AB}$	$d_{A...CP}$	$d_{B...CP}$	$\rho(\mathbf{r}_{CP})$	$\nabla^2\rho(\mathbf{r}_{CP})$	$\lambda_1$	$\lambda_2$	$\lambda_3$	$\varepsilon$
O1-	1.3659	0.8295	0.5368	1.99	-15.06	13.98	-13.82	-15.21	0.09
C2		<i>0.8285</i>	<i>0.5380</i>	<i>1.85</i>	<i>-10.37</i>	<i>14.13</i>	<i>-11.88</i>	<i>-12.62</i>	<i>0.06</i>
O1-	1.3571	0.8309	0.5263	2.02	-16.03	13.50	-14.08	-15.45	0.09
C9		<i>0.8303</i>	<i>0.5274</i>	<i>1.89</i>	<i>-12.30</i>	<i>13.33</i>	<i>-12.39</i>	<i>-13.24</i>	<i>0.06</i>
O3-	1.3562	0.8202	0.5361	2.01	-15.16	13.98	-13.88	-15.26	0.09
C3		<i>0.8106</i>	<i>0.5457</i>	<i>1.97</i>	<i>-12.70</i>	<i>14.74</i>	<i>-13.02</i>	<i>-14.43</i>	<i>0.10</i>
O4-	1.2832	0.7886	0.4946	2.59	-27.96	12.17	-18.81	-21.32	0.12
C4		<i>0.8088</i>	<i>0.4744</i>	<i>2.36</i>	<i>-22.21</i>	<i>13.10</i>	<i>-17.31</i>	<i>-18.00</i>	<i>0.04</i>
O5-	1.3544	0.8220	0.5324	2.01	-15.56	13.65	-13.90	-15.31	0.09
C5		<i>0.8207</i>	<i>0.5339</i>	<i>1.98</i>	<i>-14.68</i>	<i>12.80</i>	<i>-13.17</i>	<i>-14.31</i>	<i>0.08</i>
O7-	1.3499	0.8223	0.5278	2.02	-15.92	13.47	-14.00	-15.39	0.09
C7		<i>0.8247</i>	<i>0.5253</i>	<i>1.98</i>	<i>-15.50</i>	<i>12.17</i>	<i>-13.38</i>	<i>-14.29</i>	<i>0.06</i>
O13- C13	1.3659	0.8202	0.5458	1.98	-14.27	14.37	-13.60	-15.04	0.10
		<i>0.8226</i>	<i>0.5433</i>	<i>1.90</i>	<i>-12.01</i>	<i>13.87</i>	<i>-12.33</i>	<i>-13.55</i>	<i>0.09</i>
O14- C14	1.3511	0.8219	0.5294	2.02	-15.76	13.58	-13.97	-15.37	0.09
		<i>0.8180</i>	<i>0.5333</i>	<i>1.99</i>	<i>-14.39</i>	<i>13.28</i>	<i>-13.25</i>	<i>-14.41</i>	<i>0.08</i>
C2-	1.3830	0.6900	0.6932	2.25	-21.76	10.19	-13.91	-18.05	0.23
C3		<i>0.7023</i>	<i>0.6808</i>	<i>2.14</i>	<i>-18.85</i>	<i>9.91</i>	<i>-12.22</i>	<i>-16.53</i>	<i>0.26</i>
C2-	1.4602	0.7448	0.7155	1.92	-14.60	11.31	-11.60	-14.30	0.19
C11		<i>0.7642</i>	<i>0.6960</i>	<i>1.82</i>	<i>-13.06</i>	<i>10.49</i>	<i>-10.68</i>	<i>-12.88</i>	<i>0.17</i>
C3-	1.4347	0.7337	0.7010	2.05	-17.88	10.49	-12.74	-15.63	0.18
C4		<i>0.7114</i>	<i>0.7234</i>	<i>1.97</i>	<i>-15.83</i>	<i>10.58</i>	<i>-11.78</i>	<i>-14.63</i>	<i>0.19</i>
C6-	1.3829	0.6844	0.6985	2.18	-20.02	10.12	-13.38	-16.76	0.20
C5		<i>0.6575</i>	<i>0.7254</i>	<i>2.11</i>	<i>-17.88</i>	<i>10.23</i>	<i>-12.33</i>	<i>-15.78</i>	<i>0.22</i>

C6-	1.4090	0.6978	0.7113	2.10	-18.09	10.68	-12.76	-16.01	0.20
C7		<i>0.6744</i>	<i>0.7347</i>	<i>2.01</i>	<i>-16.05</i>	<i>10.60</i>	<i>-11.89</i>	<i>-14.76</i>	<i>0.19</i>
C8-	1.3963	0.6877	0.7086	2.15	-19.29	10.43	-13.23	-16.50	0.20
C7		<i>0.6695</i>	<i>0.7268</i>	<i>2.06</i>	<i>-17.08</i>	<i>10.36</i>	<i>-12.19</i>	<i>-15.26</i>	<i>0.20</i>
C8-	1.3878	0.6871	0.7007	2.16	-19.64	10.24	-13.27	-16.62	0.20
C9		<i>0.6589</i>	<i>0.7289</i>	<i>2.08</i>	<i>-17.31</i>	<i>10.45</i>	<i>-12.17</i>	<i>-15.58</i>	<i>0.22</i>
C10-	1.4269	0.7148	0.7121	2.00	-16.71	10.54	-12.46	-14.78	0.16
C4		<i>0.6957</i>	<i>0.7312</i>	<i>1.94</i>	<i>-14.58</i>	<i>11.16</i>	<i>-11.71</i>	<i>-14.03</i>	<i>0.16</i>
C10-	1.4211	0.6936	0.7276	2.04	-16.80	11.07	-12.35	-15.52	0.20
C5		<i>0.6924</i>	<i>0.7288</i>	<i>1.96</i>	<i>-15.18</i>	<i>10.70</i>	<i>-11.67</i>	<i>-14.21</i>	<i>0.18</i>
C10-	1.4061	0.6762	0.7299	2.05	-17.42	10.65	-12.53	-15.54	0.19
C9		<i>0.6774</i>	<i>0.7287</i>	<i>2.02</i>	<i>-15.97</i>	<i>10.72</i>	<i>-11.97</i>	<i>-14.72</i>	<i>0.19</i>
C11- C12	1.4122	0.7087	0.7034	2.05	-16.88	10.82	-12.53	-15.17	0.17
		<i>0.7096</i>	<i>0.7026</i>	<i>1.95</i>	<i>-14.38</i>	<i>11.05</i>	<i>-11.57</i>	<i>-13.86</i>	<i>0.17</i>
C11- C16	1.4072	0.7087	0.6985	2.07	-17.46	10.71	-12.80	-15.37	0.17
		<i>0.7042</i>	<i>0.7031</i>	<i>1.99</i>	<i>-15.32</i>	<i>10.64</i>	<i>-11.79</i>	<i>-14.17</i>	<i>0.17</i>
C12- C13	1.3874	0.6858	0.7017	2.16	-19.67	10.22	-13.27	-16.62	0.20
		<i>0.6731</i>	<i>0.7144</i>	<i>2.09</i>	<i>-17.47</i>	<i>10.37</i>	<i>-12.31</i>	<i>-15.52</i>	<i>0.21</i>
C13- C14	1.4099	0.7051	0.7051	2.17	-19.60	10.79	-13.23	-17.16	0.23
		<i>0.7024</i>	<i>0.7078</i>	<i>2.06</i>	<i>-18.21</i>	<i>9.72</i>	<i>-12.24</i>	<i>-15.69</i>	<i>0.22</i>
C14- C15	1.3978	0.7064	0.6915	2.13	-18.86	10.46	-13.01	-16.31	0.20
		<i>0.7227</i>	<i>0.6751</i>	<i>2.06</i>	<i>-17.08</i>	<i>10.23</i>	<i>-12.21</i>	<i>-15.10</i>	<i>0.19</i>
C16- C15	1.3895	0.6945	0.6950	2.11	-18.47	10.24	-13.05	-15.65	0.17
		<i>0.6946</i>	<i>0.6949</i>	<i>2.04</i>	<i>-15.93</i>	<i>11.12</i>	<i>-12.22</i>	<i>-14.83</i>	<i>0.18</i>
O3-	0.9832	0.7484	0.2348	2.39	-52.38	28.68	-40.15	-40.92	0.02
H3		<i>0.7396</i>	<i>0.2436</i>	<i>2.37</i>	<i>-43.43</i>	<i>32.18</i>	<i>-37.59</i>	<i>-38.02</i>	<i>0.01</i>
O5-	0.9948	0.7511	0.2437	2.34	-48.23	28.65	-38.05	-38.82	0.02
H5		<i>0.7484</i>	<i>0.2464</i>	<i>2.31</i>	<i>-41.99</i>	<i>31.81</i>	<i>-36.70</i>	<i>-37.10</i>	<i>0.01</i>

O7-	0.9864	0.7493	0.2371	2.38	-51.36	28.66	-39.59	-40.42	0.02
H7		<i>0.7463</i>	<i>0.2401</i>	<i>2.34</i>	<i>-45.89</i>	<i>31.09</i>	<i>-38.25</i>	<i>-38.74</i>	<i>0.01</i>
O13-	0.9926	0.7506	0.2420	2.35	-49.05	28.66	-38.44	-39.26	0.02
H13		<i>0.7495</i>	<i>0.2431</i>	<i>2.29</i>	<i>-42.94</i>	<i>31.13</i>	<i>-36.76</i>	<i>-37.31</i>	<i>0.01</i>
O14-	0.9788	0.7476	0.2312	2.41	-54.19	28.67	-41.01	-41.84	0.02
H14		<i>0.7368</i>	<i>0.2420</i>	<i>2.40</i>	<i>-45.67</i>	<i>32.16</i>	<i>-38.65</i>	<i>-39.18</i>	<i>0.01</i>
O1W-	0.9735	0.7385	0.2350	2.52	-53.53	30.66	-41.79	-42.40	0.01
H1W		<i>0.7314</i>	<i>0.2421</i>	<i>2.45</i>	<i>-44.39</i>	<i>31.24</i>	<i>-37.48</i>	<i>-38.15</i>	<i>0.02</i>
O1W-	0.9747	0.7387	0.2360	2.52	-53.04	30.68	-41.63	-42.09	0.01
H2W		<i>0.7352</i>	<i>0.2394</i>	<i>2.42</i>	<i>-44.18</i>	<i>31.06</i>	<i>-37.37</i>	<i>-37.88</i>	<i>0.01</i>
C6-	1.0823	0.7033	0.3789	1.85	-19.18	14.86	-16.56	-17.48	0.05
H6		<i>0.6880</i>	<i>0.3942</i>	<i>1.81</i>	<i>-15.38</i>	<i>16.69</i>	<i>-15.60</i>	<i>-16.46</i>	<i>0.05</i>
C8-	1.0798	0.7018	0.3780	1.86	-19.37	14.87	-16.66	-17.58	0.05
H8		<i>0.6846</i>	<i>0.3952</i>	<i>1.81</i>	<i>-15.40</i>	<i>16.64</i>	<i>-15.53</i>	<i>-16.50</i>	<i>0.06</i>
C12-	1.0794	0.7012	0.3782	1.86	-19.39	14.83	-16.66	-17.56	0.05
H12		<i>0.6944</i>	<i>0.3850</i>	<i>1.84</i>	<i>-16.46</i>	<i>16.81</i>	<i>-16.29</i>	<i>-16.99</i>	<i>0.04</i>
C15-	1.0832	0.7040	0.3793	1.85	-19.11	14.85	-16.53	-17.43	0.05
H15		<i>0.6883</i>	<i>0.3949</i>	<i>1.81</i>	<i>-15.62</i>	<i>16.27</i>	<i>-15.56</i>	<i>-16.34</i>	<i>0.05</i>
C16-	1.0825	0.7030	0.3795	1.85	-19.17	14.80	-16.53	-17.44	0.05
H16		<i>0.6893</i>	<i>0.3932</i>	<i>1.83</i>	<i>-16.28</i>	<i>16.05</i>	<i>-15.82</i>	<i>-16.50</i>	<i>0.04</i>

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The distances are given in Å, electron density  $\rho(\mathbf{r}_{\text{CP}})$  in  $\text{e}\text{Å}^{-3}$ . The Laplacian  $\nabla^2\rho(\mathbf{r}_{\text{CP}})$ , Hessian eigenvalues  $\lambda_1, \lambda_2, \lambda_3$  are in  $\text{e}\text{Å}^{-5}$ . The ellipticity  $\varepsilon$  is dimensionless. The distances,  $d_{\text{AB}}$ , are the same for both models because the same optimized geometry was used.

**Table S5.** Topological properties of the electron density at ring critical points of quercetin monohydrate for the TAAM\_OPT (first entry) and THEO\_OPT (second entry in italics) models. The values of  $\rho(\mathbf{r}_{\text{CP}})$  are given in  $\text{e}\text{\AA}^{-3}$  and  $\nabla^2\rho(\mathbf{r}_{\text{CP}})$  in  $\text{e}\text{\AA}^{-5}$ .

Ring label	$\rho(\mathbf{r}_{\text{CP}})$	$\nabla^2\rho(\mathbf{r}_{\text{CP}})$
A (C11...C14)	0.1210	3.59
	<i>0.1115</i>	<i>3.78</i>
B (O1...C4)	0.1169	3.62
	<i>0.1178</i>	<i>3.78</i>
C (C8...C5)	0.1037	3.56
	<i>0.1063</i>	<i>3.72</i>
D (O5...C4)	0.1065	2.85
	<i>0.1220</i>	<i>2.89</i>
E (C3...C12)	0.0687	1.69
	<i>0.0634</i>	<i>1.90</i>



**Table S6.** Monopole charges obtained for the TAAM\_OPT and THEO\_OPT models.The charge values are given in  $|e|$ .

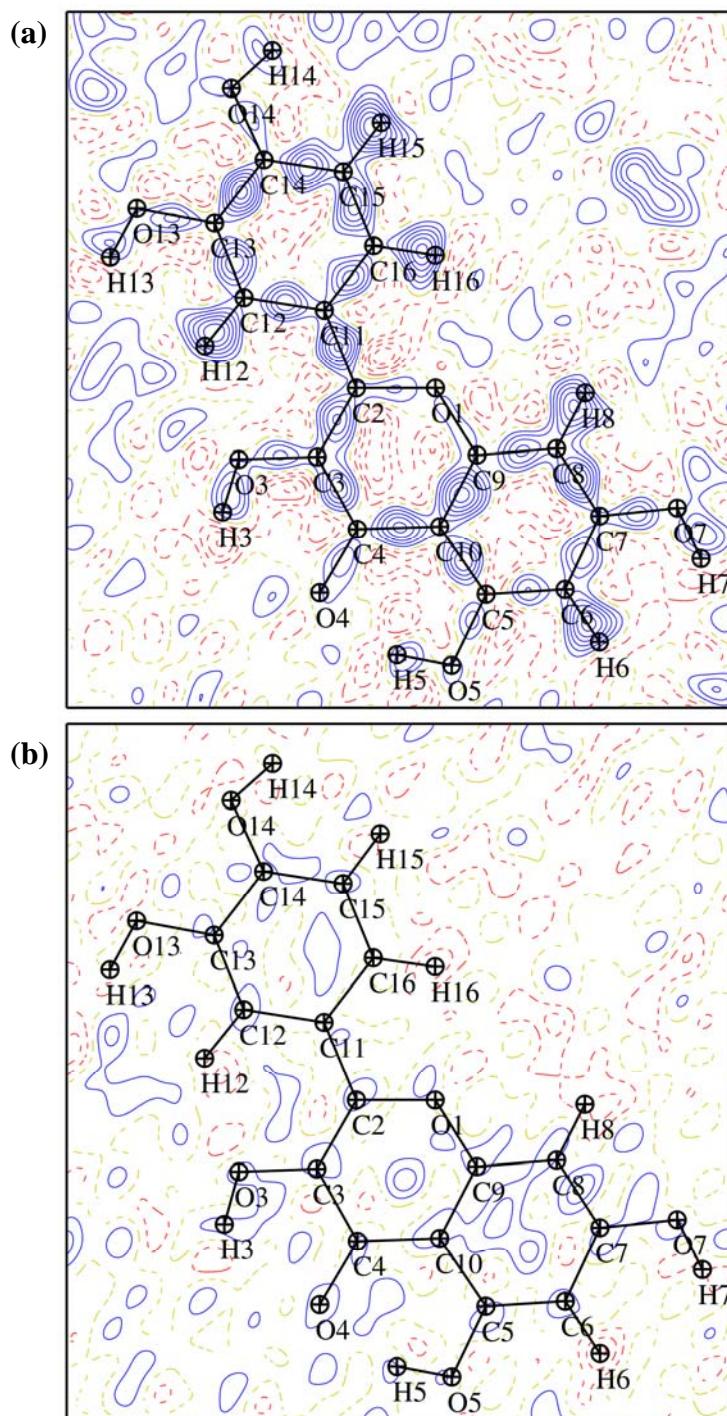
Atom Label	TAAM_OPT	THEO_OPT
O13	-0.23	-0.29
H13	0.21	0.23
O4	-0.17	-0.38
O1	-0.22	-0.20
O7	-0.23	-0.23
H7	0.21	0.23
O5	-0.23	-0.24
H5	0.21	0.21
O14	-0.23	-0.23
H14	0.21	0.20
O3	-0.23	-0.26
H3	0.21	0.21
O1W	-0.34	-0.21
H1W	0.17	0.10
H2W	0.17	0.11
C10	0.10	0.22
C2	0.04	0.04
C8	-0.03	0.18
H8	0.08	0.05
C9	0.04	-0.09
C6	-0.03	0.21
H6	0.08	0.03
C11	-0.03	-0.03
C7	0.04	-0.15
C3	0.04	0.12
C4	-0.08	0.01
C12	-0.03	0.08
H12	0.08	0.07
C13	0.04	0.01
C14	0.04	0.01
C5	0.04	-0.14
C16	-0.03	-0.04
H16	0.08	0.06
C15	-0.03	0.02
H15	0.08	0.06

**Table S7.** Computed Electrostatic Potential surface quantities for the TAAM\_OPT and THEO\_OPT models.

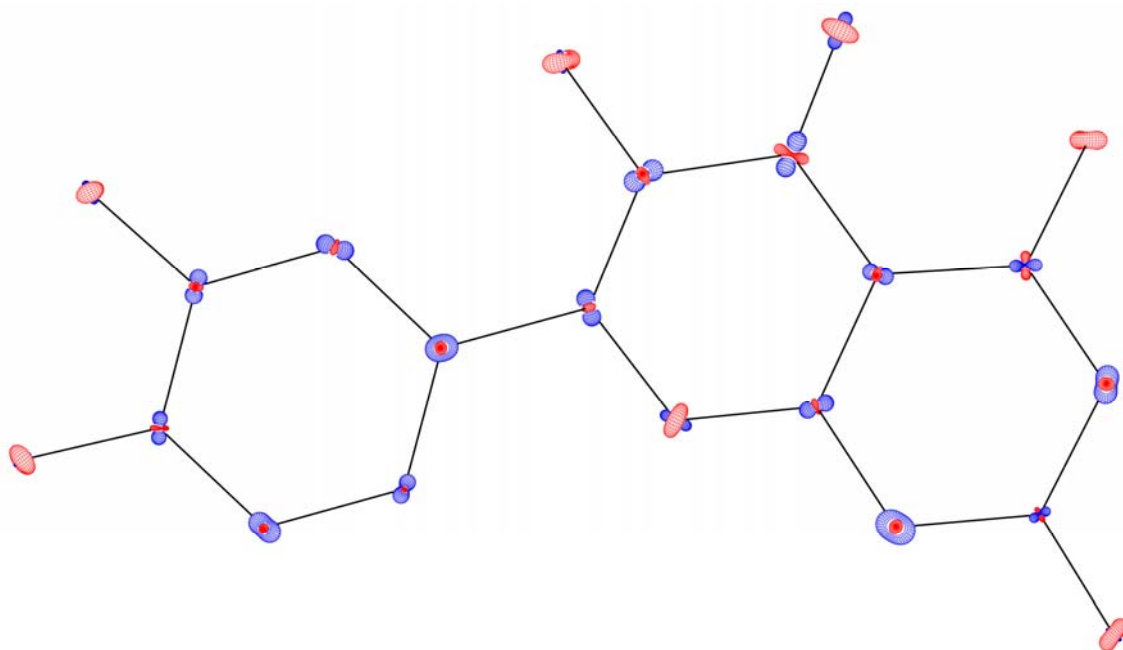
Surface quantity	TAAM_OPT	THEO_OPT
$V_{S,\max}$	50.8	45.6
$V_{S,\min}$	-48.1	-37.0
$\bar{V}_S^+$	18.2	14.2
$\bar{V}_S^-$	-11.6	-10.2
$\bar{V}_S$	1.2	2.0
$\Pi$	14.7	12.4
$\sigma_+^2$	177.4	156.7
$\sigma_-^2$	86.8	86.8
$\sigma_{\text{tot}}^2$	264.2	243.5
$\nu$	0.221	0.229

The quantities:  $V_{S,\max}$ ,  $V_{S,\min}$ ,  $\bar{V}_S^+$ ,  $\bar{V}_S^-$ ,  $\bar{V}_S$  and  $\Pi$  are given in kcal/mol;  $\sigma_+^2$ ,  $\sigma_-^2$  and  $\sigma_{\text{tot}}^2$  are in (kcal/mol)<sup>2</sup>;  $\nu$  is dimensionless. The following definitions were used to calculate surface quantities:

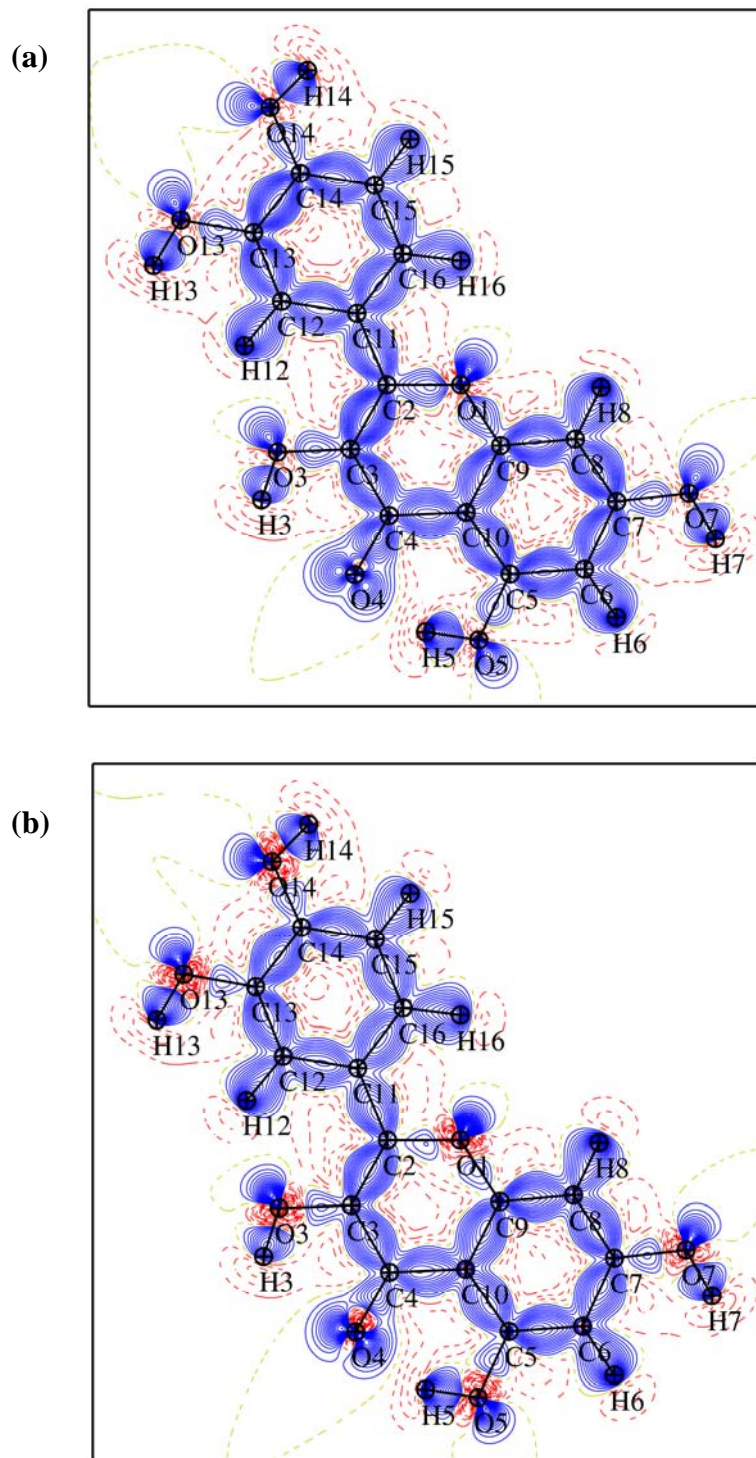
$$\begin{aligned} \bar{V}_S^+ &= \frac{1}{p} \sum_{i=1}^p V_S^+(\mathbf{r}_i), & \bar{V}_S^- &= \frac{1}{n} \sum_{i=1}^n V_S^-(\mathbf{r}_i), & \bar{V}_S &= \frac{1}{p+n} \sum_{i=1}^{p+n} V_S(\mathbf{r}_i), \\ \Pi &= \frac{1}{p+n} \sum_{i=1}^{p+n} |V_S(\mathbf{r}_i) - \bar{V}_S|, & \sigma_+^2 &= \frac{1}{p} \sum_{i=1}^p [V_S^+(\mathbf{r}_i) - \bar{V}_S^+]^2, & \sigma_-^2 &= \frac{1}{n} \sum_{i=1}^n [V_S^-(\mathbf{r}_i) - \bar{V}_S^-]^2, \\ \sigma_{\text{tot}}^2 &= \sigma_+^2 + \sigma_-^2, & \nu &= \frac{\sigma_+^2 \sigma_-^2}{(\sigma_{\text{tot}}^2)^2}. \end{aligned}$$



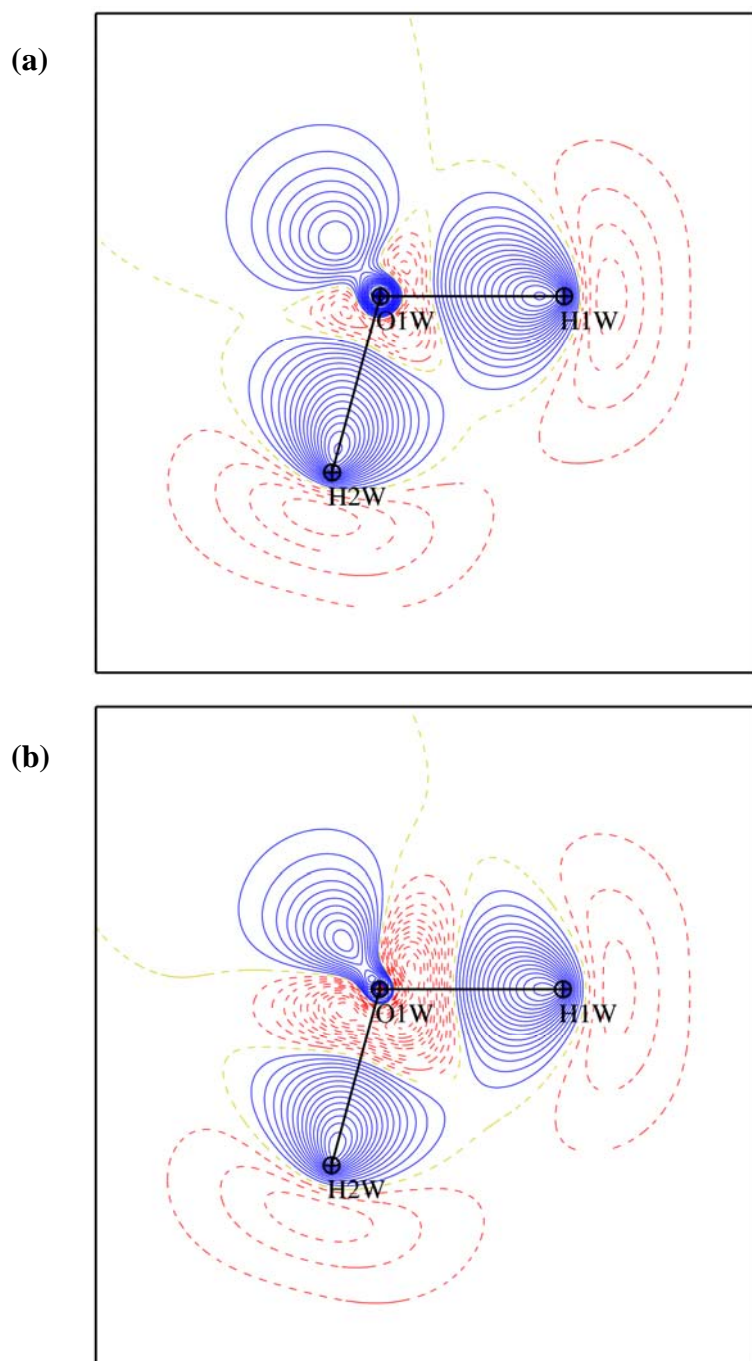
**Figure S1.** Residual experimental electron density maps in the plane of quercetin molecule for the (a) IAM\_R and (b) TAAM\_R models. Blue solid lines and red dashed lines denote positive and negative values, respectively. Contour level at  $\pm 0.05 \text{ e}\text{\AA}^{-3}$ . The zero contours are shown in yellow dashed lines.



**Figure S2.** PEANUT<sup>64</sup> representations of the ADP differences between the TAAM\_THEO\_R and TAAM\_R restrained models. The root mean square displacement differences surfaces are shown in scale of 6.15. The positive differences appear in blue, while the negative are in red. Hydrogen atoms were omitted.

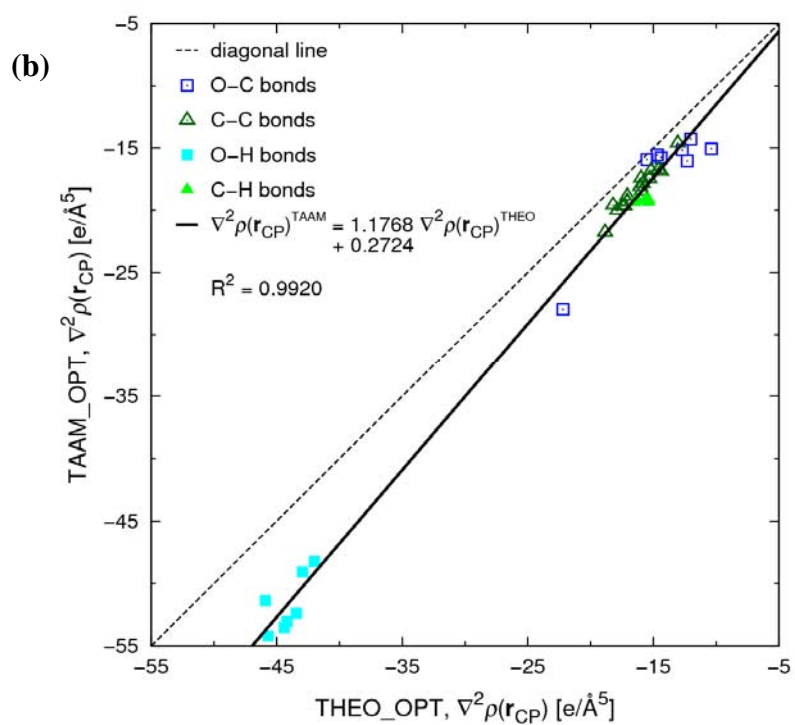
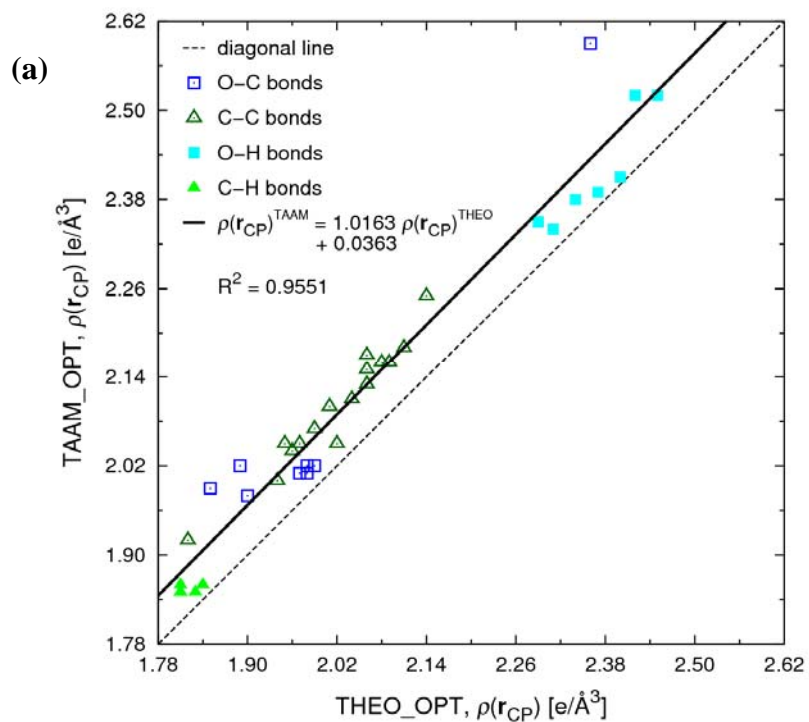


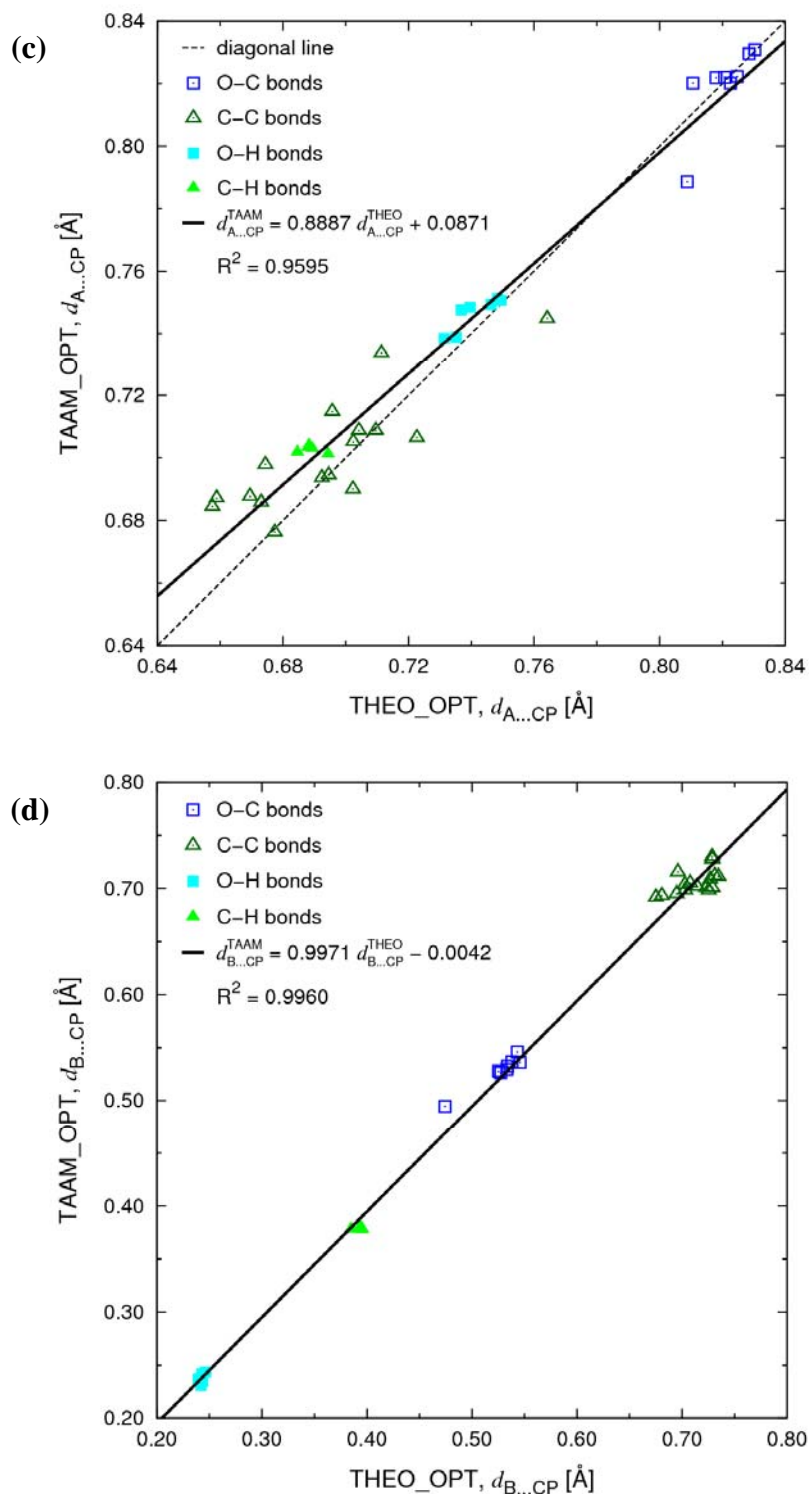
**Figure S3.** Static deformation density maps in the plane of the quercetin molecule for (a) TAAM\_OPT and (b) THEO\_OPT models. Blue solid lines and red dashed lines denote positive and negative contours, respectively. Contour level at  $\pm 0.05 \text{ e}\text{\AA}^{-3}$ . The zero contours are shown in yellow dashed lines.



**Figure S4.** Static deformation density maps for the water molecule in quercetin monohydrate presented in the plane defined by H1W, O1W and H2W atoms for (a) TAAM\_OPT and (b) THEO\_OPT models, respectively. Blue solid lines and red dashed lines denote positive and negative values, respectively. Contour level at  $\pm 0.05 \text{ e}\text{\AA}^{-3}$ . The zero contours are shown in yellow dashed lines.

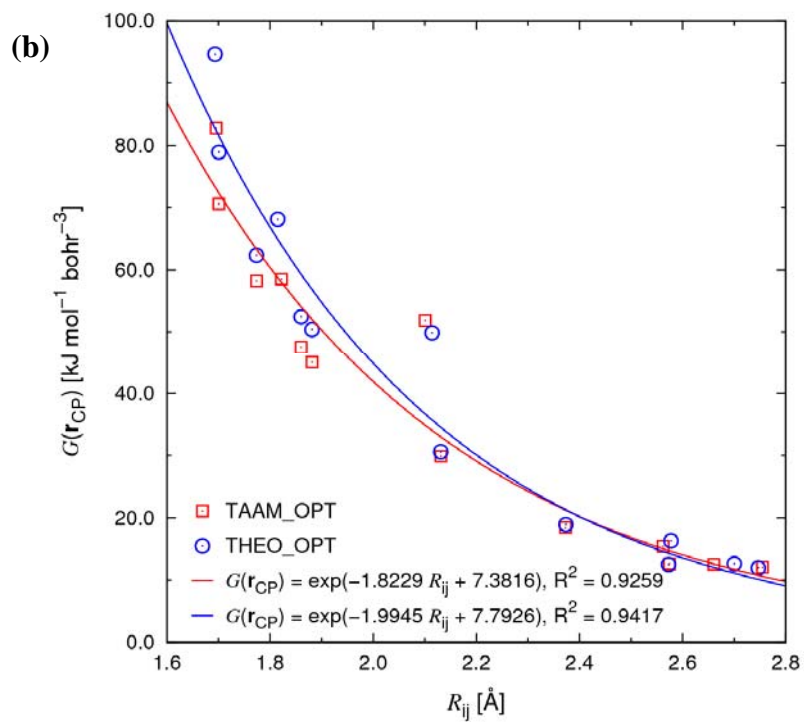
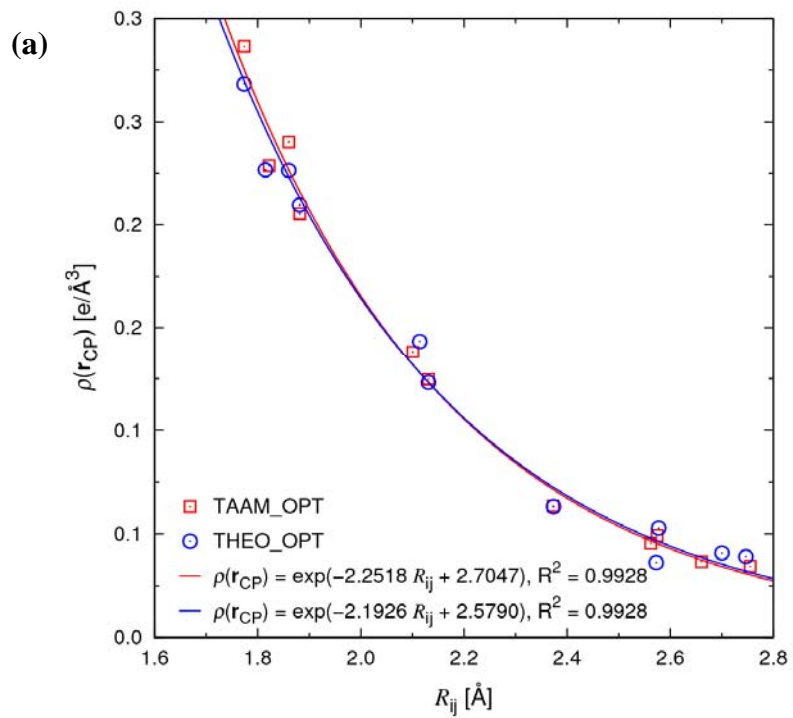


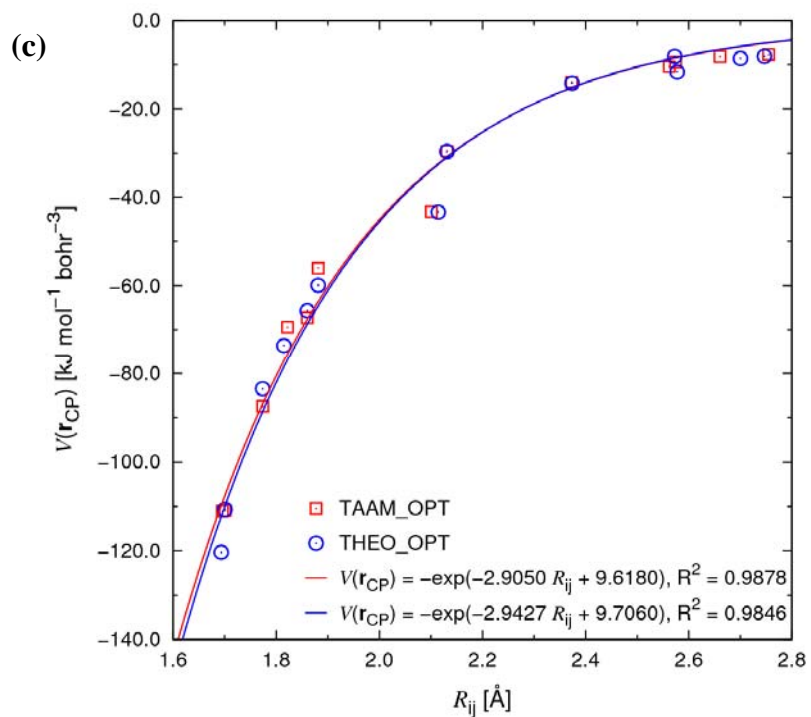




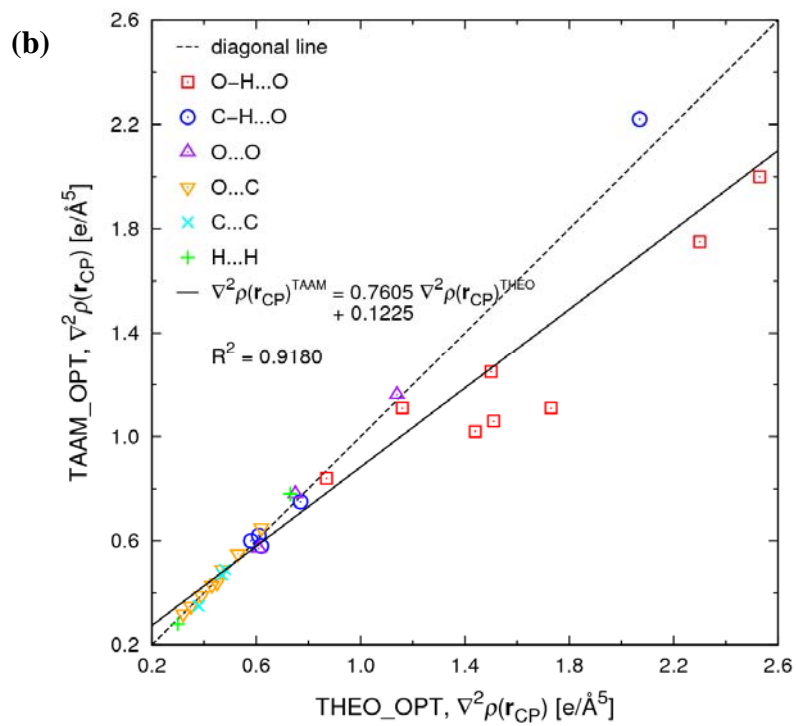
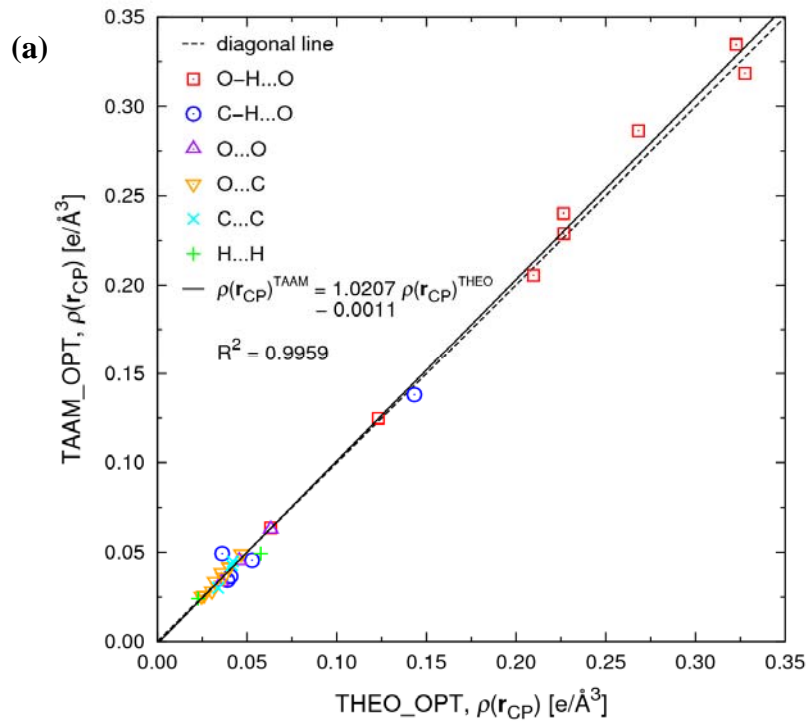
**Figure S5.** Correlation between topological properties of the total electron density of the covalent bonds for the TAAM\_OPT and THEO\_OPT models. Plots show the relative correlation of (a)  $\rho(\mathbf{r}_{CP})$ , (b)  $\nabla^2 \rho(\mathbf{r}_{CP})$  (c)  $d_{A...CP}$  and (d)  $d_{B...CP}$  values between TAAM\_OPT and THEO\_OPT.

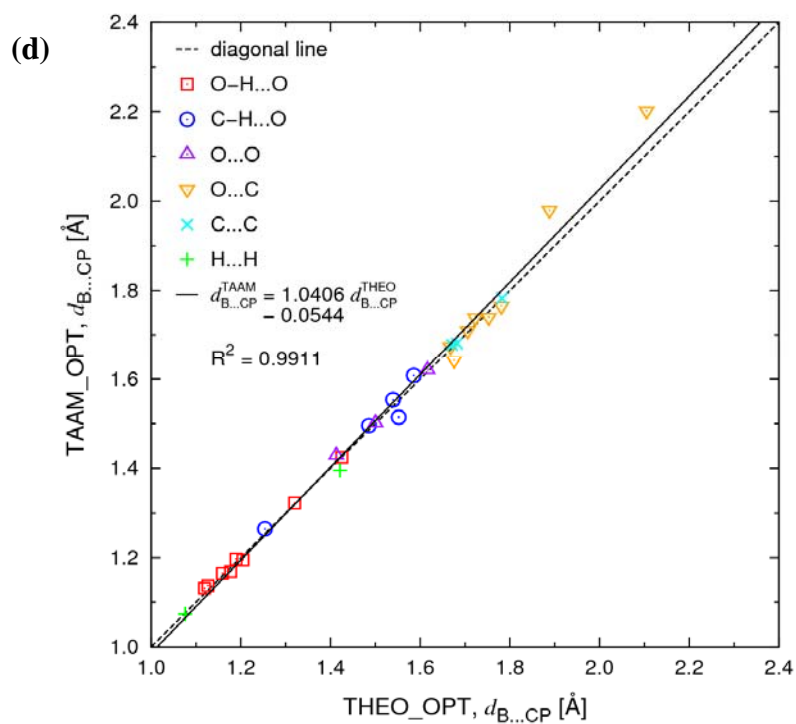
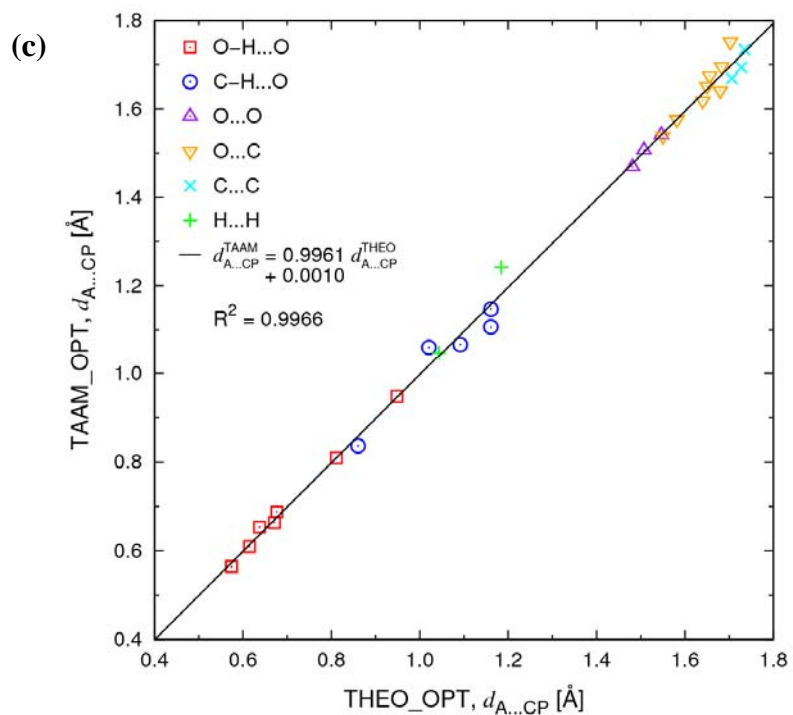




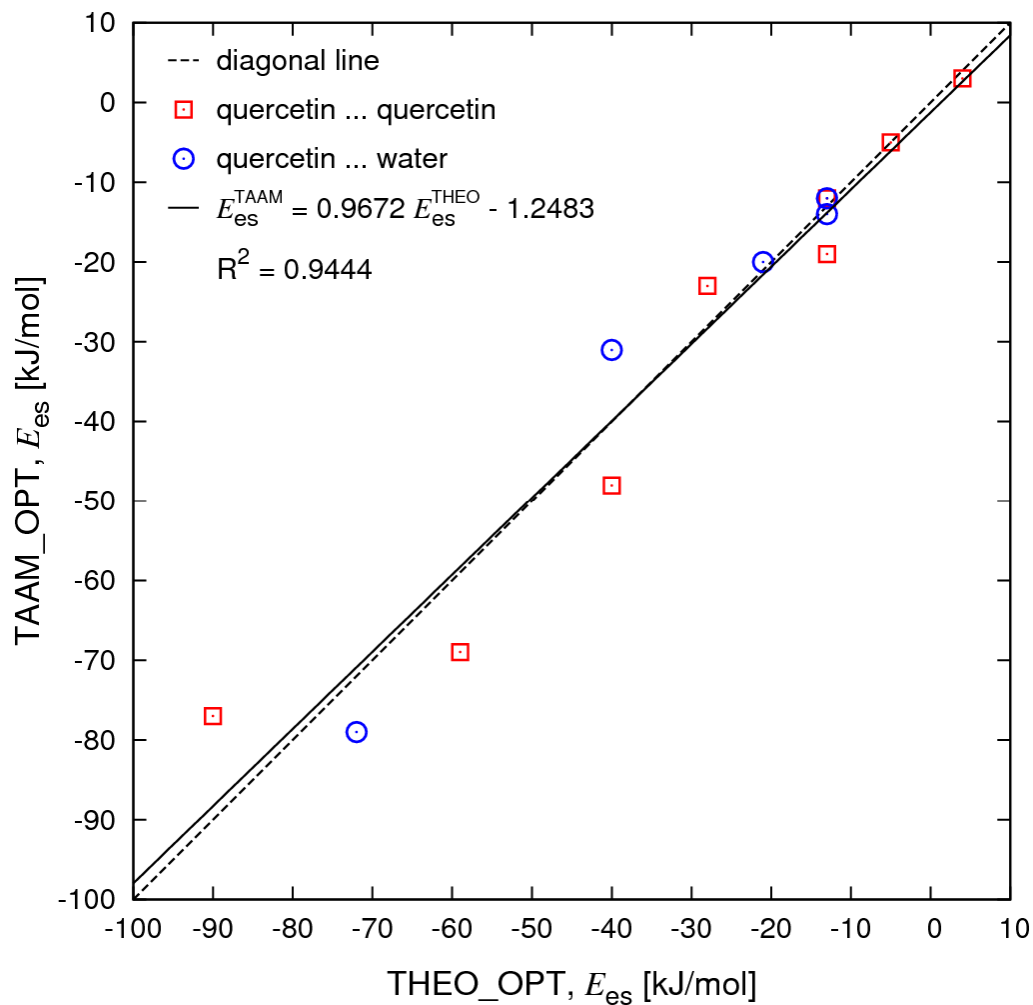


**Figure S6.** Plots of dependencies between (a)  $\rho(\mathbf{r}_{\text{CP}})$  and  $R_{ij}$ ; (b)  $G(\mathbf{r}_{\text{CP}})$  and  $R_{ij}$ ; (c)  $V(\mathbf{r}_{\text{CP}})$  and  $R_{ij}$  for X-H...O contacts. Dependencies shown for both TAAM\_OPT (blue data points) and THEO\_OPT (red data points) models.

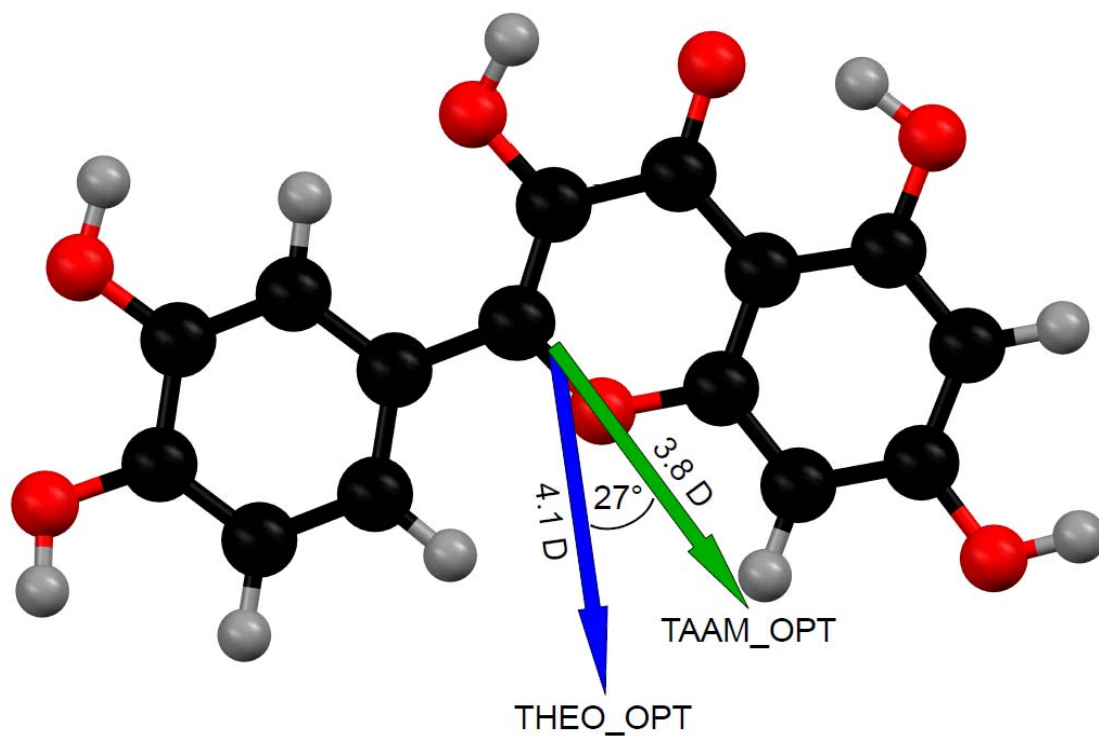




**Figure S7.** Correlation between topological properties of the total electron density for the weak interactions for the TAAM\_OPT and THEO\_OPT models. Plots show the relative correlation of (a)  $\rho(\mathbf{r}_{CP})$ , (b)  $\nabla^2 \rho(\mathbf{r}_{CP})$  (c)  $d_{A...CP}$  and (d)  $d_{B...CP}$  values between TAAM\_OPT and THEO\_OPT.



**Figure S8.** Correlation between electrostatic interaction energies obtained for different pairs of molecules for the TAAM\_OPT and THEO\_OPT models. Interactions between quercetin molecules are displayed as red squares, whereas quercetin...water short contacts are presented as blue circles.



**Figure S9.** Representation of quercetin molecule with the dipole moment vectors marked for the TAAM\_OPT (green arrow) and THEO\_OPT (blue arrow) models.