Supplementary Material

for

Mechanism of antiradical activity of coumarin-trihydroxybenzohydrazide derivatives: A comprehensive kinetic DFT study

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To quantify the molar fractions of acid-base species, the relationship between the acid constants (Ka) and the pK_a values as well as the expression for the equilibrium constant of the deprotonation process was used:

$$Ka_1 = 10^{-pKa1} \tag{1s}$$

$$Ka_2 = 10^{-pKa2} \tag{2s}$$

$$Ka_3 = 10^{-pKa3} \tag{3s}$$

$$Ka_4 = 10^{-pKa4} \tag{4s}$$

$$Ka_1 = \frac{\left[H_3A^-\right]\left[H^+\right]}{\left[H_4A\right]} \tag{5s}$$

$$Ka_2 = \frac{\left[H_2A^{2-}\right]\left[H^+\right]}{\left[H_3A^-\right]} \tag{6s}$$

$$Ka_3 = \frac{\left[HA^{3-}\right]\left[H^+\right]}{\left[H_2A^{2-}\right]} \tag{7s}$$

$$Ka_4 = \frac{\left[A^{4-}\right]\left[H^+\right]}{\left[HA^{3-}\right]} \tag{8s}$$

Molar fractions (f) of the represented acid-base species, were calculated using the following equations:

$$f(A^{4-}) = \frac{1}{1 + \beta_1\left[H^+\right] + \beta_2\left[H^+\right]^2 + \beta_2\left[H^+\right]^2 + \beta_4\left[H^+\right]^4} \tag{9s}$$

$$f(HA^{3-}) = \beta_1 \left[H^+ \right] f(A^{4-}) \tag{10s}$$

$$f(H_2A^{2-}) = \beta_2 \left[H^+ \right]^2 f(HA^{3-}) \tag{11s}$$

$$f(H_3A^-) = \beta_3 \left[H^+ \right]^3 f(H_2A^{2-}) \tag{12s}$$

where [H$^+$] represents the concentration of hydrogen ions at physiological pH in this case ([H$^+$]=3.98×10^{-8} M), while β represents global formation equilibrium constants:

$$\beta_1 = 10^{pKa4} \tag{13s}$$

$$\beta_2 = 10^{pKa4+pKa3} \tag{14s}$$

$$\beta_3 = 10^{pKa4+pKa3+pKa2} \tag{15s}$$

$$\beta_4 = 10^{pKa4+pKa3+pKa2+pKa1} \tag{16s}$$

The overall rate constants ($k_{overall}$) for the reactions between the investigated compounds and HO$^\bullet$ were calculated by the following equations:

$$k_{overall} = f(H_4A) \times k_{TOT}^{H_4A} + f(H_3A^-) \times k_{TOT}^{H_3A^-} + f(H_2A^{2-}) \times k_{TOT}^{H_2A^{2-}} + f(HA^{3-}) \times k_{TOT}^{HA^{3-}} + f(A^{4-}) \times k_{TOT}^{A^{4-}} \tag{17s}$$

Due to the very small proportion of acid-base species at physiological pH, eq. 17s can be represented as:

$$k_{overall}^{Comp1} = f(H_4A) \times k_{TOT}^{H_4A} + f(H_3A^-) \times k_{TOT}^{H_3A^-} + f(H_2A^{2-}) \times k_{TOT}^{H_2A^{2-}}$$

(18s)

$$k_{overall}^{Comp2} = f(H_4A) \times k_{TOT}^{H_4A} + f(H_3A^-) \times k_{TOT}^{H_3A^-}$$

(19s)

where k_{TOT} represent the sum of the favorable reaction pathways for the investigation species and HO$^\bullet$:

$$k_{TOT} = k_{HAT} + k_{RAF} + k_{SPL} + k_{ET}$$

(20s)

Antioxidative capacity r^T of the examined antioxidant relative to trolox (**Tx**)/gallic acid (**GA**) ($k_{overall}^{Ref}$) was calculated using the formula:

$$r^T = \frac{k_{overall}}{k_{overall}^{Ref}}$$

(21s)

The relative amounts of products (%), i.e. branching ratios (Γ_i) were determined:

$$\Gamma_i = \frac{k_i}{k_{overall}}$$

(22s)

to evaluate which of the mechanistic pathways, i, is dominant.

Standard mechanisms of antioxidant action between neutral forms of the investigated compounds and HO$^\bullet$ radical:

HAT/PCET: $H_4A + HO^\bullet \rightarrow H_3A^\bullet + H_2O$ (23s)

RAF: $H_4A + HO^\bullet \rightarrow [H_4A\text{-}OH]^\bullet$ (24s)

SET: $H_4A + HO^\bullet \rightarrow H_4A^{\bullet+} + HO^-$ (25s)

PT: $H_4A^{\bullet+} + HO^- \rightarrow H_3A^\bullet + H_2O$ (26s)

SPL: $H_4A + HO^- \rightarrow H_3A^- + H_2O$ (27s)

ET: $H_3A^- + HO^\bullet \rightarrow H_3A^\bullet + HO^-$ (28s)

where H_4A, $H_3A^\bullet$, $[HO\text{-}H_4A]^\bullet$, $H_4A^{\bullet+}$, and H_3A^- denote neutral species, its radical, radical adduct, radical cation, and anion.
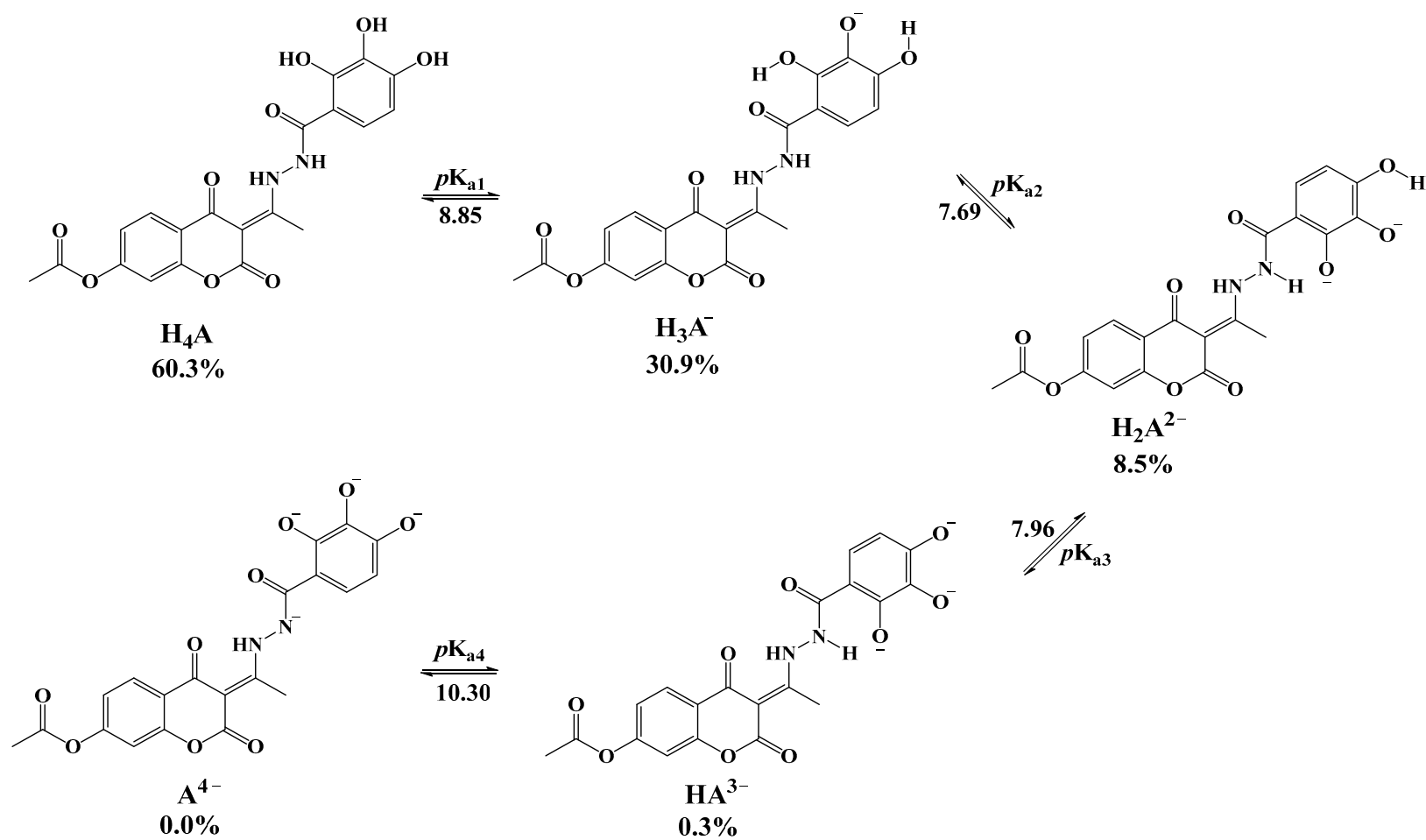
Figure S1. Deprotonation route, estimated pK_a values and molar fractions of acid-base species of **1** at physiological pH (7.4)
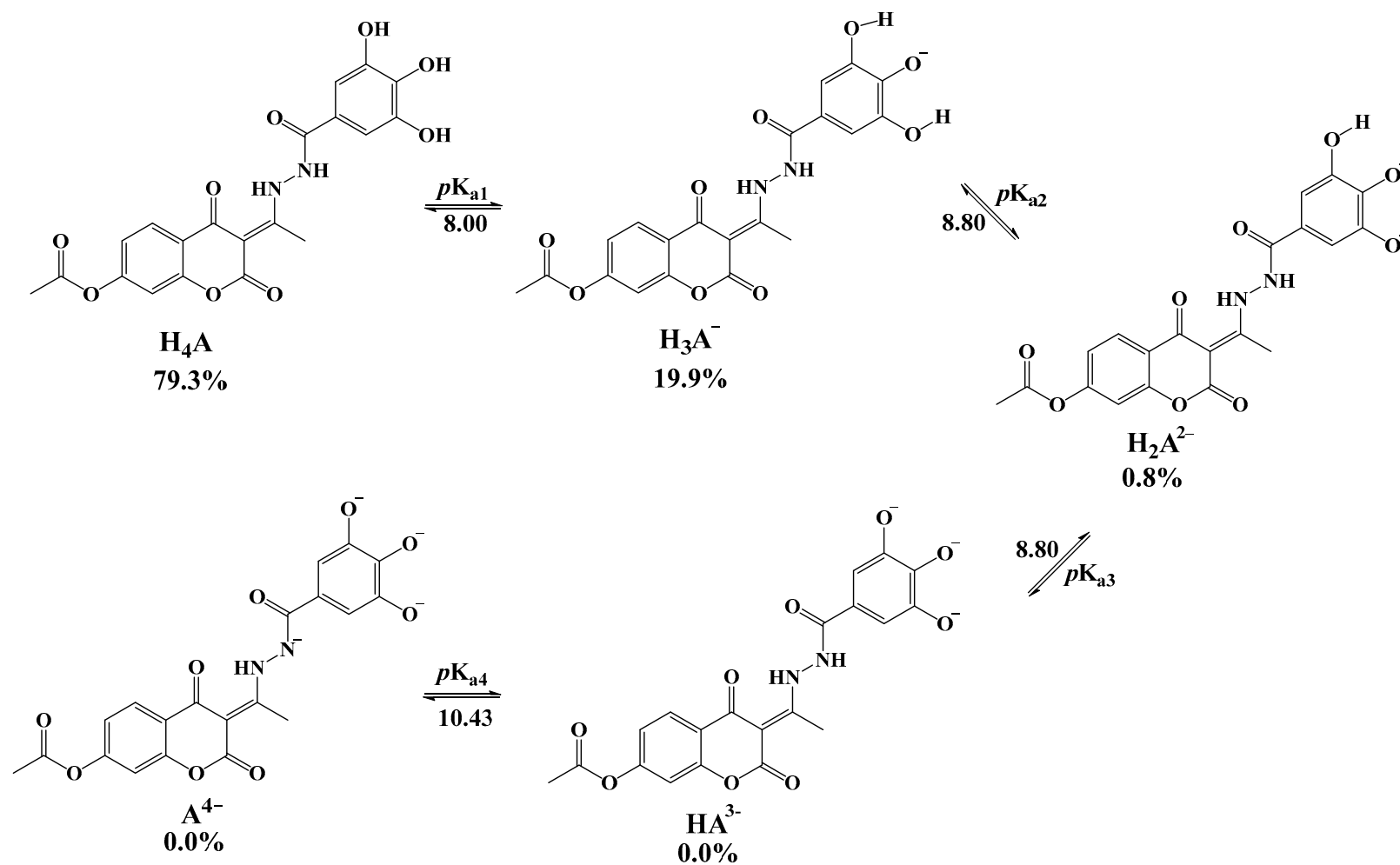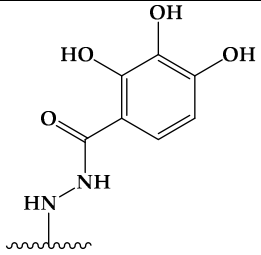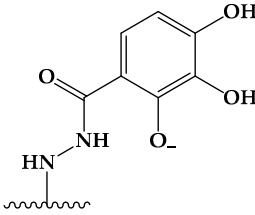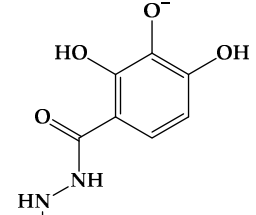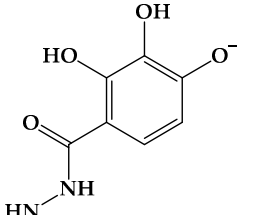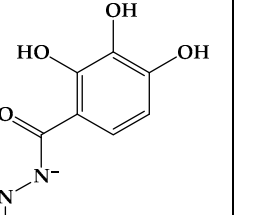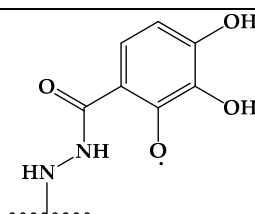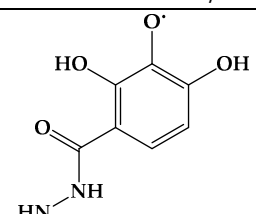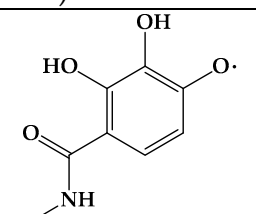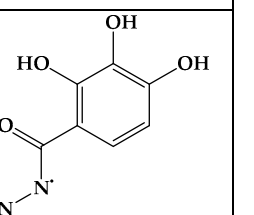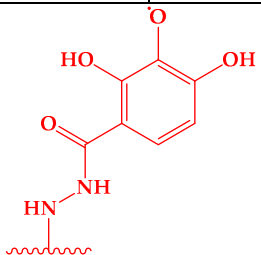
Figure S2. Deprotonation route, estimated pKa values and molar fractions of acid-base species of **2** at physiological pH (7.4)

Table S1. 2D geometries of the investigated acid-base species of **1** compound that participate in the various investigated mechanisms of antiradical action. The black color describes the thermodynamically favored products obtained between neutral species **H₄A** and HO•. The red color describes the favored products obtained between monoanionic species **H₃A⁻(H₃A•)** and HO•, while the blue color describes favored products obtained between dianionic species **H₂A²⁻(H₂A•⁻)** and HO•.

H₄A

SPLET (H₄A)			
2″-OH	3″-OH	4″-OH	N2-H

HAT/PCET (H₄A)			
2″-OH	3″-OH	4″-OH	N2-H

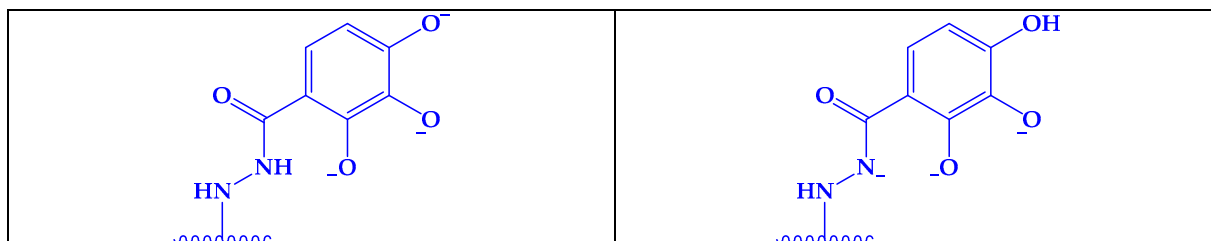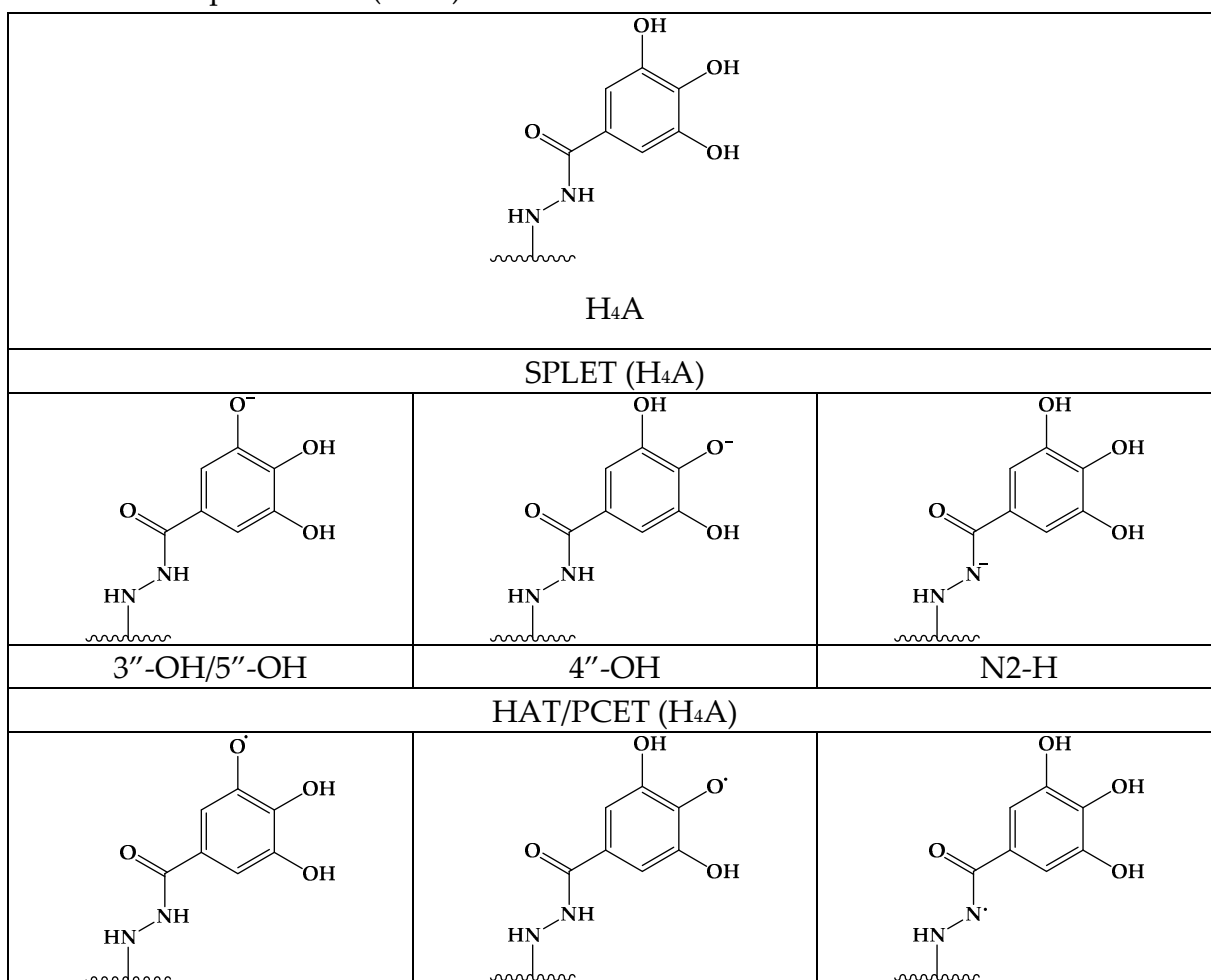| $H_3A^\bullet$ / H_3A^-		
HAT/PCET ($H_3A^\bullet$ / H_3A^-)		
2″-OH	4″-OH	N2-H
PT ($H_3A^\bullet$ / H_3A^-)		
2″-OH	4″-OH	N2-H
H_2A^{2-} / $H_2A^{\bullet-}$		
HAT/PCET (H_2A^{2-} / $H_2A^{\bullet-}$)		
4″-OH		2-NH/2″-OH
PT (H_2A^{2-} / $H_2A^{\bullet-}$)		

Table S2. 2D geometries of the investigated acid-base species of compound **2** that participate in the various investigated mechanisms of antiradical action. The black color describes the thermodynamically favored products obtained between neutral species **H₄A** and HO•. The red color describes the favored products obtained between monoanionic species **H₃A⁻**(**H₃A•**) and HO•.

H₄A

SPLET (H₄A)		
3″-OH/5″-OH	4″-OH	N2-H

HAT/PCET (H₄A)		

H_3A^-

HAT/PCET ($H_3A^{\bullet}$/ H_3A^-)

3″-OH	5″-OH	N2-H

PL ($H_3A^{\bullet}$/ H_3A^-)

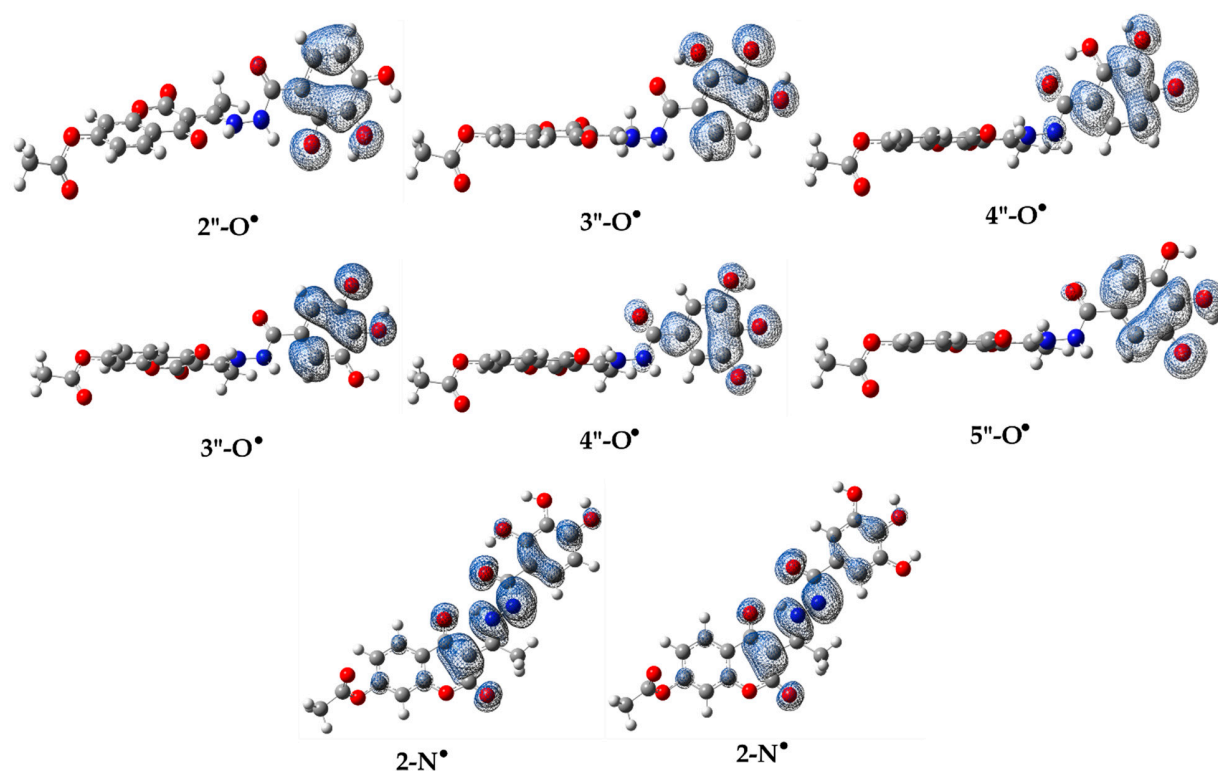| 3″-OH	5″-OH	N2-H

Figure S3. Spin density distribution maps (0.002 electrons/bohr^3) of formed radical species formed in the reaction between **H₄A** of compounds **1** (a,c) and **2** (b,d) and HO$^\bullet$ radical. Blue color represents positive spin density values.
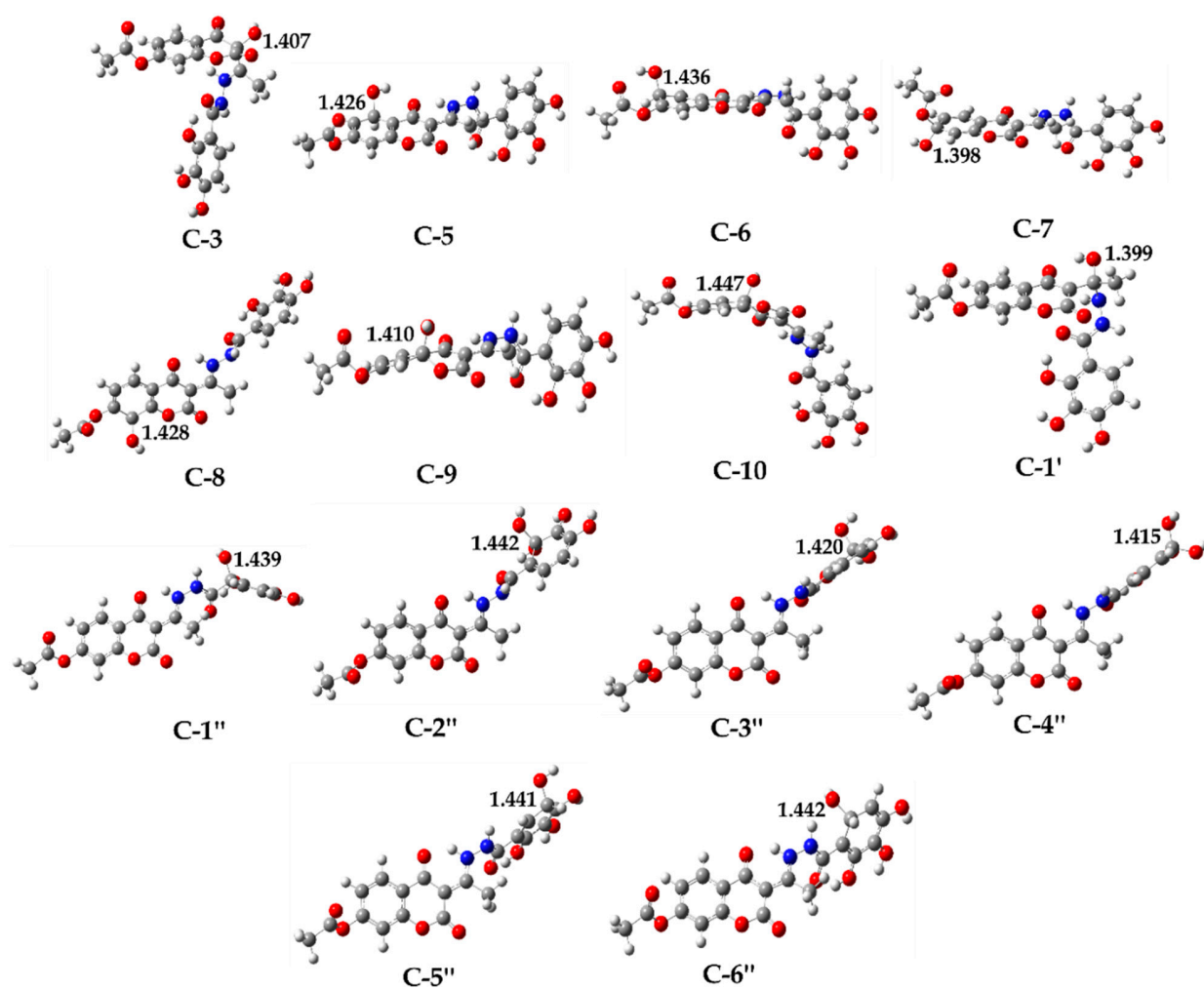
Figure S4. Optimized geometries of radical adducts with characteristic intramolecular distances (Å) formed between **H₄A** of compound **1** and HO•
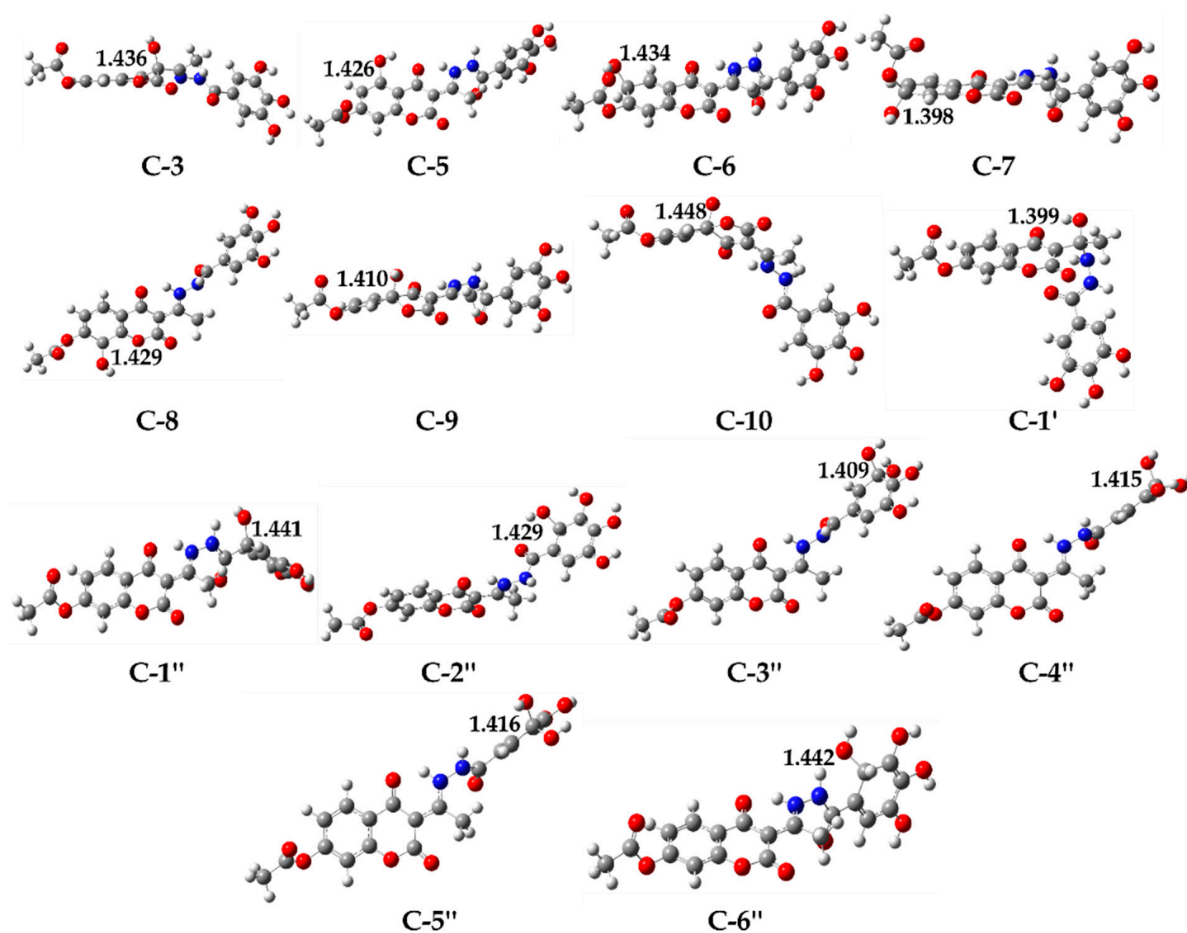
Figure S5. Optimized geometries of radical adducts with characteristic intramolecular distances (Å) formed between **H₄A** of compound **2** and HO•

a)

2"-O⁻   3"-O⁻   4"-O⁻

b)

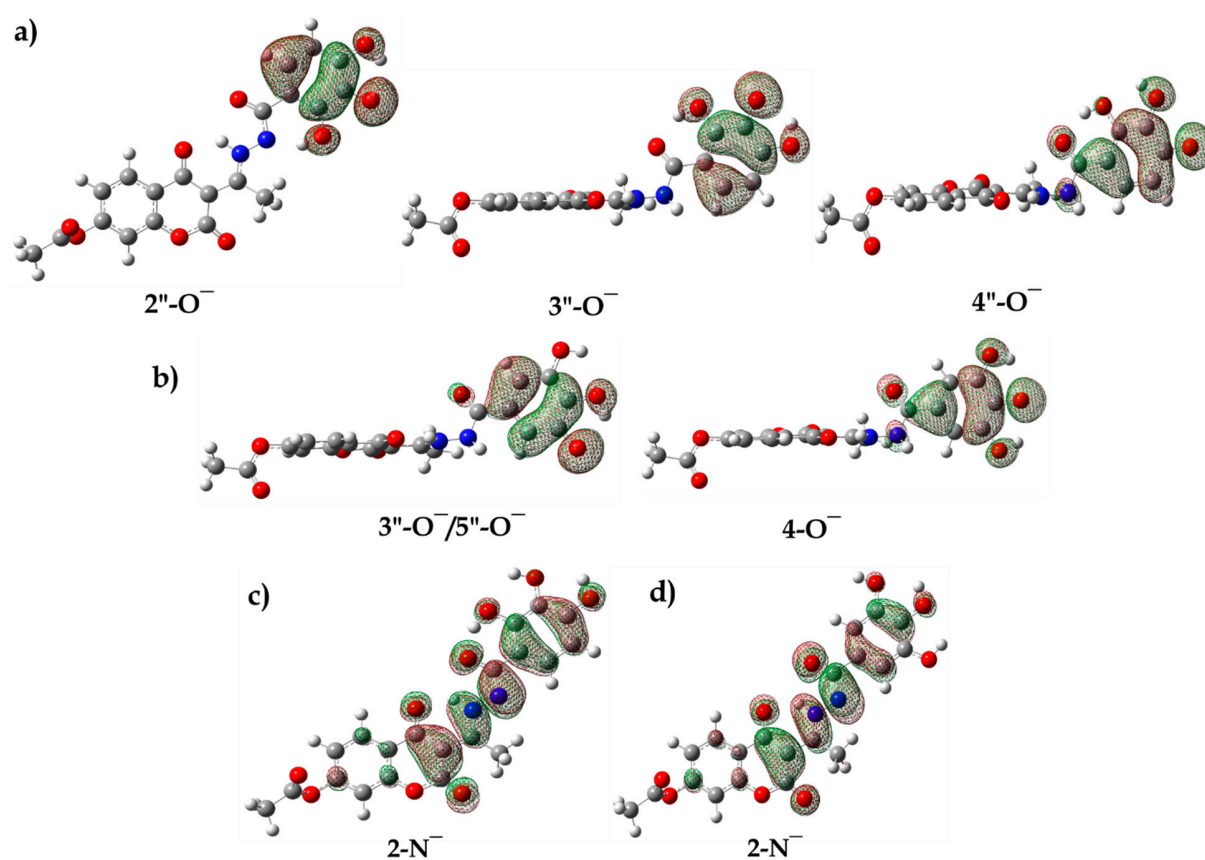3"-O⁻/5"-O⁻   4-O⁻

c)   d)

2-N⁻   2-N⁻

Figure S6. HOMO (*Highest Occupied Molecular Orbital*) orbitals of formed phenoxide anions formed in the reaction between **H₄A** of compounds **1** (a,c) and **2** (b,d) and HO⁻
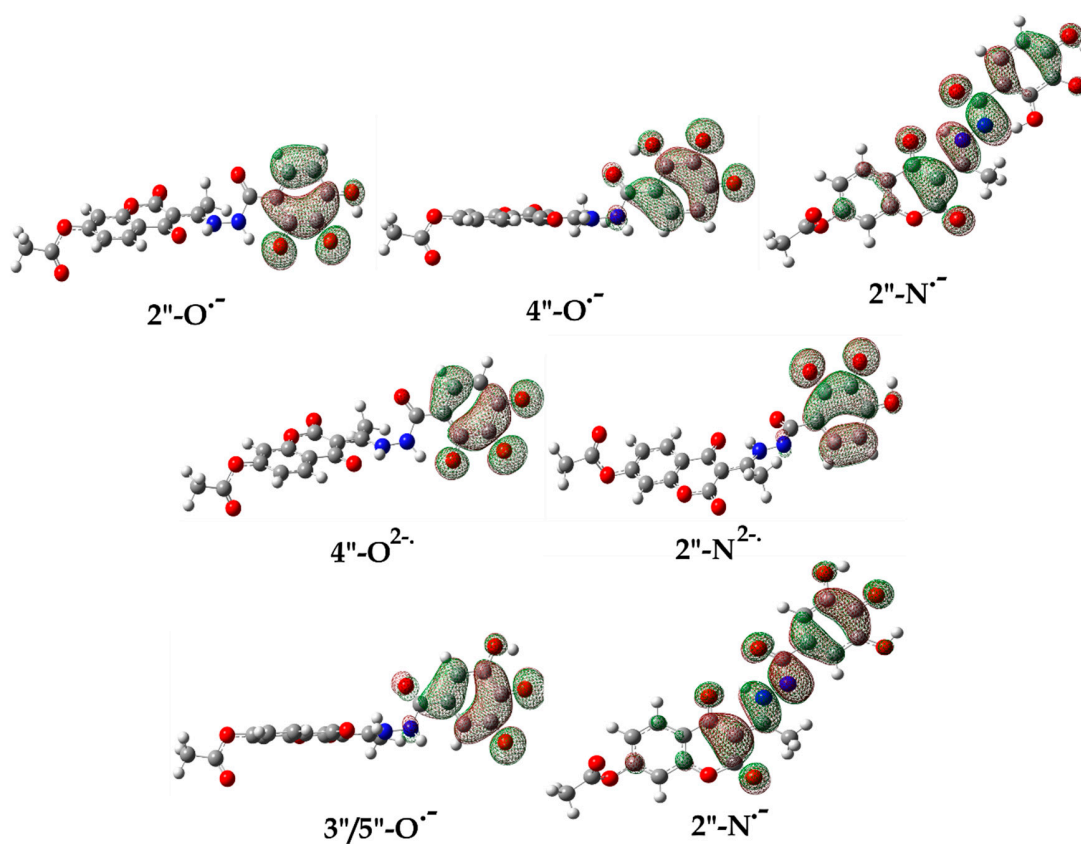
Figure S7. HOMO orbitals for the corresponding **H₂A·⁻** nad **H₂A²·⁻** species formed in the reaction of proton transfer from **H₃A·** (top) and **H₃A·⁻** (middle) of compound **1** to HO⁻ as well as from **H₃A·** of compound **2** (bottom) to HO⁻.
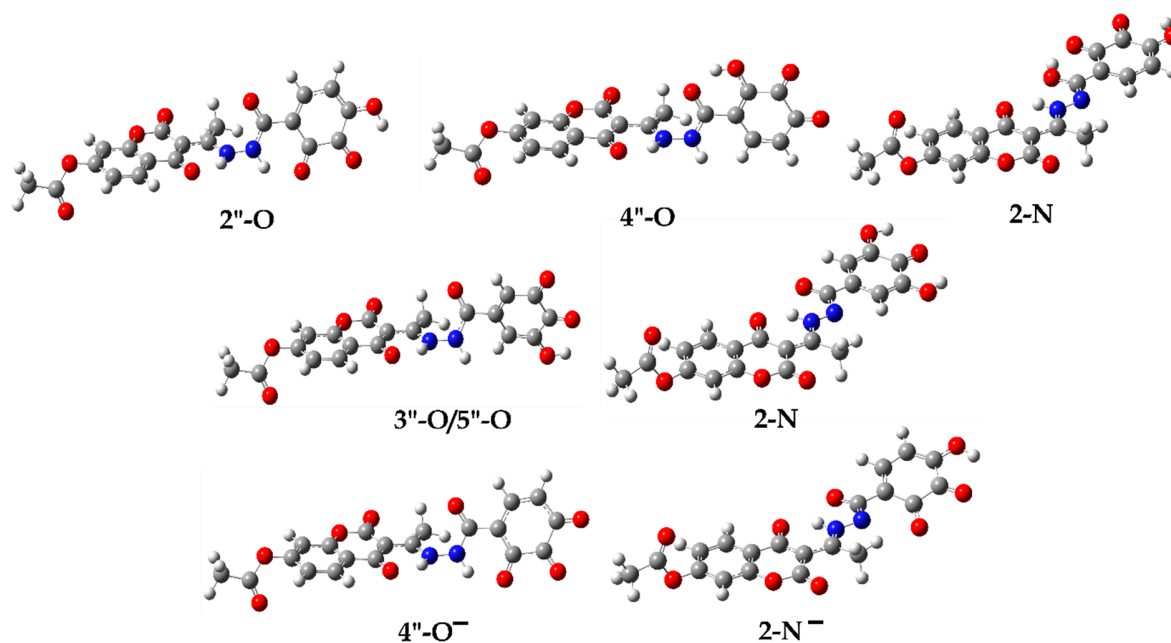
Figure S8. Optimized geometries of neutral H_2A products of compounds **1** (top) and **2** (middle) and monoanionic HA^- product (bottom) formed in the reaction between $H_3A^{\bullet}$ and $H_2A^{\bullet-}$ and $HO^{\bullet}$.

Figure S9. Optimized geometries of adducts with characteristic intramolecular distances (Å) formed between **H₃A•** of compound **1** and HO•

Figure S10. Optimized geometries of adducts with characteristic intramolecular distances (Å) formed between **H₃A•** of compound **2** and HO•
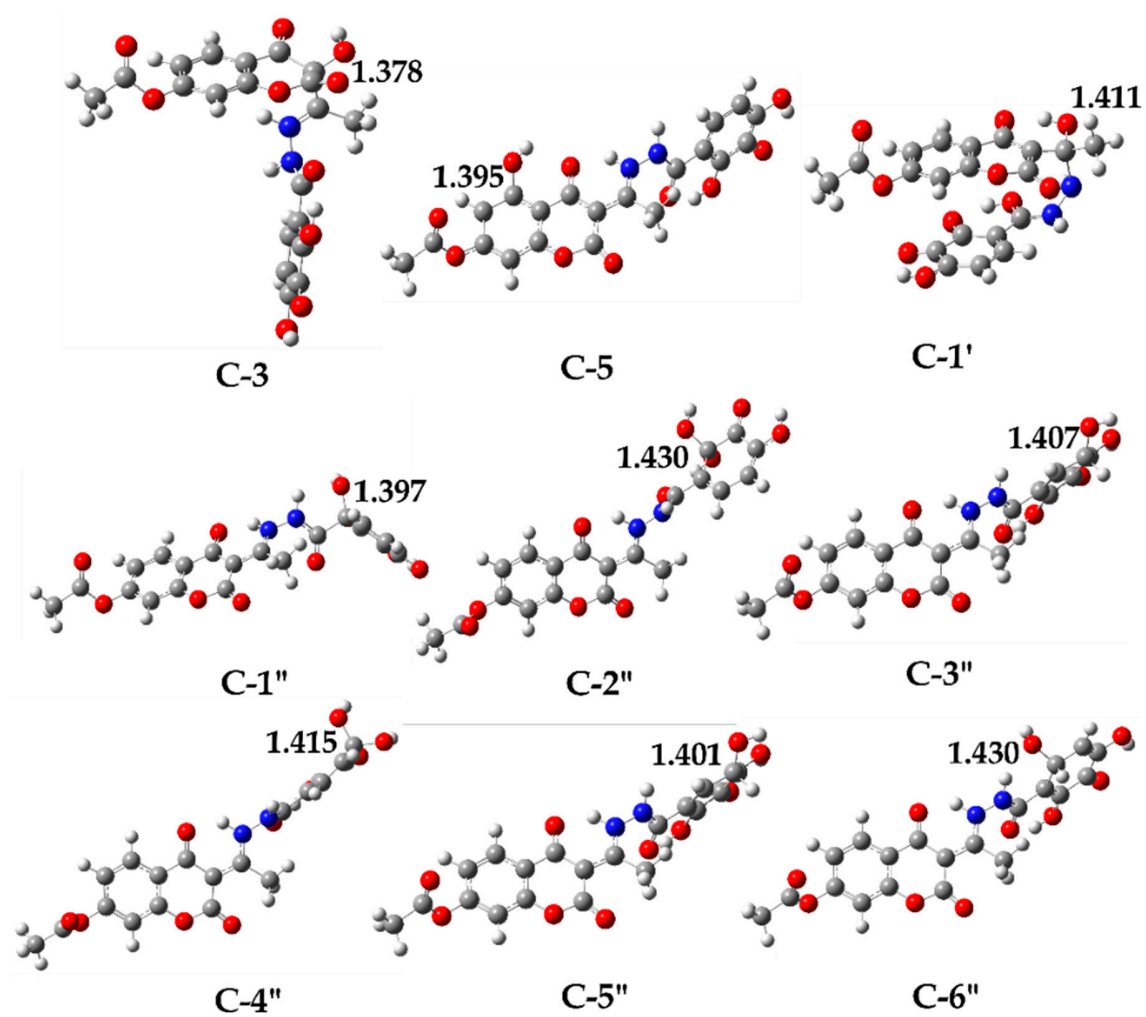
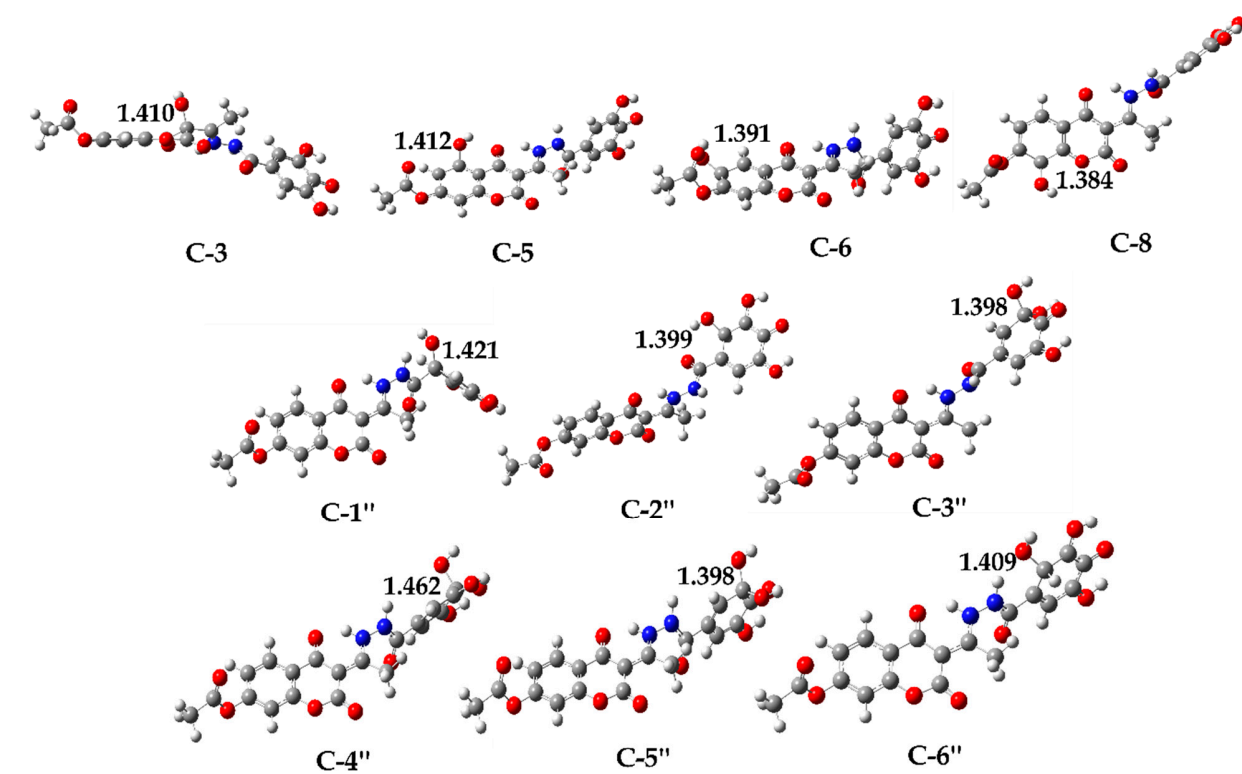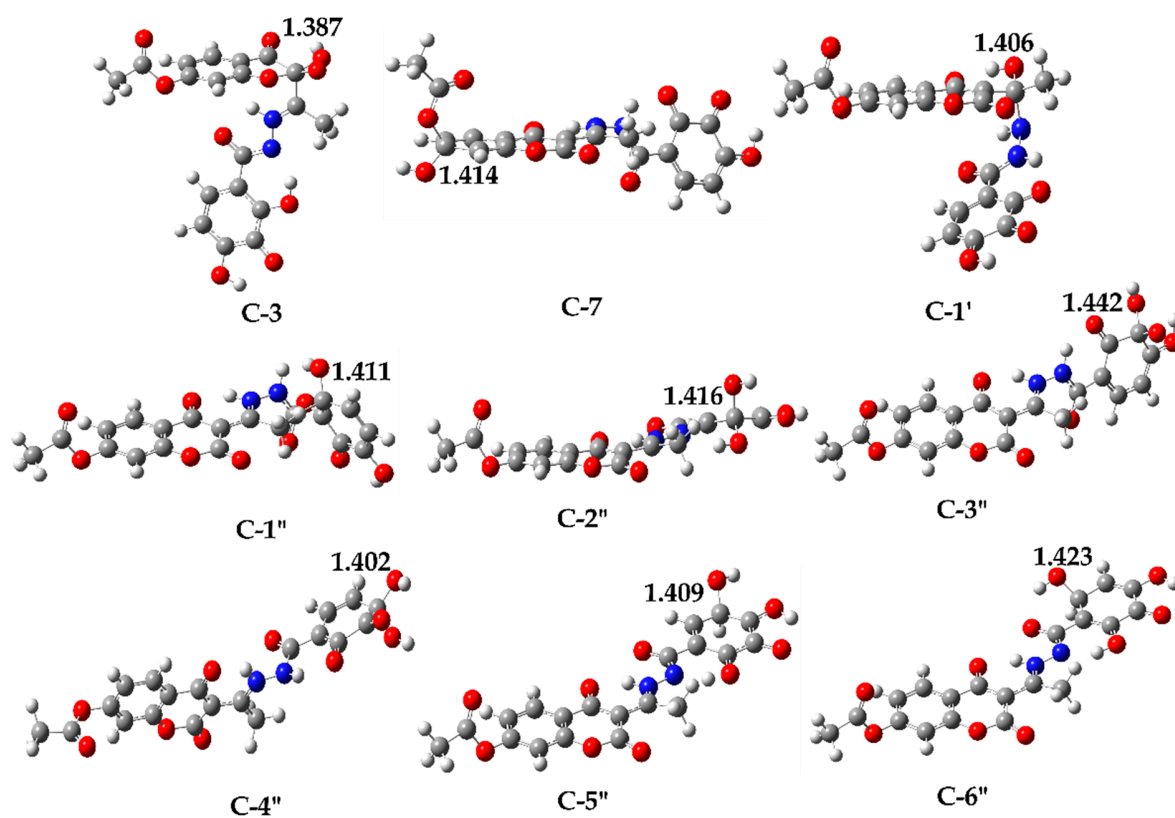**Figure S11.** Optimized geometries of adducts with characteristic intramolecular distances (Å) formed between $H_3A^{\bullet-}$ of compound **1** and $HO^{\bullet}$

Table S3. Estimated values of kinetic parameters: activation energy (ΔG_a, kJ mol^{-1}), rate constants of the bimolecular chemical reactions (M^{-1}s^{-1}) between the investigated compounds and HO$^\bullet$ estimated by the Transition State Theory (k_{TST}).

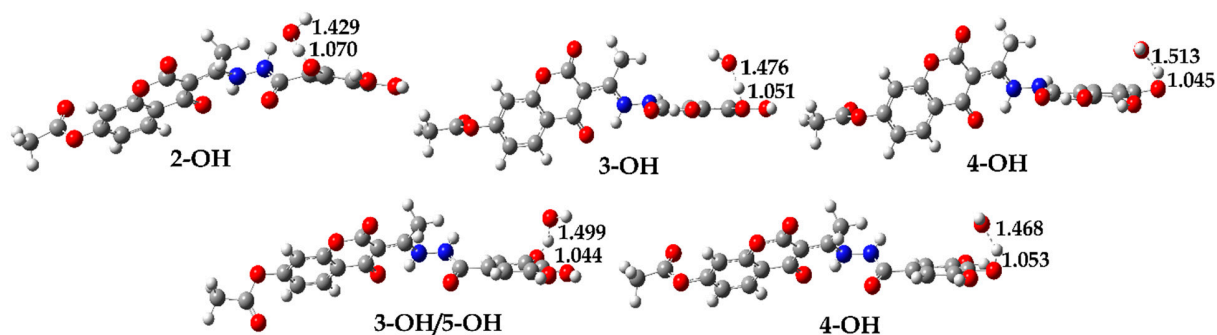| HAT/PCET				
1/2	**1 (H₄A)**		**2 (H₄A)**	
Position	ΔG_a^{HAT}	k_{TST}^{HAT}	ΔG_a^{HAT}	k_{TST}^{HAT}
2"-OH/3"-OH	58	9.53×10^3	59	6.79×10^3
3"-OH/5"-OH	41	8.26×10^6		
4"-OH/4"-OH	55	6.28×10^4	54	6.28×10^4
	RAF			
Position	**1 (H₄A)**		**2 (H₄A)**	
	ΔG_a^{RAF}	k_{TST}^{RAF}	ΔG_a^{RAF}	k_{TST}^{RAF}
C-3	38	3.29×10^7	48	6.19×10^5
C-5	51	1.88×10^5	56	2.72×10^4
C-6	38	3.26×10^7	45	2.13×10^6
C-7	60	1.60×10^3	63	1.53×10^3
C-8	39	2.59×10^7	41	9.33×10^6
C-9	50	2.27×10^5	53	7.10×10^4
C-10	43	4.01×10^6	47	9.34×10^5
C-1'	42	7.23×10^6	45	1.80×10^6
C-1"	37	5.60×10^7	35	1.02×10^8
C-2"	39	2.71×10^7	29	1.40×10^9
C-3"	22	1.91×10^{10}	47	9.00×10^5
C-4"	38	3.17×10^7	29	1.53×10^9
C-5"	38	3.38×10^7	39	2.08×10^7
C-6"	28	2.06×10^7	25	6.80×10^9

Figure S12. Optimized transition state geometries for HAT/PCET reaction pathways between neutral **H4A** of compound **1** (top) and **2** (bottom) and HO• with characteristic intraatomic distances (Å)
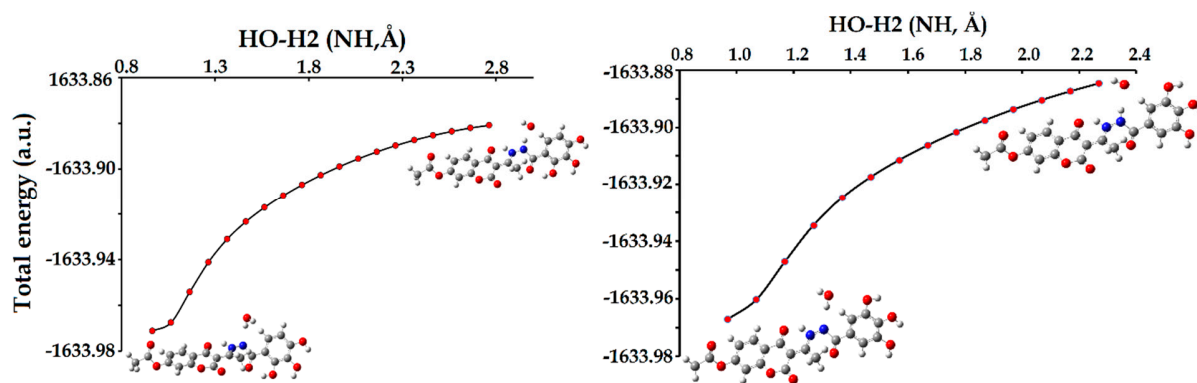

Figure S13. Dependence of the total energy (a.u.) on the HO–H2 characteristic distance (Å) for the HAT/PCET mechanism between neutral **H4A** of compound **1** (left) and **2** (right) and HO•
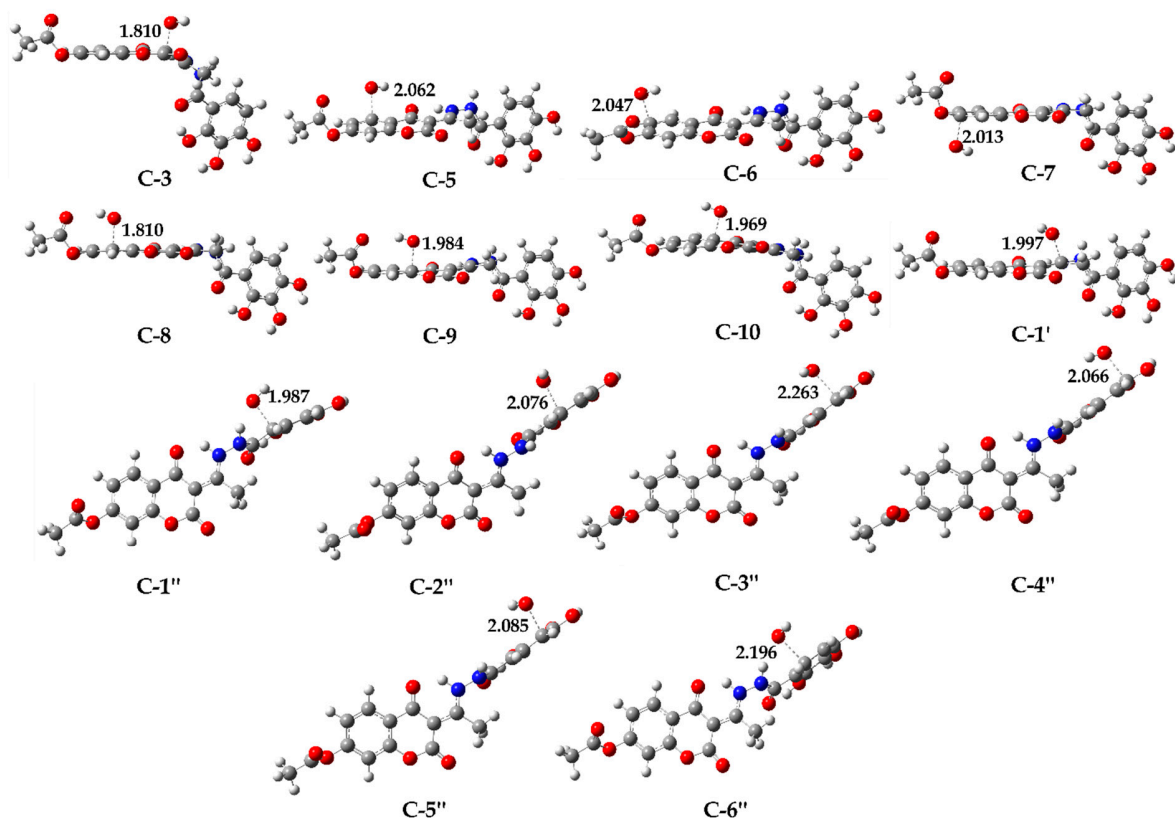
Figure S14. Optimized transition state geometries for the RAF mechanism between neutral **H₄A** of compound **1** and HO• at the M06-2X/6-311++G(d,p) level of theory with characteristic intermolecular distances (Å)
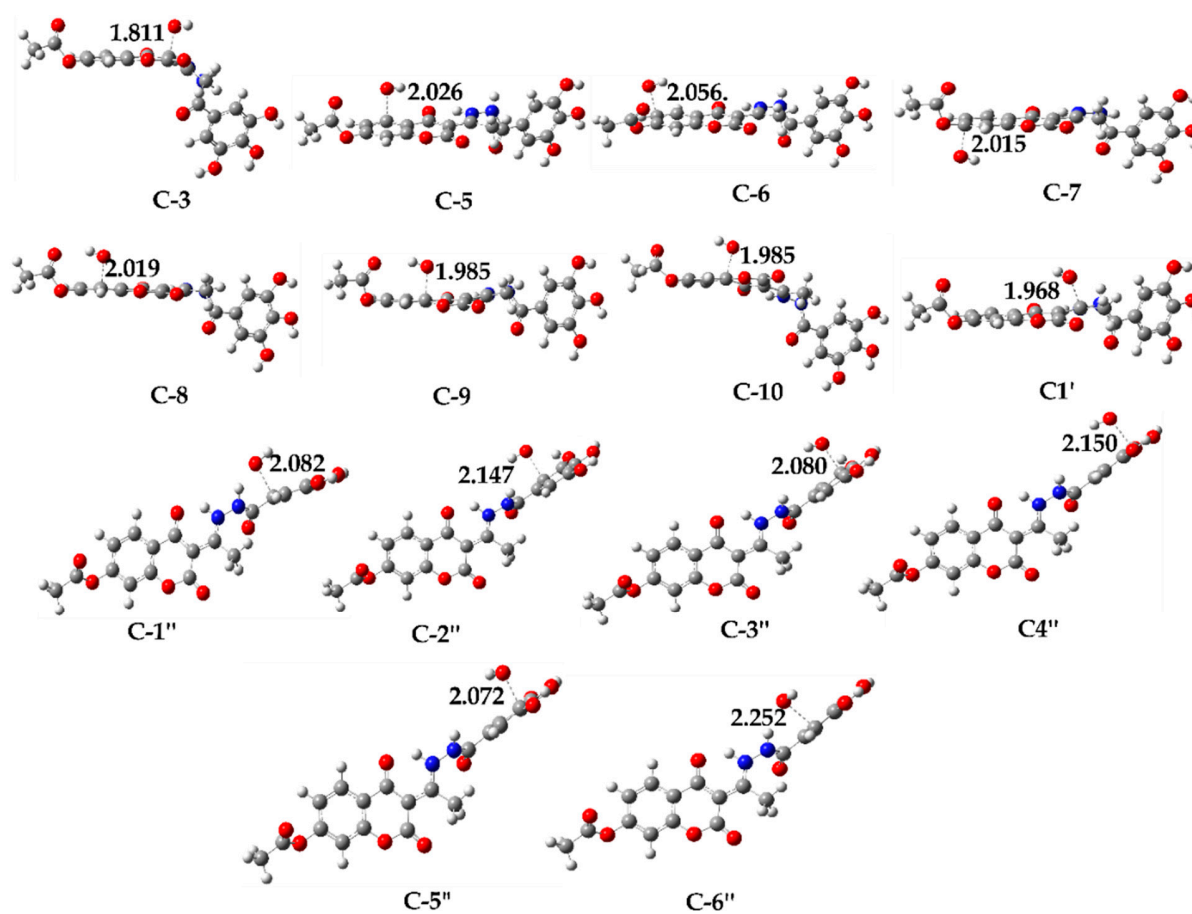
Figure S15. Optimized transition state geometries for the RAF mechanism between neutral **H₄A** of compound **2** and HO• at the M06-2X/6-311++G(d,p) level of theory with characteristic intermolecular distances (Å)
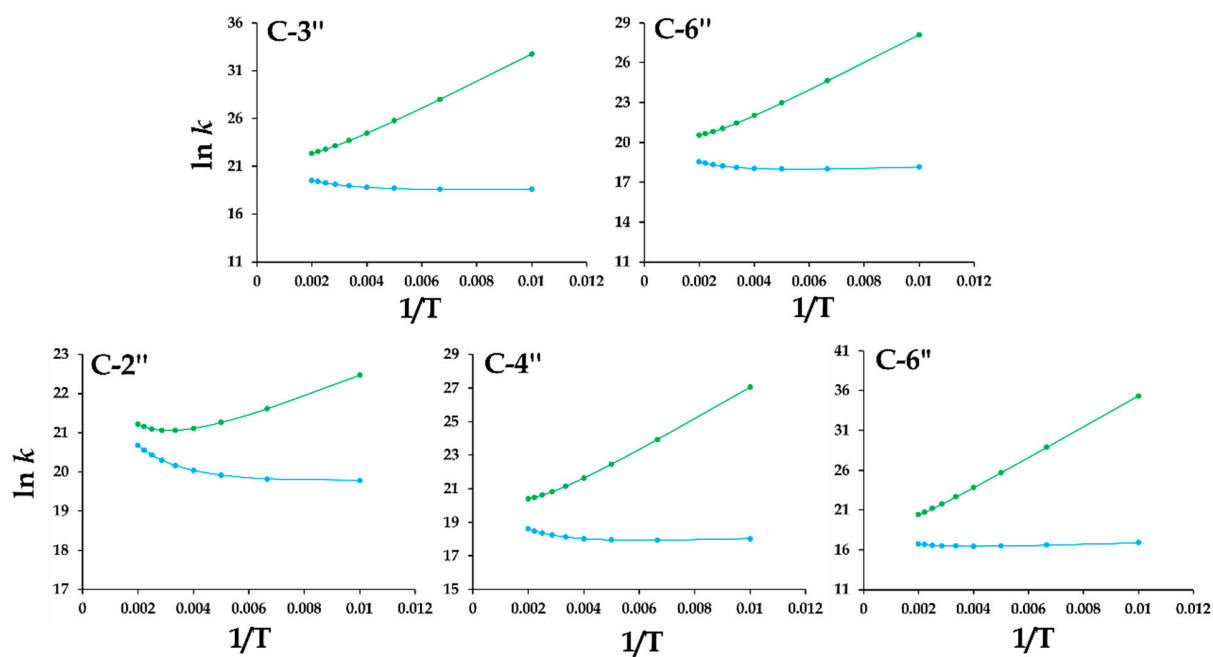
Figure S16. Graph of the dependence of lnk_{TST} (green line) and lnk_{ZCT_0} (blue line) on the reciprocal values of temperature for the RAF mechanism (**1** (top) and **2** (bottom))
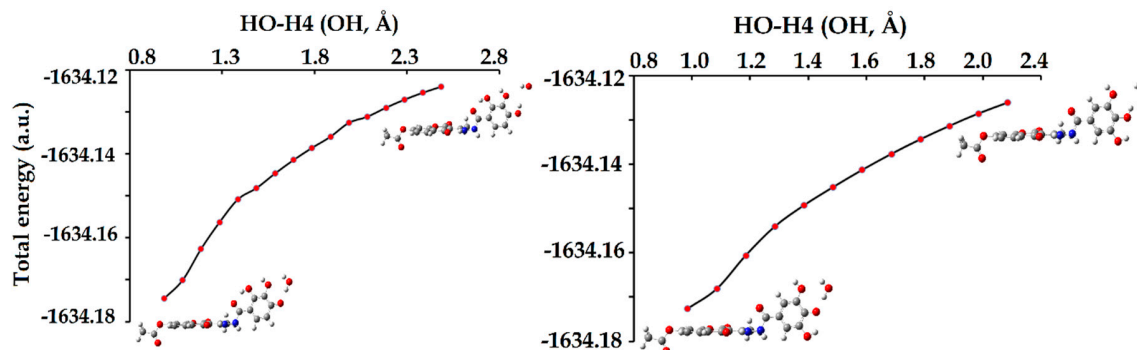

Figure S17. Dependence of the total energy (a.u.) on the HO–H4 characteristic distance (Å) for the SPL mechanism between neutral **H4A** of compound **1** (left) and **2** (right) and HO$^-$
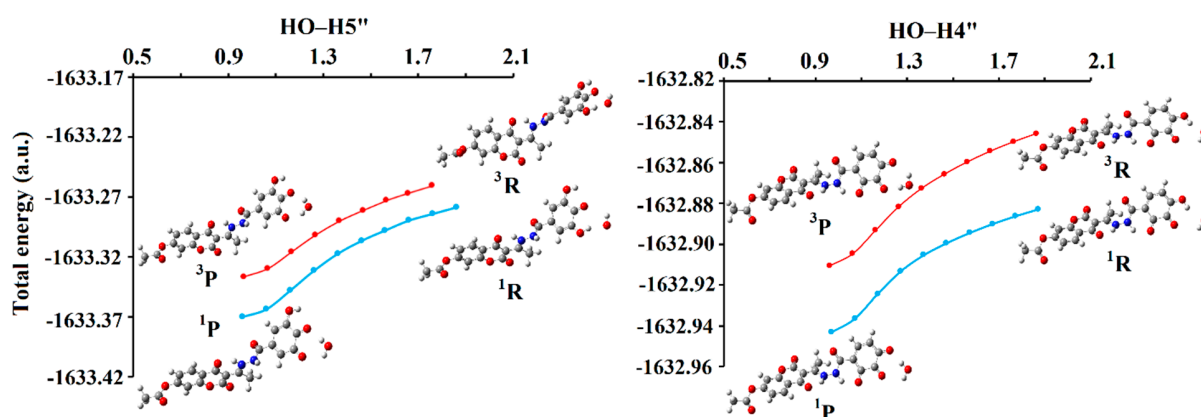
Figure S18. Energy profile for the HAT/PCET reaction pathway between **H₃A·** of compound **2** and HO• (left) as well as **H₂A·⁻** of compound **1** and HO• (right) in the singlet (blue) and triplet (red) spin states. Interatomic distances are given in angstroms [Å]

Figure S19. Energy profile for the RCF reaction path between **H₃A·** of compound **1** and HO• in the singlet (blue) and triplet (red) spin states. Interatomic distances are given in angstroms [Å]

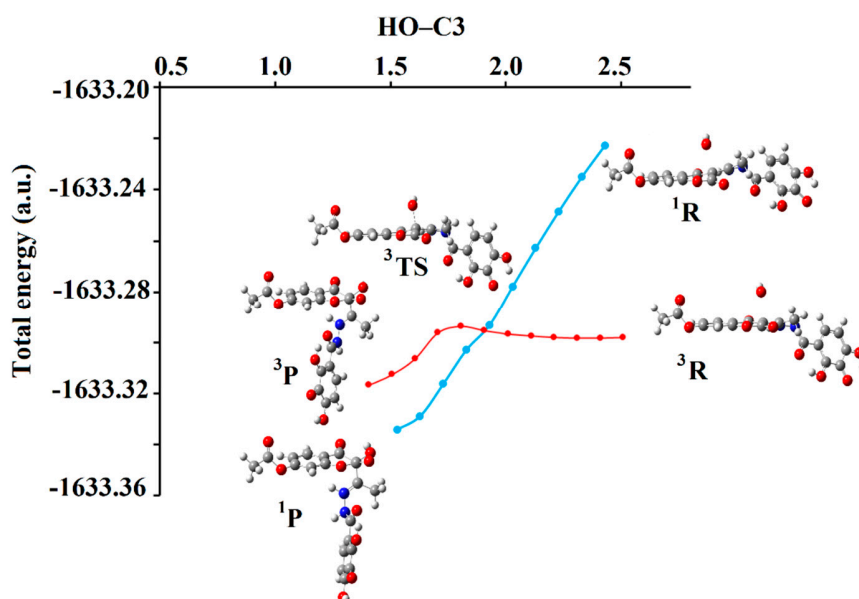**Table S4.** Estimation values of overall rate constants ($k_{overall}$), relative antiradical capacity (r^T) and branching ratios, Γ_i (%), for exergonic reaction pathways evaluated at physiological conditions, pH=7.4.

Mechanism	Position	Compound 1			Compound 2	
		H₄A	**H₃A⁻**	**H₂A²⁻**	**H₄A**	**H₃A⁻**
HAT/PCET	**2″-OH/3″-OH**	0.0	1.6	/	0.0	0.9
	3″-OH/4″-OH	0.0	/	/	0.0	/
	4″-OH/5″-OH	0.0	1.6	0.4	0.0	0.9
	2-NH	3.1	1.6	0.4	3.7	0.9
RAF/RCF	**C-3**	0.1			0.0	
	C-5	0.0			0.0	
	C-6	0.1			0.0	
	C-7	0.0			0.0	
	C-8	0.1			0.0	
	C-9	0.0			0.0	
	C-10	0.0	22.0	5.6	0.0	12.6
	C-1′	0.0			0.0	
	C-1″	0.1			0.2	
	C-2″	0.1			1.1	
	C-3″	0.3			0.0	
	C-4″	0.1			0.1	
	C-5″	0.1			0.0	
	C-6″	0.1			0.0	
SPL/SET	**n-OH/n-NH**	12.3	6.2	0.4	14.7	3.9
ET/PT	**2″-O/3″-O**	12.9	1.6	/	15.5	0.9
	3″-O/4″-O	12.2	/		15.5	/
	4″-O/5″-O	12.9	1.6	0.4	15.3	0.9
	2-N	0.2	1.6		11.3	0.9
$k_{overall}$		**3.75×10¹⁰**			**4.11×10¹⁰**	
$r^T{}_{Tx}$		**19.1**			**21.2**	
$r^T{}_{GA}$		**1.5**			**1.6**	