

Supplementary

Investigating the Antiparasitic Potential of the Marine Sesquiterpene Avarone, its reduced form Avarol, and the Novel Semisynthetic Thiazinoquinone analogue Thiazoavarone

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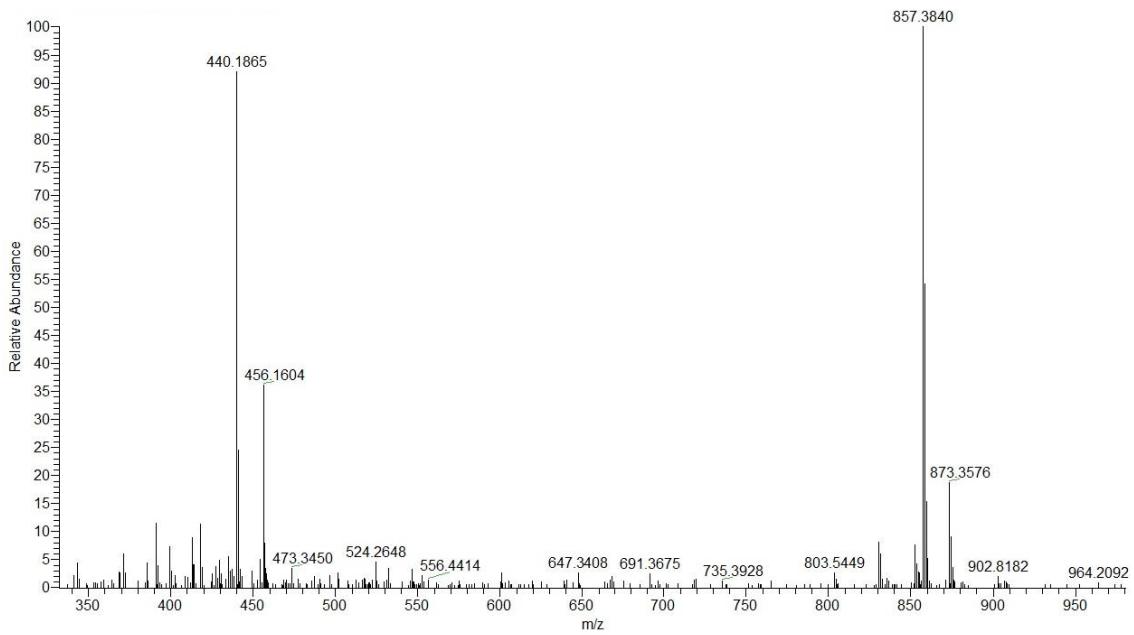


Figure S1. HRESIMS spectrum of thiazoavarone (2)

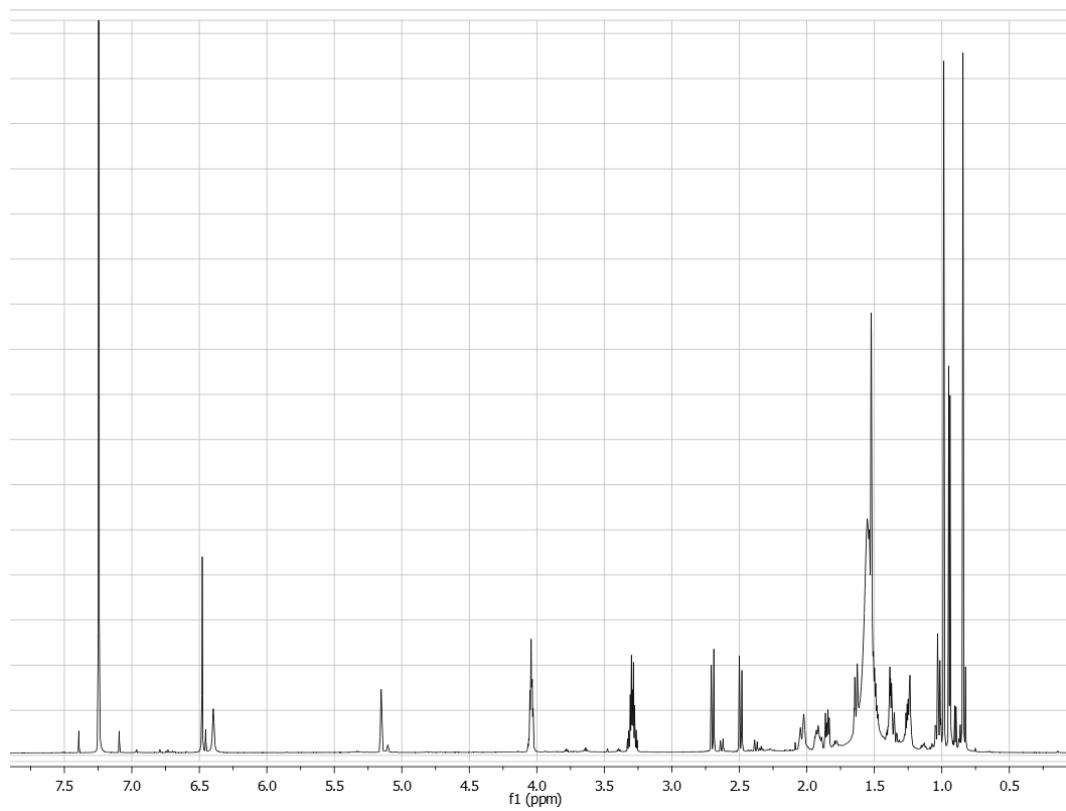


Figure S2. ^1H NMR spectrum of thiazoavarone (2) in CDCl_3

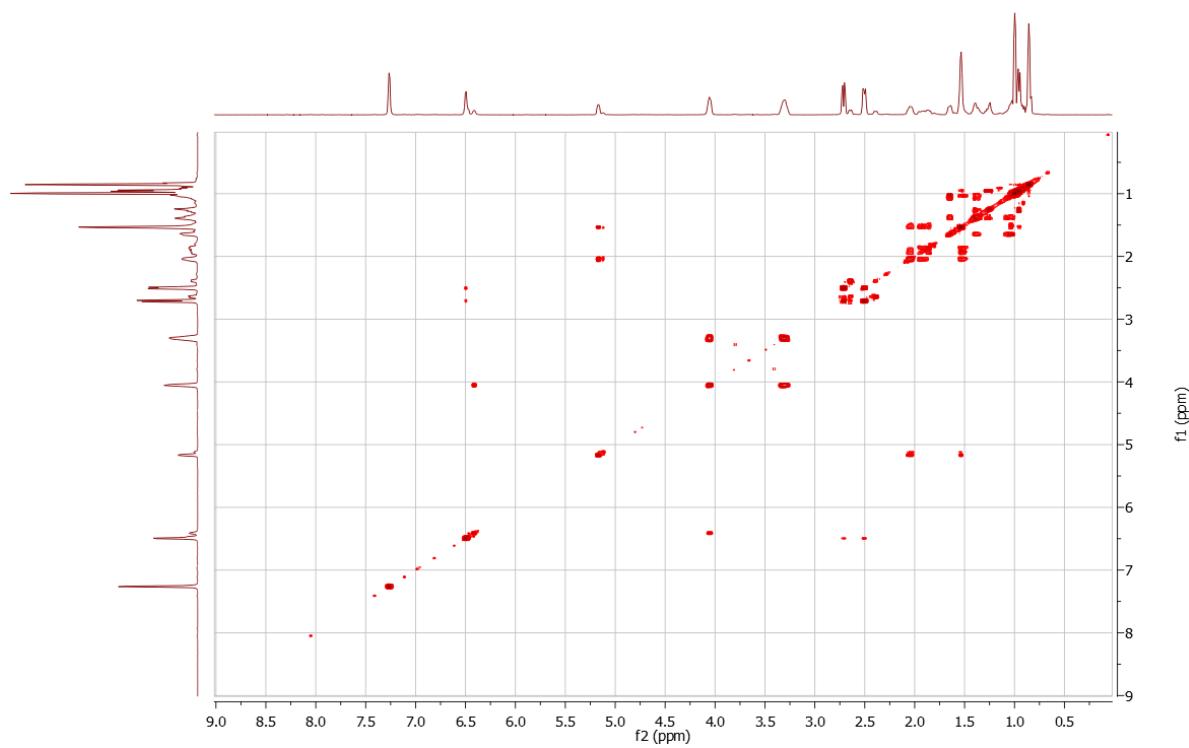


Figure S3. ^1H - ^1H COSY spectrum of thiazoavarone (**2**) in CDCl_3

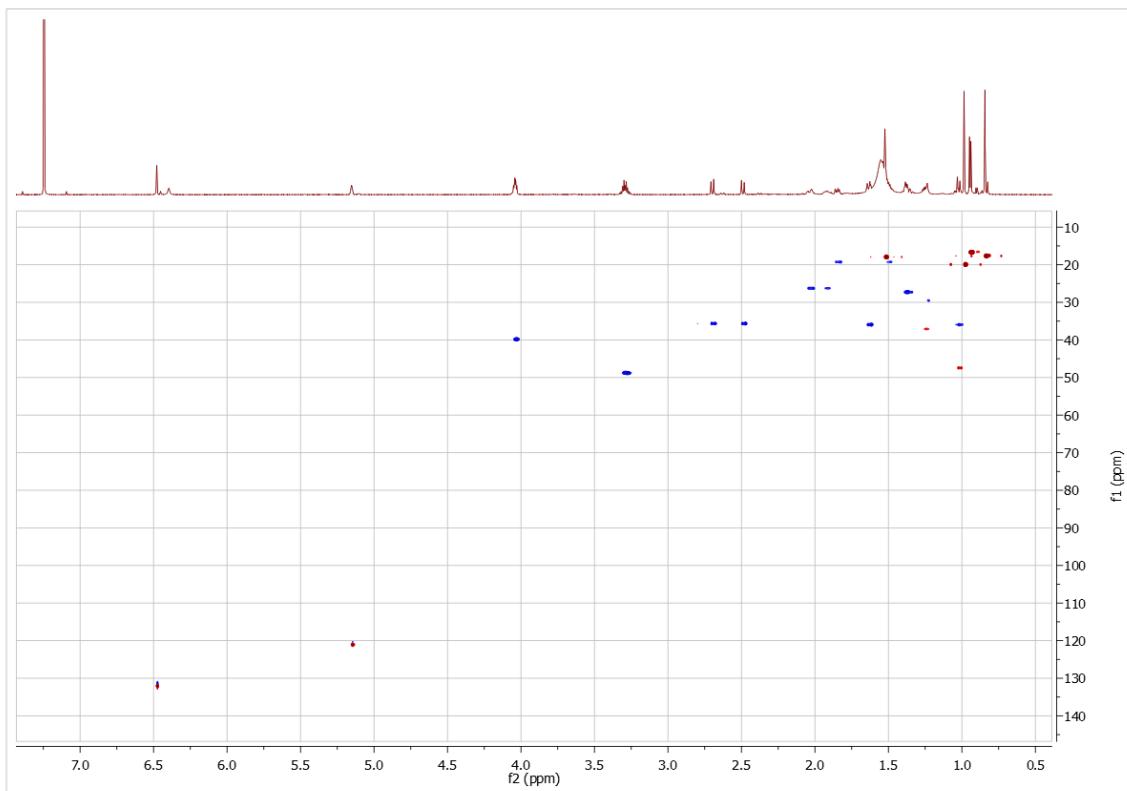


Figure S4. HSQC spectrum of thiazoavarone (**2**) in CDCl_3

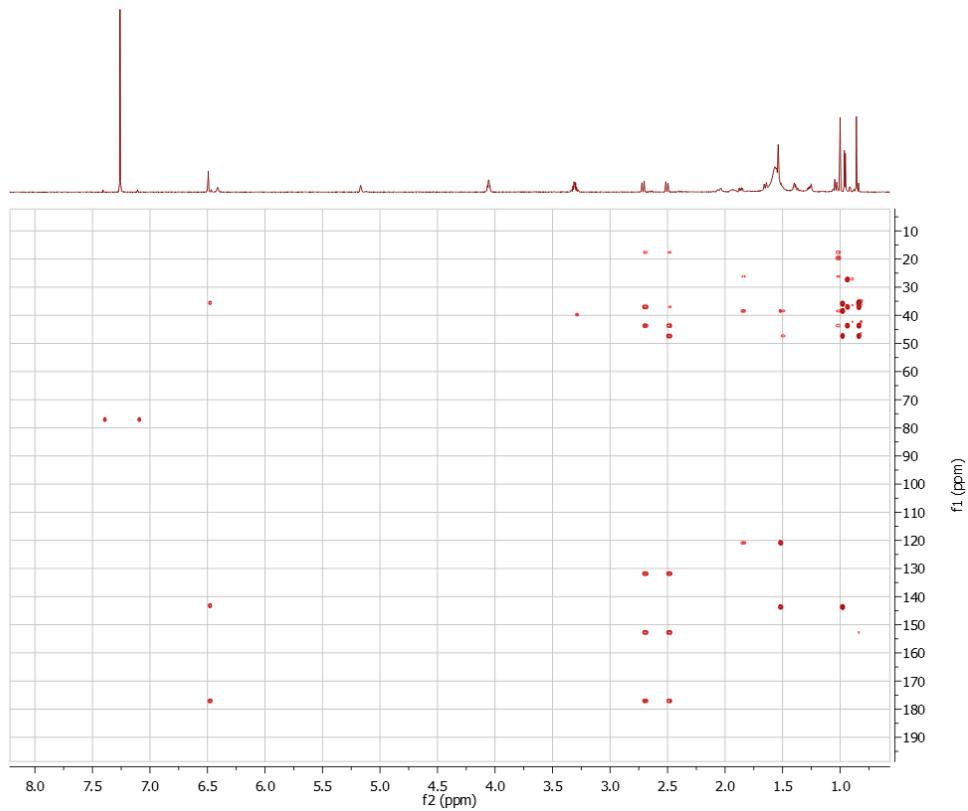


Figure S5. HMBC spectrum of thiazoavarone (**2**) in CDCl_3

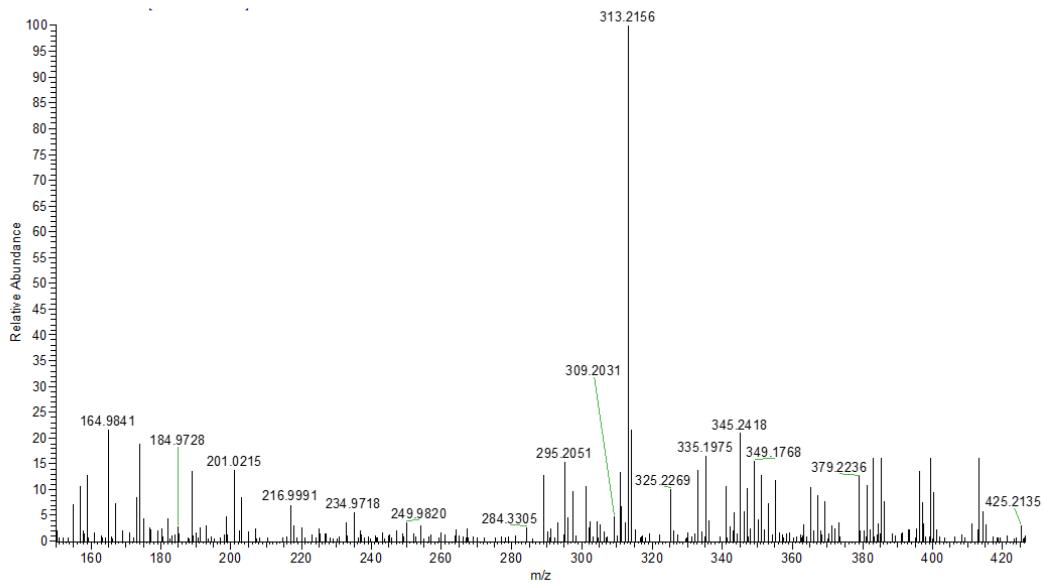


Figure S6. HRESIMS spectrum of avarone (**1**)

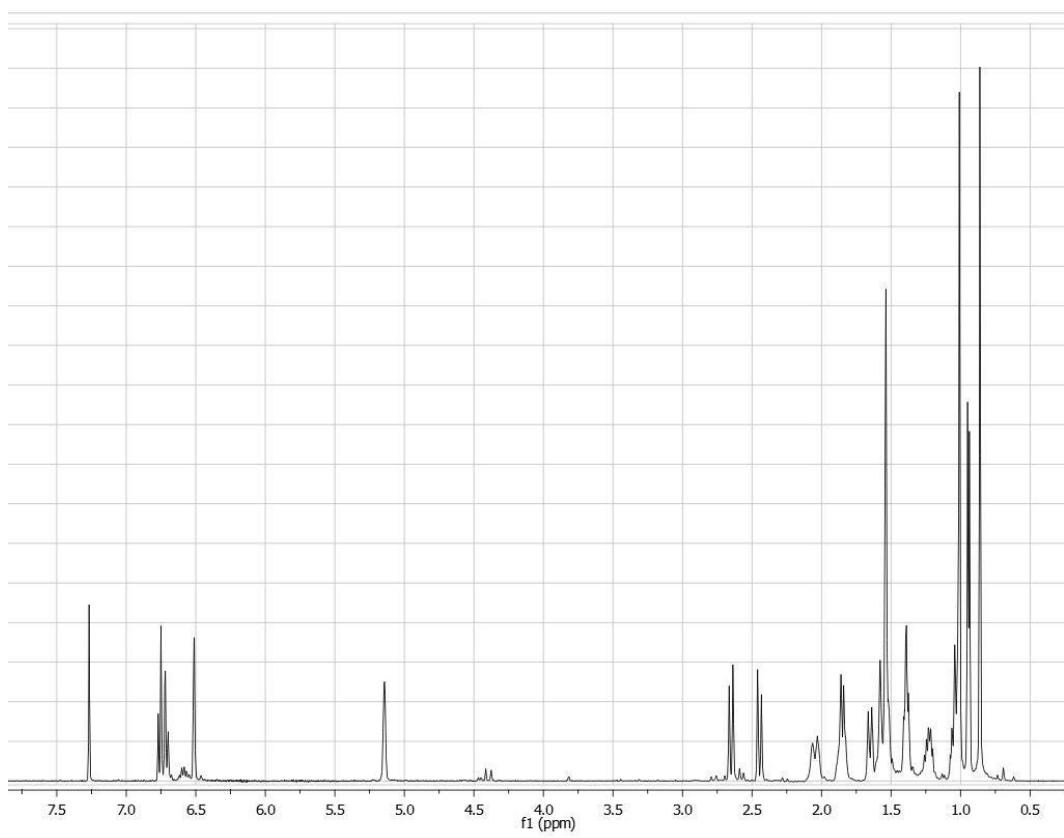


Figure S7. ^1H -NMR spectrum of avarone (**1**) in CDCl_3

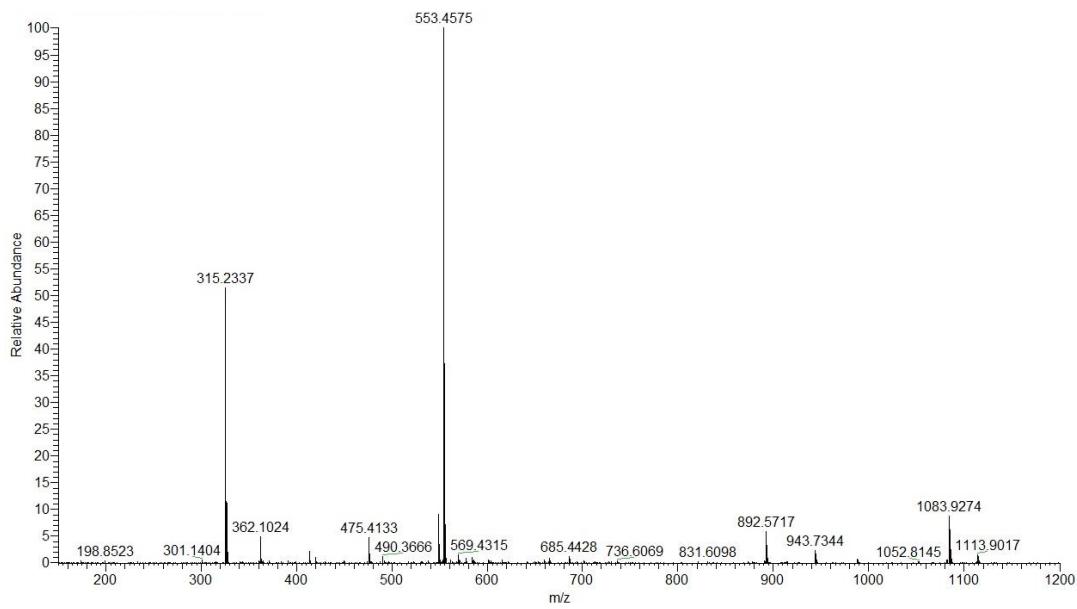


Figure S8. HRESIMS spectrum of avarol (**3**)

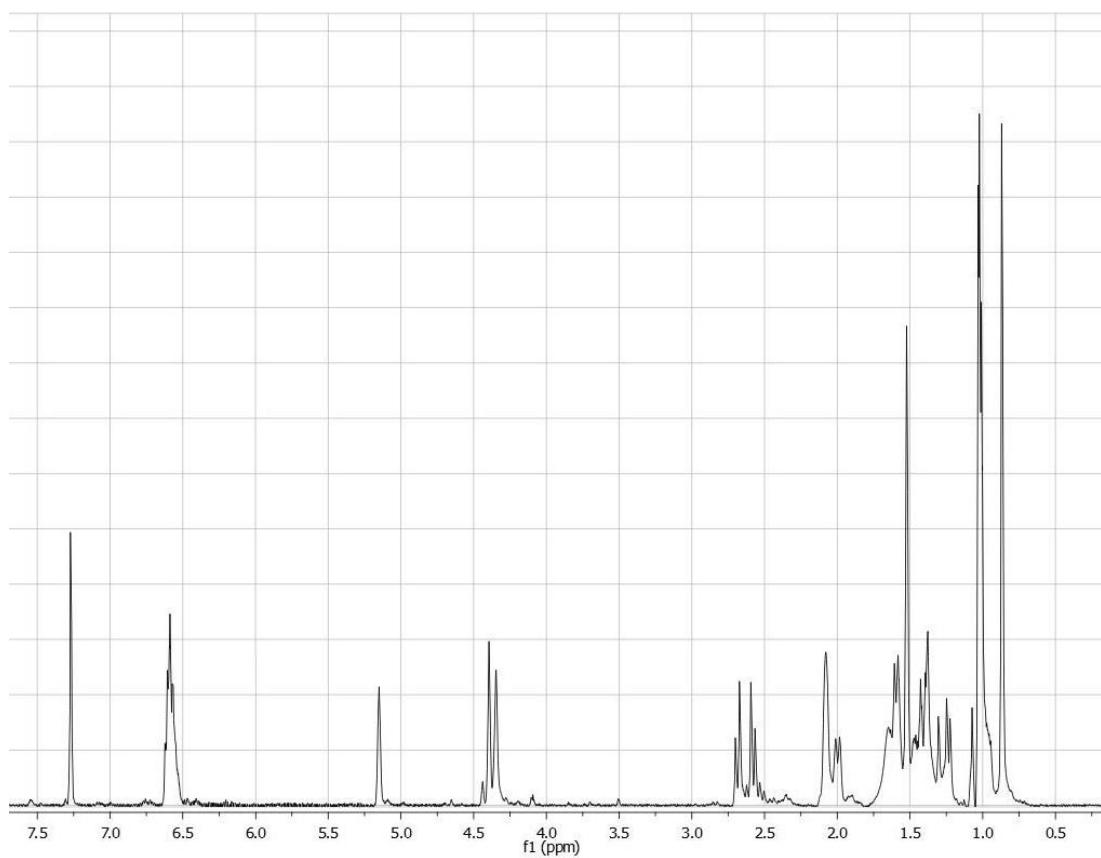


Figure S9. ^1H -NMR spectrum of avarol (**3**) in CDCl_3

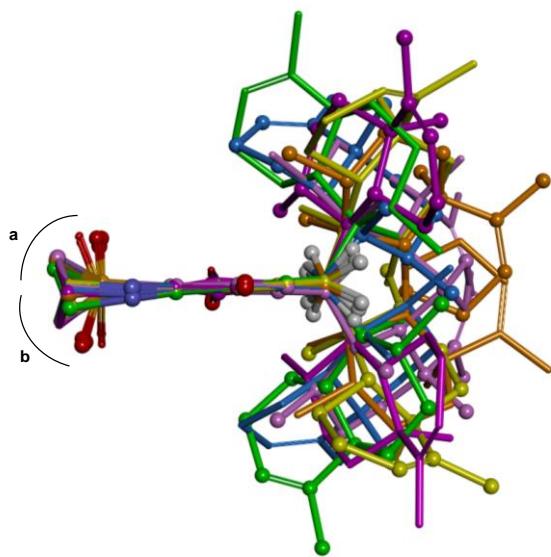


Figure S10. DFT conformers of compound **2** superimposed by the carbon atoms of quinone ring. The conformers characterized by flip a are showed in ball and stick while those characterized by the flip b are showed in stick. Carbon atoms are coloured according to family classification (I=green, II=magenta, III=pink, IV= light blue, V= orange, and VI=Yellow); heteroatoms are colored by atom type (O= red, N= blue, S= yellow). Hydrogens are omitted for sake of clarity, with the exception of those of the first R' methylene group.

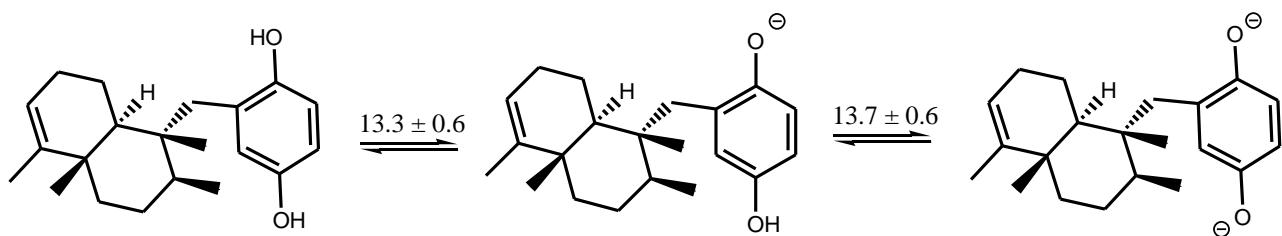


Figure S11. Structures of ionic forms and microcostant pka values calculated for **3** using the algorithm ACD/pKa Classic (ACD/Percepta software).

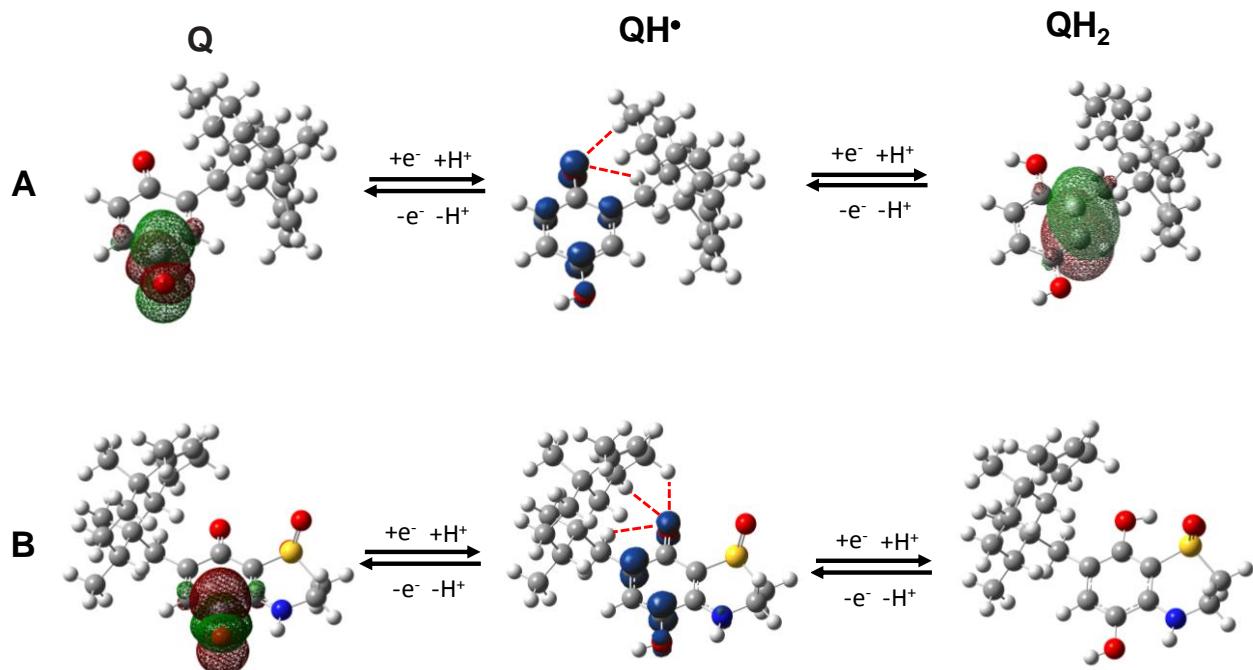
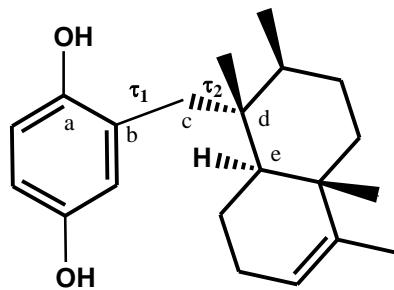


Figure S12. DFT conformer II of **1** (**Q**), **3** (**QH₂**) and their semiquinone radical (**QH[•]**). (B) DFT conformer I of **2** (**Q**) together with its one- and two-electron reduced species **QH[•]** and **QH₂**. Atoms possibly involved in an intramolecular radical shift are evidenced with red dashed lines. The LUMO of **1** and **2**, and the HOMO of **3** are visualized using GaussView with an isosurface value of 0.02 e⁻/a.u.³ The NBO spin density isosurface of the **QH[•]** species is displayed using GaussView with an isosurface value of 0.01 e⁻/a.u.³. The blue surface (positive spin density) corresponds to an excess of α -electron density.

Table S1. ΔE_{GM} values (kcal/mol) and torsion angle values (degrees) of the DFT conformers of **3**.

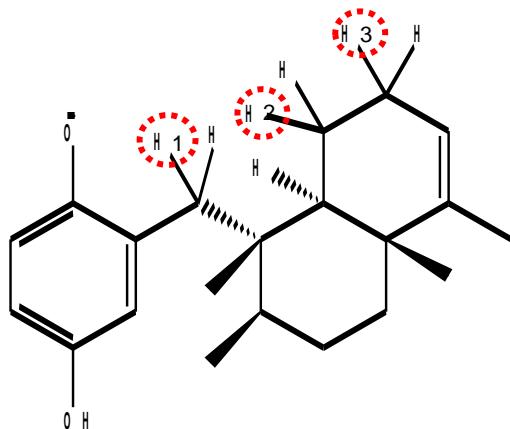


Conf.	ΔE_{GM} (kcal/mol)	Torsion Angles (°)	
		τ_1^{a}	τ_2^{b}
II	0.00	96.77	56.16
I	0.23	-104.40	59.80
III	1.45	-102.26	-179.87
IV	1.71	108.27	-69.84
V	2.12	89.20	179.85
VI	2.68	-86.45	-76.24

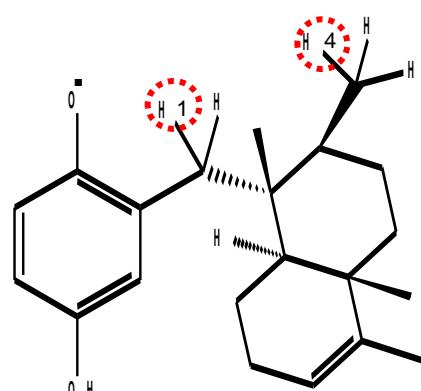
^a τ_1 torsion angle is defined by a, b, c, and d atoms. ^b τ_2 torsion angle is calculated considering b, c, d, and e atoms.

Table S2. List of the hydrogen atoms at a distance suitable for an intramolecular H-radical shift from a carbon atom of the R' substituent to the reduced/oxidized oxygen atom of the semiquinone radical. Corresponding distances in the Q, QH[·] and QH₂ forms are also reported. Values are expressed in Å.

Conf I



Conf II

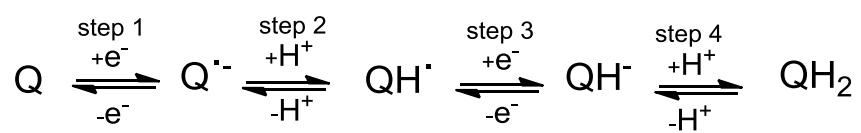


Cmp	Conf	Distance	Q species	QH [·] species	QH ₂ species
Avarone/Avarol	I	O-H1	2.43	2.41	2.43
	I	O-H2	2.45	2.40	2.45
	I	O-H3	2.92	3.02	2.77
	II	O-H1	2.43	2.42	2.46
	II	O-H4	2.49	2.45	2.54
Thiazoavarone	I	O-H1	2.35	2.39	2.41
	I	O-H2	2.39	2.39	2.44
	I	O-H3	3.04	3.02	2.82
	II	O-H1	2.38	2.40	2.45
	II	O-H4	2.47	2.47	2.56

Table S3. ClogD of compounds **1-3**.

Cmp	cLogD ^a
1	6.13
2	3.83
3	5.22

^alogD calculated considering pH values: 7.4, 7.2 and 5.5.



Scheme S1. Reduction pathway of quinone to hydroquinone in aqueous environment.