

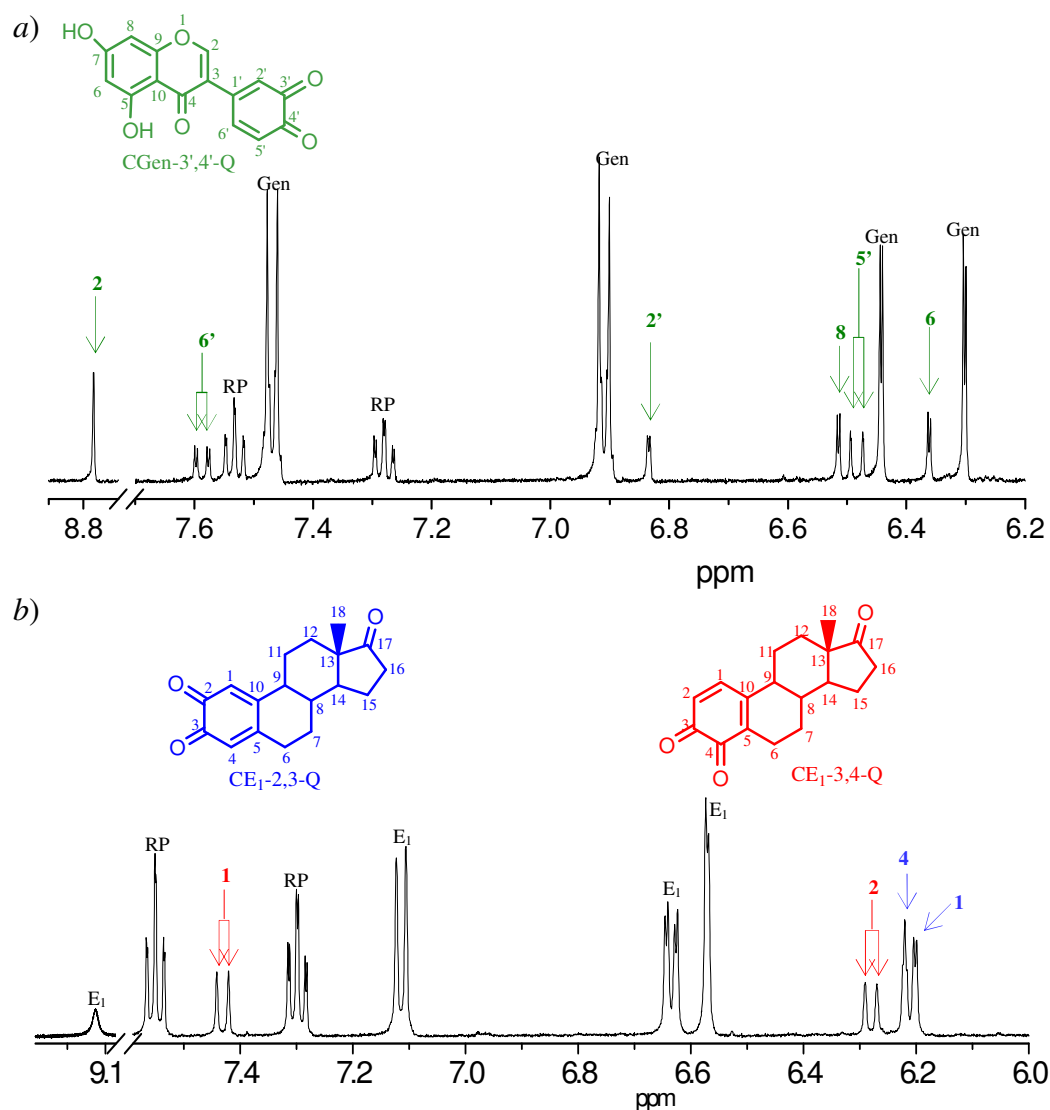
# Supporting Information

## The Balance of Beneficial and Deleterious Health Effects of Quinones: A Case Study of the Chemical Properties of Genistein and Estrone Quinones

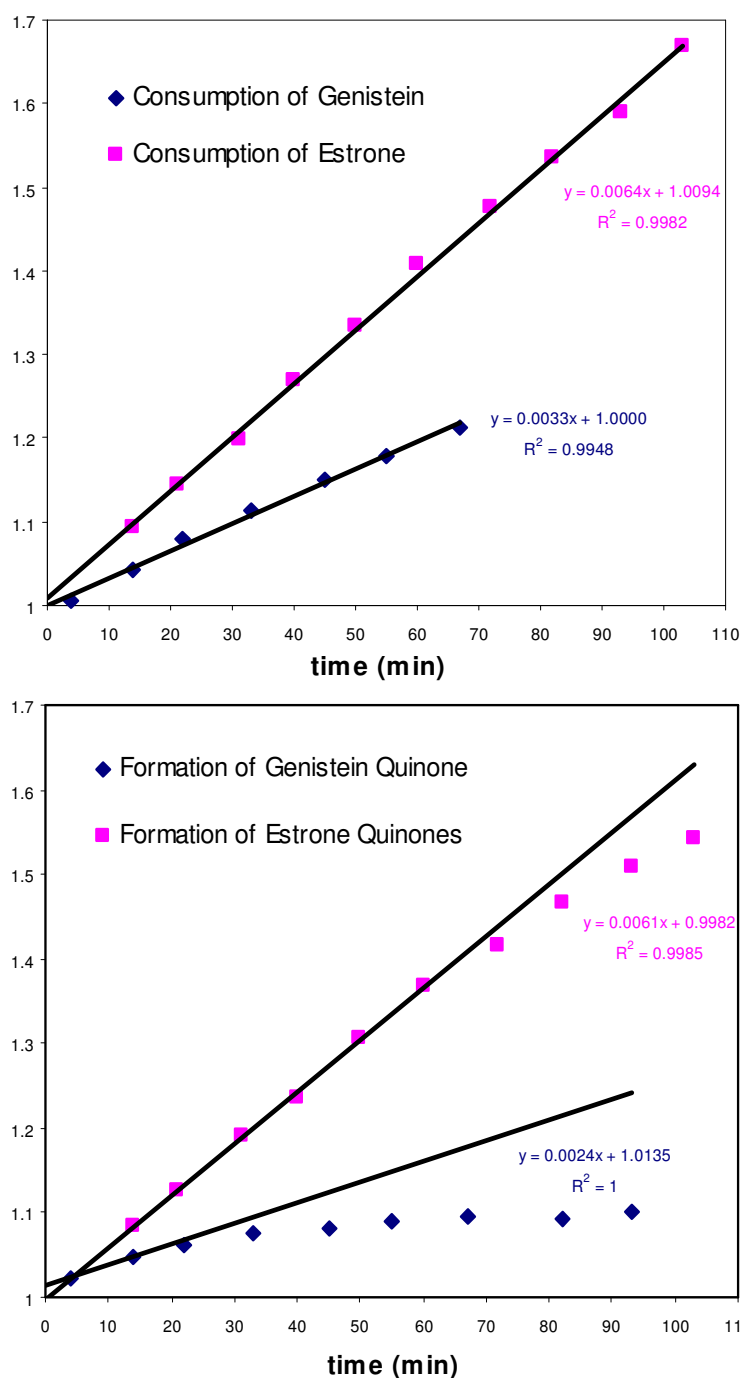
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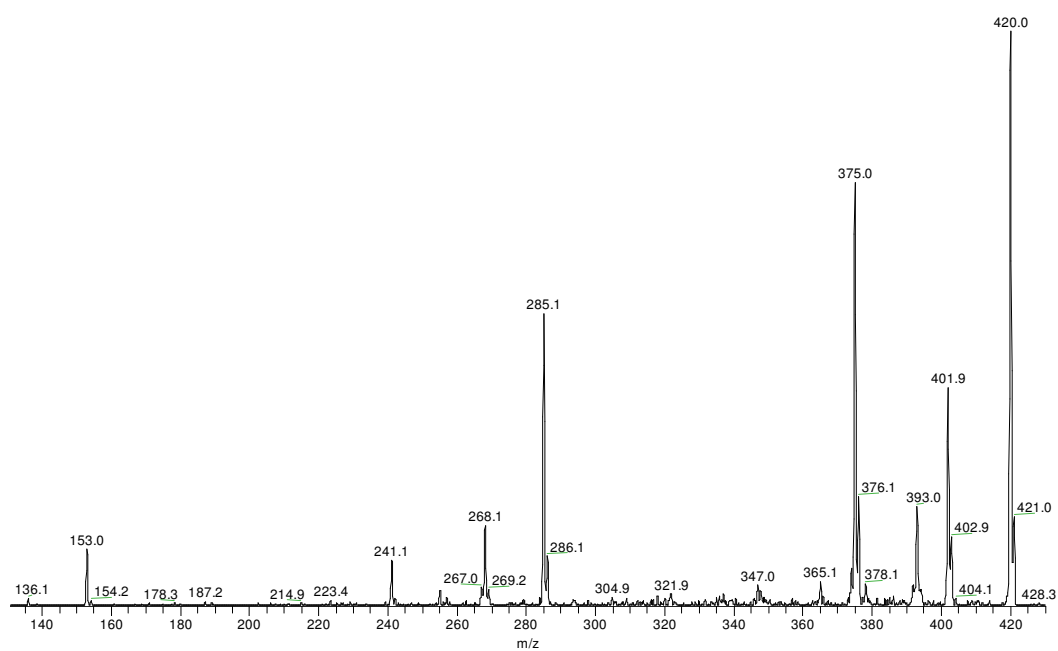
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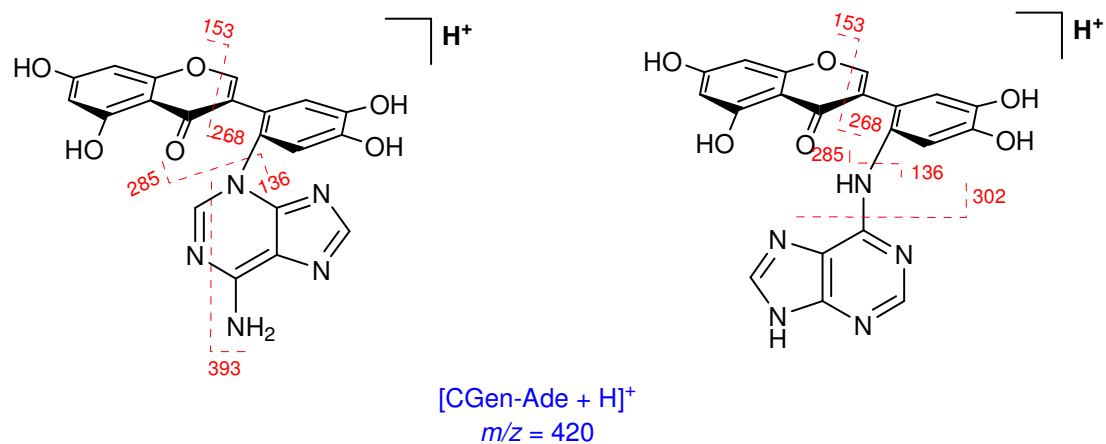
**Figure S1.**  $^1\text{H}$  NMR spectra of a) Gen + IBX with characteristic  $^1\text{H}$  of CGen-3',4'-Q product labeled in green and b)  $\text{E}_1$  + IBX with characteristic  $^1\text{H}$  of  $\text{CE}_1$ -2,3-Q and  $\text{CE}_1$ -3,4-Q products labeled in red and blue, respectively. Peaks of the reactants Gen or  $\text{E}_1$  and the reduced product from IBX (RP) in the reaction mixtures were labeled in black. The y axis is intensity.



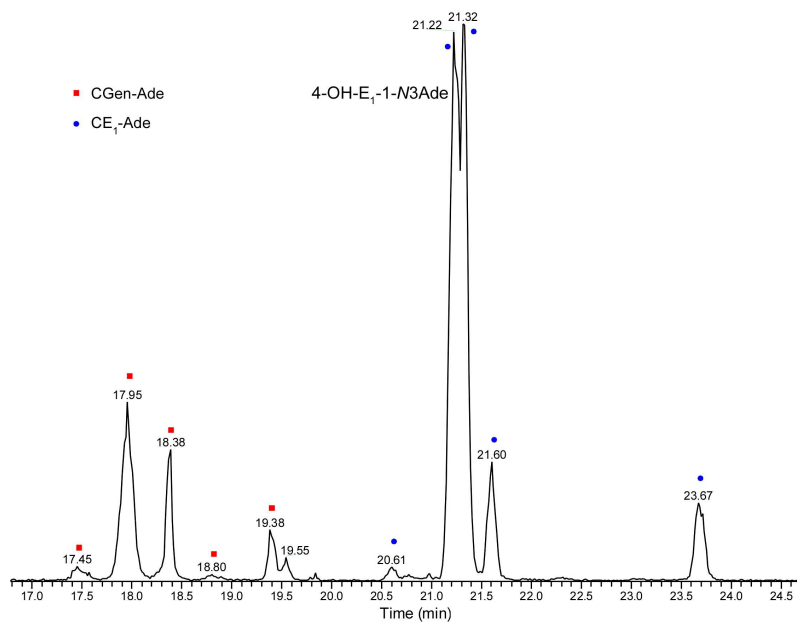
**Figure S2.** 2<sup>nd</sup>-Order kinetics of Gen + IBX and E<sub>1</sub> + IBX monitored with NMR. A) Plot of the consumption of Gen and E<sub>1</sub>, where  $y = \frac{1}{[S]/[S]_0}$  and  $[S] \equiv$  concentration of E<sub>1</sub> or Gen at time t and  $[S]_0 \equiv$  initial concentration of E<sub>1</sub> or Gen. B) Plot of the formation of CGen-3',4'-Q and CE<sub>1</sub>-2,3-Q + CE<sub>1</sub>-3,4-Q, where  $y = \frac{1}{1-[Q]/[S]_0}$  and  $[Q] \equiv$  concentration of the quinone at time t and  $[S]_0 \equiv$  initial concentration of E<sub>1</sub> or Gen. Red squares represent data for E<sub>1</sub> and blue diamond for Gen. For the formation of E<sub>1</sub>, the first six data points were used to calculate rate constant. The formation of CGen-3',4'-Q deviates from 2<sup>nd</sup>-order kinetics at ~ 30 min and only the first two data points were used to estimate the rate constant.



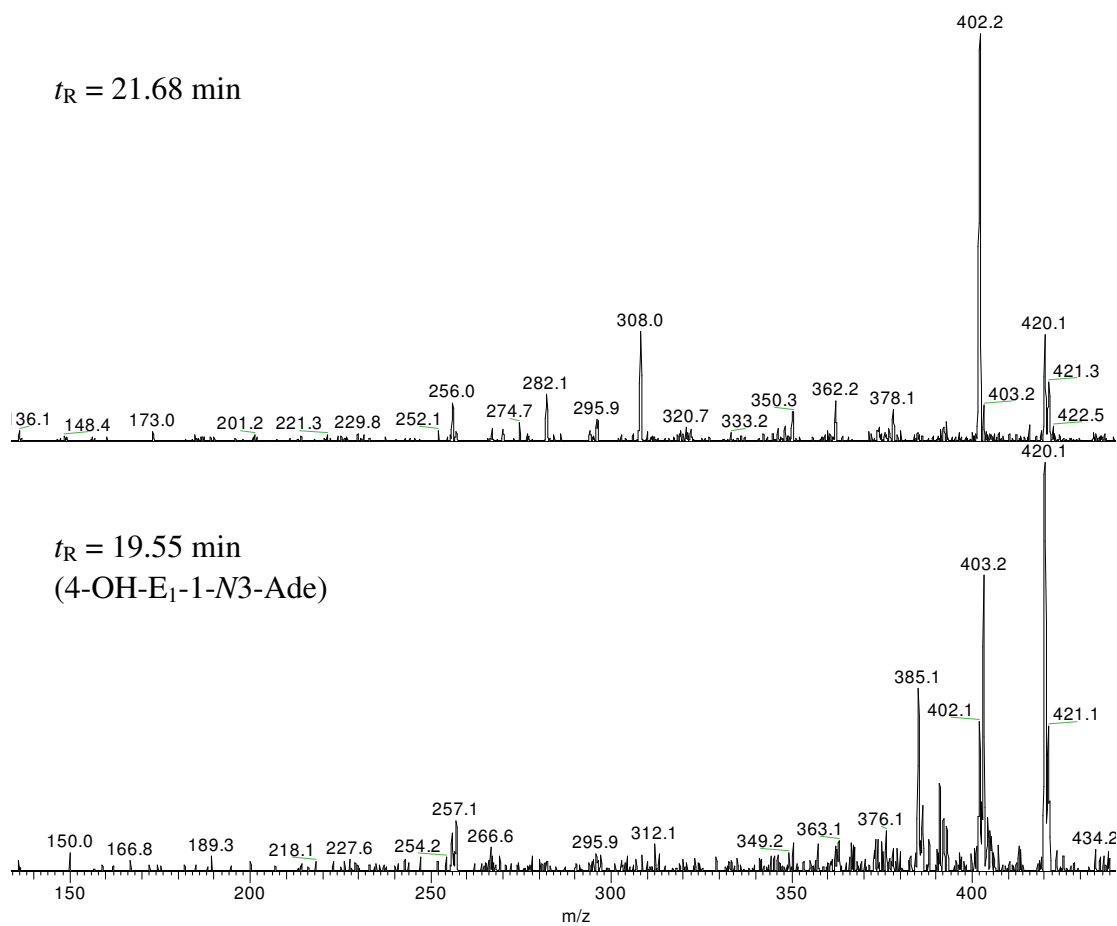
**Figure S3.** An example of MS/MS product-ion spectrum (y axis is relative abundance) of the CGen-Ade adducts in Figure 4 ( $t_R = 16.82$  min).



**Figure S4.** Representative structures of [CGen-Ade + H]<sup>+</sup> adducts. Cleavages shown are in accord with MS/MS observations. One characteristic cleavage of [CGen-Ade + H]<sup>+</sup> is C-ring cleavage that produces fragment ions of *m/z* 153 and complementary ion of *m/z* 268.



**Figure S5.** Reconstructed base-peak ion chromatogram (y axis is ion intensity) shows the formation of CGen-Ade and CE<sub>1</sub>Q-Ade adducts in DMF. The CGenQ + Ade and CE<sub>1</sub>Q + Ade were mixed at 1:1 ratio.



**Figure S6.** MS/MS Product-ion spectra of the two major CE<sub>1</sub>-Ade adducts (4-OH-E<sub>1</sub>-1-N<sup>3</sup>Ade and unknown) identified in the reaction mixture of CE<sub>1</sub>Q + calf thymus DNA at pH 7.4, 37 °C. The y axis is relative abundance.

**Table S1.** Accurate mass measurement of MS/MS product ions of genistein  $[\text{C}_{15}\text{H}_{10}\text{O}_5 + \text{H}]^+$  and catechol genistein  $[\text{C}_{15}\text{H}_{10}\text{O}_6 + \text{H}]^+$

Parent	Product	Measured Mass	Theoretical Mass	Deviation (mau)
Genistein	<b><math>\text{C}_{15}\text{H}_{11}\text{O}_5^+</math></b>	<b>271.0601</b>	<b>271.0601</b>	<b>0</b>
	$\text{C}_{15}\text{H}_9\text{O}_4^+$	253.0495	253.0495	0
	$\text{C}_{14}\text{H}_{11}\text{O}_4^+$	243.0652	243.0652	0
	$\text{C}_{14}\text{H}_{11}\text{O}_3^+$	227.0703	227.0703	0
	$\text{C}_{14}\text{H}_9\text{O}_3^+$	225.0546	225.0546	0
	$\text{C}_{13}\text{H}_{11}\text{O}_3^+$	215.0703	215.0703	0
	$\text{C}_{13}\text{H}_9\text{O}_2^+$	197.0597	197.0597	0
	$\text{C}_{11}\text{H}_7\text{O}_3^+$	187.0390	187.0390	0
	$\text{C}_{10}\text{H}_7\text{O}_2^+$	159.0441	159.0441	0
	$\text{C}_7\text{H}_5\text{O}_4^+$	153.0183	153.0183	0
	$\text{C}_8\text{H}_5\text{O}_3^+$	149.0233	149.0233	0
Catechol Genistein	<b><math>\text{C}_{15}\text{H}_{11}\text{O}_6^+</math></b>	<b>287.0551</b>	<b>287.0550</b>	<b>+0.1</b>
	$\text{C}_{15}\text{H}_9\text{O}_5^+$	269.0445	269.0444	+0.1
	$\text{C}_{14}\text{H}_{11}\text{O}_5^+$	259.0602	259.0601	+0.1
	$\text{C}_{13}\text{H}_7\text{O}_6^+$	259.0238	259.0237	+0.1
	$\text{C}_{14}\text{H}_9\text{O}_4^+$	241.0496	241.0495	+0.1
	$\text{C}_{13}\text{H}_5\text{O}_5^+$	241.0137	241.0131	+0.6
	$\text{C}_{13}\text{H}_{11}\text{O}_4^+$	231.0653	231.0652	+0.1
	$\text{C}_{12}\text{H}_7\text{O}_5^+$	231.0289	231.0288	+0.1
	$\text{C}_{13}\text{H}_9\text{O}_3^+$	213.0546	213.0546	0
	$\text{C}_{11}\text{H}_7\text{O}_4^+$	203.0339	203.0339	0
	$\text{C}_7\text{H}_5\text{O}_4^+$	153.0183	153.0183	0



**Table S2.** Accurate mass measurement for the five catechol genistein adenine  $[\text{C}_{20}\text{H}_{13}\text{N}_5\text{O}_6 + \text{H}]^+$  adducts shown in Figure S5.

Retention time (min)	Measured Mass	Deviation (mau)
17.45	420.0939	0.0
17.95	420.0938	-0.1
18.38	420.0941	+0.2
18.80	420.0937	-0.2
19.38	420.0938	-0.1