

Supplementary Materials

Table S1. Experimental design for optimizing the production of SeNPs from Kombucha fermented pollen.

Run	Factor A (SCOBY)	Factor B (Pollen)	Factor C (Na ₂ SeO ₃)	TPC (µg/mL)	TFC (µg/mL)	HAT (µg/mL)	DPPH (µM)	FRAP (µM)	CUPRAC (µM)	Si (mg/L)	Se ⁰ (mM)	Se ⁰ yield (%)
1	15	15	190	8.602	2.056	3.587	436.5	1720	695.7	5	2.043	55.80
2	30	5	190	19.03	4.509	12.84	733.6	4094	1643	4.5	0	0
3	30	15	370	8.671	2.056	3.819	462.1	1754	995.5	5	1.273	17.86
4	30	15	190	13.44	2.294	4.282	490.3	1374	1032	8	1.678	45.84
5	30	5	190	17.27	4.104	16.04	719.8	3779	1313	4	0	0
6	30	15	10	25.54	5.196	19.09	738.4	3726	1723	8	0	0
7	30	15	10	27.55	4.541	25.27	646.7	2764	1531	7	0	0
8	15	5	10	28.33	4.353	19.79	740.3	3414	1477	7	0	0
9	30	25	190	15.18	3.542	8.680	677.5	3044	1140	10	3.456	94.38
10	30	15	190	12.26	2.343	4.861	501.9	1227	1014	5	1.694	46.27
11	30	15	190	8.045	1.769	4.097	514.4	1389	984.3	5	1.980	54.07
12	30	25	190	18.19	3.692	10.67	701.9	3410	1122	13	3.638	99.36
13	15	15	190	9.262	2.468	4.050	569.8	1564	801.8	6	1.940	52.99
14	45	5	370	26.24	4.884	21.41	723.0	648.5	1840	7	0	0
15	15	5	10	28.41	4.272	24.37	654.4	2634	1339	6	0	0
16	45	5	370	26.63	5.065	25.57	719.8	864	1968	5	0	.00
17	30	15	190	11.59	1.875	4.513	642.3	1032	1204	5	2.003	54.72
18	45	25	10	20.19	4.335	12.15	712.1	2633	1248	9	0	0
19	15	25	370	12.24	2.718	4.745	492.9	1314	712.4	9	3.115	43.68
20	30	15	190	12.89	3.136	6.25	682.6	1453	1170	6	2.242	61.22
21	45	15	190	11.08	2.606	4.513	592.3	1573	742.2	8	0	0
22	45	25	10	17.20	3.829	12.15	709.6	2389	1286	9	0	0
23	30	15	370	9.193	2.194	4.166	540.3	1954	969.4	5	1.535	21.53
24	15	25	370	12.89	2.855	4.166	539.1	1139	768.3	6	2.861	40.12
25	45	15	190	12.47	2.824	3.356	600.6	1714	772.0	5	0	0

Table S2. R², Adjusted R² from the ANOVA analysis.

	R ²	Adjusted R ²
TPC	0.971	0.957
TFC	0.873	0.83
HAT	0.96	0.936
DPPH	0.69	0.628
FRAP	0.663	0.574
CUPRAC	0.952	0.924
Si	0.649	0.578
Se ⁰	0.838	0.815
Se ⁰ yield	0.698	0.655

Table S3. Coded equation.

Coded equation									
TPC = 11.81 + 1.17 × A - 0.73 × B - 8.80 × C - 6.79 × AB + 5.15 × AC - 1.56 × A² + 5.50 × B² + 5.81 × C²									
TFC = 2.53 + 0.40 × A - 0.51 × B - 1.37 × C - 1.21 × AB + 1.07 × B² + 0.60 × C²									
HAT = 5.72 + 0.05 × A - 2.38 × B - 9.09 × C - 7.52 × AB + 4.85 × AC - 2.21 × BC - 2.42 × A² + 5.75 × B² + 6.78 × C²									
DPPH = 570.68 + 52.02 × A - 38.16 × B - 60.36 × C + 106.37 × B²									
FRAP = 2078.04 - 163.624 × A - 125.28 × B - 823.85 × C - 941.83 × A² + 996.81 × B²									
CUPRAC = 1050.24 + 4.18 × A - 173.60 × B - 322.54 × C - 314.92 × AB + 152.77 × AC - 251.68 × BC - 277.92 × A² + 274.17 × B² + 274.13 × C²									
Si = 6.68 + 0.33 × A + 3.75 × B - 0.75 × C + 2.75 × AC									
Se⁰ = 1.59 + 1.08 × B + 0.87 × C + 0.91 × BC - 0.71 × C²									
Se⁰ yield = 43.43 + 19.97 × B + 12.25 × C + 12.86 × BC - 31.17 × C²									

Table S4. Pearson correlation for the responses of the RSM experimental design.

		TPC	TFC	HAT	DPPH	FRAP	CUPRAC	Si	Se ⁰	Se ⁰ yield
TPC	Pearson correlation	1	0.916**	0.960**	0.737**	0.407*	0.832**	0.192	-0.568**	-0.512**
	Sig. (2-tailed)		0.000	0.000	0.000	0.044	0.000	0.358	0.003	0.009
	N	25	25	25	25	25	25	25	25	25
TFC	Pearson correlation	0.916**	1	0.898**	0.844**	0.517**	0.843**	0.218	-0.566**	-0.516**
	Sig. (2-tailed)	0.000		0.000	0.000	0.008	0.000	0.294	0.003	0.008
	N	25	25	25	25	25	25	25	25	25
HAT	Pearson correlation	0.960**	0.898**	1	0.703**	0.380	0.861**	0.067	-0.600**	-0.537**
	Sig. (2-tailed)	0.000	0.000		0.000	0.061	0.000	0.749	0.002	0.006
	N	25	25	25	25	25	25	25	25	25
DPPH	Pearson correlation	0.737**	0.844**	0.703**	1	0.546**	0.767**	0.238	-0.466*	-0.343
	Sig. (2-tailed)	0.000	0.000	0.000		0.005	0.000	0.252	0.019	0.093
	N	25	25	25	25	25	25	25	25	25
FRAP	Pearson correlation	0.407*	0.517**	0.380	0.546**	1	0.316	0.231	-0.242	-0.164
	Sig. (2-tailed)	0.044	0.008	0.061	0.005		0.124	0.266	0.243	0.433
	N	25	25	25	25	25	25	25	25	25
CUPRAC	Pearson correlation	0.832**	0.843**	0.861**	0.767**	0.316	1	-0.031	-0.557**	-0.458*
	Sig. (2-tailed)	0.000	0.000	0.000	0.000	0.124		0.885	0.004	0.021
	N	25	25	25	25	25	25	25	25	25
Si	Pearson correlation	0.192	0.218	0.067	0.238	0.231	-0.031	1	0.325	0.353
	Sig. (2-tailed)	0.358	0.294	0.749	0.252	0.266	0.885		0.112	0.083
	N	25	25	25	25	25	25	25	25	25
Se ⁰	Pearson correlation	-0.568**	-0.566**	-0.600**	-0.466*	-0.242	-0.557**	0.325	1	0.942**
	Sig. (2-tailed)	0.003	0.003	0.002	0.019	0.243	0.004	0.112		0.000
	N	25	25	25	25	25	25	25	25	25
Se ⁰ yield	Pearson correlation	-0.512**	-0.516**	-0.537**	-0.343	-0.164	-0.458*	0.353	0.942**	1
	Sig. (2-tailed)	0.009	0.008	0.006	0.093	0.433	0.021	0.083	0.000	
	N	25	25	25	25	25	25	25	25	25

**. Correlation is significant at the 0.01 level (2-tailed); *. Correlation is significant at the 0.05 level (2-tailed).

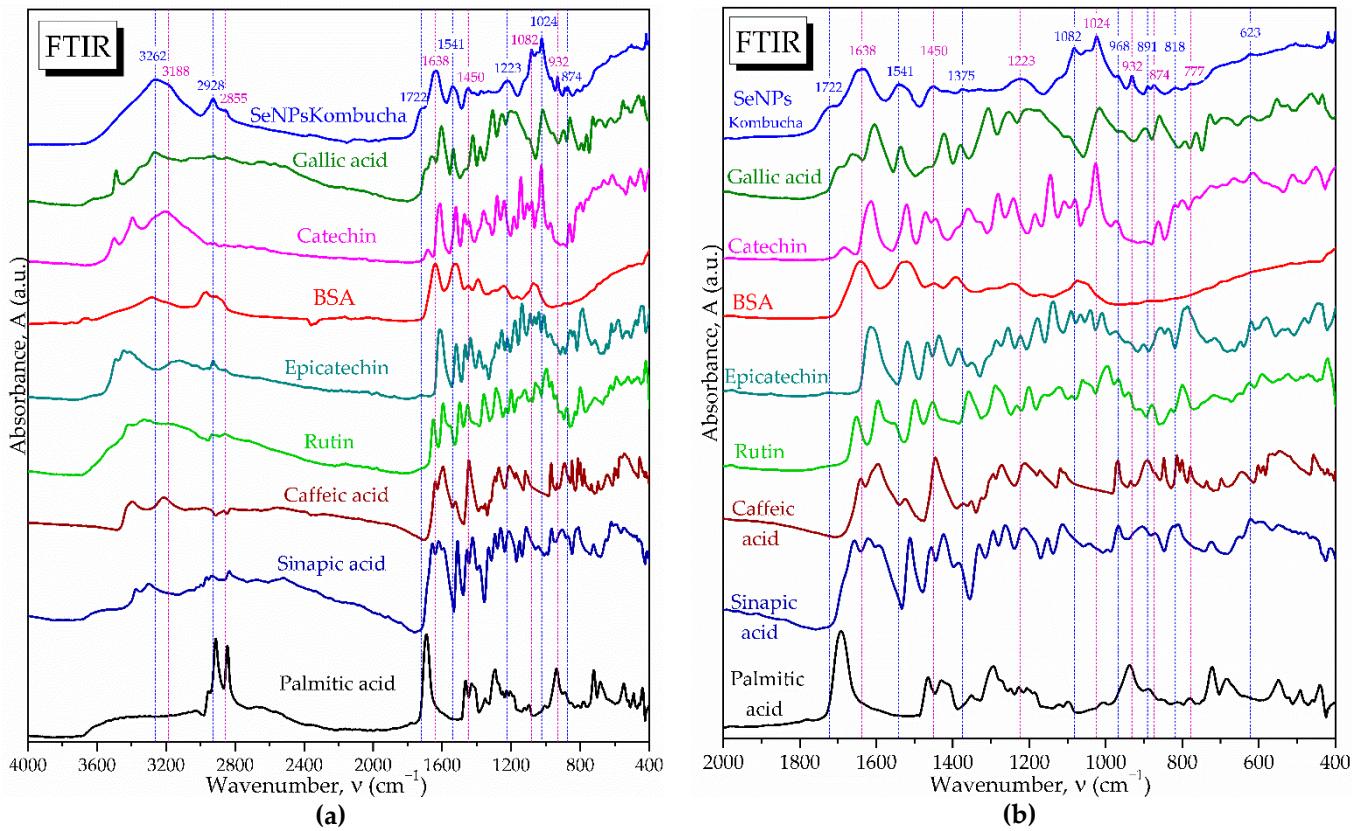


Figure S1. FTIR analysis of SeNPs Kombucha in comparison with different compounds: **(a)** Full spectral range; **(b)** Fingerprint region.

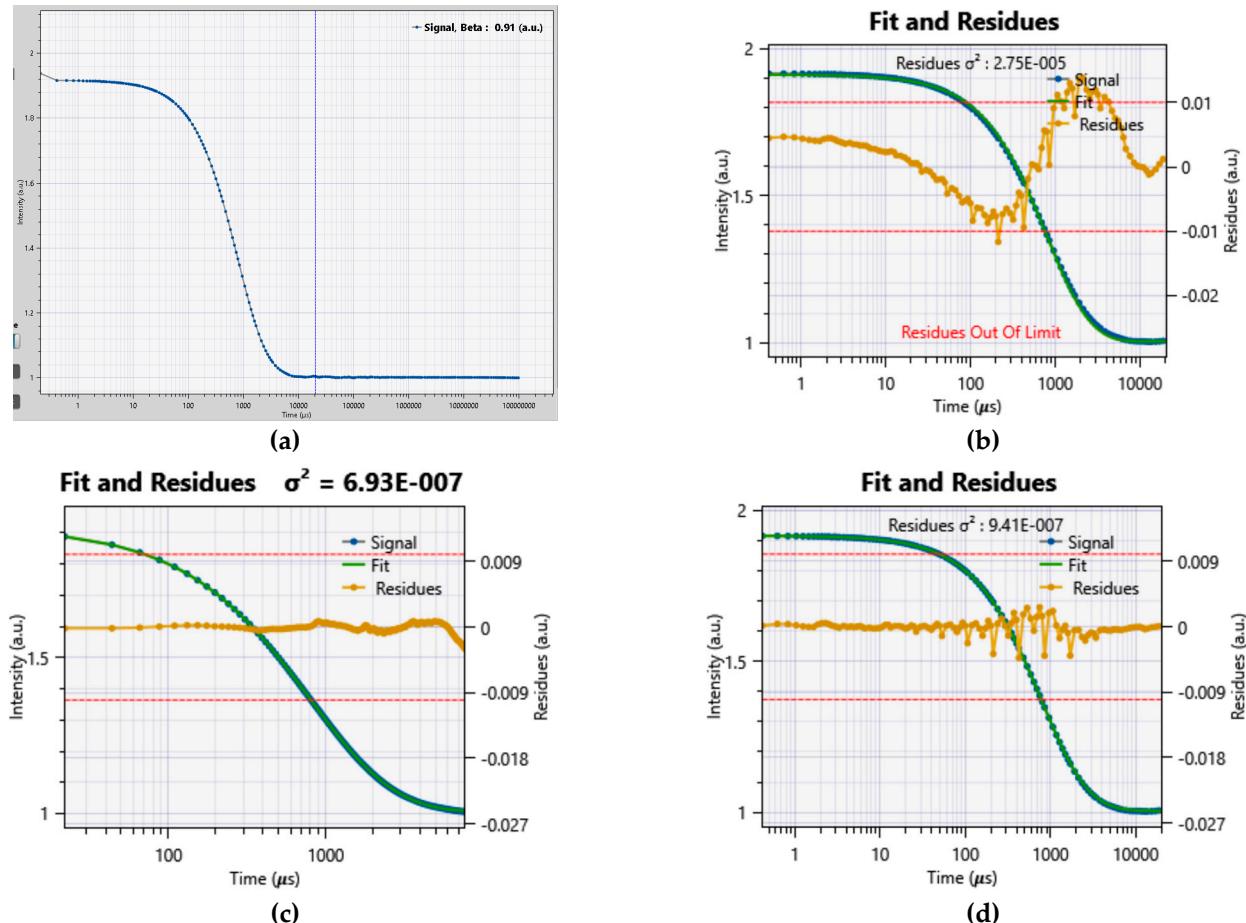
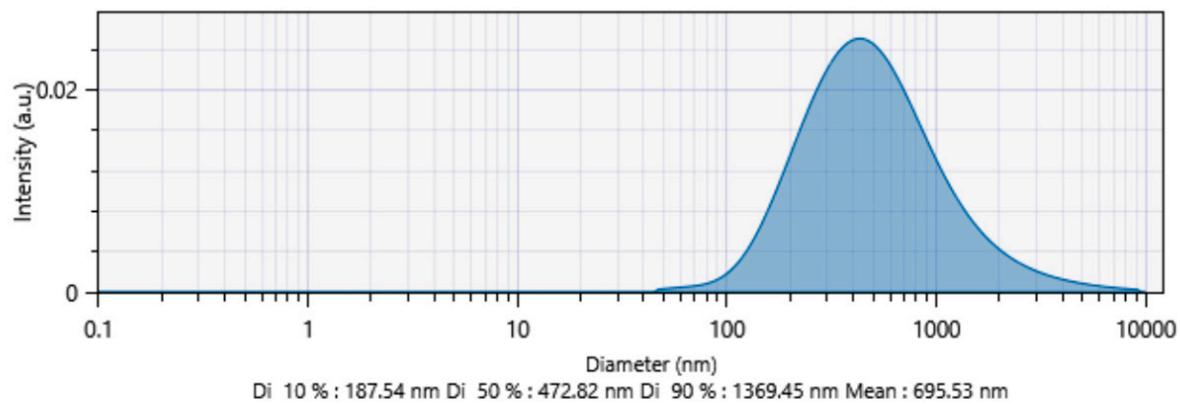
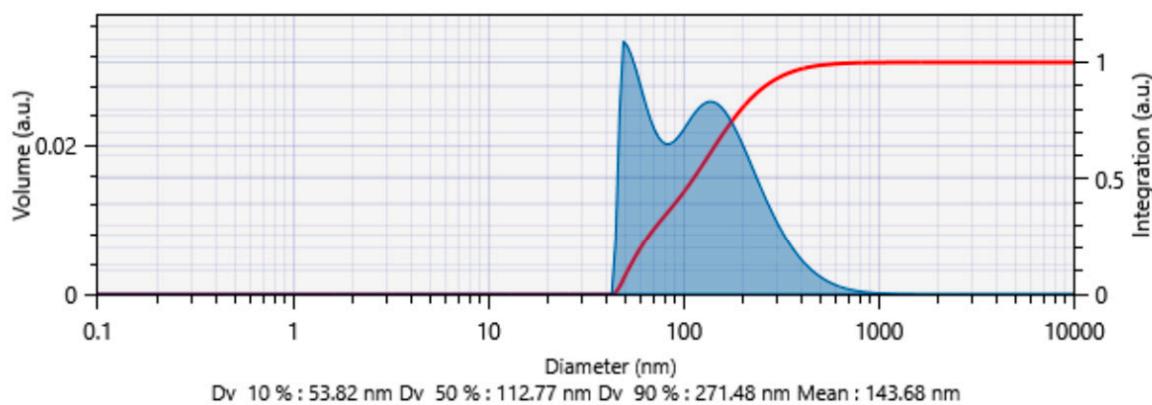


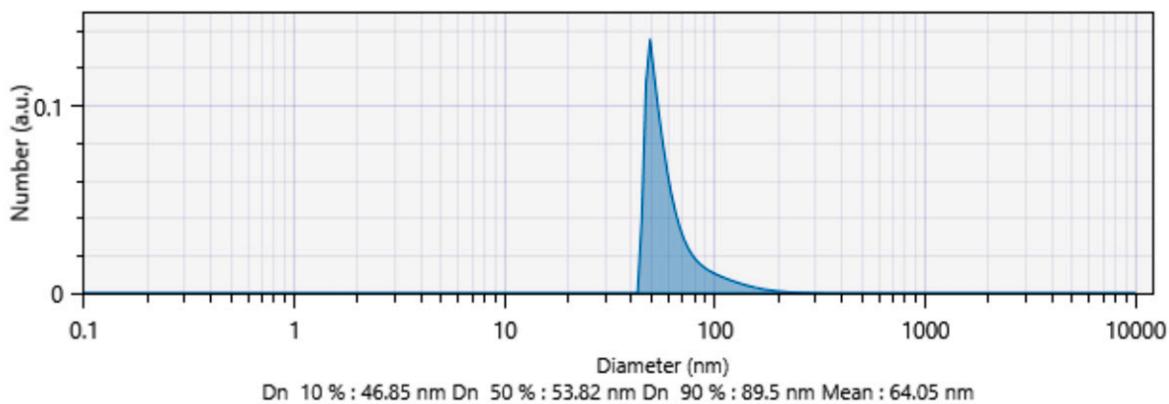
Figure S2. DLS analysis: **(a)** autocorrelation function for SeNPs Kombucha; **(b)** simulation of autocorrelation function for Cumulants method; **(c)** simulation of autocorrelation function for Pade Laplace (PL) method; **(d)** simulation of autocorrelation function for SBL method.



(a)



(b)



(c)

Figure S3. DLS analysis of SeNPs: (a) Intensity; (b) Volume; (c) Number.

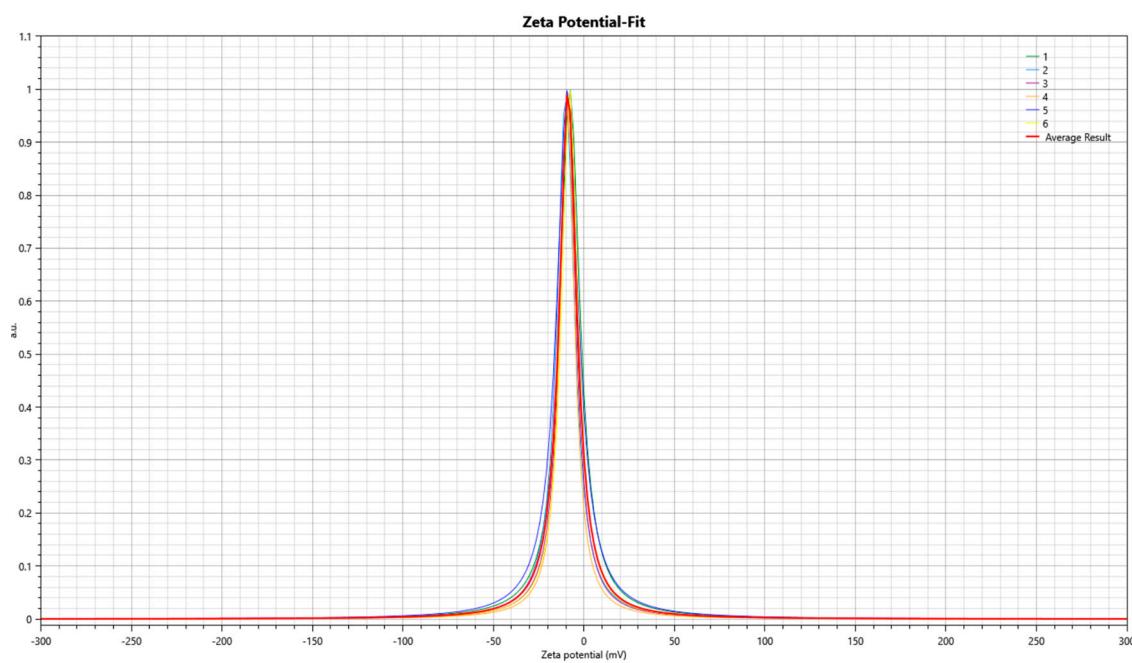


Figure S4. Zeta potential analysis of SeNPs.

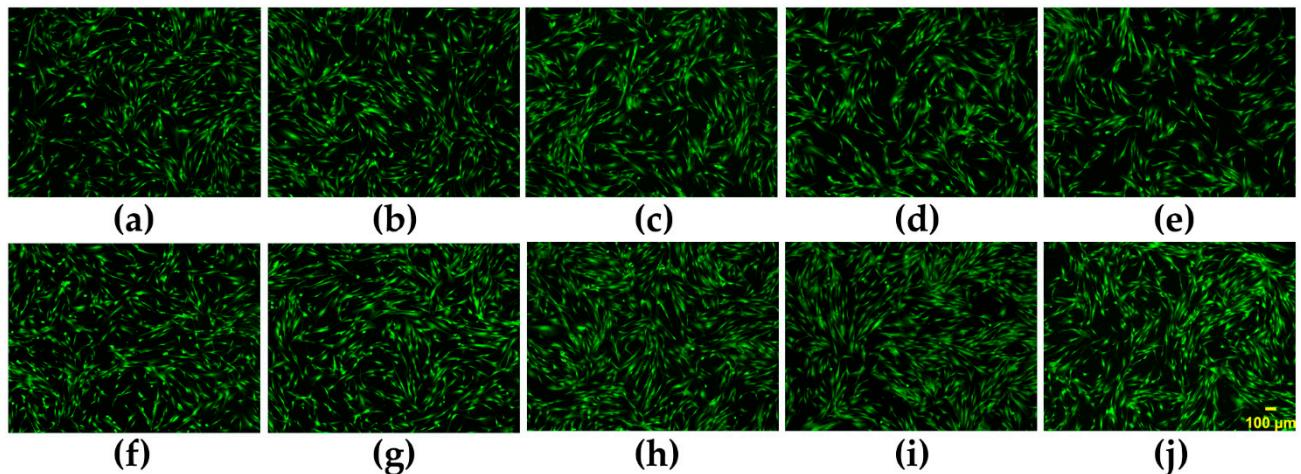


Figure S5. Biocompatibility of Kombucha beverage: (a–j) LIVE/DEAD assay (live cells – green fluorescence, dead cells – red fluorescence): (a) 0.1 mg/mL KPol5; (b) 1 mg/mL KPol5; (c) 3 mg/mL KPol5; (d) 5 mg/mL KPol5; (e) 7 mg/mL KPol5; (f) 0.1 mg/mL KPol15; (g) 1 mg/mL KPol15; (h) 3 mg/mL KPol15; (i) 5 mg/mL KPol15; (j) 7 mg/mL KPol15; KPol5 – Kombucha beverage with 30 mL SCOBY, 5 g pollen, and 190 mg sodium selenite. KPol15 – Kombucha beverage with 30 mL SCOBY, 15 g pollen, and 190 mg sodium selenite.

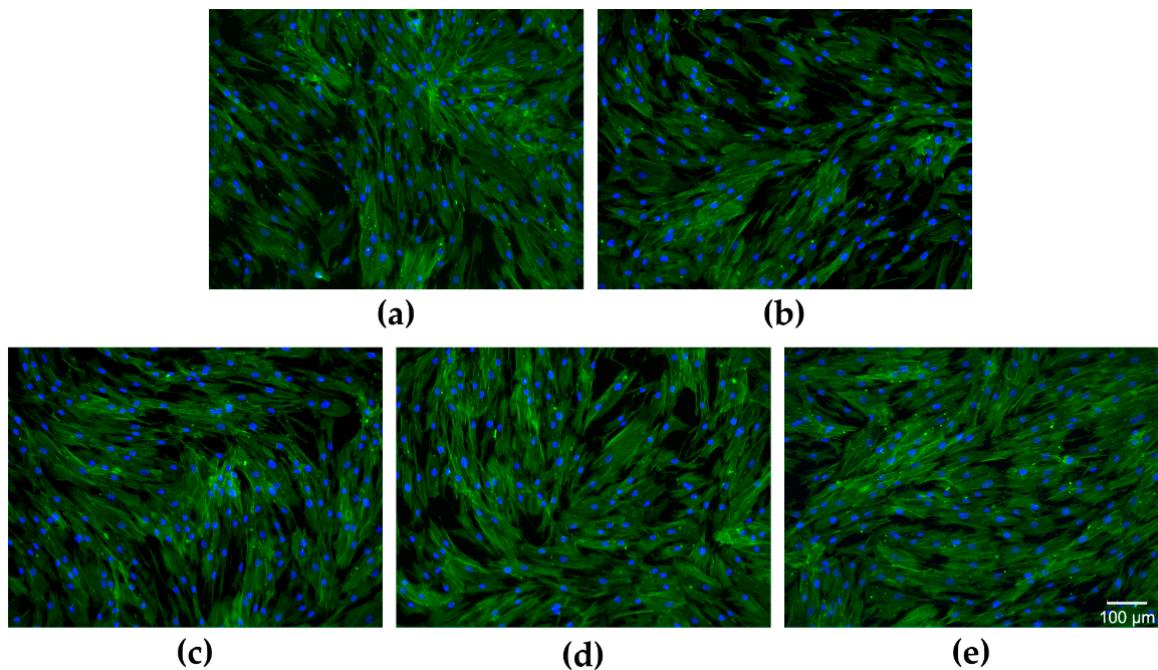


Figure S6. Cell morphology – Fluorescence microscopy images after labelling the actin cytoskeleton with Alexa Fluor 488-coupled phalloidin (green fluorescence), and the nuclei with DAPI (blue fluorescence): (a) Untreated cells (C-; Negative cytotoxicity control); HGF-1 cells treated with: (b) 3 mg/mL K; (c) 3 mg/mL KPol5; (d) 5 mg/mL KPol15; (e) 7 mg/mL KPol25; K – Kombucha beverage with 30 mL SCOBY; KPol5 – Kombucha beverage with 30 mL SCOBY, 5 g pollen, and 190 mg sodium selenite; KPol15 – Kombucha beverage with 30 mL SCOBY, 15 g pollen, and 190 mg sodium selenite, KPol25 – Kombucha beverage with 30 mL SCOBY, 25 g pollen, and 190 mg sodium selenite.

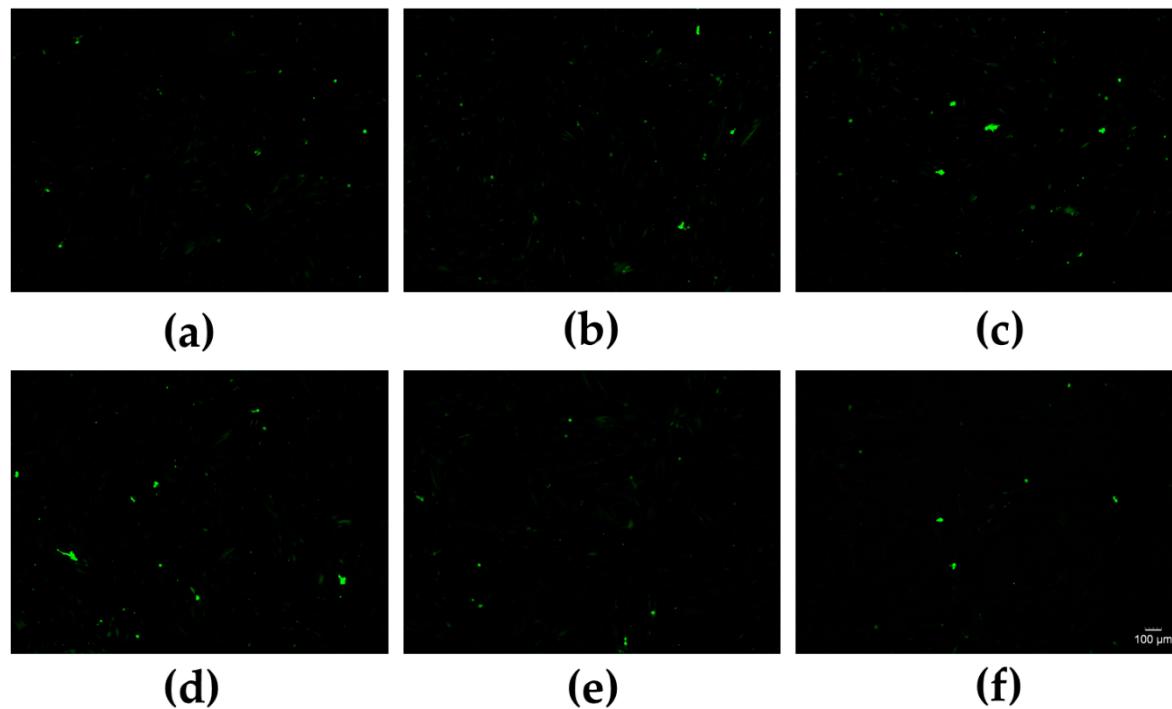


Figure S7. In vitro antioxidant activity of Kombucha beverage: (a-f) Fluorescence microscopy images after labelling total intracellular ROS with H₂DCFDA (green fluorescence): (a) 3 mg/mL KPol5; (b) 5 mg/mL KPol5; (c) 7 mg/mL; (d) 3 mg/mL KPol15; (e) 5 mg/mL KPol15; (f) 7 mg/mL KPol15; KPol5 – Kombucha beverage with 30 mL SCOBY, 5 g pollen, and 190 mg sodium selenite; KPol15 – Kombucha beverage with 30 mL SCOBY, 15 g pollen, and 190 mg sodium selenite.

Table S5. EDX analysis of fresh polyfloral pollen (P).

Element	Weight %	σ
C	62.8	0.8
O	35.6	0.8
Al	0.8	0.1
K	0.4	0.1
P	0.2	0.1

Table S6. EDX analysis of PK.

Element	Weight %	σ
C	64.8	0.7
O	32.6	0.6
N	2.6	0.8

Table S7. EDX analysis of PK25.

Element	Weight %	σ
C	67.0	0.5
O	30.6	0.4
Se	2.5	0.3

Table S8. EDX analysis of BCK.

Element	Weight %	σ
C	56.7	0.3
O	41.8	0.3
Al	0.6	0.0
P	0.5	0.0
S	0.2	0.0
Ca	0.2	0.0
K	0.1	0.0

Table S9. EDX analysis of BCKPol5.

Element	Weight %	σ
O	54.2	0.4
C	45.3	0.4
Al	0.4	0.0

Table S10. EDX analysis of BCKPol15.

Element	Weight %	σ
C	52.4	0.5
O	45.9	0.5
P	0.5	0.0
Al	0.4	0.0
K	0.3	0.0
S	0.2	0.0
Ca	0.2	0.0
Na	0.2	0.0

Table S11. EDX analysis of BCKPol25.

Element	Weight %	σ
C	51.2	0.4
O	46.7	0.4
K	0.6	0.0
Al	0.6	0.0
P	0.4	0.0
Ca	0.2	0.0
Se	0.2	0.1
S	0.2	0.0

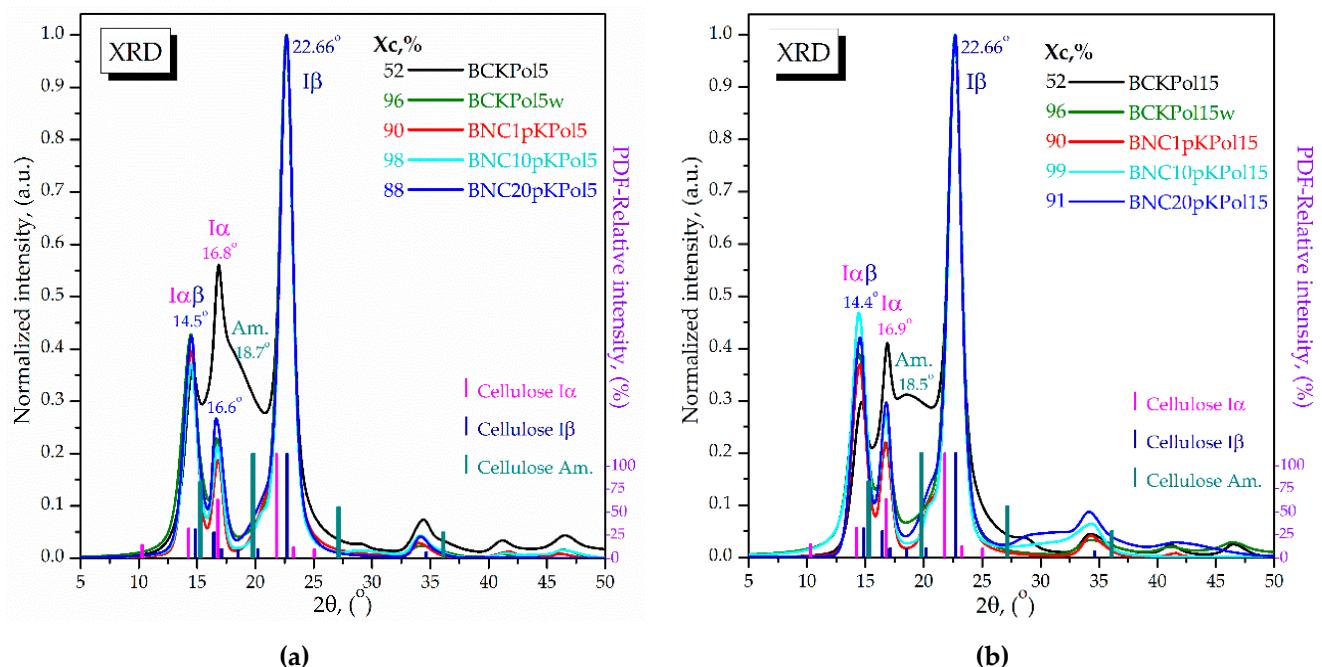


Figure S8. XRD analyses of bacterial celluloses membranes (BC) obtained by purification and microfluidization: (a) BCKPol5 – Bacterial cellulose from Kombucha beverage prepared with 30 mL SCOBY, 5 g pollen, and 190 mg sodium selenite; BCKPol5w – Purified/washed bacterial cellulose; BNC1pKPol5 – Bacterial nanocellulose after 1 pass of microfluidization; BNC10pKPol5 – Bacterial nanocellulose after 10 passes of microfluidization; BNC20pKPol5 – Bacterial nanocellulose after 20 passes of microfluidization; (b) BCKPol15 – Bacterial cellulose from Kombucha beverage prepared with 30 mL SCOBY, 15 g pollen, and 190 mg of sodium selenite; BCKPol15w – Purified/washed bacterial cellulose; BNC1pKPol15 – Bacterial nanocellulose after 1 pass of microfluidization; BNC10pKPol15 – Bacterial nanocellulose after 10 passes of microfluidization; BNC20pKPol15 – Bacterial nanocellulose after 20 passes of microfluidization. The purification of BCKPol5 and BCKPol15 increases the crystallinity degree from 52% to 96% in washed membranes (w), while 10 passes of microfluidization induce a partial arrangement of amorphous cellulose into a $I_{\alpha\beta}$ structure, as suggested by the increased intensity of the $I_{\alpha\beta}$ peak around 14.5° and increased crystallinity up to 98-99%. Subsequent microfluidization up to 20 passes shortens the cellulose microfibrils and partially disintegrate the I_{α} and $I_{\alpha\beta}$ structure into amorphous cellulose, decreasing the crystallinity degree to 88% for BNC20pKPol5, respectively 91% for BNC20pKPol15.

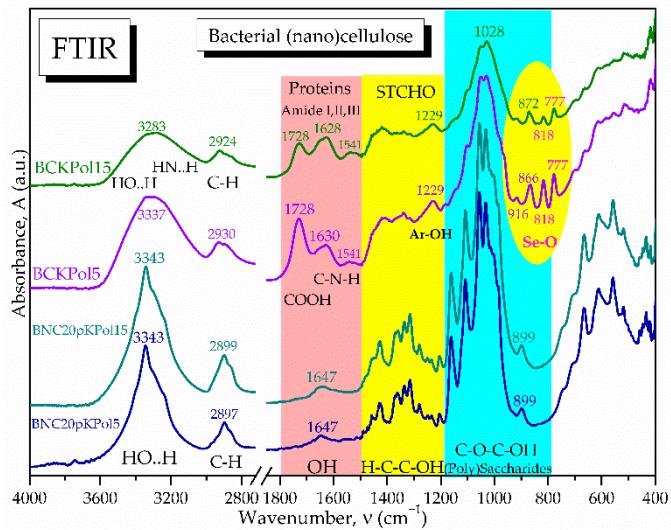


Figure S9. ATR-FTIR spectra of initial membranes of bacterial celluloses (BC) compared with the corresponding bacterial nanocelluloses (BNC) obtained by purification and microfluidization; BCKPol5 – Bacterial cellulose from Kombucha beverage prepared with 30 mL SCOBY, 5 g pollen, and 190 mg sodium selenite; BNC20pKPol5 – Bacterial nanocellulose after 20 passes of microfluidization; BCKPol15 – Bacterial cellulose from Kombucha beverage prepared with 30 mL SCOBY, 15 g pollen, and 190 mg of sodium selenite; BNC20pKPol15 – Bacterial nanocellulose after 20 passes of microfluidization.

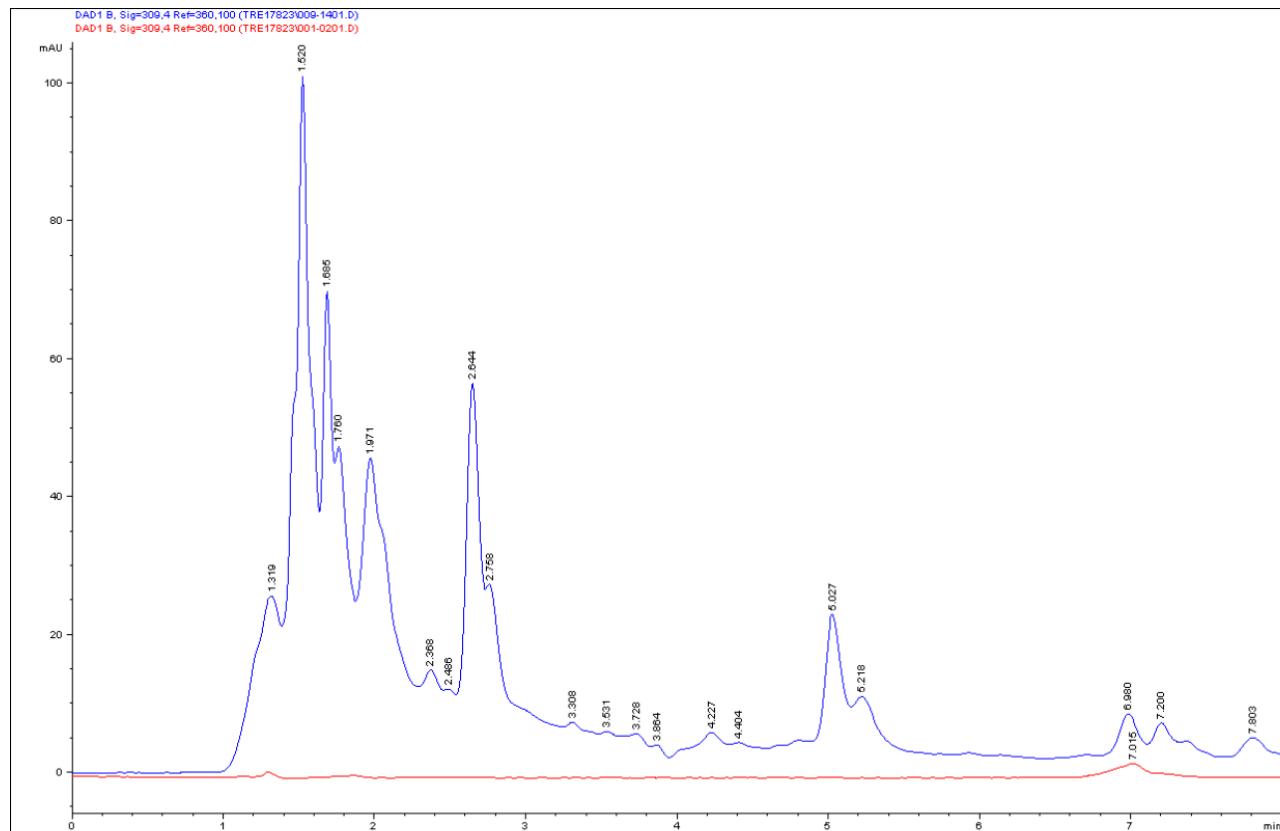


Figure S10. HPLC-DAD chromatogram analysis of black tea (blue) and t-resveratrol reference material (red); t-resveratrol ($t_r = 7.015$ min for reference material, $t_r = 6.980$ min. for sample; $\lambda = 309$ nm)

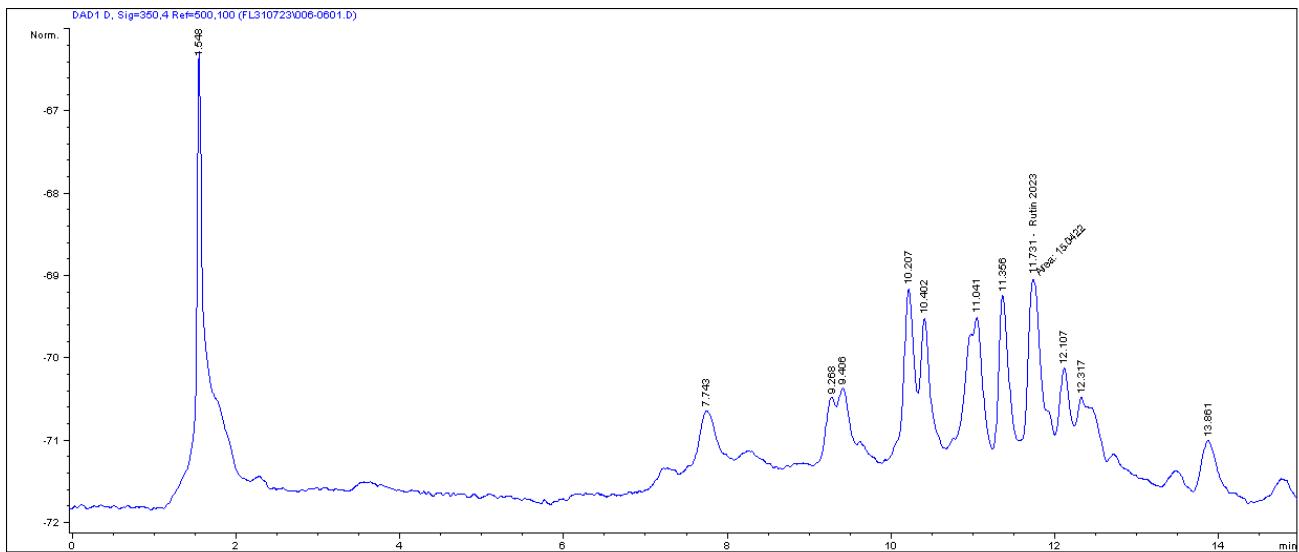


Figure 11. HPLC-DAD chromatogram analysis of black tea sample: quercetin-3-rutinoside ($tr = 11.73 \text{ min}$, $\lambda = 350 \text{ nm}$)