



Supplementary Materials: Fabrication, Optimization, and Characterization of Antibacterial Electrospun Shellac Fibers Loaded with *Kaempferia parviflora* Extract

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Table S1. Experimental design matrix and corresponding responses based on experimental runs proposed by BBD design.

Run	Factor 1 (x ₁)	Factor 2 (x ₂)	Factor 3 (x ₃)	Factor 4 (x ₄)	Response 1	Response 2
	Shellac concentration (% w/w)	Feed rate (mL/h)	Electrical voltage (kV)	Extract concentration (% w/w)	Fiber diameters (nm), n = 3 Mean ± SD	Bead-to-fiber ratio, n = 3 Mean ± SD
1	38	0.8	18	3	832 ± 6.07	0.24 ± 0.06
2	38	0.8	24	1	824 ± 7.56	0.32 ± 0.07
3	38	0.4	18	1	635 ± 8.40	0.41 ± 0.03
4	36	0.8	18	5	724 ± 8.88	0.42 ± 0.01
5	36	1.2	18	3	495 ± 9.14	0.58 ± 0.02
6	40	0.4	18	3	915 ± 9.23	0.04 ± 0.03
7	38	0.8	12	1	606 ± 9.20	0.30 ± 0.02
8	40	0.8	18	1	819 ± 9.06	0.14 ± 0.01
9	40	0.8	18	5	1149 ± 16.65	0.07 ± 0.01
10	40	0.8	24	3	962 ± 9.29	0.12 ± 0.02
11	40	0.8	12	3	864 ± 12.72	0.10 ± 0.01
12	38	0.4	24	3	807 ± 5.81	0.26 ± 0.02
13	40	1.2	18	3	878 ± 6.92	0.08 ± 0.00
14	38	0.8	18	3	766 ± 8.11	0.27 ± 0.03
15	36	0.4	18	3	475 ± 9.12	0.42 ± 0.06
16	38	1.2	18	1	473 ± 3.02	0.23 ± 0.04
17	38	1.2	24	3	788 ± 6.94	0.11 ± 0.02
18	38	1.2	18	5	954 ± 12.20	0.15 ± 0.03
19	38	1.2	12	3	728 ± 10.21	0.25 ± 0.04
20	38	0.4	18	5	1015 ± 29.34	0.28 ± 0.02
21	38	0.8	24	5	902 ± 25.32	0.21 ± 0.04
22	36	0.8	18	1	433 ± 4.02	0.86 ± 0.02
23	38	0.8	18	3	785 ± 4.86	0.30 ± 0.03
24	36	0.8	24	3	527 ± 7.43	0.81 ± 0.05
25	36	0.8	12	3	524 ± 8.83	0.78 ± 0.06
26	38	0.8	12	5	545 ± 9.19	0.29 ± 0.02
27	38	0.4	12	3	667 ± 14.45	0.38 ± 0.04

Table S2. ANOVA results for the prediction of fiber diameter and bead-to-fiber ratio of electrospun shellac fibers using the BBD matrix.

Source	Coefficients	p-value	Standard deviation	% CV	Adequate precision	R ²	F-value
Y ₁ (diameter)			89.83	12.07	18.836	0.7983	
x ₁	200.66	< 0.0001					
x ₂	72.96	0.0099					
x ₄	124.97	< 0.0001					
Intercept	744.07						
Model		< 0.0001				30.34	
Lack of fit		0.1245				7.47	
Y ₂ (bead-to-fiber ratio)			0.11	35.76	18.598	0.7652	
x ₁	-0.28	< 0.0001					
x ₄	-0.07	0.0408					
Intercept	0.31						
Model		< 0.0001				39.10	
Lack of fit		0.0643				14.99	

Table S3. Verification of experimental results in relation to the predicted values generated by the designed models.

Run	Fiber diameters (nm)			Bead-to-fiber ratios		
	Predictive value	Acceptable interval (95% confident level)		Predictive value	Acceptable interval (95% confident level)	
		Experimental 1 value	Experimental 1 value		Experimental 1 value	Experimental 1 value
1			582.00			0.43
2	574.71	504.83-633.30	589.00	0.34	0.10-0.57	0.29
3			573.00			0.33

Table S4. Hot-stage microscope (HSM) images of shellac flakes and fibers at elevated temperatures.

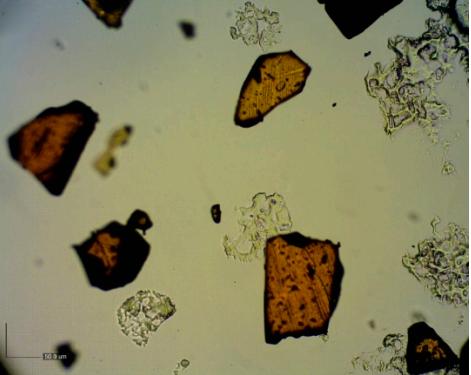
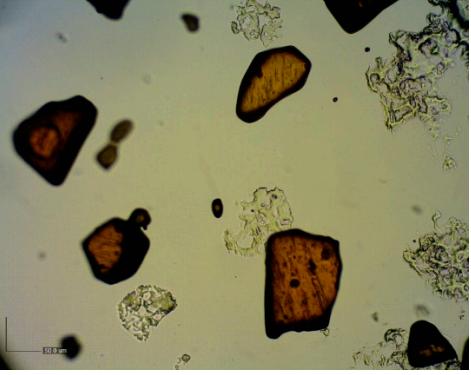
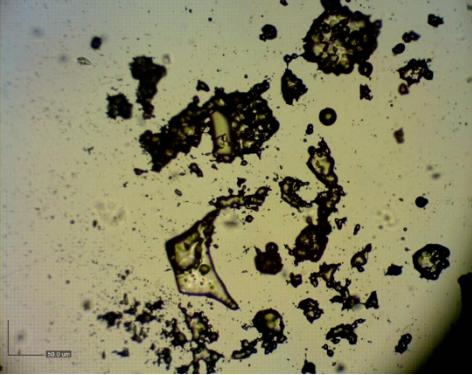
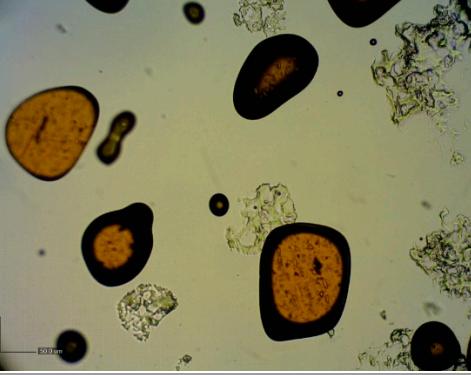
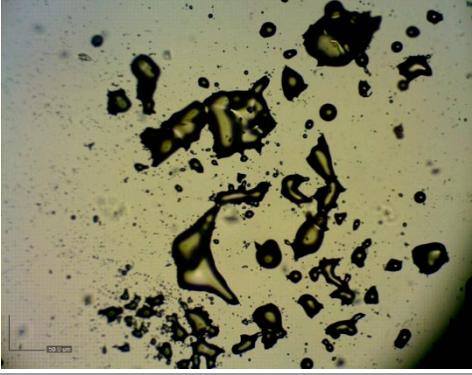
Temperature (°C)	HSM images	
	Shellac flakes	Shellac fibers
50		
60		
70		

Table S5. Total methoxyflavones release kinetics models of KP extract-loaded electrospun shellac fibers with various formulations.

Modelling	Formulation	Parameters	R ² Adjusted	AIC	MSC
Zero-order	36% (w/w) Shellac (Run 5)	k ₀ = 0.658	0.6261	33.9314	0.5837
	38% (w/w) Shellac (Run 1)	k ₀ = 0.362	0.8586	29.3385	1.5559
	40% (w/w) Shellac (Run 10)	k ₀ = 0.279	0.7723	23.8660	1.0799
	1% (w/w) KP extract (Run 8)	k ₀ = 0.707	0.7320	32.7120	0.9168
	3% (w/w) KP extract (Run 13)	k ₀ = 0.717	0.9303	27.1213	2.2633
	5% (w/w) KP extract (Run 9)	k ₀ = 0.548	0.1348	34.5053	-0.2552
First-order	Run optimized	k ₀ = 0.615	0.7643	31.5195	1.0452
	36% (w/w) Shellac (Run 5)	k ₁ = 0.011	0.8456	30.0709	1.3558
	38% (w/w) Shellac (Run 1)	k ₁ = 0.005	0.8301	30.8168	1.2602
	40% (w/w) Shellac (Run 10)	k ₁ = 0.003	0.8474	21.8649	1.4802
	1% (w/w) KP extract (Run 8)	k ₁ = 0.012	0.9643	22.6377	2.9317
	3% (w/w) KP extract (Run 13)	k ₁ = 0.012	0.9413	25.6829	2.3226
	5% (w/w) KP extract (Run 9)	k ₁ = 0.008	0.6404	30.1149	0.6228
	Run optimized	k ₁ = 0.009	0.9037	27.6057	1.8279
Higuchi	36% (w/w) Shellac (Run 5)	k _H = 6.166	0.8683	28.7122	1.6275
	38% (w/w) Shellac (Run 1)	k _H = 3.948	0.7251	32.6617	0.8913
	40% (w/w) Shellac (Run 10)	k _H = 2.594	0.8548	21.6160	1.5299
	1% (w/w) KP extract (Run 8)	k _H = 6.605	0.9303	25.9781	2.2336
	3% (w/w) KP extract (Run 13)	k _H = 6.606	0.9096	28.4231	2.0030
	5% (w/w) KP extract (Run 9)	k _H = 5.224	0.8553	25.5361	1.5332
	Run optimized	k _H = 5.724	0.8888	27.7630	1.7965
	36% (w/w) Shellac (Run 5)	k_{KP} = 4.217 n = 0.698	0.8995	27.3629	1.8974
Kosmeyer-Peppas	38% (w/w) Shellac (Run 1)	k_{KP} = 0.830 n = 0.612	0.8859	28.2652	1.7706
	40% (w/w) Shellac (Run 10)	k_{KP} = 1.242 n = 0.669	0.8684	21.6875	1.5156
	1% (w/w) KP extract (Run 8)	k_{KP} = 4.060 n = 0.612	0.9643	20.8333	3.5210
	3% (w/w) KP extract (Run 13)	k_{KP} = 2.245 n = 0.747	0.9957	13.7837	4.9309
	5% (w/w) KP extract (Run 9)	k_{KP} = 6.387 n = 0.554	0.9815	13.2558	1.1850
	Run optimized	k_{KP} = 3.012 n = 0.647	0.9519	23.5706	2.6350

Note: The significance of **bold** reveals the estimated parameters derived from the best-fit mathematical model.

Table S6. Antimicrobial activity of KP extract-loaded electrospun shellac fibers as determined by time-kill kinetics assay.

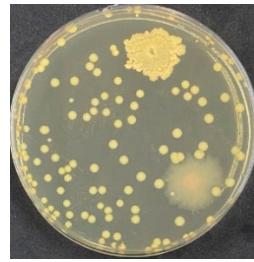
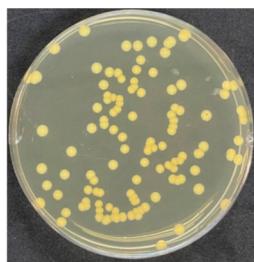
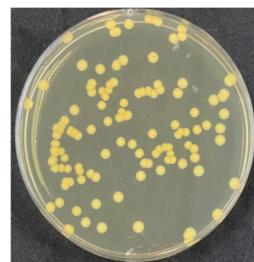
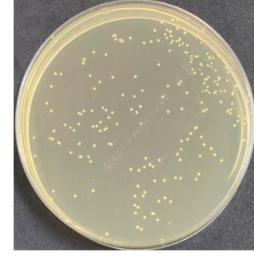
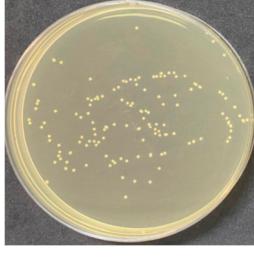
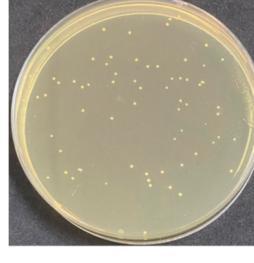
	Time (min)		
	15	30	60
Digital image			
Microbial load (CFU/mL)	1.09×10^6	1.87×10^6	1.18×10^6
R (%)	-	-	-
<i>S. aureus</i>			
Digital image			
Microbial load (CFU/mL)	7.80×10^5	1.60×10^5	1.30×10^5
R (%)	28.44	85.32	88.07
	Time (min)		
	15	30	60
Digital image			
Microbial load (CFU/mL)	1.09×10^6	1.62×10^6	2.62×10^6
R (%)	-	-	-
<i>E. coli</i>			
Digital image			
Microbial load (CFU/mL)	1.34×10^6	9.99×10^5	8.60×10^5
R (%)	-	9.17	21.10

Table S6 (continue). Antimicrobial activity of KP extract-loaded electrospun shellac fibers as determined by time-kill kinetics assay.

	Time (min)		
	15	30	60
Digital image			
Microbial load (CFU/mL)	3.24×10^7	3.56×10^7	3.60×10^7
P. <i>aeruginosa</i>	R (%) -	-	-
	Time (min)		
	120	240	360
Digital image			
Microbial load (CFU/mL)	5.71×10^7	5.75×10^7	5.82×10^7
R (%)	-	-	-

R, the value of antimicrobial activity (%).

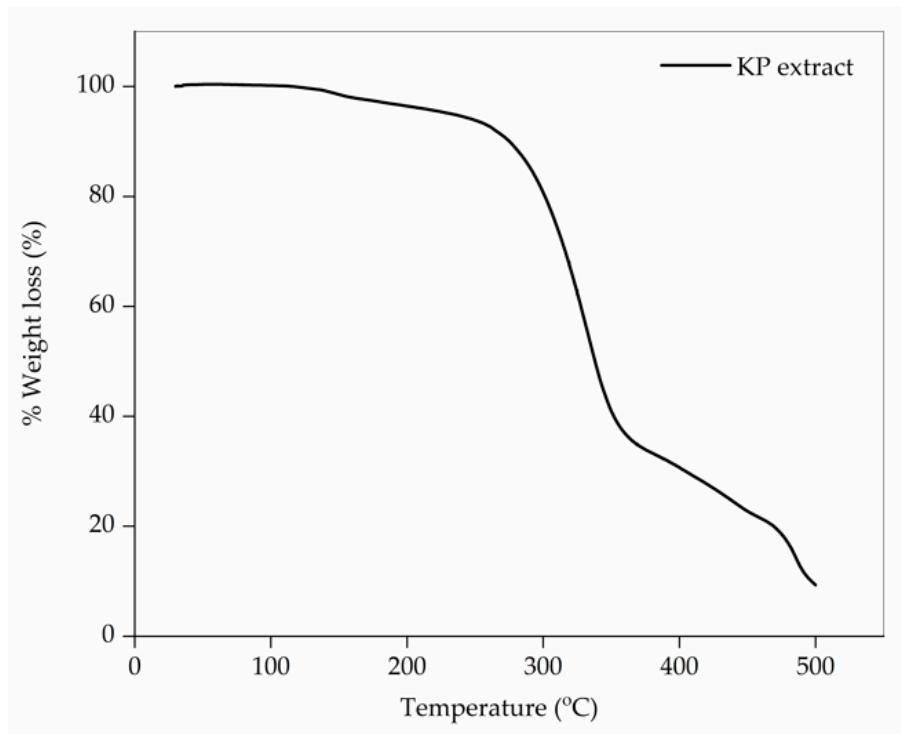


Figure S1. TGA thermogram of KP extract.

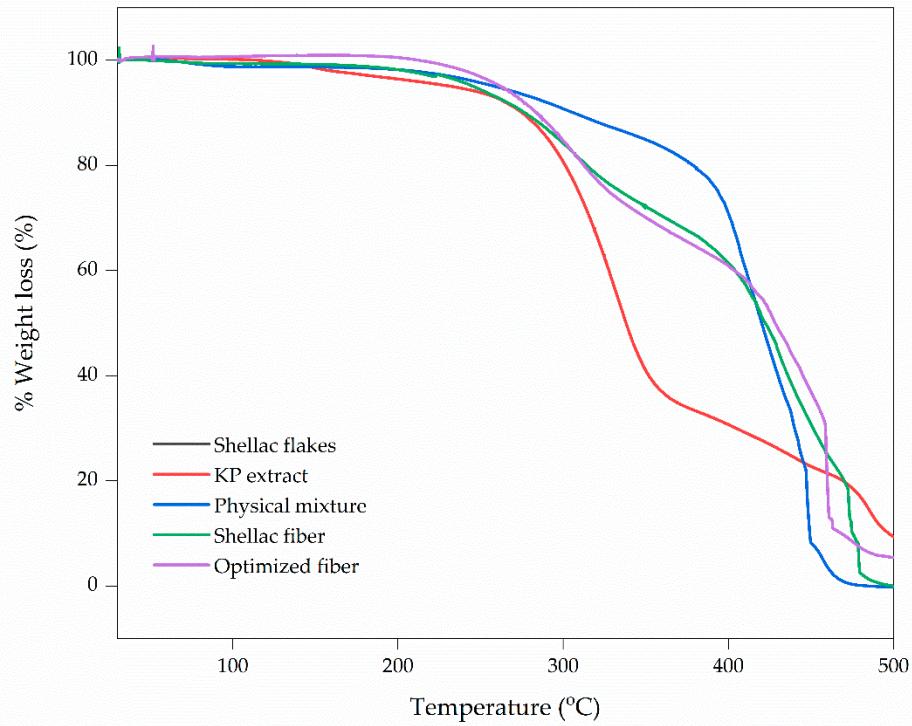


Figure S2. TGA thermograms of samples.

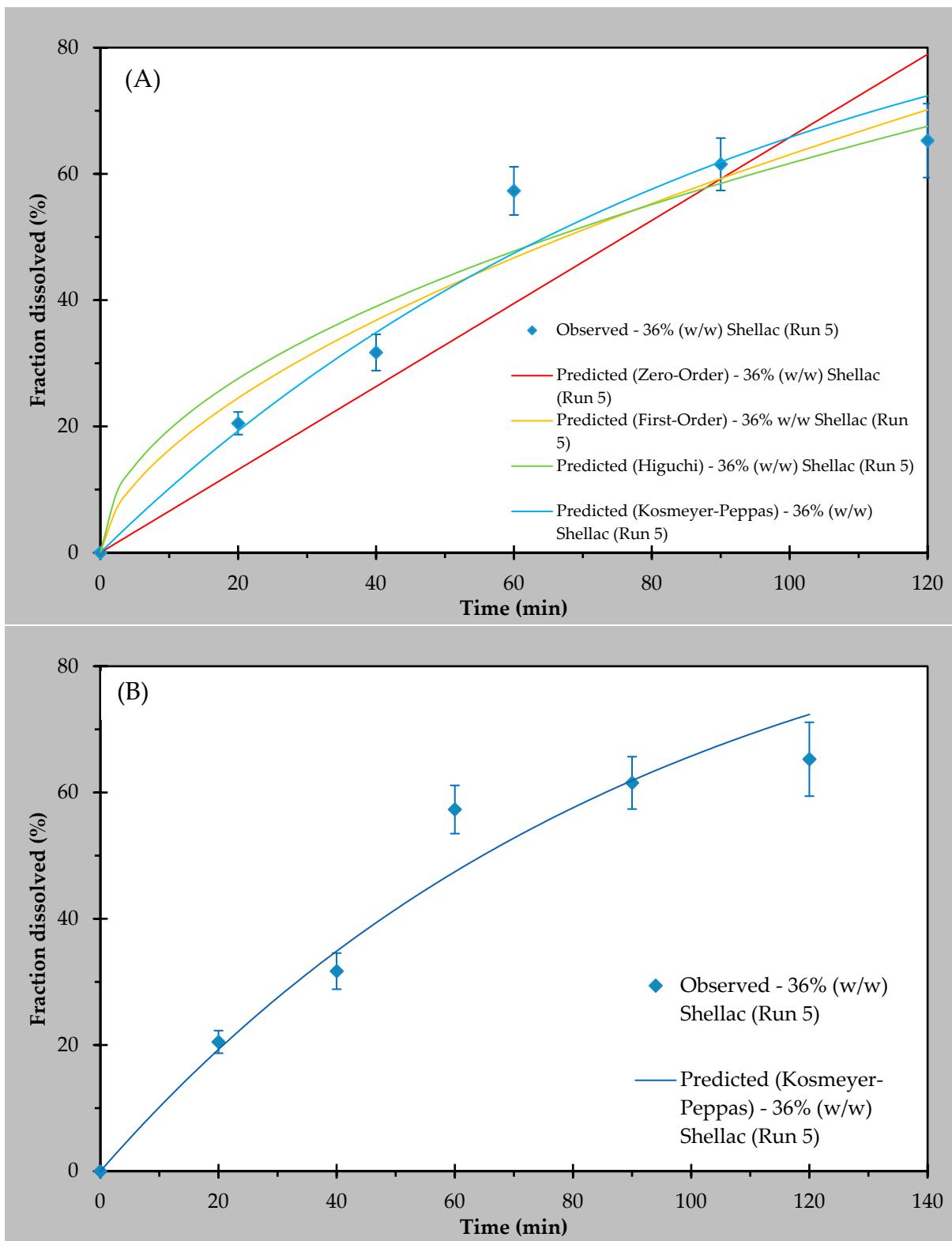


Figure S3. A mathematical model fits the KP extract release profiles of KP extract-loaded electrospun shellac fiber containing shellac and KP extract concentrations of 36% (*w/w*) and 3% (*w/w*), respectively: (A) All mathematical model fits, (B) The best fit model.

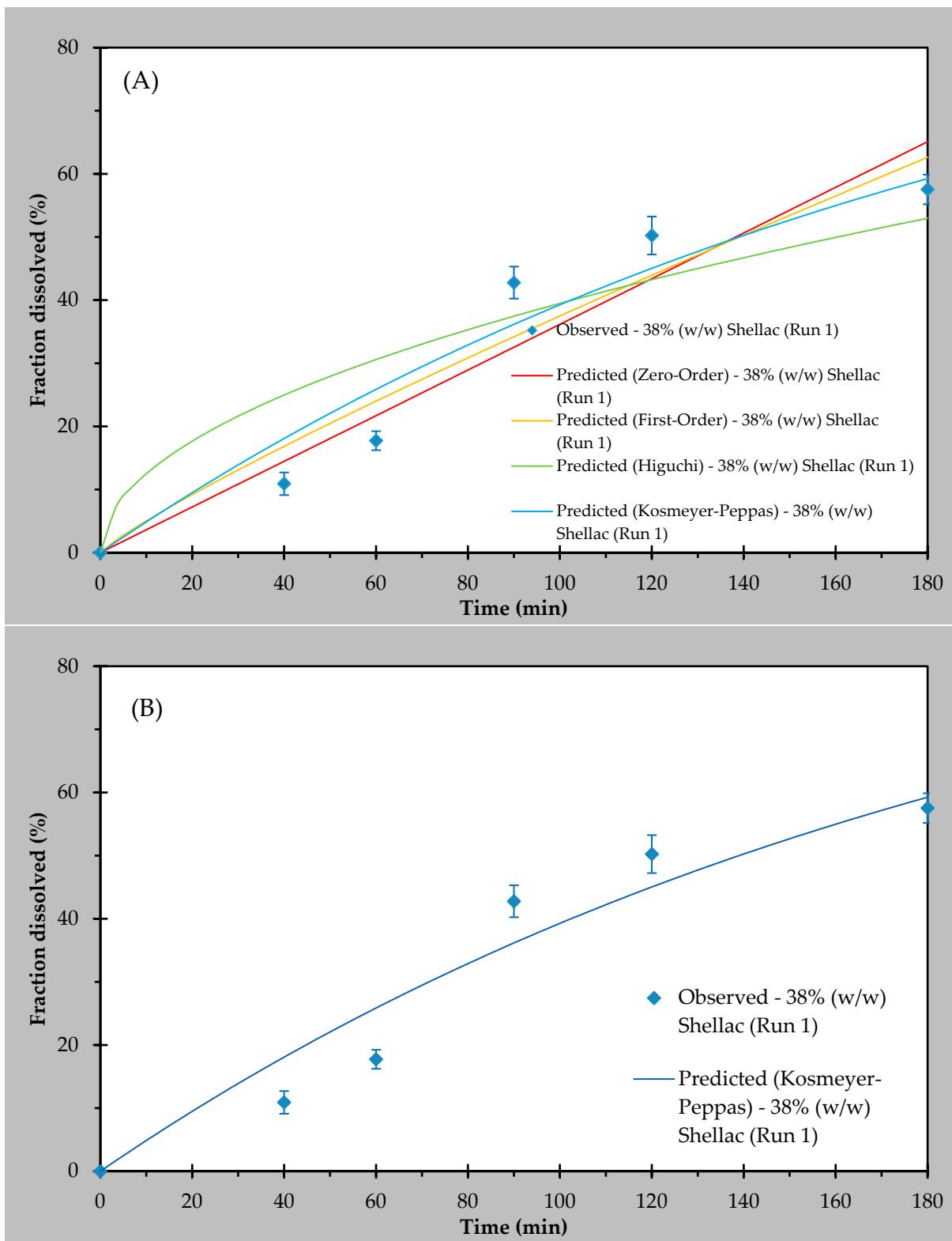


Figure S4. A mathematical model fits the KP extract release profiles of KP extract-loaded electrospun shellac fiber containing shellac and KP extract concentrations of 38% (*w/w*) and 3% (*w/w*), respectively: (A) All mathematical model fits, (B) The best fit model.

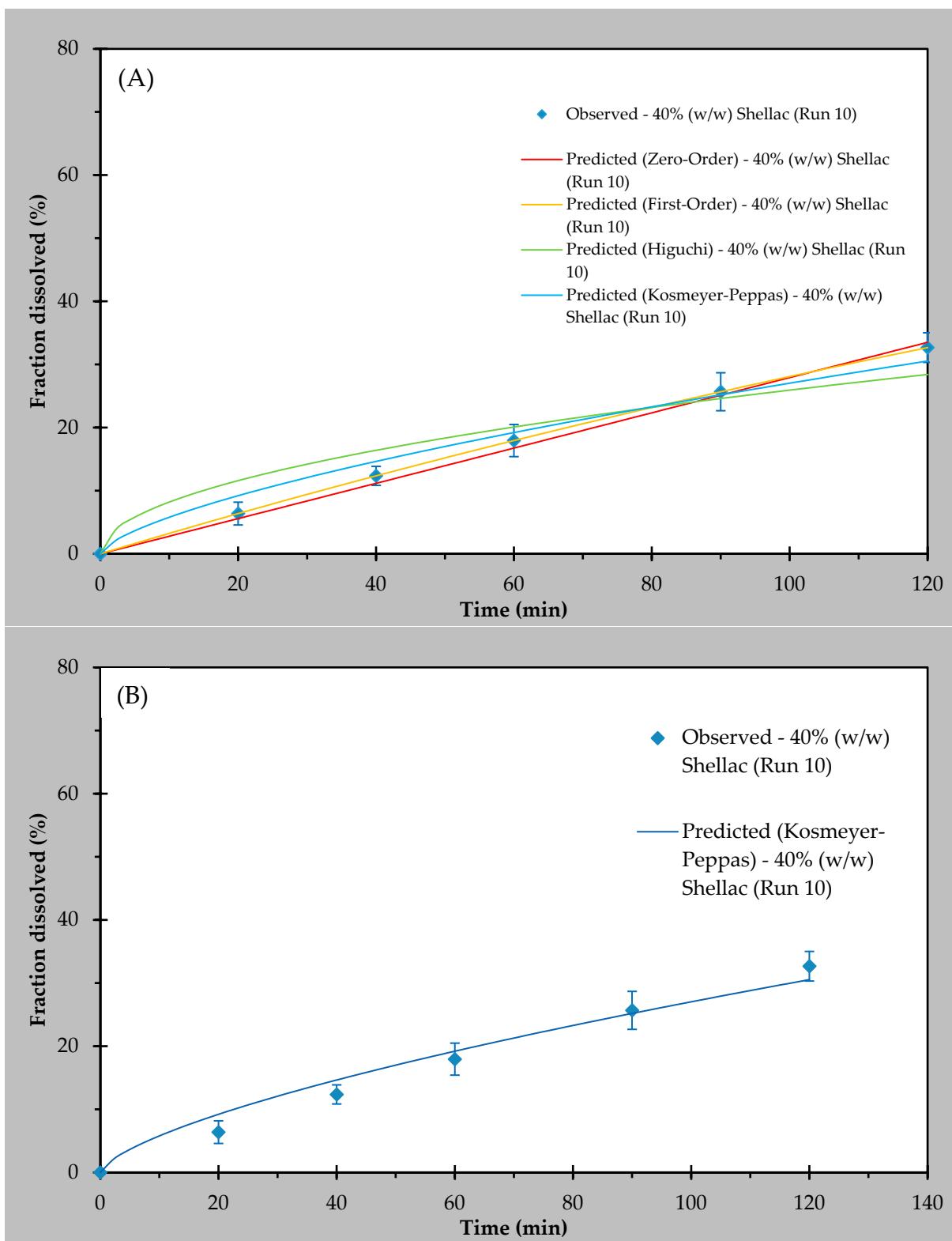


Figure S5. A mathematical model fits the KP extract release profiles of KP extract-loaded electrospun shellac fiber containing shellac and KP extract concentrations of 40% (w/w) and 3% (w/w), respectively: (A) All mathematical model fits, (B) The best fit model.

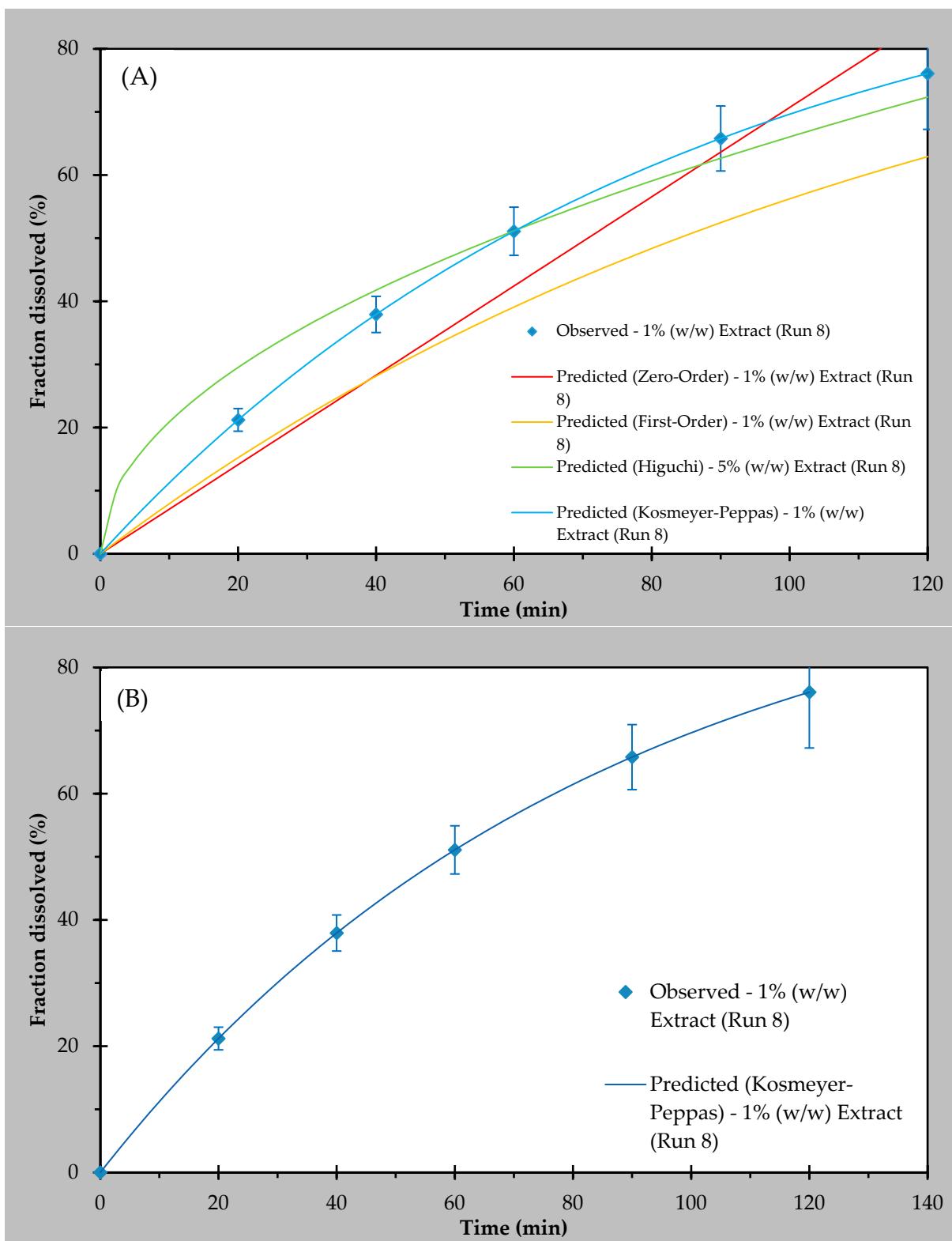


Figure S6. A mathematical model fits the KP extract release profiles of KP extract-loaded electrospun shellac fiber containing shellac and KP extract concentrations of 40% (*w/w*) and 1% (*w/w*), respectively: (A) All mathematical model fits, (B) The best fit model.

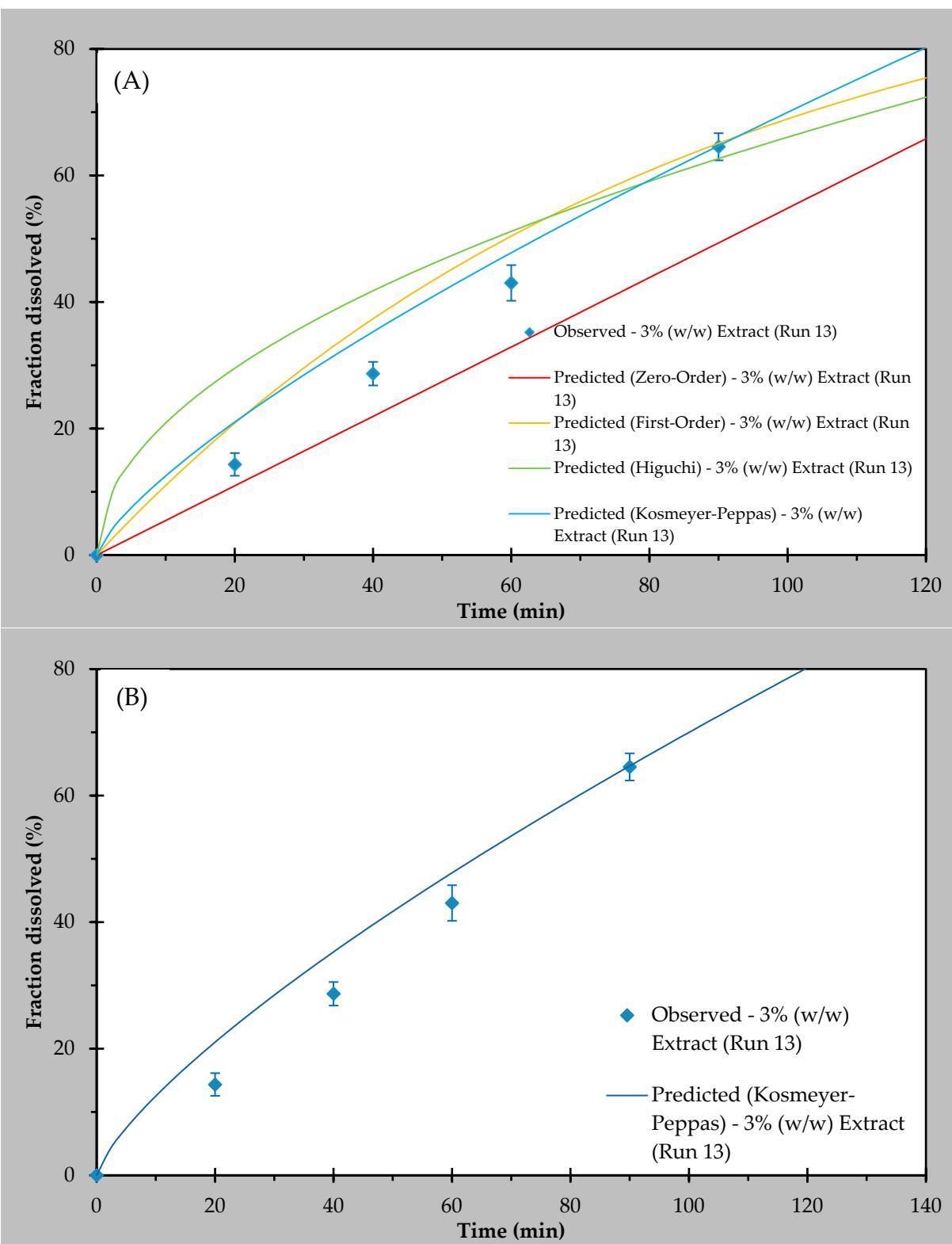


Figure S7. A mathematical model fits the KP extract release profiles of KP extract-loaded electrospun shellac fiber containing shellac and KP extract concentrations of 40% (*w/w*) and 3% (*w/w*), respectively: (A) All mathematical model fits, (B) The best fit model.

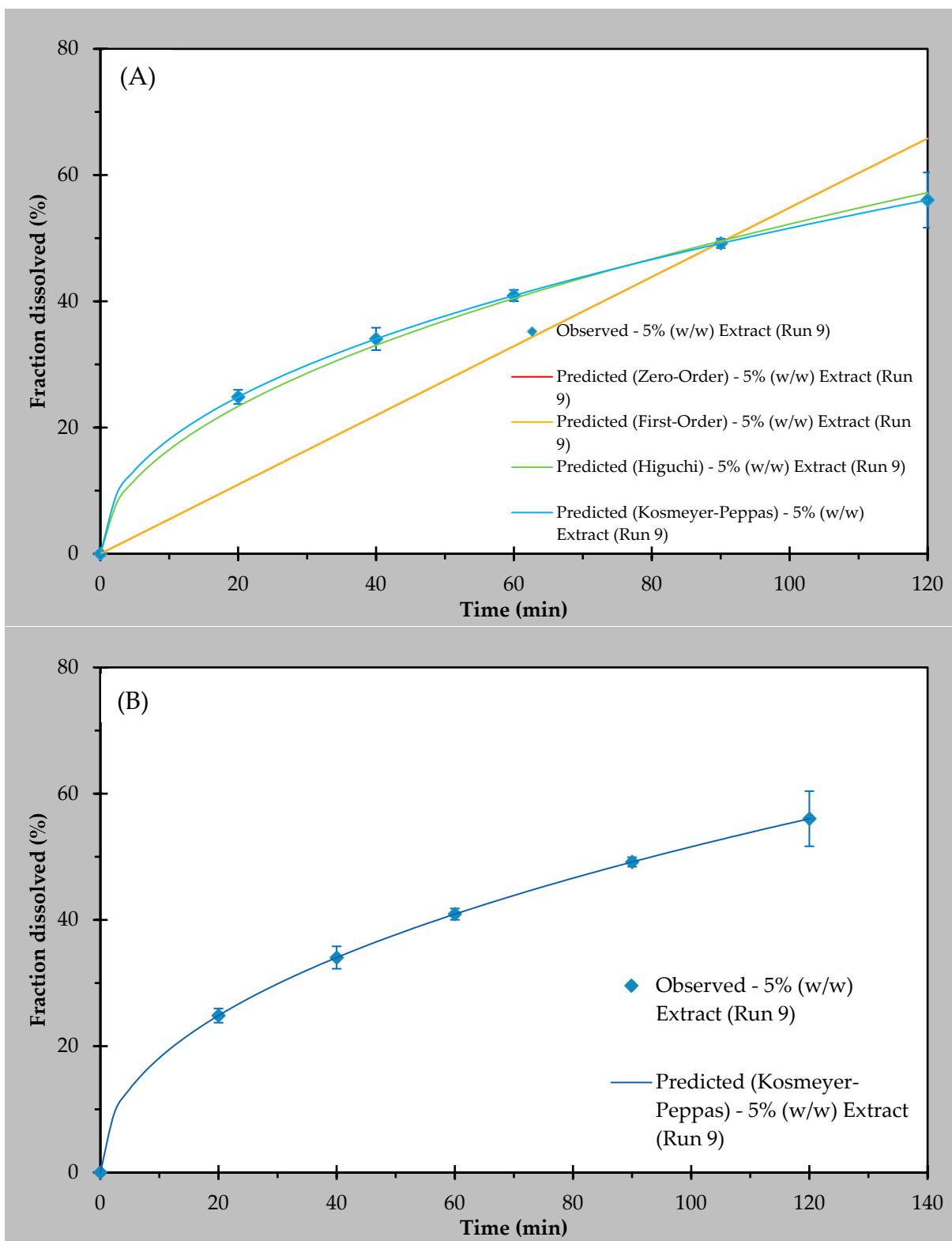


Figure S8. A mathematical model fits the KP extract release profiles of KP extract-loaded electrospun shellac fiber containing shellac and KP extract concentrations of 40% (*w/w*) and 5% (*w/w*), respectively: (A) All mathematical model fits, (B) The best fit model.

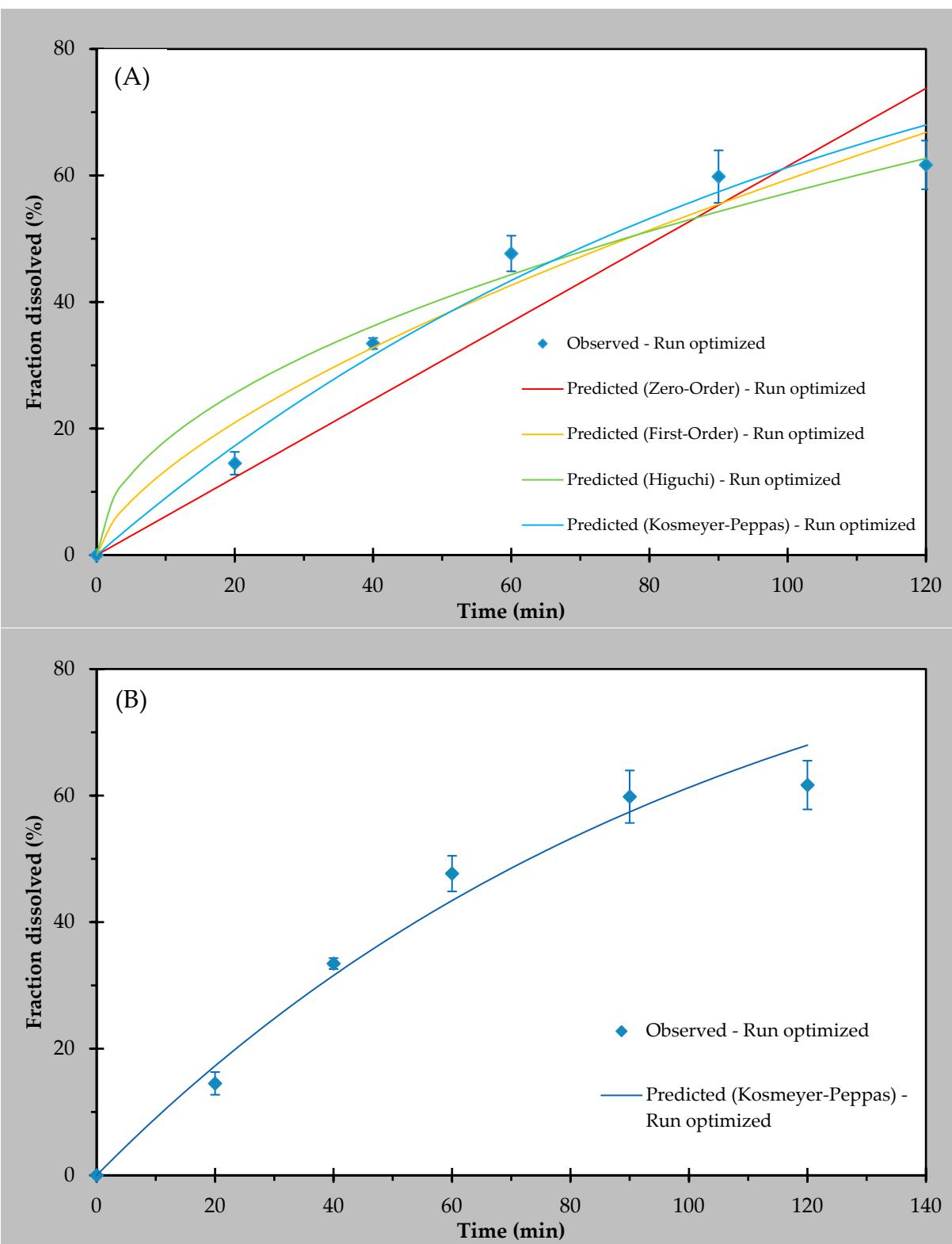


Figure S9. A mathematical model fits the KP extract release profiles of optimized KP extract-loaded electrospun shellac fiber containing shellac and KP extract concentrations of 37.25% (*w/w*) and 1.50% (*w/w*), respectively: (A) All mathematical model fits, (B) The best fit model.