

## Supplementary Material

### **Characterization of Bacterial Cellulose Produced by *Komagataeibacter maltaceti* P285 Isolated from Contaminated Honey Wine**

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### **Legends of Supplementary Figures and Tables**

**Supplementary Figure S1.** Final media pH of HS broth cultivated by *K. maltacetii* P285, *K. pomacetii* O277 and *K. nataicola* TISTR 975 and incubated at different temperatures for 7 days (A). (B) Final media pH of culture broth after cultivation in HS broth with varying initial media pH and incubated at 30°C for 7 days.

**Supplementary Figure S2.** Final media pH of modified MSM with varying carbon sources; citric acid (A), ethanol (B), glucose (C) and sucrose (D) cultivated by *K. maltacetii* P285, *K. pomacetii* O277 and *K. nataicola* TISTR 975 and incubated at 30°C for 7 days.

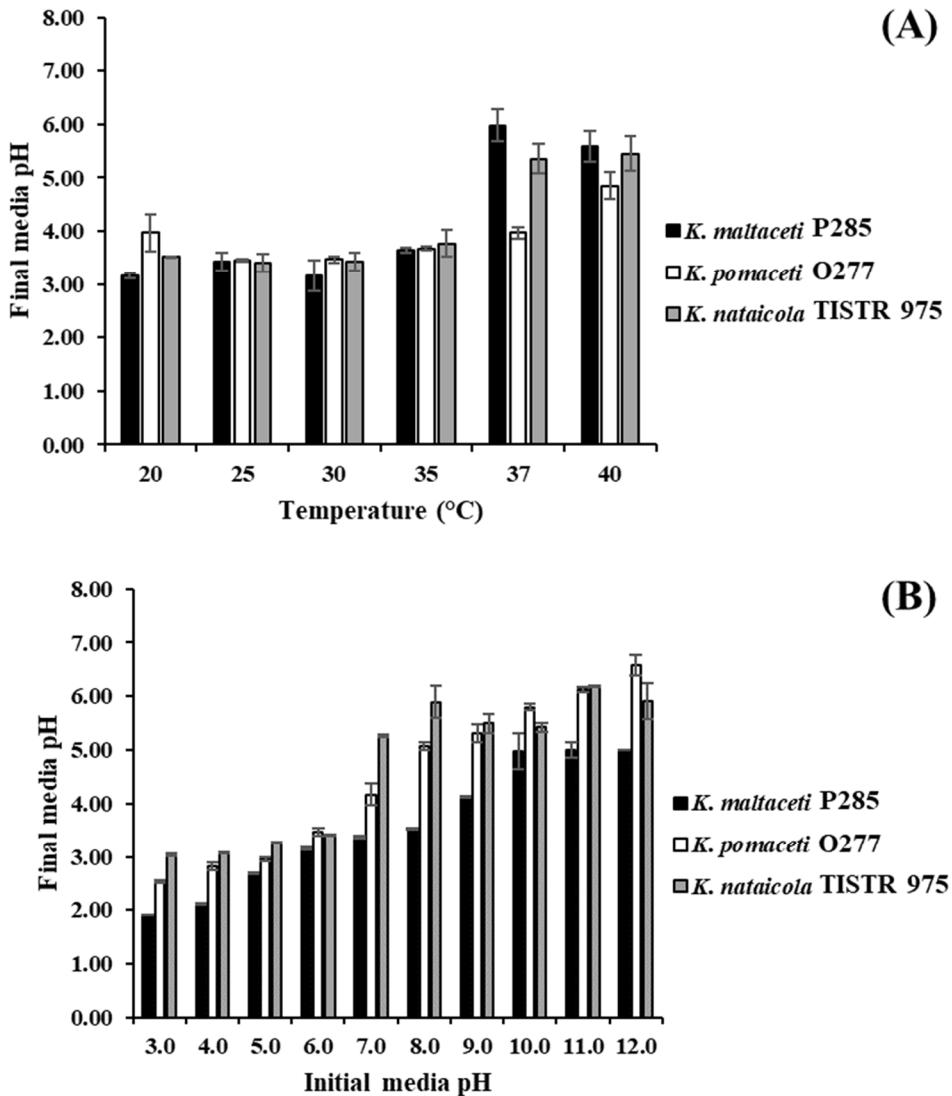
**Supplementary Figure S3.** Final media pH of modified MSM with different nitrogen sources; peptone (A), yeast extract (B),  $(\text{NH}_4)_2\text{HPO}_4$  (C) and  $\text{NaNO}_3$  (D) cultivated by *K. maltacetii* P285, *K. pomacetii* O277 and *K. nataicola* TISTR 975 and incubated at 30°C for 7 days.

**Supplementary Figure S4.** Final media pH of sugarcane (A) and honey (B) solutions when cultivated by *K. maltacetii* P285, *K. pomacetii* O277 and *K. nataicola* TISTR 975 and incubated at 30°C for 7 days.

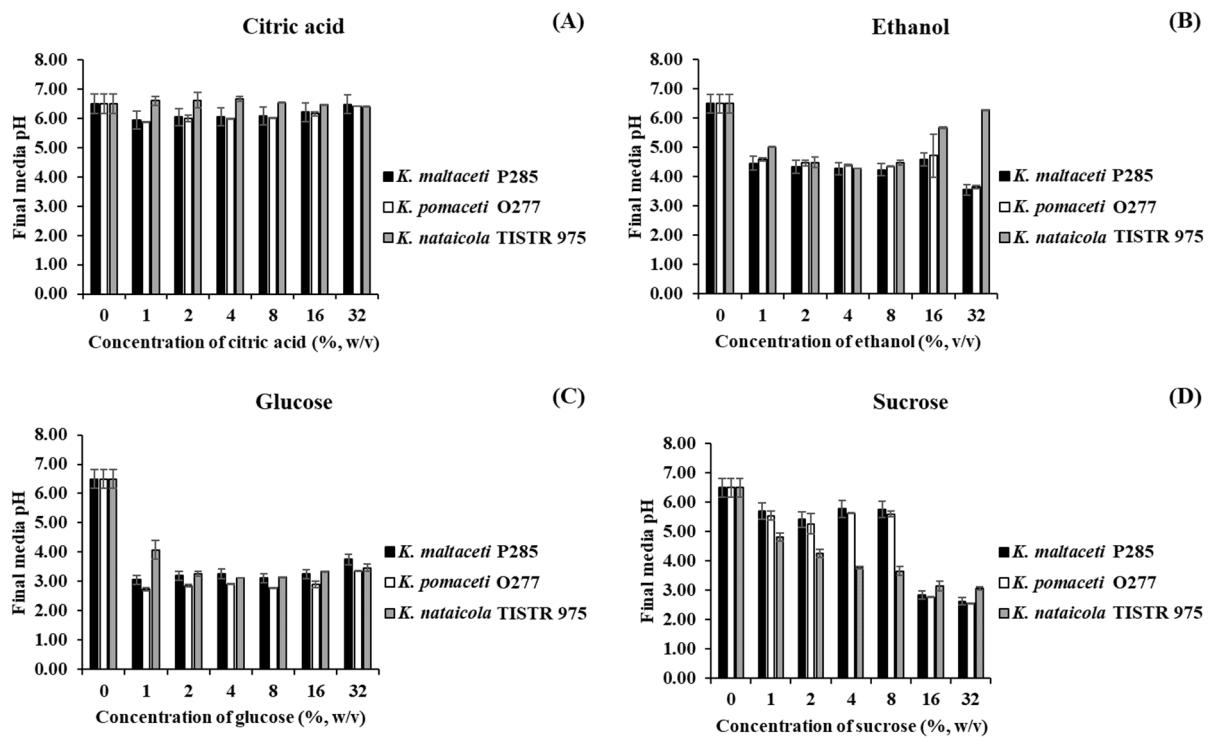
**Supplementary Figure S5.** Final media pH of modified sugarcane (S1 and S2) and honey (H1 and H2) solutions when cultivated by *K. maltacetii* P285, *K. pomacetii* O277 and *K. nataicola* TISTR 975 and incubated at 30°C for 7 days. The HS broth and the solutions of sugarcane (8%, w/v; CS) and honey (1:4 of honey : water ratio; CH) supplemented with 0.2% (w/v) yeast extract, pH 6.0 were used as controls.

**Supplementary Table S1.** Dry mass of bacterial cellulose produced by *K. maltacetii* P285 and *K. pomacetii* O277 when cultured in modified MSM with different concentrations of carbon and nitrogen sources. The *K. nataicola* TISTR 975 was used as a control. Data expressed mean  $\pm$  standard deviation of three independent experiments. The difference letters were considered statistically significant ( $p < 0.05$ ).

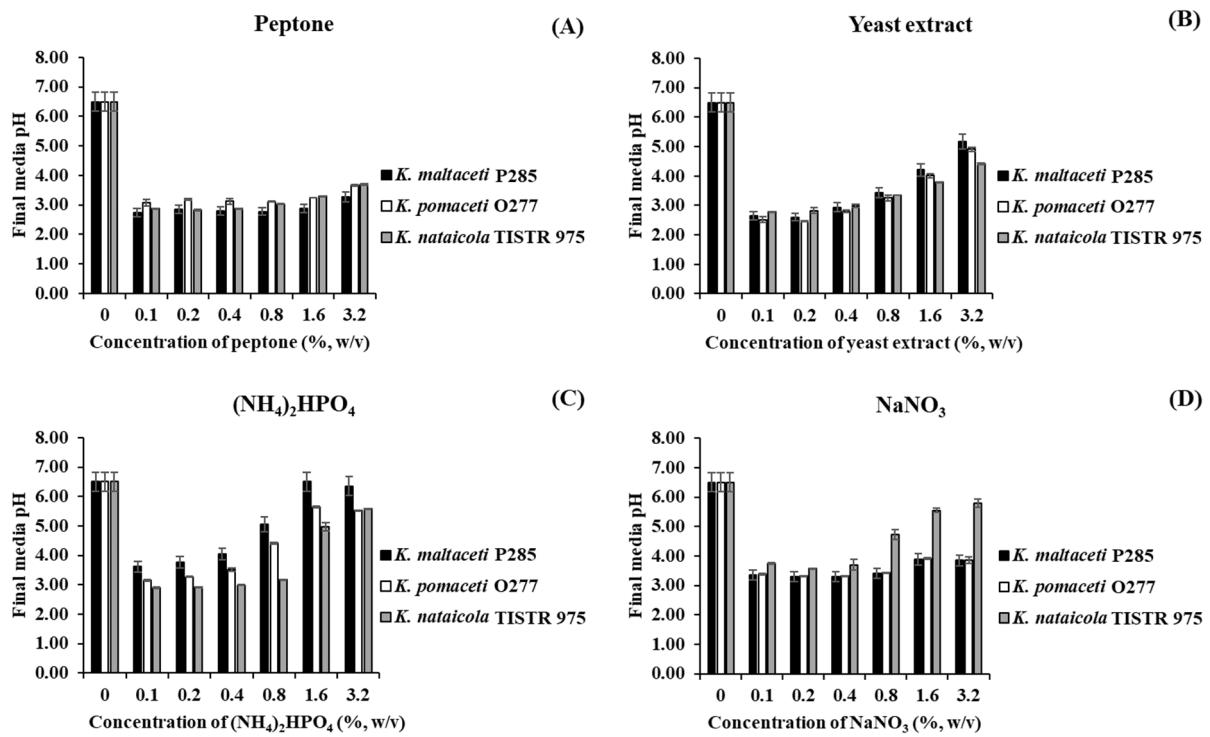
**Supplementary Figure S1**



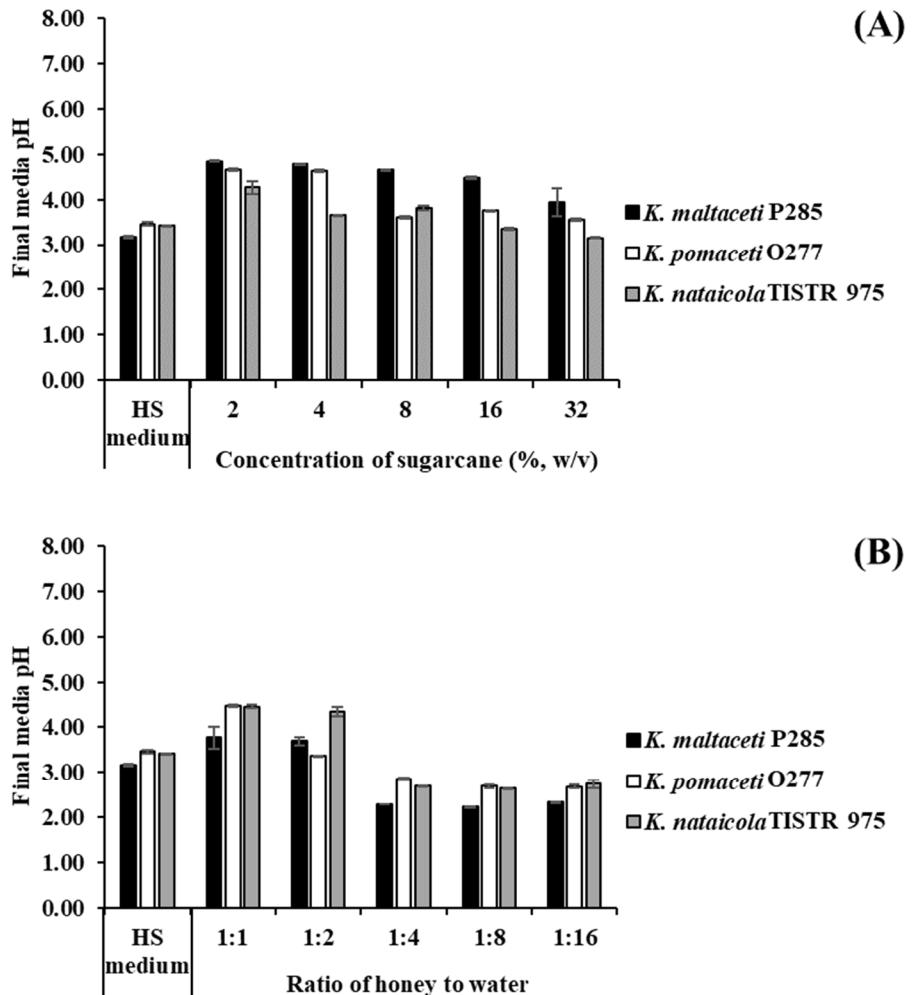
## Supplementary Figure S2



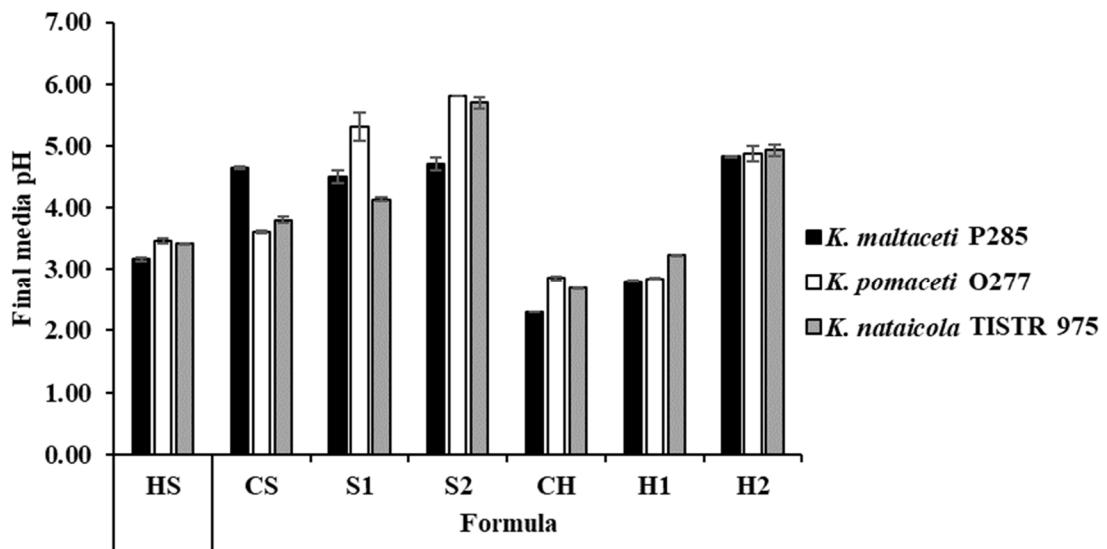
### Supplementary Figure S3



**Supplementary Figure S4**



**Supplementary Figure S5**



**Supplementary Table S1**

Carbon source concentration	Dry weight of cellulose (g/L)											
	<i>Komagataeibacter maltacei</i> P285				<i>Komagataeibacter pomacei</i> O277				<i>Komagataeibacter nataicola</i> TISTR 975			
	Citric acid	Ethanol	Glucose	Sucrose	Citric acid	Ethanol	Glucose	Sucrose	Citric acid	Ethanol	Glucose	Sucrose
0%	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>f</sup>	0 <sup>g</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>f</sup>	0 <sup>g</sup>	0 <sup>d</sup>	0 <sup>a</sup>	0 <sup>f</sup>	0 <sup>g</sup>
1%	0 <sup>a</sup>	0 <sup>a</sup>	1.2±0.1 <sup>c</sup>	0.6±0.0 <sup>f</sup>	0 <sup>a</sup>	0 <sup>a</sup>	1.2±0.2 <sup>e</sup>	1.2±0.1 <sup>f</sup>	0.2±0.1 <sup>c</sup>	0.1±0.0 <sup>a</sup>	2.9±0.0 <sup>c</sup>	0.7±0.1 <sup>f</sup>
2%	0 <sup>a</sup>	0 <sup>a</sup>	3.0±0.1 <sup>d</sup>	1.7±0.1 <sup>e</sup>	0 <sup>a</sup>	0 <sup>a</sup>	3.0±0.1 <sup>d</sup>	3.3±1.1 <sup>e</sup>	0.6±0.1 <sup>b</sup>	0 <sup>a</sup>	7.3±0.1 <sup>d</sup>	1.6±0.2 <sup>e</sup>
4%	0 <sup>a</sup>	0 <sup>a</sup>	4.2±0.2 <sup>c</sup>	3.2±0.2 <sup>d</sup>	0 <sup>a</sup>	0 <sup>a</sup>	4.4±0.9 <sup>c</sup>	6.2±1.4 <sup>d</sup>	0.9±0.1 <sup>a</sup>	0 <sup>a</sup>	15.1±0.6 <sup>c</sup>	2.1±0.1 <sup>d</sup>
8%	0 <sup>a</sup>	0 <sup>a</sup>	7.2±0.4 <sup>b</sup>	4.2±0.2 <sup>c</sup>	0 <sup>a</sup>	0 <sup>a</sup>	8.2±0.4 <sup>b</sup>	8.1±0.7 <sup>c</sup>	0.5±0.2 <sup>b</sup>	0 <sup>a</sup>	20.5±0.6 <sup>b</sup>	5.3±1.2 <sup>c</sup>
16%	0 <sup>a</sup>	0 <sup>a</sup>	11.3±0.6 <sup>a</sup>	4.9±0.2 <sup>bc</sup>	0 <sup>a</sup>	0 <sup>a</sup>	11.9±1.6 <sup>a</sup>	9.4±0.2 <sup>b</sup>	0 <sup>d</sup>	0 <sup>a</sup>	21.7±0.6 <sup>a</sup>	8.0±1.1 <sup>b</sup>
32%	0 <sup>a</sup>	0 <sup>a</sup>	1.2±0.1 <sup>c</sup>	6.7±0.3 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	1.2±0.4 <sup>e</sup>	12.8±1.3 <sup>a</sup>	0 <sup>d</sup>	0 <sup>a</sup>	16.0±0.5 <sup>c</sup>	16.7±1.2 <sup>a</sup>
Nitrogen source concentration	<i>Komagataeibacter maltacei</i> P285				<i>Komagataeibacter pomacei</i> O277				<i>Komagataeibacter nataicola</i> TISTR 975			
Peptone	Yeast extract	(NH <sub>4</sub> ) <sub>2</sub> HPO <sub>4</sub>	NaNO <sub>3</sub>	Peptone	Yeast extract	(NH <sub>4</sub> ) <sub>2</sub> HPO <sub>4</sub>	NaNO <sub>3</sub>	Peptone	Yeast extract	(NH <sub>4</sub> ) <sub>2</sub> HPO <sub>4</sub>	NaNO <sub>3</sub>	
0%	0 <sup>d</sup>	0 <sup>e</sup>	0 <sup>d</sup>	0 <sup>c</sup>	0 <sup>c</sup>	0 <sup>e</sup>	0 <sup>e</sup>	0 <sup>g</sup>	0 <sup>f</sup>	0 <sup>e</sup>	0 <sup>e</sup>	
0.1%	1.6±0.0 <sup>c</sup>	1.3±0.1 <sup>d</sup>	1.3±0.0 <sup>b</sup>	2.8±0.1 <sup>a</sup>	2.0±0.0 <sup>b</sup>	1.0±0.0 <sup>d</sup>	2.9±0.0 <sup>b</sup>	2.8±0.1 <sup>a</sup>	3.9±0.2 <sup>f</sup>	5.9±0.5 <sup>e</sup>	5.0±0.6 <sup>c</sup>	6.5±0.0 <sup>b</sup>
0.2%	1.7±0.2 <sup>c</sup>	1.2±0.0 <sup>d</sup>	1.7±0.1 <sup>a</sup>	3.0±0.1 <sup>a</sup>	2.0±0.2 <sup>b</sup>	1.0±0.0 <sup>d</sup>	3.9±0.9 <sup>a</sup>	3.0±0.0 <sup>a</sup>	5.0±0.3 <sup>e</sup>	6.3±0.3 <sup>e</sup>	3.9±0.1 <sup>d</sup>	7.3±0.5 <sup>a</sup>
0.4%	2.3±0.0 <sup>b</sup>	1.3±0.1 <sup>d</sup>	1.6±0.2 <sup>a</sup>	3.1±0.2 <sup>a</sup>	2.8±0.8 <sup>ab</sup>	1.0±0.2 <sup>d</sup>	3.6±0.0 <sup>a</sup>	3.1±0.2 <sup>a</sup>	6.1±0.2 <sup>d</sup>	7.6±0.8 <sup>d</sup>	6.5±0.5 <sup>b</sup>	7.5±0.1 <sup>a</sup>
0.8%	1.7±0.0 <sup>c</sup>	3.8±0.2 <sup>c</sup>	1.1±0.1 <sup>b</sup>	1.6±0.1 <sup>b</sup>	2.0±0.3 <sup>b</sup>	3.1±0.2 <sup>c</sup>	2.6±0.1 <sup>c</sup>	1.6±0.1 <sup>b</sup>	9.6±1.5 <sup>e</sup>	9.8±0.3 <sup>c</sup>	11.1±0.2 <sup>a</sup>	2.2±0.3 <sup>c</sup>
1.6%	1.7±0.1 <sup>c</sup>	7.1±0.4 <sup>b</sup>	0.9±0.0 <sup>c</sup>	1.5±0.1 <sup>b</sup>	2.1±0.1 <sup>b</sup>	5.7±0.0 <sup>b</sup>	2.0±0.0 <sup>d</sup>	1.6±0.0 <sup>b</sup>	15.6±0.8 <sup>b</sup>	13.0±0.6 <sup>b</sup>	10.7±0.2 <sup>a</sup>	2.1±0.1 <sup>c</sup>
3.2%	2.7±0.1 <sup>a</sup>	11.9±0.6 <sup>a</sup>	0 <sup>d</sup>	0 <sup>c</sup>	3.3±0.3 <sup>a</sup>	9.6±0.1 <sup>a</sup>	0.3±0.0 <sup>e</sup>	0 <sup>c</sup>	18.1±0.5 <sup>a</sup>	21.1±0.2 <sup>a</sup>	0 <sup>e</sup>	0.7±0.0 <sup>d</sup>