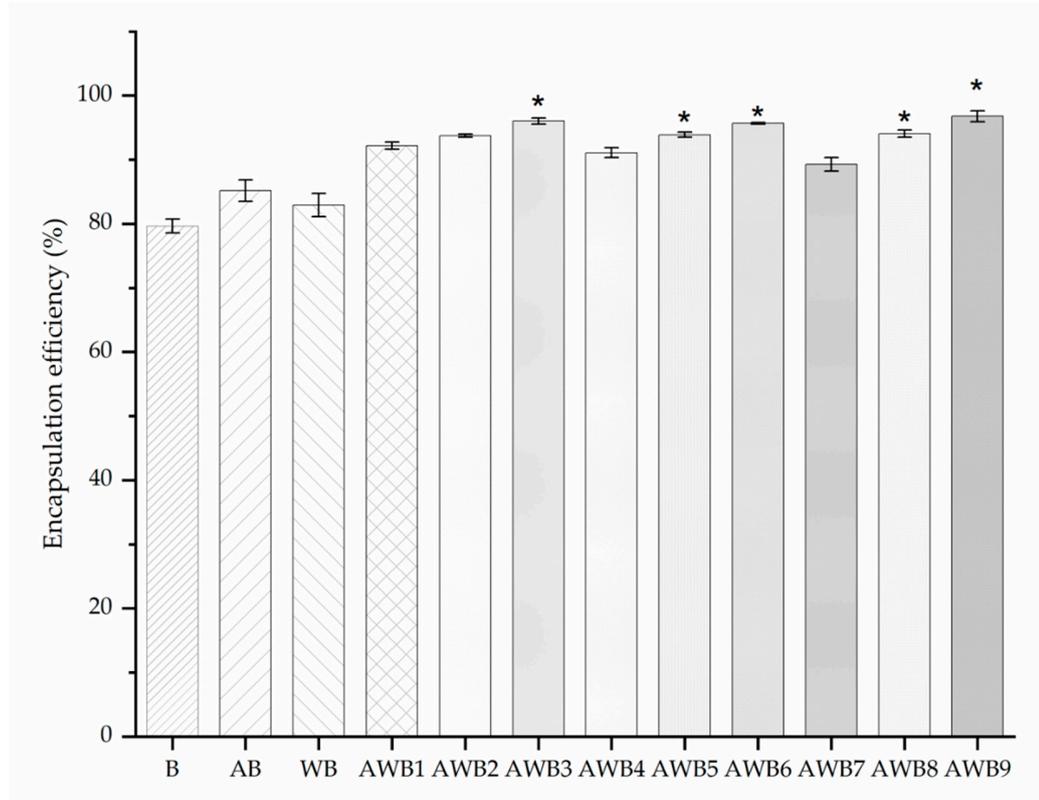
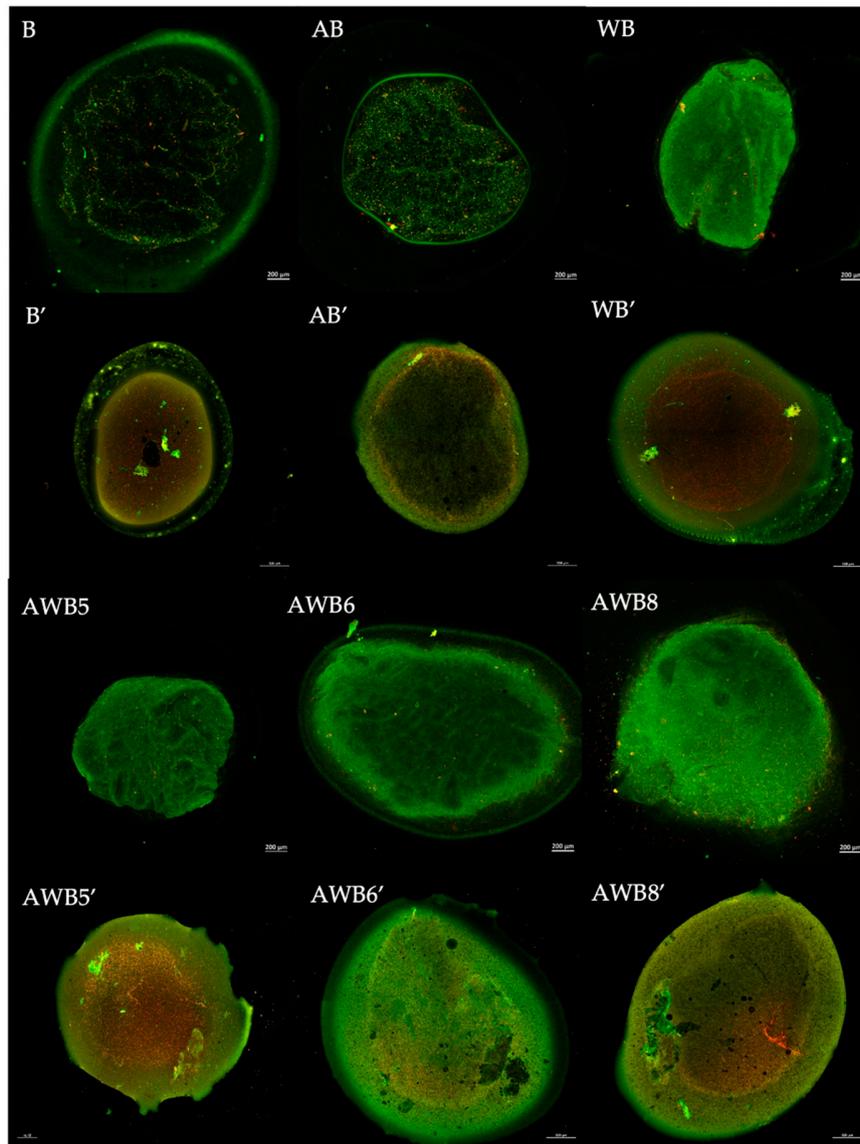


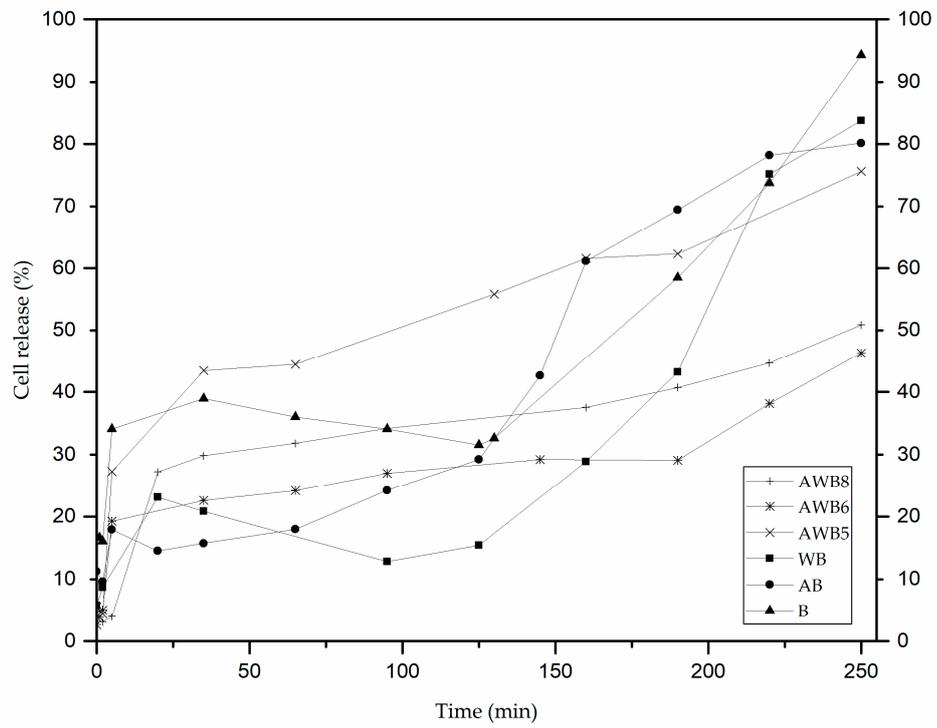
## 1. Supplementary material



**Figure S1.** Encapsulation efficiency of *S. boulardii* for beads using agavins and whey protein as wall materials at different concentrations.



**Figure S2.** Confocal micrographs of optimized beads of *S. boulardii* under *in vitro* gastrointestinal digestion conditions for viability staining. Nomenclature without apostrophe indicates micrographs taken at minute 0 (salivary phase). Nomenclature with apostrophe indicates micrographs taken at minute 250 (intestinal phase).



**Figure S3.** Percentage release of *S. boulardii* encapsulates during gastrointestinal digestion *in vitro*.

**Table S1.** Comparison of shape and dimensional parameters of *S. boulardii* encapsulates.

		Type of Bead							
		B	AB	WB	AWB3	AWB5	AWB6	AWB8	AWB9
<b>Area (mm<sup>2</sup>)</b>	Mean	4.849 <sub>d</sub>	5.540 <sub>a</sub>	5.920 <sub>c</sub>	5.405 <sub>a,b</sub>	5.893 <sub>c</sub>	5.431 <sub>a</sub>	5.805 <sub>c</sub>	5.430 <sub>a</sub>
	SE	±0.04	±0.05	±0.07	±0.05	±0.05	±0.06	±0.06	±0.02
<b>Perimeter (mm)</b>	Mean	9.439 <sub>c,j</sub>	9.396 <sub>c,h</sub>	9.516 <sub>c,i</sub>	9.168 <sub>a,c,e,f,g</sub>	9.441 <sub>c,d,e</sub>	9.082 <sub>b,e,h,i,j</sub>	9.152 <sub>a,c,e,f,g</sub>	8.931 <sub>b,g</sub>
	SE	±0.14	±0.06	±0.09	±0.07	±0.06	±0.07	±0.06	±0.03
<b>Circularity</b>	Mean	0.692 <sub>h</sub>	0.789 <sub>b,e,g</sub>	0.824 <sub>a,f,g</sub>	0.808 <sub>a,e</sub>	0.831 <sub>a,c,e</sub>	0.828 <sub>a,c,e</sub>	0.871 <sub>c</sub>	0.856 <sub>c,d,f</sub>
	SE	±0.01	±0.00	±0.01	±0.01	±0.00	±0.01	±0.00	±0.00
<b>Solidity</b>	Mean	0.983 <sub>a</sub>	0.990 <sub>b</sub>	0.989 <sub>b</sub>	0.989 <sub>b</sub>	0.990 <sub>b</sub>	0.990 <sub>b</sub>	0.993 <sub>b</sub>	0.991 <sub>b</sub>
	SE	±0.00	±0.00	±0.00	±0.00	±0.00	±0.00	±0.00	±0.00

SE: Standard error of the mean. Note: values from the same row and subtable that do not share the same subscript are significantly different in the multiple comparison test Tukey,  $p < 0.05$ .

**Table S2.** Comparison of internal structure parameters of *S. boulardii* encapsulates.

		Type of Bead					
		B	AB	WB	AWB5	AWB6	AWB8
<b>Second angular momentum</b>	Mean	0.964 <sub>b,c</sub>	0.944 <sub>a</sub>	0.963 <sub>b</sub>	0.945 <sub>a</sub>	0.950 <sub>a,b</sub>	0.956 <sub>a,b</sub>
	SE	±0.00	±0.00	±0.00	±0.00	±0.01	±0.00
<b>Contrast</b>	Mean	1008.881 <sub>b,c</sub>	1584.216 <sub>a</sub>	1051.047 <sub>b</sub>	1541.675 <sub>a</sub>	1424.428 <sub>a,b</sub>	1239.540 <sub>a,b</sub>
	SE	±53.69	±55.19	±20.13	±194.70	±132.27	±49.68
<b>Correlation</b>	Mean	0.000370 <sub>b,c</sub>	0.000232 <sub>a</sub>	0.000354 <sub>b</sub>	0.000253 <sub>a</sub>	0.000264 <sub>a</sub>	0.000301 <sub>a,b</sub>
	SE	±0.00	±0.00	±0.00	±0.00	±0.00	±0.00
<b>Entropy</b>	Mean	0.109 <sub>b,c</sub>	0.158 <sub>a</sub>	0.113 <sub>b</sub>	0.155 <sub>a</sub>	0.144 <sub>a,b</sub>	0.130 <sub>a,b</sub>
	SE	±0.00	±0.01	±0.00	±0.02	±0.01	±0.01
<b>Fractal dimension</b>	Mean	1.561 <sub>c</sub>	1.638 <sub>a</sub>	1.574 <sub>b,c</sub>	1.627 <sub>a</sub>	1.615 <sub>a,b</sub>	1.602 <sub>a,b,c</sub>
	SE	±0.01	±0.01	±0.00	±0.02	±0.01	±0.01
<b>Lacunarity</b>	Mean	0.039 <sub>a,b</sub>	0.028 <sub>b</sub>	0.031 <sub>a,b</sub>	0.043 <sub>a</sub>	0.034 <sub>a,b</sub>	0.032 <sub>a,b</sub>
	SE	±0.00	±0.00	±0.00	±0.01	±0.00	±0.00

SE: Standard error of the mean. Note: values from the same row and subtable that do not share the same subscript are significantly different in the multiple comparison test Tukey,  $p < 0.05$ .

**Table S3.** Comparison of internal structure parameters of *S. boulardii* encapsulates.

		Types of Beads					
		B	AB	WB	AWB5	AWB6	AWB8
<b>Second angular momentum</b>	Mean	0.0003 <sub>b</sub>	0.0002 <sub>b</sub>	0.0003 <sub>b</sub>	0.0008 <sub>a</sub>	0.0007 <sub>a</sub>	0.0004 <sub>b</sub>
	SE	±0.000	±0.000	±0.000	±0.001	±0.000	±0.000
<b>Contrast</b>	Mean	420.942 <sub>a</sub>	426.818 <sub>a</sub>	315.955 <sub>a</sub>	332.509 <sub>a</sub>	169.706 <sub>b</sub>	324.542 <sub>a</sub>
	SE	±29.102	±31.257	±29.355	±38.740	±11.348	±24.209
<b>Correlation</b>	Mean	0.000970 <sub>c,e</sub>	0.000862 <sub>c</sub>	0.001055 <sub>c,d</sub>	0.001713 <sub>a</sub>	0.002480 <sub>b</sub>	0.001259 <sub>a,c</sub>
	SE	±0.000	±0.000	±0.000	±0.000	±0.000	±0.000
<b>Inverse differential momentum</b>	Mean	0.097 <sub>b</sub>	0.080 <sub>b</sub>	0.096 <sub>b</sub>	0.148 <sub>a</sub>	0.143 <sub>a</sub>	0.094 <sub>b</sub>
	SE	±0.005	±0.003	±0.005	±0.011	±0.004	±0.003
<b>Entropy</b>	Mean	8.407 <sub>b,c</sub>	8.590 <sub>c</sub>	8.410 <sub>b,c</sub>	7.588 <sub>a</sub>	7.534 <sub>a</sub>	8.212 <sub>b</sub>
	SE	±0.088	±0.050	±0.074	±0.117	±0.060	±0.064
<b>SDBC</b>	Mean	2.345 <sub>a</sub>	2.411 <sub>b</sub>	2.357 <sub>a</sub>	2.359 <sub>a</sub>	2.348 <sub>a</sub>	2.398 <sub>b</sub>
	SE	±0.005	±0.006	±0.006	±0.009	±0.005	±0.005

SE: Standard error of the mean. Note: values from the same row and subtable that do not share the same subscript are significantly different in the multiple comparison test Tukey,  $p < 0.05$ .