

# Saccharomyces cerevisiae

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1 Research Article

## 2 **The effect of *Saccharomyces cerevisiae* on the growth of yeast and** 3 **fungi, $\beta$ -glucan formation, and antibacterial activities during** 4 **soybean fermentation for tempeh**

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### 12 **Abstract**

13 Generally, the microorganism involved in tempeh fermentation is *Rhizopus oligosporus*.  
14 However, *Saccharomyces cerevisiae*, a type of  $\beta$ -glucan-producing yeast, is known to be  
15 present and able to grow during tempeh fermentation. We observed the dynamics of yeast  
16 and fungi growth and  $\beta$ -glucan production during tempeh fermentation after  
17 adding *Saccharomyces cerevisiae* as an inoculum. This study used a Factorial Randomized  
18 Complete Block Design (RCBD) with two factors and three replications. The first factor  
19 was types of starter cultures as follows: *Saccharomyces cerevisiae*, *Rhizopus oligosporus*,  
20 and *Saccharomyces cerevisiae* + *Rhizopus oligosporus*. The second factor was  
21 fermentation time at room temperature ( $30\pm 2^\circ\text{C}$ ) as follows: 0, 8, 16, 24, 32, and 40 hours.  
22 The growth of yeast and fungi, the formation of  $\beta$ -glucan, and antibacterial  
23 activity to *Escherichia coli* during soybean fermentation were observed every eight hours.  
24 The results showed that yeast grew during soybean fermentation from a single culture  
25 of *Saccharomyces cerevisiae* and a mixed culture of *Rhizopus*  
26 *oligosporus* and *Saccharomyces cerevisiae*, but not from *Rhizopus*  
27 *oligosporus* culture alone. During fermentation, the growth of yeast increased until the end  
28 of the fermentation period. However, in the soybean fermentation with a mixed culture  
29 of *Rhizopus oligosporus* and *Saccharomyces cerevisiae*, yeast growth decreased at 32  
30 hours of fermentation time.  $\beta$ -glucan formed in tempeh with all types of inoculum. In all kinds of  
31 cultures, the antimicrobial activity of tempeh to *Escherichia coli* increased along with  
32 fermentation time. The highest  $\beta$ -glucan content and antibacterial activity of tempeh came  
33 from the mixed culture of *Rhizopus oligosporus* and *Saccharomyces cerevisiae*.

34 **Keywords:** soybean tempeh; yeast growth; *Saccharomyces cerevisiae*; *Rhizopus*  
35 *oligosporus*;  $\beta$ -glucan production; fermentation time.

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### 38 **Introduction**

39 Tempeh is a traditional Indonesian food made from soybeans fermented by *Rhizopus*  
40 sp. Tempeh is classified as a functional food because it contains bioactive compounds such as  
41 isoflavones which are good for the health of the body. They have nutritional advantages,  
42 unique textures, and pleasant flavors (Kustyawati et al., 2014). The quality of tempeh is  
43 influenced by the raw material and type of inoculum or starter culture used. The type of

44 inoculum plays an important role in making tempeh because it affects the quality of the  
45 tempeh produced.

46 Generally, tempeh uses an inoculum containing *Rhizopus oligosporus* (O'Toole,  
47 2016). Other important microorganisms involved in fermenting soybean to form tempeh are  
48 *R. oryzae* and *R. stolonifer*. All three microorganisms ferment soybeans into tempeh.  
49 *Rhizopus oligosporus* retains most of the nutrients contained in soybeans, increases protein  
50 digestibility, and increases the levels of some vitamins from the vitamin B family (Muchtadi,  
51 2010). *Rhizopus oligosporus* synthesizes more protease enzyme, whereas *R. oryzae* favors  
52  $\alpha$ -amylase enzyme (Triwibowo, 2011).

53 Previous researches showed that the microflora in tempeh were not just fungi.  
54 Seumahu *et al.*, (2013) and Efriwati *et al.* (2013) found lactic acid bacteria (BAL) and yeasts  
55 in tempeh. Besides *rhizopus oligosporus*, yeast and bacteria are also involved during  
56 fermentation and contribute significantly in producing functional metabolites (Kustyawati *et*  
57 *al.* 2020). One type of yeast found in tempeh fermentation was *Saccharomyces cerevisiae*  
58 (Kustyawati *et al.*, 2016), which was known as a  $\beta$ -glucan producing microorganism  
59 (Pengkumsri *et al.*, 2017).

60 The *S. cerevisiae* cell wall is composed of  $\beta$ -(1,3) and  $\beta$ -(1,6)-glucan, mannan, chitin  
61 (1-2%), and mannoproteins composing about 20-30% of the dry weight of the cell wall  
62 (Naruemon *et al.*, 2013).  $\beta$ -glucan is a polysaccharide compound that has health benefits, one  
63 of which as a biological response modifier (Corno *et al.* 2020).  $\beta$ -glucans have anti-aging  
64 effect and antibiotic activity against bacteria, fungi, viruses, and parasites (Hetland *et al.*  
65 2013). The presence of  $\beta$ -glucans in tempeh is expected to improve the functional properties  
66 of tempeh as a healthy food.

67 A research conducted by Rizal *et al.* (2020) showed that yeast could grow alongside  
68 fungi during soybean fermentation when a carbon source was added for yeast growth, thus  
69 resulting in  $\beta$ -glucans in the tempeh produced. In this study, *S. cerevisiae* was added  
70 intentionally to the soybean fermentation process without the addition of carbon sources to  
71 produce tempeh with high  $\beta$ -glucan content. It was important to examine if the addition of  
72 *Saccharomyces cerevisiae* without any carbon source in soybean fermentation could  
73 produce yeast and  $\beta$ -glucan in tempeh. In addition, the presence of  $\beta$ -glucan in tempeh  
74 produced by the addition of yeast was expected to provide added value for tempeh due to its  
75 health benefits including antibacterial activity. Therefore, this study aimed to observe the  
76 effect of adding *Saccharomyces cerevisiae* to the growth dynamics of yeast and fungi,  
77  $\beta$ -glucan formation, and antibacterial activities during soybean fermentation to produce  
78 tempeh.

## 79 **Materials and Methods**

80 The main ingredients used in this study were pure cultures of *Rhizopus oligosporus*  
81 FNCC 6010 and *Saccharomyces cerevisiae* FNCC 3012, *Eschericia coli*, soybeans (brand  
82 "Soybean USA No. 1"), Nutrient Broth (NB), Nutrient Agar (NA), Malt Extract Agar  
83 (MEA), and Potato Dextrose Agar (PDA). The study was arranged in a Factorial  
84 Randomized Block Design with three replications. The first factor was the addition of three  
85 levels of inoculum: *S. cerevisiae* (negative control), *R. oligosporus* (positive control), and *R.*  
86 *oligosporus* + *S. cerevisiae* (the main treatment). The second factor was fermentation time  
87 with six levels: 0, 8, 16, 24, 32, and 40 hours. During fermentation, samples are analyzed for  
88 microbial content (number of *S. cerevisiae* and *R. oligosporus*),  $\beta$ -glucan content and  
89 antibacterial activity to *E. coli* after each level (0, 8, 16, 24, 32, and 40 hours) of fermentation  
90 time.

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92 **Preparation of *S. cerevisiae* culture**

93 Isolate of *S. cerevisiae* was cultured into sterile Malt Extract Agar (MEA) medium using  
94 a sterilized inoculating loop needle with a scratch plate, then incubated for 24 to 48 hours at  
95 28°C to form colonies. The colonies were then harvested by adding 5 or 10 mL of distilled  
96 water into the plate disk. *S. cerevisiae* cells were harvested and poured into a 50 mL  
97 centrifuge tube. The tube was weighed and spun at 3000 rpm for 10 minutes to obtain a  
98 separate solid from the supernatant. The supernatant was then discarded and the remaining  
99 solids were diluted with 25 to 30 mL of distilled water. The *S. cerevisiae* cells were  
100 transferred into a test tube containing 9 mL of physiological saline solution and then  
101 homogenized using a vortex. The number of *S. cerevisiae* cells was calculated using a  
102 hemocytometer. The required concentration was 10<sup>7</sup> cells/mL.

103 **Preparation of *Rhizopus oligosporus* culture**

104 *Rhizopus oligosporus* from tilted agar was cultured onto a sterile medium of Potato  
105 Dextrose Agar (PDA) using a sterilized inoculating loop needle and a scratch plate. It was  
106 then incubated for five to seven days at 30 to 35°C to obtain pure colonies, harvested in the  
107 same way as the *S. cerevisiae*. The required concentration was 10<sup>5</sup> cells/mL, 100 times less  
108 than *S. cerevisiae*.

109 **Production of Soybean Tempeh**

110 Three hundred grams of soybeans were soaked at room temperature overnight. The  
111 soybean husks were removed. Soybeans were then boiled in three times their weight of water  
112 for 30 minutes, drained, and cooled until they reached 32°C (ambient temperature) and were  
113 ready to be inoculated.

114 Three separate 100 g samples of boiled soybeans received these inoculums:

- 115 (1) 1 mL suspension of 10<sup>5</sup> spores/mL of *R. oligosporus*,  
116 (2) 1 mL suspension of 10<sup>7</sup> cells/mL of *S. cerevisiae*, and  
117 (3) 1 mL suspension of 10<sup>5</sup> spores/mL of *R. oligosporus* + 1 mL suspension of 10<sup>7</sup>  
118 cells/mL of *S. cerevisiae*.

119  
120 The samples were packaged in plastics that had been perforated for ventilation then  
121 incubated at 32°C for 40 hours, observed every eight hours.

122 **Enumeration of Microorganisms**

123 Microbiological analysis followed Lay (1994), culturing on Potato Dextrose Agar  
124 (PDA) for fungi and Malt Extract Agar (MEA) for yeasts. Immediately at 0 hour, then at 8,  
125 16, 24, 32, and 40 hours, each tempeh was sampled and diluted following the method of  
126 Kustyawati et al. (2009). Ten grams of sample and 90 mL of peptone water were  
127 homogenized with a stomacher paddle blender for five minutes, then diluted into the  
128 concentration series. One mL of each dilution was planted with the appropriate surface plate  
129 calculation method on the media. Incubation continued for 24 to 48 hours at 32°C to grow  
130 fungi and 30°C to grow yeast.

131 **Analysis of β-Glucan**

132 The β-glucan analysis every eight hours during fermentation followed Kusmiati *et al.*  
133 (2007). One gram of sample and 30 mL of NaOH 0.7 N was hydrolyzed for six hours at 75°C  
134 then centrifuged at 10,000 rpm at 25°C for 30 minutes. The supernatant was removed and the

135 residue was washed with 30 ml of 0.5 M acetic acid solution, then centrifuged again at 10,000  
136 rpm and 25°C for 30 minutes. This process was repeated three times. The obtained residue  
137 was then twice-washed with 20 mL of water and centrifuged at 5,000 rpm for 10 minutes.

138 The residue with 20 mL of ethanol was centrifuged at 5,000 rpm for 10 minutes,  
139 resulting in wet  $\beta$ -glucan (crude). This biomass was dehydrated at 45°C oven for 24 hours  
140 and weighed to obtain the dry weight of  $\beta$ -glucan (crude). The dry residue with 4 mL of 1M  
141 NaOH was left for one hour. The sample was then diluted with 10 mL of sterile distilled  
142 water and shaken with an orbital shaker. After that, the sample was added with 2 mL of  
143 Pb-Acetate and left to stand for 30 minutes. Finally, one gram of sodium oxalate clears the  
144 solution, and two mL of it with 0.5 mL of phenol 5% and 2.5 mL of sulfuric acid 5N was  
145 tested using a sugar free content spectrophotometer with a wavelength of 490 A.

## 146 **Assessment of Antibacterial Activities**

### 147 ***Preparation of Target Bacteria (E. coli)***

148 Twenty  $\mu$ l of pure *Escherichia coli* was grown on Mac Conkey Agar (MCA) media,  
149 then incubated at 37°C for 24 hours. From the MCA media, *Escherichia coli* was taken with  
150 an inoculating loop needle and put into the Nutrient Broth (NB) media and incubated at 37°C  
151 for 24 hours. One mL of *E. coli* grown in Nutrient Broth for 24 hours was then diluted in 9  
152 mL of physiological NaCl 0.85% in a sterile test tube, then homogenized using vortex for 15  
153 seconds.

### 154 ***Antibacterial Testing***

155 A total of 100  $\mu$ L of target bacteria (*E. coli*) was poured evenly on the surface of the  
156 Nutrient Agar (NA) medium using the spread plate method, then let dry. A total of 2 grams of  
157 the sample from each treatment was dissolved in 8 mL of sterile distilled water. Disc paper  
158 with a diameter of 5.5 mm was inserted into each of these treatments, then allowed to stand  
159 for 10 minutes. After that, disc paper was placed on the surface of the NA medium containing  
160 the target bacteria, then incubated at 37°C for 24 hours. After 24 hours, the diameter of the  
161 inhibitory area that formed around the disc paper was measured using a slide. The sample's  
162 antibacterial activity was expressed by the diameter of the inhibition zone that formed in the  
163 clear area around the disc paper.

## 164 **Results and Discussion**

### 165 **Growth of Yeast and Fungi during Fermentation of Tempeh**

166 Results showed that yeast (*S. cerevisiae*) grew during soybean fermentation with *S.*  
167 *cerevisiae* culture and mixed culture of *R. oligosporus* and *S. cerevisiae*, but did not grow  
168 during soybean fermentation with *R. oligosporus* culture alone. Meanwhile, fungi grew  
169 during soybean fermentation with *R. oligosporus* inoculum and mixed culture of *R.*  
170 *oligosporus* and *S. cerevisiae* but did not grow during soybean fermentation with *S.*  
171 *cerevisiae* culture alone. See Figure 1.

172 Figure 1A shows the growth curve of *S. cerevisiae* in tempeh inoculated with only *S.*  
173 *cerevisiae*. The adaptation phase occurred in zero up to 8 hours of fermentation with a  
174 population of  $10^7$  CFU/g. For comparison, Sugoro *et al.* (200616) stated that the adaptation  
175 phase of *S. cerevisiae* in modified 1% tapioca solution medium containing 10.21% glucose  
176 was at the sixth hour of fermentation. Kusmiati *et al.* (2011) in media using carbon sources of  
177 glucose had the adaptation phase of *S. cerevisiae* at four hours of fermentation. On YNB  
178 medium containing 30% of glucose, Ishmayana *et al.* (2012) had it at six hours of

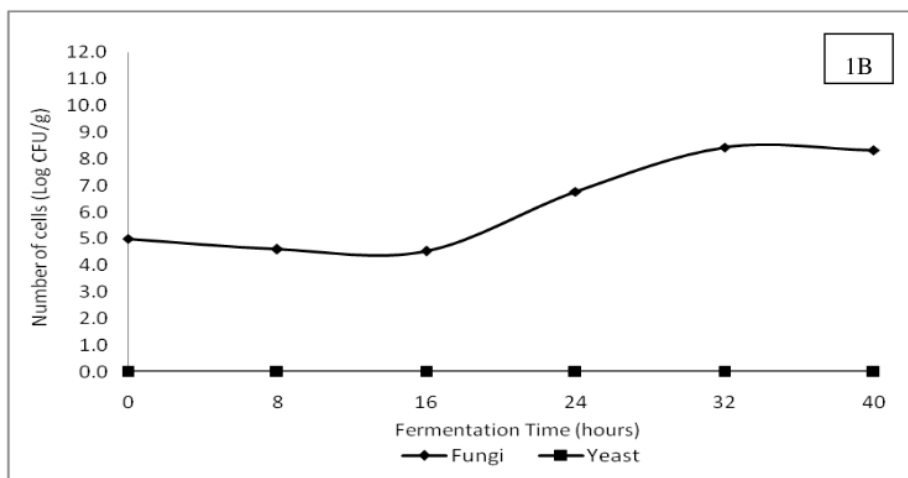
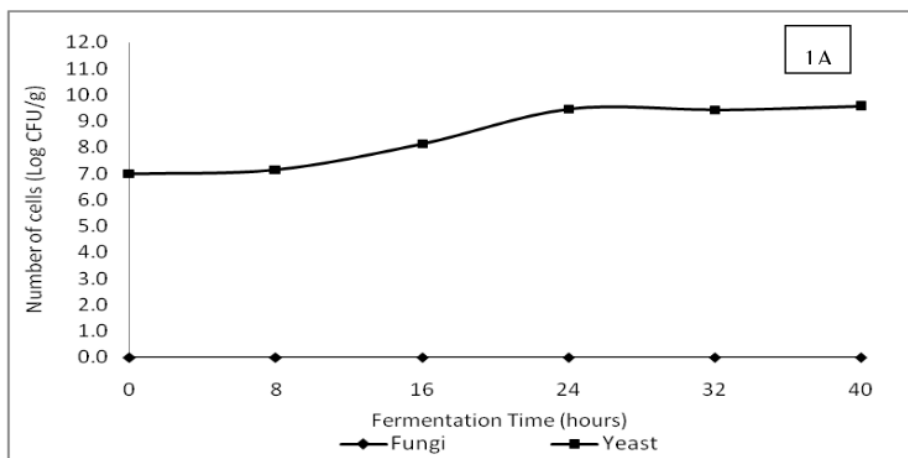
179 fermentation. Our adaptation phase of *S. cerevisiae* in this research lasted longer than those  
180 of Sugoro *et al.* (2006), Kusmiati *et al.* (2011), and Ishmayana *et al.* (2012) because in this  
181 study, there was no carbon source on the substrate, which was needed for the growth of *S.*  
182 *cerevisiae* during fermentation.

183 Figure 1A also shows that after eight hours of fermentation, yeast experiences a sharp  
184 increase in the number of cells from  $1.73 \times 10^8$  CFU/g at 16 hours of fermentation to  $3.33 \times$   
185  $10^9$  CFU/g at 24 hours of fermentation. This increase indicated that yeast (*S. cerevisiae*)  
186 entered an exponential phase after eight hours of fermentation time. Kavanagh (2005) stated  
187 that in the exponential phase, yeast reproduced by budding. Based on the exponential phase,  
188 the maximum specific growth rate ( $\mu_{max}$ ) of yeast was 0.012 cells/hour. Furthermore, yeast  
189 experienced a stationary phase from 24 hours to 40 hours of fermentation time with a  
190 population of  $4.82 \times 10^9$  CFU/g. The death phase of yeast (*S. cerevisiae*) appeared to occur  
191 after 40 hours of fermentation time.

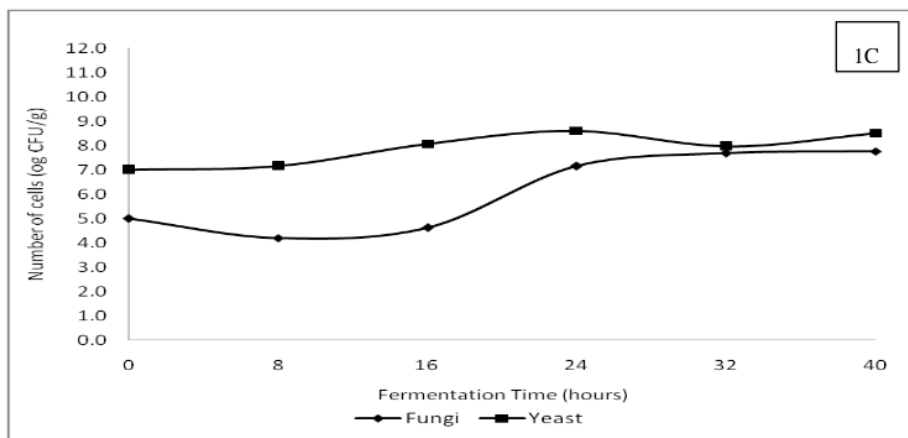
192 Yeast can grow during the fermentation process of soybeans inoculated with only *S.*  
193 *cerevisiae* even though tempeh is not formed. *Saccharomyces cerevisiae* as a sole culture  
194 (without the addition of the main tempeh fungus) in 40 hours of fermentation does not form  
195 tempeh (Figure 2A). According to Sarwono (2004), to make tempeh, a tempeh inoculum is  
196 needed. Without an inoculum of tempeh (*R. oligosporus*), soybeans will simply decay.  
197 *Saccharomyces cerevisiae* increases, but there is no presence of *R. oligosporus* unless it is  
198 inoculated. This result is in line with Wahono *et al.* (2011), who reported that during the  
199 fermentation of sorghum seeds in bioethanol production, there was an increase in the growth  
200 rate of *S. cerevisiae*. Yeast can grow by utilizing the nutrients present in the soybean  
201 substrate. According to Kustyawati (2010), almost all foods provide sufficient nutrition to  
202 support yeast growth.

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Figure 1. The growth curves of yeast and fungi during soybean fermentation inoculated with a single culture of *S. cerevisiae* (1A), single culture of *R. oligosporus* (1B), and mixed culture of *R. oligosporus* and *S. cerevisiae* (1C).

233 Figure 1B shows no growth of yeast during tempeh fermentation with *R. oligosporus* as  
234 a single inoculum. Unless *S. cerevisiae* is inoculated, there will be no yeast growth. In  
235 soybean fermentation using *R. oligosporus* as a single culture, there was no yeast growth, but  
236 tempeh was still formed due to hyphae from *R. oligosporus* (Figure 2B). This result was in  
237 line with Kustyawati (2009), which stated that yeast was not found during tempeh  
238 fermentation using *R. oligosporus*. Thus, this study revealed that the presence of yeast in  
239 tempeh could be found only if fermented soybeans were added with yeast.

240 The growth dynamics of yeast and the appearance of soybean during tempeh  
241 fermentation inoculated with the mixed culture of *R. oligosporus* and *S. cerevisiae* are  
242 presented in Figure 1C dan Figure 2C. Figure 1C shows that the adaptation phase of yeast  
243 occurs between zero and eight hours while the adaptation phase of fungi occurs between zero  
244 and 16 hours. In this sample, *S. cerevisiae* and *R. oligosporus* grew at the same time and  
245 continued growing until the end of the experiment at 40 hours of fermentation time. The  
246 appearance of tempeh inoculated with a mixed culture of *R. oligosporus* and *S. cerevisiae*  
247 during fermentation showed that between zero and 16 hours, there was no significant fungi  
248 (*R. oligosporus*) growth, as the soybeans remained intact (Figure 2C). After 16 hours of  
249 fermentation, fungi entered exponential growth phase marked by an increase in the number  
250 of *R. oligosporus* spores up to  $7.67 \times 10^6$  CFU/g at 24 hours of fermentation time and  $2.73 \times$   
251  $10^7$  CFU/g at 32 hours of fermentation time.

252 This growth pattern is in line with the growth pattern of *S. boulardi* which was  
253 inoculated together with *R. oligosporus* for tempeh fermentation in a study conducted by  
254 Kustyawati (2009). The growth pattern of yeast in this treatment was similar to that of  
255 soybean inoculated with *S. cerevisiae* alone. This indicates that *S. cerevisiae* utilizes the  
256 nutrients present in soybeans for growth and there is a mutually beneficial symbiosis between  
257 *R. oligosporus* and *S. cerevisiae* during fermentation. According to Kustyawati (2009), there  
258 may be a mutually beneficial symbiosis in terms of nutrient availability between *R.*  
259 *oligosporus* and *S. cerevisiae* during tempeh fermentation to achieve synergistic growth.  
260 *Rhizopus oligosporus* breaks down carbohydrate, fat, and protein into simple forms, and *S.*  
261 *cerevisiae* absorbs the elements C, H, O, and N from them. In turn, enzymatic activity by *S.*  
262 *cerevisiae* benefits *R. oligosporus* (Kustyawati, 2009).

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2A



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2B

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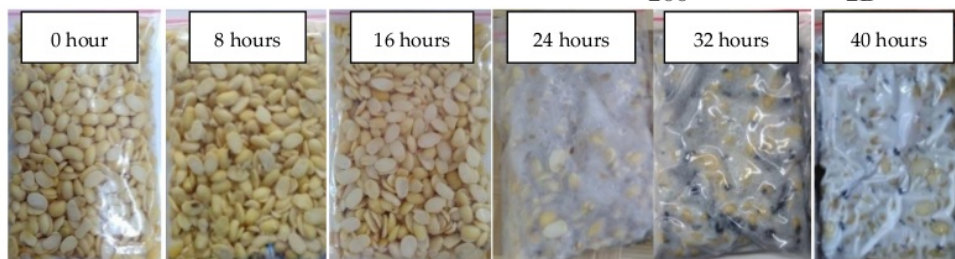
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2C

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Figure 2. The appearance of soybean inoculated with a single culture of *S. cerevisiae* (2A), a single culture of *R. oligosporus* (2B), and mixed culture of *S. cerevisiae* and *R. oligosporus* (2C) during tempeh fermentation.

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### Formation of $\beta$ -Glucan during Tempeh Fermentation

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All types of starter cultures increased  $\beta$ -glucan content of tempeh over time (Table 1). The  $\beta$ -glucan content of tempeh was higher than that – 0.05% (w/w) – of soybean without inoculum. *Saccharomyces cerevisiae* and mixed *R. oligosporus* and *S. cerevisiae* inoculum produced higher  $\beta$ -glucan content in resulting tempeh than soybeans without inoculum.

$\beta$ -glucan can be taken from the cell wall of *S. cerevisiae* through alkaline extraction, but further purification is needed (Lee *et al.*, 2001; Javmen *et al.*, 2012). Commercial tempeh inoculum contains not only *R. oligosporus* but also other microorganisms and fillers such as rice flour (Sukardi and Purwaningsih, 2008). The  $\beta$ -glucan content as a result of the addition of *S. cerevisiae* increases with the concentration of *S. cerevisiae* (Kusmiati, 2007), because the cell wall of *S. cerevisiae* contains  $\beta$ -(1,3) and  $\beta$ -(1,6) glucans (Naruemon *et al.*, 2013).

Based on careful measurement, it was found that the highest  $\beta$ -glucan content was found in tempeh inoculated with a mixed cultures of *R. oligosporus* and *S. cerevisiae* at 40 hours of fermentation time – 0.578% (w/w).

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293 Table 1. The  $\beta$ -glucan content of soybeans inoculated with various starter cultures during tempeh fermentation.  
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Fermentation time (hours)	% $\beta$ -glucan content (w/w) $\pm$ DS of tempeh on various types of starter culture		
	<i>S. cerevisiae</i>	<i>R. oligosporus</i>	<i>R. oligosporus</i> + <i>S. cerevisiae</i>
0	0.173 $\pm$ 0.001	0.207 $\pm$ 0.013	0.179 $\pm$ 0.002
8	0.189 $\pm$ 0.004	0.168 $\pm$ 0.005	0.295 $\pm$ 0.006
16	0.227 $\pm$ 0.008	0.159 $\pm$ 0.006	0.330 $\pm$ 0.008
24	0.663 $\pm$ 0.006	0.264 $\pm$ 0.008	0.593 $\pm$ 0.005
32	0.506 $\pm$ 0.004	0.413 $\pm$ 0.019	0.486 $\pm$ 0.010
40	0.515 $\pm$ 0.008	0.311 $\pm$ 0.009	0.578 $\pm$ 0.010
Soybean without inoculum	0.050 $\pm$ 0.021		

295  
296 Soybeans inoculated with *S. cerevisiae* contained more  $\beta$ -glucan than those without the  
297 addition of *S. cerevisiae* (Table 1). Our results agreed with Thontowi *et al.* (2007), who stated  
298 that the  $\beta$ -glucan content of *S. cerevisiae* in cultures with N peptone sources tended to  
299 increase along with fermentation time and was relatively fixed by the end of fermentation  
300 time (84 hours). Kusmiati *et al.* (2006) also reported an increase in  $\beta$ -glucan production with  
301 the use of different carbon sources, for example by utilizing sugar mill waste (molasses) as a  
302 fermentation medium. Increased  $\beta$ -glucan production follows the increasing number of *S.*  
303 *cerevisiae* cells. Formation of  $\beta$ -glucans continues until *S. cerevisiae* reaches a stationary  
304 growth phase. Kim *et al.* (2014) reported that the  $\beta$ -glucan content of polysaccharides in  
305 black rice bran fermented by *L. edodes* increases with time.

306 From Table 1,  $\beta$ -glucan content in this study is higher than that of Rizal *et al.* (2018) with  
307 0.578% and 0.076% respectively. Shokri *et al.* (2008) obtained  $\beta$ -glucan from *S. cerevisiae*  
308 cell walls using NaOH with 27.5% of  $\beta$ -glucan, whereas Varelas *et al.* (2016) obtained 40%  
309 of  $\beta$ -glucan. Meanwhile, our percentages ranged from 0.05% to 0.663%, significantly lower  
310 than the numbers previously mentioned. This difference was caused by the different methods  
311 used to extract  $\beta$ -glucan. In this study, the  $\beta$ -glucan content was investigated from the  
312 resulting fermented soybean flour, while the  $\beta$ -glucan content studied by Shokri *et al.* (2008)  
313 and Varelas *et al.* (2016) was directly isolated from the cell wall of *S. cerevisiae*.

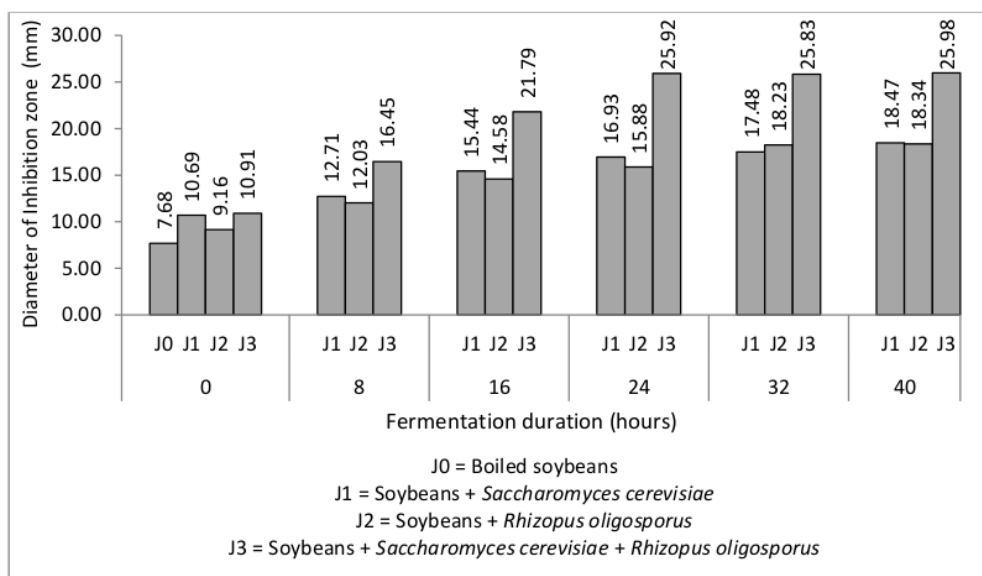
314 This study showed that the addition of *S. cerevisiae* in making of tempeh could increase  
315 yeast growth and  $\beta$ -glucan content of tempeh. The highest content of  $\beta$ -glucan was found in  
316 tempeh which was made by adding a mixed culture of *R. oligosporus* and *S. cerevisiae*  
317 inoculum at 40 hours of fermentation time with 0.578% (w/w) (Table 1).

### 318 Antimicrobial Activities of Tempeh during Fermentation

319 Antibacterial activity testing was carried out during the fermentation process of  
320 soybeans added with various types of culture (soybeans + *S. cerevisiae*, soybeans + *R.*  
321 *oligosporus*, soybeans + *S. cerevisiae* + *R. oligosporus*). In this study, the antibacterial  
322 activity of tempeh was determined by measuring the inhibitory zone's diameter in the form of

323 a clear area around the disc paper. The results showed that the antibacterial activity of tempeh  
 324 increased along with fermentation time for all treatments of starter culture types. The highest  
 325 inhibitory zone was shown in tempeh with the addition of *R. oligosporus* and *S. cerevisiae*  
 326 mixed starter culture at 40 hours of fermentation time, which was  $25.98 \pm 0.56$  mm.  
 327 Meanwhile, the diameter of the lowest inhibition area was shown in soybeans without added  
 328 inoculum with  $7.68 \pm 0.39$  mm. The diameters of the tempeh inhibition area against *E. coli* in  
 329 various types of culture at different stages of fermentation time are shown on Figure 3.

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332 Figure 3. Antibacterial activities of soybean inoculated by a culture of *Saccharomyces cerevisiae*, a culture of  
 333 *Rhizopus oligosporus*, and a mixed culture of *Saccharomyces cerevisiae* and *Rhizopus oligosporus* during  
 334 tempeh fermentation.

335 Figure 3 shows that the boiled soybeans without any starter culture addition could still  
 336 inhibit the growth of *Escherichia coli* with inhibitory area diameter of  $7.68 \pm 0.39$  mm. The  
 337 content of isoflavones in soybeans causes the antibacterial activity of soy. According to  
 338 Kustyawati (2009), antibacterial activity happens because soybeans alone contained  
 339 isoflavones in the form of genistein ( $0.25 \pm 0.60$ ) and daidzein ( $0.69 \pm 0.20$ ). Additionally,  
 340 according to Dhayakaran et al. (2015), soy isoflavones shows antibacterial activity against  
 341 several pathogens such as *Listeria monocytogenes*, *Escherichia coli*, and *Pseudomonas*  
 342 *aeruginosa*.

343 The addition of soybeans with all types of starter cultures (*S. cerevisiae*, *R.*  
 344 *oligosporus*, and a mixed culture of *S. cerevisiae* and *R. oligosporus*) caused an improvement  
 345 in antibacterial activity during fermentation that continued to increase along with  
 346 fermentation time. Both *S. cerevisiae* and *R. oligosporus* contribute to improving  
 347 antibacterial activity during tempeh fermentation. The highest antibacterial activity was  
 348 found in tempeh added with mixed cultures of *S. cerevisiae* and *R. oligosporus* after 40 hours  
 349 of fermentation. The increase in tempeh antibacterial activity during soybean fermentation  
 350 by *S. cerevisiae* and *R. oligosporus* was related to tempeh  $\beta$ -glucan content, which also  
 351 increased (Table 1). These results are consistent with research conducted by Rizal et al.  
 352 (2020). According to Rizal et al. (2020), increasing the number of *S. cerevisiae* and *R.*  
 353 *oligosporus* in tempeh can cause tempeh  $\beta$ -glucan content to increase, thus increasing the

354 antibacterial activity of tempeh. As stated by Hetland et al. (2013),  $\beta$ -glucans are compounds  
355 that are antagonistic to several microorganisms including bacteria, mold, yeast, and viruses.

356 The increased antibacterial activity of tempeh during fermentation is also caused by the  
357 increase in the number of soy isoflavones. Kustyawati et al. (2020) showed that soybean  
358 added with *S. cerevisiae* and *R. oligosporus* contained daidzein and genistein of  $224.37 \pm$   
359  $0.20$  and  $465.12 \pm 0.90$ , respectively. Increasing the amount of isoflavones can increase the  
360 inhibitory activity against bacteria because isoflavones act as antimicrobials (Mambang et  
361 al., 2014).

## 362 **Conclusions**

363 This study showed that the addition of *S. cerevisiae* in tempeh fermentation resulted in  
364 growth of yeast and fungi, production of beta-glucans, and increase in the antibacterial  
365 activity of tempeh. The highest content of  $\beta$ -glucan and diameter of inhibition (antibacterial  
366 activity to *E. coli*) was found in tempeh with the addition of a mixed culture of *R. oligosporus*  
367 and *S. cerevisiae* at 40 hours of fermentation time, with 0.578% (w / w) and 25.98 mm,  
368 respectively. Tempeh with the addition of *S. cerevisiae* inoculum has the potential of added  
369 functional properties due to the health benefits of  $\beta$ -glucan. Future research is needed to see  
370 the effect of tempeh with additional *S. cerevisiae* inoculum on body resistance in vivo.

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381

382 **References**

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