1	Optimization of enzymes addition to improve whole wheat
2	bread making quality by response surface methodology and
3	optimization technique
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18 Highlights

- 19 Optimal amounts of enzymes required to improve bread making qualities (BMQ) were
- 20 determined by response surface methodology (RSM).
- 21 Adding optimal amounts of enzymes maximized the BMQ of WWF dough and bread.
- 22 Combining RSM and optimization technique is an effective method to calculate the
- 23 optimal amounts of enzymes to add.
- 24
- 25 Abstract

1 The functional ingredients in whole wheat flour, such as dietary fiber, vitamins, and 2 minerals, have beneficial health effects. However, the excessive amount of dietary fiber 3 in whole wheat flour inhibits gluten network formation and diminishes bread making 4 qualities (BMQ). Adding appropriate amounts of enzymes, α -amylase (AM) and 5 hemicellulase (HC), could be a solution to these problems. In this study, response 6 surface methodology (RSM) created a RS model and Solver (Excel add-in software) 7 calculated the optimal amounts of the enzymes. Adding optimum concentrations of AM 8 and HC drastically improved BMQ (gas retention of dough, specific loaf volume, and 9 bread staling) of whole wheat flour dough and bread compared to whole wheat flour 10 dough and bread without the enzymes. These results show that combining RSM and 11 Solver is an effective and reasonably easy method to determine optimal concentrations 12 of enzymes to obtain the highest quality bread when using whole wheat flour.

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14 Key words: whole wheat flour, bread making quality, enzymes, response surface15 methodology

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18 Introduction

Whole wheat flour is derived by milling or grinding whole grain of wheat, which contains several functional compounds, such as dietary fiber (DF), vitamins, and minerals. These functional compounds have various positive effects on health such as reduced risk of cardiovascular diseases (Tucker et al. 2010), diabetes (Murtaugh et al. 2003), and some cancers (Schatzkin et al. 2008; Nimptsch et al. 2011). In order to enhance the functionality of bread, whole wheat flour has been used for bread making. As the characteristics of whole wheat flour are substantially different from those of white wheat flour, whole wheat flour bread exhibits increased crumb firmness, dark crumb appearance and, in some cases, alters the taste of the bread (Bruckner et al. 2001; Hung et al. 2007). In addition, an excessive amount of dietary fiber (DF), especially insoluble DF, inhibits the formation of the gluten network and decreases loaf volume (Lai et al. 1989). Thus, it is necessary to modify the bread making method or use certain additives to offset the disadvantages of making bread with whole wheat flour.

8 In this study, two kinds of enzymes, α -amylase (AM) and hemicellulase (HC), were 9 used as improvers (Caballero et al. 2007). These enzymes act on the damaged starch 10 (DS) and insoluble DF. DS is generated by the physical damage that occurs during the 11 milling process, which has a negative effect on the final bread quality. The most 12 evident effects are the reduction of loaf volume and an increase in bread staling rate. 13 On the other hand, it is possible to improve the bread making quality (BMQ) by adding 14 α -amylase, which decomposes DS. In addition, insoluble DF plays a role in disrupting 15 the formation of the gluten network, which diminishes BMQ, resulting in smaller and 16 firmer bread (Wang et al. 2002). Hemicellulase decomposes insoluble DF, which also 17 inhibits the formation of the gluten network, thus improving BMQ (Santiago et al. 18 2015a).

A combination of several enzymes that specifically counteract each negative factor is more effective in improving BMQ compared to using an individual enzyme (Caballero et al. 2007; Santiago et al. 2015a; Santiago et al. 2015b; Matsushita et al. 2017). However, determining the optimal concentration of each enzyme is difficult and a time and labor intensive task due to the complex nature of the interactions among multiple enzymes. It requires the comparison of an enormous amount of data obtained for bread making quality parameters using various enzyme combinations.

1 Therefore, in this study, we adopted a central composite face-centered design (CCF) 2 (Flander et al., 2007) as a reasonable and effective method to acquire the evaluation 3 data to determine the optimum amounts of multiple enzymes that would maximum the 4 BMQ of dough with whole wheat flour. A response surface model (RSMd) was created 5 using the data acquired, based on the CCF, and then the optimal amounts of multiple 6 enzymes were determined by using an optimization technique (OT) with Solver (Excel 7 add-in software). Finally, in order to validate the effectiveness of these methods, bread 8 making experiments, with the optimal amounts of multiple enzymes, were conducted, 9 and the effectiveness of each combination was verified from the bread making qualities 10 of the dough and various evaluations of the bread.

11

12 Materials and Methods

13 Flour and enzymes used

Camellia (Nisshin Flour Milling Co., Ltd., Tokyo, Japan) and Zenryufun Kyoriki
(Ebetsu Flour Milling Co., Ltd., Ebetsu, Japan) were used in this study. Two
commercial enzymes were used: AM (Sumizyme AS) containing 1500 α-amylase U/g,
and HC (Sumizyme SNX) containing 14,000 xylanase U/g. Both were manufactured
by Shin Nihon Chemical Co., Ltd. (Anjo, Japan).

19

20 **Optimal concentrations of added enzymes**

A central composite face-centered design (CCF), as reported by Flander et al. (2007), was used with two variables to determine optimal concentrations of enzymes. This CCF was composed of twelve experiments with four replicates at the center point (Table 1). The two variables optimized were AM (g/100 g flour) and HC (g/100 g flour). Experimental conditions (amounts of added enzymes) at the center point were

1 0.1 g/100 g flour for both AM and HC. Concentrations of both enzymes ranged from 0 2 to 0.2 g/ 100 g flour. Then random bread making tests were done using various 3 combinations of the amounts of the enzymes. In this study, specific loaf volume (SLV) 4 was adopted as the response and the amounts of added enzymes (AM and HC) were 5 the factors in analysis of RSM. The reason for choosing SLV as a response trait is that 6 it is representative of BMQ. From the results of twelve CCF experiments, a RSMd, for 7 a response and factors, was derived by multiple regression analysis. Selection of the 8 explanatory variables of the RSMd was determined by the stepwise back selection 9 method with a 2.0 F value as an index. Effectiveness of the model was assessed by 10 verifying the factor effect and lack of fit with the analysis of variance (ANOVA). 11 Optimal amounts of added enzymes were also determined with the model by using the 12 Excel add-in software Solver. After the CCF experiments, bread making tests were 13 conducted using a Control and whole wheat flour doughs with and without enzymes, 14 and the effects on BMQ were evaluated in detail.

15

16 **Dough preparation and bread making**

The Control and whole wheat flour doughs were prepared according to the formula described by Matsushita et al. 2017. The optimal amount of water was determined using a Farinograph at 500 BU according to the method used by the AACC (1991). Forty percent of the standard white wheat flour formulation used for the Control was replaced with whole wheat flour because it is the maximum percentage at which the BMQ can be improved with enzymes (Matsushita et al. 2017). The no-time method and the standard wheat bread formulation were employed (Yamauchi et al. 2001).

24

25 **Dough properties and bread evaluation**

1 The gas retention of dough (GRD) was evaluated by measuring the maximum 2 expansion volume of 20 g of dough proofed at 38° C and 85° % relative humidity (RH) in 3 a cylinder subjected to 0 to 75 cmHg (Yamauchi et al. 2000). The gassing power (GP) 4 of 20 g of dough after bench time was measured at 30°C for 1, 2, and 3 h using a 5 Fermograph II (ATTO Co., Ltd.) (Santiago et al. (2015a). The SLV of bread, cooled at 6 room temperature for 1 h after baking, was measured by the rapeseed-displacement 7 method according to the AACCI (2000). Replicates of three doughs and loaves were 8 prepared in a single bread making test to measure the GRD, GP and SLV, respectively. 9 Photographs and images of the breads were recorded using the method reported by 10 Santiago et al. (2015a). The color of the top bread crust and crumb was measured with a 11 colorimeter (CR-400, Konica Minolta Sensing, Inc., Tokyo, Japan). Moisture content of 12 the bread crumb samples, stored for 1 day in polyethylene bags, was measured using the 13 official method of the AOAC (2000). The color values and moisture content of bread 14 crumbs were measured from eight and ten slices of bread, respectively, from two loaves 15 of the same replicate.

16

17 **DS and DF analysis**

18 Sample preparation, before DS and DF analysis, was done according to the method 19 reported by Santiago et al. (2015a). The DS content in dough was measured with a 20 Megazyme assay kit (Megazyme International Ireland Ltd., Wicklow, Ireland) based on 21 the method of Gibson et al. (1991). Neutral detergent fiber (NDF), which are cellulose, 22 hemicellulose and lignin content and acid detergent fiber (ADF), which are cellulose 23 and lignin content, were measured using the official AOAC (2000). The difference 24 between NDF and ADF was calculated and used as a rough number for hemicellulose 25 content. The DS and DF of doughs, after final proofing, were measured using eight and 1 four samples, respectively, of the same replicate.

2

3 **Bread staling evaluation**

The temporal changes of crumb hardness were measured at 1, 2, and 3 days of storage (Yamauchi et al. 2001). The loaves were cut into 2 cm thick slices and a 3 x 3 cm square crumb was cut from the center. Using a creep meter (RE2-33005C; Yamaden Co., Ltd., Tokyo, Japan), the changes in temporal hardness of the bread crumbs were measured by compressing them with a special cube plunger (6 cm length x 6 cm width x 2 cm height).

10

11 Statistical analysis

12 The samples were prepared from the replicated bread making tests for all data 13 measurements except, for water absorption. Significant differences, except for water 14 absorption, were evaluated using Tukey's multiple range test at 5% significance level 15 with Excel 2012.

16

17 **Results and Discussion**

18 **Optimal concentrations of added enzymes**

19 The RSMd with SLV as the response and AM and HC as the factors, shown below was 20 derived by multiple regression analysis based on the results of the twelve, bread 21 making CCF experiments.

22 $Y=5.60X_1+4.35X_1-15.62X_1^2-16.07X_1X_2+4.85$

where Y is SLV (ml/g); X_1 is concentration of AM (g/100 g flour); X_2 is concentration of HC (g/100 g flour). R^2 and adjusted R^2 in the model showed high values, 0.841 and

25 0.751, respectively. Using ANOVA, the effectiveness and lack of fit were also assessed

and those were significant at 1% significance level and not significant at 5% 1 2 significance level, respectively. These results clarified that this RSMd sufficiently 3 estimates SLV when using two levels of added enzyme concentrations. Furthermore, the partial regression coefficients of X_1^2 and X_1X_2 explanatory variable on RSMd show 4 minus values. Therefore, especially, when both enzymes are added to the dough in a 5 6 large excess, these explanatory variables have the effect of largely lowing the BMQ 7 (SLV). Since the magnitude of the partial regression coefficient on these explanatory 8 variables is nearly same, it shows that when the enzymes are added excessively, the 9 effect of decreasing SLV of both enzymes and AM is large.

The optimal concentrations of AM and HC, calculated using Solver, an Excel add-in software, were 0.128 and 0.1 g/100 g flour, respectively. In whole wheat flour dough, SLV increased with the amount of HC added, but the dough became very sticky and extremely difficult to handle, and the improving effect plateaued when added HC exceeded 0.1 g/100 g flour. Therefore, the optimum concentration of HC is 0.1 g/100 g flour.

16

17 **BMQ evaluation**

BMQ of the Control dough, 40% of whole wheat flour (WWF) dough, and 40% of
whole wheat flour with enzyme (WWF+E) dough are presented in Table 2. Although
the WWF dough showed a lower GRD compared to the others, the GRD of the
WWF+E dough was significantly the highest among all the samples.

Initially, GP of WWF and WWF+E doughs were lower than the Control at 1 h
fermentation. At more than 2 h fermentation, the GP of these doughs were nearly same
or significantly higher compared to the Control, respectively.

25 The WWF bread had significantly lower SLV than the others. On the other hand, the

SLV of the WWF+E bread was significantly the highest among all breads. The 5.66
 value of SLV of the WWF+E bread was very close to the 5.54 value calculated using
 the RSMd. The experiments verified the effectiveness of this model.

In terms of moisture content, there was no large difference among the samples, but
WWF+E bread was significantly lower than the others. The main reason seems to be the
large reduction in weight when baking WWF+E dough, which is related to the dough's
significant expansion from the addition of enzymes.

8 Lower GRD and SLV of dough and WWF bread can be due to the higher amounts of 9 DS and DF compared to the WWF+E (Table 4). It suggests that the excessive DF in 10 whole wheat flour disrupts the gluten network formation in dough, resulting in a 11 weaker gluten network (Lai et al. 1989; Wang et al. 2002; Ozboy and Koksel 1997).

12 In terms of GRD and SLV, the WWF+E dough and bread had significantly the highest 13 values among all the samples. This might be attributed to the combined catalytic 14 activities of AM and HC that decreases DS and insoluble hemicellulose (equivalent to 15 NDF-ADF) in the dough (Table 4). The Control dough and bread had higher GRD and 16 SLV values despite having high DS content (4.16%) which might be attributed to the 17 lower values of total DF (equivalent to NDF), especially hemicellulose (equivalent to 18 NDF-ADF) compared to the WWF dough. The GP of WWF and WWF+E doughs were 19 significantly higher than the Control at 3 h fermentation. This may be related to that 20 high concentrations of various nutrients in WWF promote fermentation of yeast.

Regarding the effect of each added enzyme, AM hydrolyzes damaged and gelatinized starch in to maltose and dextrin in dough. Kim et al. (2006) reported that the high amounts of DS and DF decreased SLV of bread made with polished wheat flour, but SLV was increased by the addition of AM. Patel et al. (2012) also had a similar observation that the addition of fungal AM increased SLV in chemically leavened

1	bread. Likewise, Jiang et al. (2005) reported that HC catalyzes the degradation of
2	polysaccharides into mono-sugars and short chain saccharides, resulting in superior
3	gluten network formation. The catalytic activity of HC may have led to higher GRD
4	and SLV in WWF+E dough and bread compared to those with WWF. The addition of
5	xylanase, a kind of HC enzyme, improved SLV of whole wheat flour bread (Shah et al.
6	2006), and a millet/wheat composite bread (Schoenlechner et al. 2013).
7	From these findings, it is reasonable to expect drastic improvements of GRD and SLV
8	in WWF+E dough and bread.
9	
10	Bread color and appearance
11	Table 3 shows the results of the bread color measurement. In terms of crust color, the
12	Control bread had the highest values of L*, a* and b*among all samples. The addition
13	of WWF decreased the values of L*, a* and b*. In addition, all values of the WWF+E
14	were significantly lower than the Control.
15	In terms of crumb color, the addition of WWF significantly decreased the value of L*,
16	while it significantly increased the values of a* and b*. L* values of crumb significantly
17	decreased in descending order of the Control, WWF, and WWF+E. The a* value of the
18	Control crumb was significantly lower compared to WWF and WWF+E breads. The b*
19	values significantly increased in the order of Control, WWF+E, and WWF.
20	Figure 1 shows the bread and crumb images. The addition of whole wheat flour made
21	the external color darker; especially the color of WWF+E bread was darker compared
22	to the Control. The crumbs of WWF and WWF+E breads were darker compared to the
23	Control crumb. The loaf size of WWF bread was smaller than the Control, while the
24	WWF+E bread was obviously larger than the Control. These results were congruent
25	with the SLV data presented in Table 2.

1 The crust color of WWF bread was darker than the Control. In addition, the WWF+E 2 bread was darker compared to the Control and WWF breads (Fig. 1), which 3 corresponded with its lower L* values (Table 3). The values of redness and yellowness 4 in crust were also significantly decreased by the addition of enzymes compared to WWF bread, which is evidenced by the lower a* and b* values of crust (Table 3). 5 6 These results show that bread with WWF+E was inferior in regard to excessive 7 darkness of the crust. Goesaert et al. (2009) reported that the addition of AM increased 8 concentrations of reducing sugars, such as glucose and fructose, resulting in the 9 enhancement of the Maillard reaction.

The natural dark brown color of wheat bran makes bread crumb color darker in WWF and WWF+E breads, which results in the reduction in the L* value and the increase in a* and b* (Table 3). However, the L* value of WWF+E bread crumb was significantly lower compared to that of WWF bread crumb. These results show that WWF+E has decreased L* values of the bread crumb which makes it slightly inferior to the WWF bread crumb.

16

17 **DS and DF contents of dough**

Table 4 shows the DS content and DF composition of doughs from different treatments.
The Control had significantly higher DS content than the others. The WWF dough had
a lower value than the Control but significantly higher than the WWF+E dough. The
addition of an optimal amount of enzymes decreased the amounts of DS in dough,
therefore the WWF+E dough had significantly lower DS content than those of other
samples.

Table 4 also shows the DF content of the various doughs. The WWF and WWF+E doughs had significantly higher values than the Control dough except for the NDF-ADF of WWF+E dough. Furthermore, the values of WWF+E dough were lower
 than that of WWF dough except for ADF. In addition, NDF-ADF of WWF+E dough
 was significantly lower compared to that of WWF dough.

The higher DS contents of dough without the enzymes can be associated with the
amounts of DS generated due to the physical damages during the milling process.
Excess amounts of DS causes undesirable effects on BMQ (Santiago et al. 2015a;
Yamauchi et al. 2014). The WWF+E dough had significantly lower DS than the others,
which can be related to the enzymatic activity of AM.

9 From Table 4, WWF dough had higher DF content (NDF, ADF, and NDF-ADF), since 10 whole wheat flour contains high amounts of DF. Generally, excess DF negatively 11 effects the formation of the optimal gluten network, resulting in the reduction of GRD 12 and SLV. Conversely, WWF+E dough showed lower DF content, except for ADF, 13 which was attributable to the xylanase activity of HC, compared to WWF dough. HC 14 hydrolyzes DF, such as xylan and arabinoxylan, resulting in low NDF content and 15 crude hemicellulose (NDF-ADF) in the WWF+E dough (Stojceska and Ainsworth 16 2008; Jiang et al. 2005).

Ultimately, the improvement of GRD and SLV of bread treated with the optimal
amount of enzymes can be associated with the reduction of the amounts of DS and DF
(mainly pentosan, an insoluble hemicellulose).

20

21 Hardness of bread

Figure 2 shows staling of breads from different treatments during 3-day storage. The WWF bread showed a significantly higher value than that of WWF+E bread at 1-day storage. The Control and WWF+E breads showed similar values. The hardness of the Control and the WWF breads had similar values and were significantly higher than 1 WWF+E bread at 2-day storage. WWF bread had significantly the highest value of
2 hardness among all samples at 3-day storage, while the WWF+E bread had a
3 significantly lower value than the others.

4 There are various factors which relate to the temporal changes in crumb hardness
5 during the storage: retrogradation rate of gelatinized starch gel (GSG), the contents of
6 DS and insoluble pentosan, and SLV.

7 The AM mainly breaks down DS and GSG in dough into low molecular weight 8 dextrins, and oligo-saccharides during bread making. In addition, the endogenous 9 β -amylase in wheat flour converts the saccharides into maltose. These complementary 10 functions during the bread making process bring about partial decompositions of DS 11 and GSG. As a result, AM increases the content of low molecular weight saccharides 12 (LMWSs) in bread. It was reported that these LMWSs retard the retrogradation of 13 GSG and reduce the amount of available starch for the retrogradation in bread (Duran 14 et al. 2001; Palacios et al. 2004; Goesaert et al. 2009). Caballero et al. (2007) and 15 Palacios et al. (2004) also reported that the AM has an anti-staling effect on bread 16 during the storage. Martin and Hoseney (1991) and Palacios et al. (2004) suggested 17 that the partially decomposed starch gel has a lower retrogradation rate. Moreover, the 18 LMWSs produced by the AM hydrolysis in the dough interfere with the starch-protein 19 interactions, resulting in few and weak crosslinks between the starch and protein, and a 20 reduction of hardening rate of the bread (Martin and Hoseney 1991; Martin et al. 1991). 21 The SLV of WWF+E bread was significantly larger than the others (Table 2 and Fig. 22 2). It has also been reported that the staling rate clearly decreases when there is a large 23 SLV (Maleki et al. 1980).

The insoluble pentosan in dough interferes with the formation of a desirable gluten network, and HC attacks the insoluble pentosan, resulting in the improvement of BMQ.

It was reported that the addition of HC improved SLV and increased LMWSs in dough
 (Caballero et al. 2007; Matsushita et al. 2017; Ghoshal et al. 2013). The WWF+E
 dough had significantly lower amounts of crude hemicellulose (NDF-ADF) than WWF
 dough (Table 4).

5 From these findings, it seems the main factors concerning the suppression of staling in 6 the WWF+E bread is that the enzymes decompose DS and insoluble pentosan and 7 strengthen the gluten network , which promote high SLV, and the enzymes produce the 8 LMWSs that retard starch gel retrogradation in the bread.

9

10 Overall BMQ

11 This study established that a treatment with an optimal amount of AM and HC 12 drastically improves BMQ of whole wheat flour dough and bread. The most improved 13 properties of BMQ were GRD and SLV, which increased, and the suppression of bread 14 staling (Table 2 and Fig. 2). These WWF+E dough and bread properties were 15 significantly improved compared to those of WWF dough and bread, which were also 16 significantly better than the Control. On the other hand, a negative effect of the 17 treatment was a reduction in the bread color evaluation, especially a decrease in L* 18 value of crust, (Table 3). In the WWF bread, the decrease in L* value of the bread was 19 comparable to the Control and was considered to be an acceptable characteristic. 20 However, the addition of enzymes resulted in increased browning of the bread crust, an 21 effect of promoting the Maillard reaction during the baking process, and lowering the 22 bread color evaluation. This seems to be a negative effect of adding enzymes. In this 23 study, the optimal amounts of enzymes (AM and HC) were derived using SLV as a 24 response in an RSM and OT. The optimal value calculated for SLV using the RSMd 25 was 5.54, which corresponded to the actual experimental value of 5.66, which validates this model to some extent. There is a limit to optimizing bread making conditions using SLV as an index of optimum bread quality because degradation in crust color, a negative trait, was obtained when using enzymes in this study. Based the findings, combining RSM and OT is effective method for the optimizing bread making conditions. To more effectively use this method in the future, it will be necessary to create an overall index that integrates SLV, bread color, and staling suppression as indicators of BMQ.

8

9 Conclusion

10 Although the high amounts of DF in whole wheat flour have good functionality, it 11 decreases bread making properties. The insoluble pentosan of DF interferes with the 12 formation of the gluten network, resulting in the reduction of GRD and SLV, and the 13 acceleration of staling rate during the storage. The addition of optimal amounts of 14 enzymes (AM and HC) solved these problems. These changes can be attributed to the 15 degradation of DS and hemicellulose (mainly insoluble pentosan) into soluble low 16 molecular weight saccharides which do not negatively influence the formation of the 17 gluten network. As a result, the addition of optimal amounts of enzymes enables the 18 production of satisfactory whole wheat flour bread which has a large amount of DF 19 and several desirable BMQ, such as high GRD and SLV, and a suppressed staling rate. 20 The findings suggest that the combination of RSM and OT (Solver) are an effective 21 method for establishing optimum conditions for bread making with whole wheat flour.

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18	Figure Caption
19	Figure 1. The appearance and scanned crumb images of breads: the Control, WWF, and
20	WWF+E ^a
21	^a WWF: whole wheat flour, E, enzymes. Optimal amounts of enzymes, α -amylase and
22	hemicellulase, were added to WWF+E dough.
23	Figure 2. Temporal hardness changes of bread crumbs: the Control, WWF, and
24	WWF+E during storage ^a
25	^a WWF: whole wheat flour, E: enzymes. Optimal amounts of enzymes, α -amylase and

- 1 hemicellulase, were added to WWF+E dough. The vertical bar is the standard deviation
- 2 of each value (n=8). The symbols followed by different letters are significantly different
- 3 (p<0.05). \bigcirc : Control, \triangle : WWF, \Box : WWF+E.
- 4

	Scaled value ^b		Actual concentration			
Run"		X ₂	AM	НС		
	X_1		(g/100 g flour)	(g/100 g flour)		
1	0.0	0.0	0.1	0.1		
2	0.0	-1.0	0.1	0.0		
3	-1.0	-1.0	0.0	0.0		
4	0.0	+1.0	0.1	0.2		
5	0.0	0.0	0.1	0.1		
6	-1.0	0.0	0.0	0.1		
7	+1.0	+1.0	0.2	0.2		
8	0.0	0.0	0.1	0.1		
9	-1.0	+1.0	0.0	0.2		
10	0.0	0.0	0.1	0.1		
11	+1.0	0.0	0.2	0.1		
12	+1.0	-1.0	0.2	0.0		

Table 1. Central composite face-centered design on scaled values and actual
 concentrations of AM and HC ^a

^a Scaled values and actual concentrations of AM and HC are shown.

4 AM : α -amylase, HC : hemicellulase.

5 ${}^{b}X_{1} = (AM-0.1)/0.1$, where the actual concentration of AM ranged from 0.0 to 0.2/100 6 g flour, X₂= (HC-0.1)/0.1, where the actual concentration of HC ranged from 0.0 to 7 0.2/100 g flour

	Bread making		GRD (ml)		GP (ml)		SLV (ml/g)	Moisture content
	treatments	Water absorption (%)		1h	2h	3h		of crumb ^b (%)
	Control	68	$105.0\pm10.0~\text{b}$	$28.4\pm0.6~a$	62.5 ± 1.6 a	$92.9\pm0.8~\text{b}$	$4.95\pm0.14\ b$	42.05 ± 1.22 a
	WWF	69	$95.6\pm7.7\ b$	27.1 ± 1.8 a	62.2 ± 3.2 a	99.4 ± 3.1 a	$4.59\pm0.11\ c$	42.10 ± 0.51 a
	WWF+E	69	121.9 ± 6.1 a	26.6 ± 1.4 a	61.2 ± 3.2 a	99.6 ± 2.5 a	$5.66\pm0.24~a$	$41.56\pm0.42\ b$
2	^a GRD: gas retention of	dough, GP: gassing power	of dough, SLV: s	pecific loaf volu	me, WWF: whole	e wheat flour, E:	enzymes. Optima	l amounts of enzymes,
3	α -amylase and hemicellul	lase, were added to WWF+F	E dough. Each valu	e, except for wat	er absorption, is t	he mean \pm SD (th	e others: n=6, moi	isture content of crumb:
4	n=10). The values follows	ed by different letters within	a column are signi	ficantly different	(p<0.05).			
5	^b Moisture content of crur	nb was measured with the sa	imples stored 1 day	into polyethylen	e bags after bakin	g.		
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Table 2. Bread making qualities of doughs: the Control, WWF, and WWF+E a

1 **Table 3.** Color of the breads: the Control, WWF, and WWF+E^a

Bread making	Bread making				Bread crumb color		
treatments	L* (-)	a* (-)	b* (-)	L* (-)	a* (-)	b* (-)	
Control	49.31 ± 1.16 a	16.71 ± 0.12 a	31.22 ± 1.53 a	81.12 ± 1.21 a	-2.48 ± 0.09 b	9.32 ± 0.17 c	
WWF	48.67 ± 0.79 a	$15.26\pm0.31~b$	29.28 ± 1.11 a	75.14 ± 1.33 b	-0.41 ± 0.29 a	11.65 ± 0.51 a	
WWF+E	$41.75\pm0.60\ b$	$14.83\pm0.26~\text{c}$	$22.12\pm0.70\ b$	70.11 ± 1.56 c	-0.51 ± 0.13 a	$10.76\pm0.55~b$	

Table 3. Color of breads made from doughs of Control, WWF and WWF+ $E^{(1)}$

^a WWF: whole wheat flour, E: enzymes, L*: level of lightness, a*: level of redness, b*: level of yellowness. Optimal amounts of enzymes,

3 α -amylase and hemicellulase, were added to WWF+E dough. Each value is the mean \pm SD (n=8). The values followed by different letters

4 within a column are significantly different (p < 0.05).

	Bread making	DS (%)	NDF (%)	ADF (%)	NDF-ADF (%)
	treatments				
	Control	4.16 ± 0.48 a	$0.59\pm0.17~b$	$0.44\pm0.13~\text{b}$	$0.15 \pm 0.09 \text{ c}$
	WWF	$3.73\pm0.39~b$	$2.29\pm0.28~a$	1.19 ± 0.08 a	1.10 ± 0.21 a
	WWF+E	$2.01\pm0.07\ c$	1.85 ± 0.28 a	1.24 ± 0.08 a	$0.62\pm0.22~\text{b}$
2	^a WWF: whole w	heat flour, E: en	zymes, DS: dama	ged starch, NDF	neutral detergent
3	fiber, ADF: acid	detergent fiber	, NDF-ADF: cru	de hemicellulose	content. Optimal
4	amounts of enzy	mes, α-amylase	and hemicellulas	e, were added to	WWF+E dough.
5	Each value is the	e mean ± SD (I	DS: n=8, the othe	ers: n=4). The va	alues followed by
6	different letters w	ithin a column a	re significantly di	fferent (p<0.05).	
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Table 4. DS and DF contents of doughs: the Control, WWF, and WWF+E^a



Control

WWF

WWF+E

Figure 1

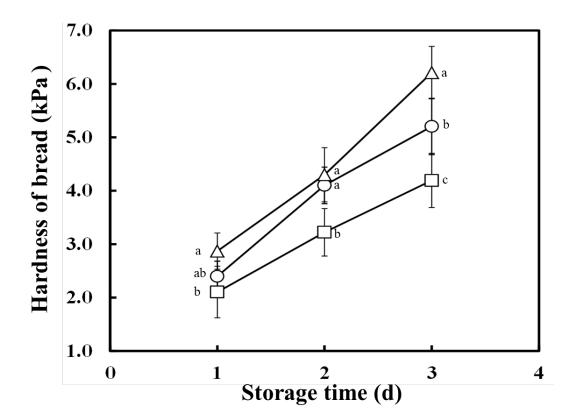


Figure 2