

Rice (*Oryza sativa* L.) breeding for novel black short grain associated with yield, cooking quality, high nutrition and antioxidant potential derived from indica black rice x japonica white rice

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26 unpolished rice, two breeding lines (69-1-1 and 72-4-3) showed higher scores than their parents.
27 However, only 69-1-1 was identified as japonica type according to its genetic background. Therefore,
28 this breeding program, involving the crossing of a temperate japonica white rice with a tropical indica
29 black rice, can create novel black short grain rice variety adapted to a tropical environment, similar to
30 japonica-type rice.

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32 **Introduction**

33 Japanese food has been widespread in Thailand for more than three decades. The Japan
34 External Trade Organization Bangkok (2020) reported that the number of Japanese restaurants
35 increased dramatically in Thailand from 1,307 to 4,094 restaurants over ten years (2010-2020),
36 demonstrating the high market demand for japonica rice in Thailand. As a result, Thailand has the
37 largest Japanese restaurant business in Asian. However, the cultivation of temperate japonica varieties
38 in tropical regions such as Thailand is very limited because these varieties are sensitive to high
39 temperatures. Thus, in countries where japonica varieties do not traditionally grow, consumers need to
40 pay a high price for this commodity (Kang 2010).

41 Recently, growing consumer interest in health-promoting food products has generated a
42 substantial market for rice with higher nutritional value (Mbanjo et al. 2020). Thus, individuals who eat
43 rice prefer unpolished rice that has a black- or red-colored pericarp. The pericarp of colored rice grains
44 accumulates proanthocyanidins (“red rice”) or anthocyanins (“black rice”). These compounds have
45 antioxidant activity and health benefits (Maeda et al. 2014). Anthocyanins are the flavonoid pigments
46 of black rice and are a source of antioxidants that have the ability to inhibit the formation or to reduce
47 the concentrations of reactive, cell-damaging free radicals (Adom and Liu 2002). In addition, black rice
48 is high in fiber, vitamins B and E, iron, thiamine, magnesium, niacin and phosphorous (Qiu et al. 1993;
49 Zhang et al. 2004). Thus, black rice is becoming popular among rice consumers and dieticians due to
50 its high nutritive and medicinal value. Black rice can be found in several varieties and has a long history

51 of cultivation in Asian countries such as China, India and Thailand (Kong et al. 2008), but most of black
52 rice varieties have been identified in indica-type rice.

53 Due to the high demand for Japanese food and healthy functional foods, many short grain
54 black rice varieties have been bred with improved agronomical traits in Japan, but their eating quality
55 and yields still need to be improved (Saka et al. 2007; Takita et al. 2001; Maeda et al. 2014). In general,
56 consumers who eat japonica rice prefer cooked rice with a short grain and a sticky and chewy texture
57 and dislike rice with a dry and crisp texture. In fact, high stickiness is one of the most important features
58 of japonica rice (Wada et al. 2008; Hossain et al. 2009). At present, the premium short grain rice variety
59 and the most widely produced variety in Japan is Koshihikari (Kobayashi et al. 2018). Therefore, many
60 researchers have tried to take advantage of Koshihikari's good eating quality while adding a black- or
61 red-colored pericarp (Maeda et al. 2014; Yamaguchi et al. 2015), containing anthocyanins or tannins
62 (both healthful nutraceuticals), respectively.

63 The cultivation of temperate japonica varieties, such as Koshihikari, is very limited in tropical
64 climates and results in low yields and low cooking quality (Kang 2010); however, japonica rice
65 varieties, Akitakomachi is more suitable than other japonica rice varieties for growth in northern parts
66 of Thailand due to their resistance to hot weather (Warinrak 2013; Seemanon et al 2015). In addition,
67 Akitakomachi has been developed from Koshihikari and is being gradually accepted by consumers,
68 continuously maintaining the fourth position in Japan. (Kobayashi et al. 2018). Therefore, it can be used
69 in short grain breeding programs in tropical climates.

70 Thailand occupies the ninth position in black rice cultivation worldwide (Ichikawa et al. 2001;
71 Sompong et al. 2011). Recently, the premium long grain black rice variety "Riceberry" has become a
72 registered rice variety in Thailand. Researchers initially developed Riceberry with the aim of boosting
73 the nutritional value, fragrance and taste of rice (Vanavichit 2021). Therefore, Riceberry is the most
74 popular black rice variety and is known for its health-promoting properties. In clinical studies, Riceberry
75 has been found to be enriched in water-soluble antioxidants, particularly anthocyanins and lipid-soluble
76 antioxidants, such as carotenoids, gamma oryzanol, and vitamin E. All nutritive properties of Riceberry

77 are contained in its rice bran, with only a small fraction accumulating in its endosperm (Leardkamolkarn
78 et al. 2011; Prangthip et al. 2013).

79 The successes of rice breeders at incorporating desirable genetic traits from indica rice into
80 japonica rice for breeding program have been limited. However, combining the desired characteristics
81 from indica and japonica rice should give an elite offspring in breeding program (Karla 2020). Thus,
82 the present research focused on the improvement of a novel black short grain rice (japonica-like) variety
83 that is suitable for growth in tropical regions to produce high yield and has increased nutritional quality,
84 with improved micronutrient and antioxidant contents. The genetic combination of temperate japonica
85 white rice, Akitakamachi and tropical indica black rice, Riceberry should be introgressed into the
86 progenies that performed by pedigree selection associated with MAS.

87 **Materials and Methods**

88 **Rice growth conditions**

89 This experiment was conducted from 2015 to 2020 at Tana Grain Polish, Ltd., Phan District,
90 Chiang Rai Province (19°35'09.4" N, 99°44'42.7" E, 413 m above sea level). Weather parameters,
91 including the air temperature, relative humidity, and the amount of rain in the field, were measured
92 every 3 h each year (2016-2020) by a data logger (WatchDog 2000 Series Micro Stations, Spectrum
93 Technologies, Inc., USA). The mean day/night temperatures over the five years were 26.8/23.1°C, and
94 the mean maximum/minimum temperatures over the five years were 31.1/20.2°C. The mean relative
95 humidity during the day/night over the five years was 72.1/87.8% RH, and the average rainfall over the
96 five years was 4.3 mm (Fig. S1). The rice plants in every generation were seeded in a field nursery.
97 After 30 d, the rice seedlings were transplanted into breeding plots. The soil in the Phan District
98 consisted of 1.56% organic matter, 0.07% total N, 26.70 mg/kg available (avail) P, 75.54 mg/kg
99 exchangeable (exch) K, 629.0 mg/kg exchangeable Ca, and 76.50 mg/kg exchangeable Mg and had a
100 pH of 5.40. Additionally, basal fertilizer was applied 15 d after planting at rates of 33.7 kg N/ha
101 (diammonium phosphate) and 41.3 kg of P₂O₅/ha. The second split of fertilizer was applied at the

102 booting stage (65 d) at a rate of 57.5 kg N/ha (Rice Department, 2020). Other management practices
103 were in accordance with conventional approaches for high-yield japonica rice cultivation.

104 **Breeding schemes**

105 An indica black long grain, Riceberry plant was used as the female parent, and a japonica
106 white short grain, Akitakomachi plant was used as the male parent. The breeding program is shown in
107 Fig. 1. The two parents were crossed to obtain F₁ seeds, and the progeny were then selfed and selected
108 from F₂ until F₇ by the pedigree method. Grain color and japonica grain shape (Juliano and Villareal
109 1993; Codex Alimentarius Commission 1990) were used as criteria for phenotypic selection. In F₆, the
110 selected lines were grown for the first yield trial with parents and control varieties in the dry season
111 from December 2019 – March 2020 (DS19/20). The experiment was conducted as an RCBD, with
112 three replications. The plot size for each treatment was 2.5 x 2.5 m (6.25 m²), with a spacing of 25 x 25
113 cm. After that, the validation of candidate lines (F₇) was conducted in the second yield trial wet season
114 from June – September 2020 (WS20). The RCBD with three replications was applied, and the plot size
115 for each treatment was 2.5 m x 3.5 m (8.75 m²), with a spacing of 25 x 25 cm. The grain yield, grain
116 quality, cooking quality, nutrition, phytochemical, antioxidation and enzymatic analysis were evaluated
117 in WS20.

118 **Phylogenetic analysis based on genotyping by sequencing (GBS)**

119 A phylogenetic analysis was performed among breeding lines (F₄), parents and control
120 varieties. The gDNA from the leaves was isolated according to the DNeasy Plant Mini Kit (Qiagen)
121 protocol. The gDNA was then sequenced on an Illumina HiSeq X by Novogene AIT, Singapore. The
122 Bowtie 2 program was subsequently used to align the nucleotides (Langmead and Salzberg 2012), and
123 the GATK program was used to analyze the single-nucleotide polymorphisms (SNPs) in each sample
124 (McKenna et al. 2010). Finally, the nucleotide sequences from the candidate lines and control varieties
125 were used to construct a phylogenetic tree using the MEGA X program.

126 **Screening SNP markers by KASP genotyping technology**

127 The screening of MAS in breeding lines was conducted in F₃ and F₄ using SNP markers,
128 including markers for starch (*wx^b*), gelatinization temperature (*SSIIa*), short grain (*GS3*) and blast
129 resistance (*Pi-ta*; TBGI453598) (Table S1). All KASP assay genotyping was performed using the LGC
130 SNP line system following standard KASP protocols (LGC Group 2016).

131 **Agronomic trait collection in yield trials**

132 Agronomic traits were collected, including days to 100% flowering, number of tillers per
133 plant, number of panicles per plant, plant height, 1000 grain weight and grain yield. The days to
134 flowering were recorded when 100% of the individual plants in each plot flowered. The number of
135 tillers per plant, number of panicles per plant and plant height were measured at maturity. The grain
136 yield in each plot was determined per harvested area of 6.25 or 8.75 m². The grain yield moisture was
137 adjusted to 14% and then extrapolated to kg per ha. After threshing, the grains were weighed to obtain
138 a 1000 grain weight.

139 **Grain and cooking quality assessments**

140 The dried grains were stored at room temperature for one month prior to grain quality trait
141 evaluation. The paddy grains were dehulled using a mini-polisher. Three physical grain qualities,
142 namely, grain length, grain width and the grain length to width ratios of both paddy grain and unpolished
143 grain, were measured using a two-decimal-point digital Vernier caliper. Three chemical grain qualities,
144 including amylose (AC) was determined based on the colorimetric procedure as described by Juliano
145 (1985). Protein (PC) was calculated by estimating nitrogen content with the Kjeldahl method and then
146 using a conversion factor of 5.95 to calculate PC in the rice sample (AOAC 2000). Fat was determined
147 using the soxhet distillation method which involved repeated fat extractions with petroleum ether
148 (AOAC 2000).

149 The cooking times of the unpolished rice samples were determined according to the method
150 described by Juliano (1985). The pasting properties of the rice flours were evaluated according to the
151 AACC method (AACC 2000). Rice flour (3 g) was suspended in water (25 g) in a canister, and the

152 viscosity changes were measured using a Rapid Visco Analyzer (RVA, Model 4-D, Newport Scientific,
153 Australia).

154 **Evaluation of the sensory qualities of cooked rice**

155 The rice cooking procedure by Xu et al. (2018) was applied. Unpolished rice samples (500 g,
156 14% moisture content) were washed three times with distilled water and soaked for 12 h. Distilled water
157 was added to the rice samples to obtain an optimum rice:water weight ratio (1.0:1.4). The samples were
158 cooked using the preset cooking setting of a rice cooker (Sharp model KS-ZT18, Thailand). Seven
159 panelists who had been well trained in the principles and concepts of descriptive sensory analysis
160 participated in the sensory quality evaluation. The sensory items included smell (score 1–5), appearance
161 (score 1–5), stickiness (score 1–5), softness (score 1–5) and taste (score 1–5). A comprehensive
162 assessment was performed based on the above factors. Using a relative scale, the panelists gave a score
163 for each attribute compared with the reference sample attributes, and the overall quality was the sum of
164 the scores for all the attributes.

165 **Pericarp thickness and pigment color analysis**

166 The pigment intensity of unpolished rice was measured with a colorimeter meter (model CR-
167 300, Minolta, Japan). L^* , a^* and b^* values were calculated to determine the color of the rice pericarp
168 of each line/variety, where ' L^* ' indicates the degree of lightness or darkness ($L^* = 0$ indicates perfect
169 black and $L^* = 100$ indicates most perfect white; hue chart); ' a^* ' indicates the degree of redness (+)
170 and greenness (-); and ' b^* ' indicates the degree of yellowness (+) and blueness (Lamberts et al. 2006).

171 The pericarp thickness was measured in each line/variety. The unpolished rice grains were
172 soaked in water for 24 h and dissected into longitudinal sections using a razor blade. The dissected grain
173 was placed on agar in a Petri dish and observed under a light stereoscope (Leica model EZ4,
174 Switzerland). The pericarp thickness was measured at 5 points on the pericarp using the program GIMP
175 2.10.12.

176 **Evaluation of nutritional and phytochemical contents**

177 Nutritional values, including iron, zinc vitamin E (α -tocopherol), vitamin B6 and folic acid
178 contents, were analyzed by the Institute of Nutrition, Mahidol University, Thailand, which followed the
179 protocol of AOAC (2000). The anthocyanin content in pericarp grains was measured according to the
180 procedures described by Rahman et al. (2015). The absorbance was measured at 535 nm using a
181 spectrophotometer (SpectraMax M2 Multilabel Microplate Reader).

182 The total flavonoid content was determined following the method of Djeridane et al. (2006).
183 The total flavonoid content was calculated from the calibration curve of rutin equivalents (RUE)
184 according to the formula $Y = 0.0083x + 0.0511$, $R^2 = 0.999$. In addition, the total phenolic content was
185 evaluated using Folin-Ciocalteu reagent (Kähkönen et al. 1999). The total phenolic content was
186 calculated from the calibration curve of gallic acid equivalents (GAE) according to the formula $Y =$
187 $0.0094x + 0.0028$, $R^2 = 0.998$.

188 **Antioxidant and enzymatic assays**

189 The inhibitory effect of rice grain water extracts on DPPH radical scavenging was determined
190 following the method of Boskou et al. (2006). In addition, the antioxidant properties of rice grain water
191 extracts on ABTS radical scavenging were investigated using the method of Hsu et al. (2011).

192 The α -amylase inhibition assay was performed following the method of Kwon et al. (2006),
193 and the α -glucosidase inhibition assay was performed following the method of Ahmad et al. (2011).
194 The percent inhibition of free radical scavenging and enzymatic activities was calculated from the
195 absorbance data using the formula: inhibition (%) = $[(Abs_{control} - Abs_{test}) / Abs_{control}] \times 100$.

196 **Statistical analysis**

197 All the data were analyzed using R program version 3.6.1 (R Core Team 2014) to test the
198 significance of the results in terms of agronomic traits and cooking quality. The means were separated
199 using Duncan's test at alpha levels of 0.01 and 0.05. If there was a significant difference among the
200 experiments for a given parameter, then the values from all of the experiments for that parameter were
201 used to obtain the mean and standard error. Chi-square (χ^2) was used to analyze the relationship of the
202 segregation ratio of grain color in the F₂ generation. In addition, correlation analysis (r) was used to

203 evaluate color intensity, nutritional and phytochemical values, and antioxidation and antidiabetic
204 activities.

205 **Results**

206 **Breeding results during early generations**

207 Among 156 F₂ plants obtained from a cross between the indica black long grain variety
208 “Riceberry” and the japonica white short grain variety “Akitakomachi”, the seed pericarps of 85, 34
209 and 37 plants were classified as showing black, brown and white coloration, respectively (Fig. 2a and
210 2S). Therefore, the color of the pericarp was identified as a qualitative trait with a segregation ratio of
211 9 black:3 brown:4 white ($\chi^2 = 2.62, P > 0.05$). The Mendelian ratio 9:3:4 fit to recessive epistasis of two
212 gene interactions. However, the black seed group consisted of two types, black and purple, depending
213 on the degree of color intensity. Conversely, the paddy grain shape of the F₂ population segregated
214 according to a normal distribution, as shown in Fig. 2b and c. The paddy grain shapes of Akitakomachi
215 (length/width 7.20/3.90 mm) and Riceberry (length/width 11.20/2.60 mm) were clearly identified as the
216 shortest and longest grain phenotypes, respectively, whereas the paddy grain shapes of their F₂
217 progenies ranged from 6.60-11.00 mm in grain length and 2.60-3.90 mm in grain width, respectively.
218 Therefore, 65 of 156 F₃ seeds that had short grains with black color were selected and grown as F₃
219 plants. After that, 26 of the F₄ seeds were selected according to the same criteria to grow F₄ plants as
220 the family lines. In this generation, 7 lines were selected, and their growth was continued to obtain the
221 F₅ generation (Fig. 1).

222 In addition, MAS for cooking quality and blast resistance (*wx^b*, *SSIIa*, *GS3* and *Pi-ta*) was
223 used to detect the target genes/QTLs at the seedling stage in F₃ and F₄. The results indicated that 17 of
224 26 plants in F₃ were successfully fixed in terms of the homozygosity of all target genes. Moreover, the
225 homozygosity of these genes was confirmed again in the 7 selected lines of the F₄ generation. When
226 considering these genes in the parents, it was found that Akitakomachi was homozygous for *wx^b*, *SSIIa*,
227 *GS3* and *Pi-ta*, while Riceberry was homozygous for *wx^a* and *Pi-ta*.

228 **Phylogenetic relationships of the F₄ selected lines**

229 In the F₄ generation, the seven selected lines were analyzed by GBS for their genetic
230 backgrounds together with their parents and control varieties. The phylogenetic tree constructed from
231 these lines was divided into two groups (Fig. 3). Group I contained one selected line (69-1-1) together
232 with Akitakomachi, Sasanishiki, Koshihikari and DOA1. This group was clearly identified as a japonica
233 type. In addition, the genetic background of Akitakomachi had closer relationships with 69-1-1. Group
234 II contained Riceberry and the control indica varieties together with six selected lines. However, the
235 coloration of six selected lines, 69-3-4, 72-4-1, 72-4-2, 72-4-3, 72-4-9 and 62-2-17, was clarified as
236 black to purple. In addition, grains 69-3-4, 72-4-1, 72-4-2, 72-4-3, and 72-4-9 were identified as having
237 short-medium grains. In contrast, only one line among the selected lines, 62-2-17, had a slender shape.
238 Finally, the growth of three lines, 69-1-1, 72-4-3, and 72-4-9, was continued through the F₅ generation
239 (data not shown) according to the criteria of genetic background, grain shape, grain color and agronomic
240 traits. These three candidate lines were used for yield trial experiments in the F₆ and F₇ generations.

241 **Evaluations of agronomic traits and grain yield**

242 The yield trial experiment was conducted in DS19/20 (F₆) and WS20 (F₇). The plant types of
243 the candidate lines with their parents are shown in Fig. 4a and b. The three candidate lines (69-1-1, 72-
244 4-3 and 72-4-9) were grown along with control varieties including Akitakomachi, Koshihikari and
245 Riceberry. The three candidate lines in F₆ (DS19/20) and F₇ (WS20) generations exhibited significant
246 grain yield and agronomic traits among three lines and control varieties ($P < 0.05$), as shown in Table 1.
247 The days to flowering periods of the candidate lines and japonica varieties were significantly shorter
248 than that of Riceberry in both seasons. In addition, the other agronomic traits varied among the
249 lines/varieties and seasons.

250 The grain yield of 69-1-1 was the highest in both DS19/20 (7.12 kg/ha) and WS20 (6.67
251 kg/ha), with a significant difference when compared with the other two lines, and the japonica control
252 varieties. However, the grain yield of 69-1-1 was not significantly different from that of Riceberry. In
253 addition, the grain yields of 72-4-3 and 72-4-9 were lower than that of Akitakomachi in both seasons
254 (Table 1). Hence, the grain yield of 69-1-1 showed the highest performance, similar to tropical indica,
255 which can be grown in tropical climates, especially in Chiang Rai Province. However, the grain weight

256 of 69-1-1 was the lowest and was significantly different from those of 72-4-3, 72-4-9, Akitakomachi
257 and Koshihikari, while it was not significantly different from Riceberry (Table 1). Moreover, it was
258 confirmed that the grain size of 69-1-1 was smaller than those of other lines/varieties (Table 2; Fig 4C
259 and D).

260 The length-to-width ratios of unpolished grains in DS19/20, 69-1-1 and 72-4-3 were not
261 significantly different from those in Akitakomachi and Koshihikari, and the ratios of these candidate
262 lines were within the standard range for short grain rice (ratio<2) (Codex Alimentarius Commission,
263 1990). However, the length-to-width ratios of 69-1-1 and 72-7-3 in WS20 were above the standard for
264 short grain rice (ratio=2.05), but the difference was not significant when compared with Akitakomachi
265 and Koshihikari. Conversely, the length-to-width ratio of 72-4-9 was over the short grain standard and
266 was significantly different from 69-1-1, 72-7-3, Akitakomachi and Koshihikari in both seasons (Table
267 2). Therefore, in terms of grain yield, grain color and the standard of short grain, 69-1-1 was identified
268 as a promising black short grain line. However, the cooking and sensory qualities, nutritional value,
269 antioxidation inhibition and antidiabetic activity of 69-1-1 were evaluated to ensure confidence before
270 its release to the public.

271 **Color intensity and pericarp thickness**

272 The color intensity on the rice pericarp differed among the candidate lines and their parents,
273 as shown in Table 3 and Fig. 4f. Lines 72-4-3 and 72-4-9 were not significantly different from Riceberry
274 and had the most intense color, while 69-1-1 was slightly brighter than the other candidate lines and
275 Riceberry. Conversely, the nonpigmented Akitakomachi showed the brightest color and was identified
276 as yellow brown. When considering the pericarp thickness, the results showed that the pericarp of 69-
277 1-1 was thinner than those of other lines/varieties, while the pericarp of Akitakomachi was the thickest
278 (Table 3).

279 **Sensory test and cooking quality**

280 The sensory test of unpolished rice samples in WS20 (Fig. 5) revealed the highest overall
281 scores for 69-1-1 (overall score=34.75) and 72-4-3 (overall score=33.00). When each sensory item was

282 considered, it was found that 69-1-1 and 72-4-3 had high scores and differed significantly from 72-4-9
283 and Riceberry in terms of smell, glossiness, stickiness, elasticity, taste and texture of cold rice,
284 respectively, while the hardness scores of 69-1-1 and 72-4-3 were lower than that of 72-4-9 and were
285 not significantly different from that of Akitakomachi. In addition, the color scores of the three candidate
286 lines were not significantly different from that of Riceberry (Fig. 5).

287 The cooking times of unpolished grains of the candidate lines and control varieties ranged
288 from 15-25 min. The three candidate lines and Riceberry, had the longest cooking times (23-25 min),
289 while Akitakomachi (white color grain) had a shorter cooking time (15 min). In addition, the amylose
290 content of 69-1-1 (19.10%) was within the standard range for japonica rice (<20). However, the amylose
291 content of 69-1-1 was significantly higher than those of Akitakomachi (17.29%) and Riceberry
292 (17.50%) (Table 4).

293 The pasting temperature (PT), peak viscosity (PV), breakdown (BD), final viscosity (FV) and
294 setback (SB) of the rice flour samples were significantly different ($P<0.05$) among the lines/varieties
295 (Table 4). The PV, FV and SB of Akitakomachi were the highest, followed by those of the three
296 candidate lines and Riceberry. Regarding PT values, the candidate lines had higher values than
297 Akitakomachi but lower values than Riceberry. BD is usually related to the tendency of gelatinized
298 starch granules to break when holding at high temperature with continuous shearing. The BDs of two
299 candidate lines (72-4-9 and 72-4-3) were not significantly different (59.79-61.12 RVU), and those two
300 lines had the highest BD values among the examined lines/varieties. Conversely, the BD of 69-1-1
301 (44.29 RVU) was the lowest among the candidate lines but higher than that of Riceberry.

302 The two principal components (PCA) (Fig. 6) explained a total of 91.12% of the variation.
303 The first principal component (PC1) accounted for 54.35% of the variation and seemed to differentiate
304 lines/varieties according to their protein content and FV and SB values, and these values were
305 negatively correlated with the CT values. The second principal component (PC2) accounted for 36.77%
306 of the variation and primarily explained why the fat content and PT properties of rice were negatively
307 correlated with the amylose content and BD value. In this study, PCA showed the clustering pattern of
308 the three groups. The two candidate lines 72-4-9 and 72-4-3 seemed to have similar properties in terms

309 of high amylose content and BD, while both 69-1-1 and Riceberry had high fat and PT values.
310 Akitakomachi showed a high protein content and high FV, PV, and SB property values.

311 **Nutritional values and antioxidant and antidiabetic activities**

312 The protein contents of the candidate lines and control varieties were significantly different
313 ($P<0.05$). The rice flour from Akitakomachi and 69-1-1 had the highest protein content (6.87 and
314 6.58%) compared to the other lines/varieties, though the candidates and varieties did not differ
315 significantly in terms of the protein content (5-8%). In addition, the protein contents of all lines/varieties
316 in this study were within ranges previously reported for rice.

317 The results of the nutrition analysis are shown in Table 5. The unpolished grain of 72-4-9 had
318 the highest total anthocyanin content (78 mg/100 g), which was significantly different from that of
319 Riceberry (68 mg/100 g), while the anthocyanin contents of 72-4-3 and 69-1-1 were 62 and 59 mg/100
320 g, respectively. In addition, 69-1-1 showed the highest nutrient concentrations in two categories (Fe,
321 and Vit B6), while 72-4-9 had the highest nutrient concentrations for Zn category. Conversely,
322 Riceberry had the highest folic acid content (70 $\mu\text{g}/100$ g), while 69-1-1 had less folic acid (48 $\mu\text{g}/100$
323 g) than the other two candidate lines and Riceberry. However, when considering all parameters
324 evaluated in the WS20 yield trial, 69-1-1 showed a higher in grain yield, low amylose (<20%), the
325 highest score of sensory tests and higher in Fe and Vit B6 than 72-4-3 and 72-4-9.

326 Candidate line 72-4-9 showed the highest contents of flavonoids and phenolics (71.65 \pm 1.79
327 mg RUE/100 g extract and 176.45 \pm 16.30 mg GAE/100 g extract, respectively), followed by 69-1-1
328 (61.03 \pm 1.14 mg RUE/100 g extract and 141.88 \pm 6.62 mg GAE/100 g extract, respectively). Moreover,
329 72-4-3 also showed higher concentrations of flavonoids and phenolics than Riceberry and
330 Akitakomachi (Table 6).

331 The results of antioxidant activity assays showed that the candidate line 72-4-9 presented the
332 strongest inhibition of DPPH (11.99%) among the lines/varieties, while 69-1-1 (8.13%) and 72-4-3
333 (8.24%) had an inhibitory effect on DPPH similar to that of Riceberry (7.04%). Conversely,
334 Akitakomachi did not show the antioxidant property of DPPH radical scavenging. In terms of ABTS

335 radical scavenging inhibition, all candidate lines, 69-1-1, 72-4-3 and 72-4-9, showed strong inhibitory
336 activities of more than 80% and a similar inhibitory effect on ABTS to Riceberry. In contrast, the water
337 extract of Akitakomachi showed weaker ABTS inhibition than the other extracts (Table 6).

338 The α -amylase activity assay showed that the water extracts of Riceberry and 72-4-9 (13.08%
339 and 11.10%) had higher capacity to inhibit α -amylase than the two other candidate lines and
340 Akitakomachi. In the case of α -glucosidase inhibition, it was found that all candidate lines (51.52%-
341 60.93%) showed stronger inhibitory activity than Riceberry and Akitakomachi (Table 6).

342 When considering the correlation analysis (Fig. 7), it was found that the intensity of grain
343 color (L^*) was highly negatively correlated with anthocyanin, zinc, flavonoid, and phenolic levels and
344 antioxidation activities (DPPH, ABTS). Conversely, the anthocyanin content was highly positively
345 correlated with zinc, folic acid, flavonoid, and phenolic levels and α -amylase and antioxidation
346 activities.

347 **Discussion**

348 **The segregation of color in pericarp and grain size**

349 In this research, color segregation analysis in F_2 progenies from Riceberry (black color) x
350 Akitakomachi (white color) showed the same results as many researchers reported that the character of
351 black pigment is controlled by two complementary dominant genes, *Pb* and *Pp*, with recessive epistasis
352 (9:3:4) (Lee et al. 2018; Kristantini et al. 2019). However, in the black and brown color groups, there
353 was also variation in color intensity. This may suggest that the color of the rice pericarp is controlled
354 by other genes in the pathway of anthocyanin synthesis (Yamuangmorn and Prom-u-Thai 2021).

355 The segregation of grain size was controlled by quantitative genetics that combines grain
356 length, grain width, and grain thickness characteristics (Ponce et al. 2020). Li et al. (2018) reported that
357 the size of rice grains is coordinately controlled by cell proliferation and cell expansion in the spikelet
358 hull and identified several quantitative trait loci and a number of genes as key grain size regulators.
359 However, the *GRAIN SIZE3* (*GS3*) gene was the first molecularly characterized QTL for grain size and
360 is used as the major QTL to identify grain length differences between indica and japonica types (Fan et

361 al. 2006, Saichompoo et al. 2021). Thus, *GS3* could be used to identify grain size in this breeding
362 program.

363 **Agronomic and environmental factors**

364 Japonica rice is suitable for cultivation between 53° N and 36° S latitudes at lower
365 temperatures and under longer days than in tropical areas (Khush 1997). Moreover, japonica rice can
366 produce grain yields in temperate regions higher than that in tropical regions due to the photoperiod
367 sensitivity and spikelet sterility caused by high temperatures (Kobayashi et al. 2018; Lee et al. 2018;
368 Yoshida 1983). In this research, the breeding lines were bred at 19° N and 99° E latitude under
369 26.8/23.1°C mean day/night temperatures over five years. The 69-1-1 showed a high grain yield similar
370 to that of Riceberry, although the flowering day was the same as those of temperate japonica control
371 varieties. Therefore, rice plants genetically belonging to temperate japonica varieties can be bred to
372 adapt to tropical region by introgressing indica genetically into breeding lines (Negrao et al. 2008).

373 **Grain and cooking quality**

374 Amylose is one of the components of rice starch that greatly affects cooking and eating
375 qualities. Low-amylose rice generally becomes soft and sticky after cooking (Okadome et al. 1999).
376 Amylose content between 15–20% is identified as low amylose rice and contain nearly all temperate
377 japonica varieties (Juliano 1971). In this study, an amylose content of less than 20% among the
378 candidate lines was found only in 69-1-1 (19.10%) and the unpolished grain size ratio of this line is
379 within the standard for japonica grain shape (ratio<2) (Codex Alimentarius Commission 1990).
380 However, the grain weights of the three breeding lines were lower than those of japonica white rice
381 because anthocyanin deposition in the pericarp of black rice reduced the photosynthetic rate (Rahman
382 et al. 2015).

383 The sensory quality of cooked rice is an important factor in determining its market price, as
384 well as consumer acceptance and breeding efforts to improve rice grain quality (Xu et al. 2018). In this
385 study, the 69-1-1 and 72-4-3 had higher scores for overall sensory quality which were lower protein
386 content (6.23-6.58%) than the mean of unpolished rice grain (8%) (Mahender et al. 2016). In addition,

387 both lines showed low hardness and high stickiness scores in the sensory analysis. The results were
388 consistent with the findings of Xu et al. (2018), who suggested that the overall sensory quality is
389 negatively correlated with protein content and positively correlated with hardness and stickiness. In
390 addition, the stickiness after cooking is due to the low amylose content (Juliano 1985) and the wx^b gene
391 that is critical gene to control amylose content (Shao et al. 2020) was presented in the candidate lines.
392 Thus, the cooking and eating quality characteristics of breeding lines were derived from both indica and
393 japonica parents which contained wx^b and *SSIa* genes.

394 The pasting properties of rice flour are related to the cooking and eating quality of rice. The
395 differences in pasting properties among rice varieties in this research could be attributed to differences
396 in the amounts of amylose, lipids, and branch chain-length distribution of amylopectin present in rice
397 starch (Singh et al. 2006). Due to the lipid content, the paste viscosity of 69-1-1 was found to be lower
398 than that of the other lines. Starches with a high lipid content can form a rigid network of structures in
399 granules due to amylose-lipid complexation, causing a restriction of granule swelling and resulting in a
400 higher pasting temperature (Becker et al. 2001). Moreover, the formation of amylose-lipid complexes
401 may have prevented amylose from leaching out, resulting in a reduction in the hot paste viscosity
402 (Richardson et al. 2004).

403 The pasting and chemical properties of Akitakomachi were quite different from those of the
404 breeding lines. This observation might be explained by a characteristic difference between japonica-
405 and indica-type rice in terms of the different branch-chain lengths of their amylopectin molecules. The
406 amylopectin of japonica rice has a larger proportion of short-branch chains than that of indica rice (Kang
407 et al. 2006), and short-branch chains in amylopectin behave in a manner similar to amylose by
408 restricting starch swelling, resulting in high PV, FV and SB values (Jane et al. 1999). Overall, the
409 differences in pasting properties among different genotypes are attributed to protein, lipid and amylose
410 contents, granule rigidity, and starch crystallinity. Therefore, pasting profiles associated with chemical
411 content will provide a baseline for the selection of cooking and eating quality grain for further
412 development. In addition, the pasting properties of 69-1-1 was closer with indica parent than those of
413 japonica parent.

414 **Nutritional value and antioxidant and antidiabetic activities**

415 The anthocyanins content in black rice pericarp also varies depending on the rice cultivars
416 (Jiamyangyuen et al. 2017; Ji et al. 2012). Kushwaha et al. (2020) reported that temperate climate could
417 favor the rise in grain's anthocyanin content than the warm climate. In addition, the increase in the color
418 intensity was increase the total anthocyanin content (Mackon et al. 2021). Moreover, the color intensity
419 and pericarp thickness had affected to anthocyanin content in rice pericarp. Therefore, the breeding
420 procedure for anthocyanin content from indica black rice x japonica white rice by phenotypic selection
421 using color intensity was successful in obtaining new breeding lines that had similar total anthocyanin
422 contents to the black-grained parent.

423 Most of the nutrients found in rice grain accumulate in the outer aleurone layer and the embryo
424 (Mbanjo et al. 2020). The standardization of unpolished rice grains has resulted in ranges of 1.4-5.2
425 mg/100 g of iron, 1.9-2.8 mg/100 g of zinc, 0.80-2.50 mg/100 g of vitamin E, 0.50-0.70 mg/100 g of
426 vitamin B6 and 16-20 µg/100 g of folate (Juliano 2016). Therefore, the three candidate lines in this
427 study were within the standard levels. In addition, the correlation analysis in this study confirmed the
428 results of Gao (2011) who reported that the color intensity was positively correlated with iron, zinc and
429 folic acid contents. Other studies have suggested that pigmented rice contains higher levels of iron and
430 zinc than white grain (Hurtada et al. 2018; Shao et al. 2018).

431 Total flavonoids and phenolics are natural bioactive compounds in plants that can indicate
432 their potential for use as therapeutic agents and in controlling the quality of medicinal sources. They
433 can also function as free radical-scavenging and reducing agents. (Ghasemzadeh and Ghasemzadeh
434 2011; Tungmannithum et al. 2018). In this research, the contents of phenolics, flavonoids, and
435 anthocyanins were negatively correlated with L* color intensity values, in accord with previous findings
436 (Shao et al. 2018). However, the total antioxidant activity of black rice bran was correlated with the
437 contents of total anthocyanins, total phenolics and total flavonoids (Zhang et al. 2010; Goufo and
438 Trindade 2014), which were determined in the same way as in this research.

439 Extracts of black rice grain have been shown to effectively inhibit the activities of endogenous
440 α -amylase and α -glucosidase, thereby inhibiting the conversion of starch to glucose in the small
441 intestine, which acts as a source of resistant starch utilized by the gut microbiota in the colon (Boue et
442 al. 2016; Chiou et al. 2018). All black and red bran extracts inhibit α -glucosidase activity; however,
443 only red rice bran extracts inhibit α -amylase activity (Boue et al. 2016). In this study, α -glucosidase had
444 the greatest correlation with anthocyanin content (Yao et al. 2010; Bae et al. 2017). However, α -amylase
445 activity was not correlated with anthocyanin content that was different result from those of Wongsu et
446 al. (2019).

447 **Conclusion**

448 A breeding program for obtaining black color, short grain with high nutrition and
449 phytochemical can provide a novel rice variety for growth in Thailand that combines tropical
450 adaptability with high yield and high nutritional value from tropical indica rice and short grain with
451 good cooking and eating quality from temperate japonica rice. Finally, the promising 69-1-1 line has
452 undergone short grain rice breeding with the same yield as indica rice and resistance to blast disease
453 together with high anthocyanin content and good cooking and eating quality according to Japanese food
454 standards. In addition, it has already been registered as a new short grain rice variety in Thailand named
455 – Fuji-Murasaki (Murasaki means purple color).

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639 Table 1 Agronomic traits of the three candidate short grain lines compared with commercial varieties and their parents in DS19/20 (F₆) and WS20 (F₇).

Lines/Varieties	DF (days)		PH (cm)		NTP		NPP		TGW (g)		GY (t/ha)	
	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20
69-1-1	89b ^{1/}	63c	98b	110a	22ab	36a	22ab	33a	22.13d	22.60c	7.12a	6.67a
72-4-3	90b	72b	121a	117a	21ab	27bc	21ab	25bc	25.73c	28.13b	5.63bc	5.95bc
72-4-9	90b	74b	107ab	108a	25a	30ab	25a	29ab	27.37b	28.50b	4.85c	4.65c
Akitakomachi	91b	65c	100b	92b	15b	17d	15b	17d	25.57a	30.60a	5.93b	6.50b
Koshihikari	91b	64c	99b	83c	21ab	21cd	21ab	20cd	29.83a	30.67a	4.38d	4.40cd
Riceberry	113a	107a	94c	110a	27a	27bc	27a	27abc	20.73d	21.77c	7.38a	6.96a
F-test			**	**	**	**	*	**	**	**	**	**
(P<0.01)	**	**										
CV%	3.17	4.59	3.17	4.59	2.23	4.59	2.23	4.59	4.59	4.59	5.24	4.85

640 ^{1/} Different letters in the same column indicate significant differences at the 0.05 level using LSD.

641 DF=Days to flower, PH=Plant height, NTP=Number of tillers per plant, NPP=Number of panicles per plant, TGW=1000 grain weight, GY=Grain yield

642

643 Table 2 Grain length, grain width and grain length/width ratio of paddy and unpolished grain of the three candidate short grain lines compared with
 644 commercial varieties and their parents in DS19/20 (F₆) and WS20 (F₇).

Lines/Varieties	Paddy grain						Unpolished grain					
	Length (mm)		Width (mm)		Ratio		Length (mm)		Width (mm)		Ratio	
					(length/width)						(length/width)	
	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20	DS19/20	WS20
69-1-1	7.23d	8.13cd	3.00c	3.45bc	2.28c	2.35c	5.29c	6.01bc	2.81b	2.94a	1.88bc	2.05cd
72-4-3	8.29bc	8.57bc	3.49a	3.36bc	2.23c	2.55bc	5.76c	6.15b	3.15a	2.88a	1.83c	2.07cd
72-4-9	9.10ab	8.94b	3.47ab	3.26c	2.68b	2.74b	6.91b	6.05b	3.13a	2.93a	2.20b	2.14b
Akitakomachi	7.37cd	7.52d	3.87a	3.96a	1.84d	1.90d	5.36c	5.51d	3.16a	3.16a	1.70c	1.75cd
Koshihikari	7.53cd	7.57d	3.75a	3.65b	2.08cd	2.08d	5.55c	5.58cd	3.20a	3.12a	1.73c	1.79cd
Riceberry	9.96a	10.52a	2.88c	2.50d	3.89a	4.22a	7.71a	7.39a	1.98c	1.96b	3.91a	3.77a
F-test ($P<0.01$)	**	**	**	**	**	**	**	**	**	**	**	**
CV%	4.59	4.59	4.59	4.59	4.59	4.59	4.59	4.59	4.59	4.59	4.59	4.59

645 ^{1/} Different letters in the same column indicate significant differences at the 0.05 level using LSD.

646 Table 3 Pericarp thickness, pigmentation intensity and pericarp color of the three candidate lines compared with their parents in WS20.

Lines/Varieties	Pericarp thickness (mm)	Pigmentation intensity			Pericarp color
		L^*	a^*	b^*	
		69-1-1	0.020c ^{1/}	10.35b	

CV% 0.85 2.25 1.54 3.21 2.85 1.20 0.51 0.65 0.75

650 ^{1/} Different letters in the same column indicate significant differences at the 0.05 level using LSD.

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654 Table 5 Nutrition concentration of the three candidate lines compared with their parents in WS20.

Lines/Varieties	Iron	Zinc	Vitamin E	Vitamin B6	Folic acid	Anthocyanin
	(mg/100 g)	(mg/100 g)	(mg/100 g)	(mg/100 g)	(µg/100 g)	(mg/100 g)
69-1-1	1.69a ^{1/}	1.92b	1.49a	0.23a	48c	59bc
72-4-3	1.39b	1.88b	1.48a	0.17b	52b	62b
72-4-9	1.19c	2.15a	1.43ab	0.17b	54b	78a
Akitakomachi	1.02d	1.38c	0.41c	0.15bc	41d	0e
Riceberry	1.61a	1.99b	1.34b	0.15bc	70a	68b
F-test (<i>P</i> <0.01)	**	**	**	**	**	**

CV% 4.23 3.89 4.18 3.52 4.01 5.96

655 ^{1/} Different letters in the same column indicate significant differences at the 0.05 level using LSD.

656

657 Table 6 Phytochemical contents, antioxidant inhibitory effects and enzyme inhibitory activities of the three candidate lines compared with their parents in
658 WS20.

Lines/Varieties	Phytochemical contents		Antioxidants inhibitory effects		Enzymes inhibitory activities	
			(%)		(%)	
	Flavonoids (mg RU/100 g extract)	Phenolics (mg GA/100 g extract)	DPPH	ABTS	α -amylase	α -glucosidase
69-1-1	61.03±1.14b ^{1/}	141.88±9.62b	8.13±3.91b	86.59±0.39a	3.35±0.22b	51.52±5.41a
72-4-3	51.13±1.25c	123.50±3.06c	8.24±3.97b	87.18±0.60a	2.99±0.23b	56.76±9.77a
72-4-9	71.65±1.79a	176.45±16.30a	11.99±2.89a	86.72±0.25a	11.10±1.38a	60.93±4.66a
Akitakomachi	1.05±0.58e	4.08±0.24e	0.00±0.00c	38.69±5.02b	3.86±0.75b	33.57±8.86b
Riceberry	31.56±0.51d	89.02±10.41d	7.04±3.21b	86.72±0.34a	13.08±0.61a	28.99±5.04b
F-test (<i>P</i> <0.05)	*	*	*	*	*	*
CV%	2.67	9.12	14.70	7.42	8.66	25.52

659 ^{1/} Different letters in the same column indicate significant difference at the 0.05 level using LSD.

660

661 Table S1 Gene-based/linked markers used for foreground selection for blast resistance and cooking quality for validation in the breeding lines.

Target Traits	Gene	Chr.	Marker name	SNP/Indel	Homozygous target SNP/Indel	LGC code ^{1/}	Aliquot ID ^{1/}
Amylose content	<i>wx^b</i>	6	wx_5UTR_G/T	G/T	T:T	002-0052.1	1107690034
Gelatinization temperature	<i>SSIIa</i>	6	ALK_ex8_SNP_A/G	A/G	A:A	002-0049.1	1103441167
Short grain	<i>GS3</i>	3	GS3	C/A	C:C	002-0755.1	1132528814
Blast resistance	<i>Pi-ta</i>	11	TBGI453598	T/C	T:T	002-0821.1	1140749274

662 ^{1/} LGC group, 2015

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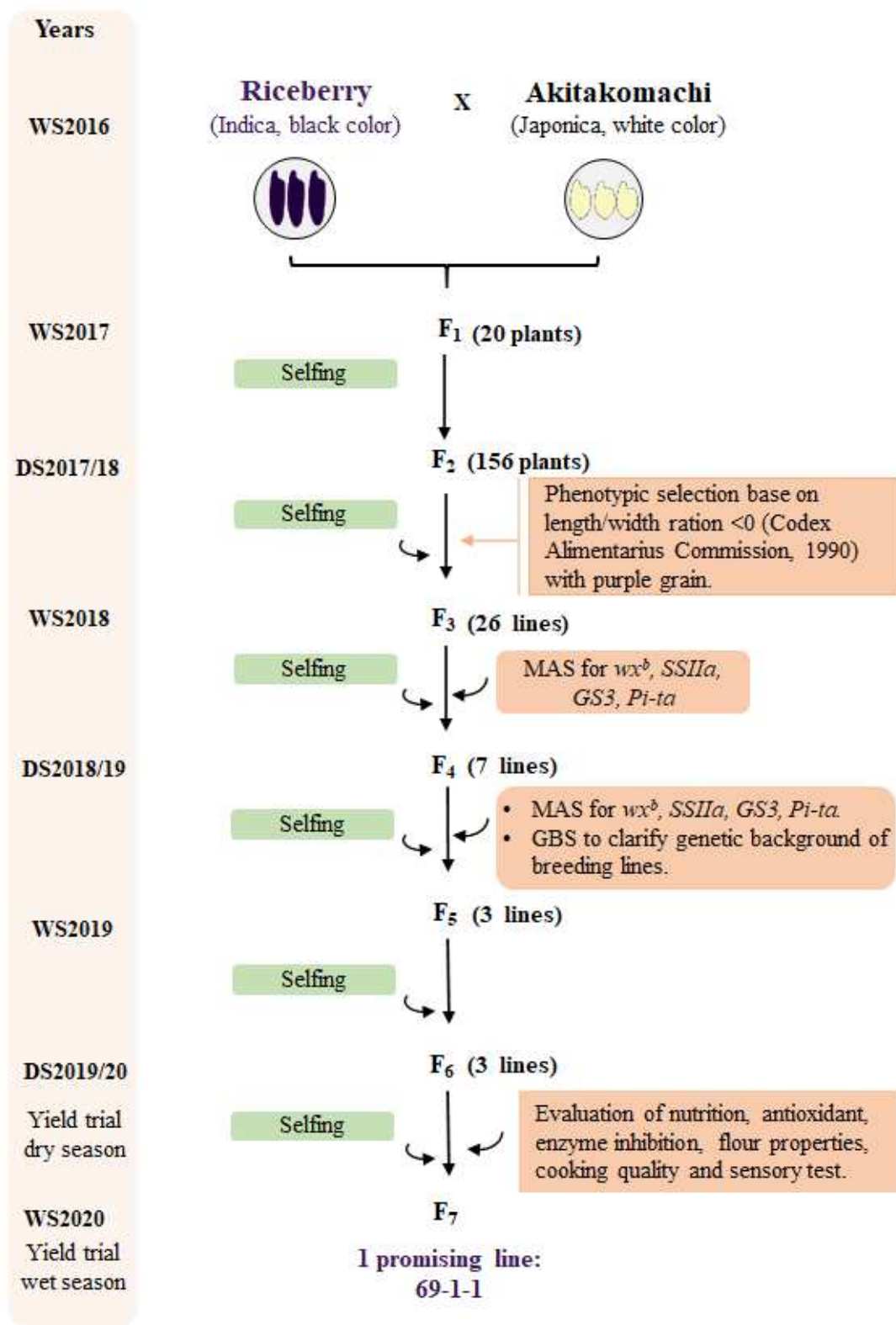
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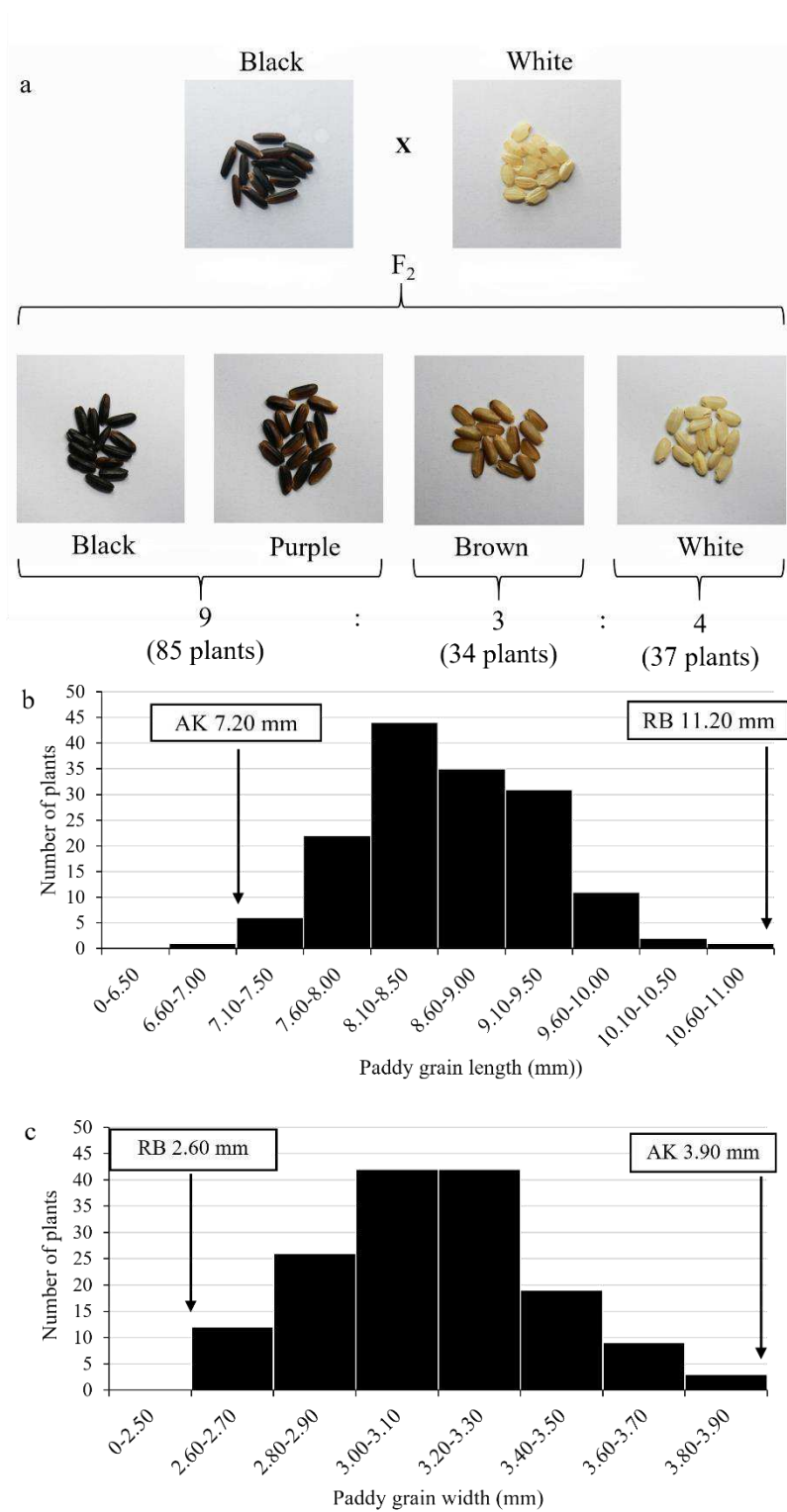


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672 Figure 1 Scheme of the breeding program for black short grain rice derived from Akitakomachi x

673 Riceberry from WS16 to WS21 in the Phan District, Chiang Rai Province, Thailand.

674 WS, Wet season; DS, Dry season; MAS, Marker-assisted selection; GBS, Genotype by sequencing

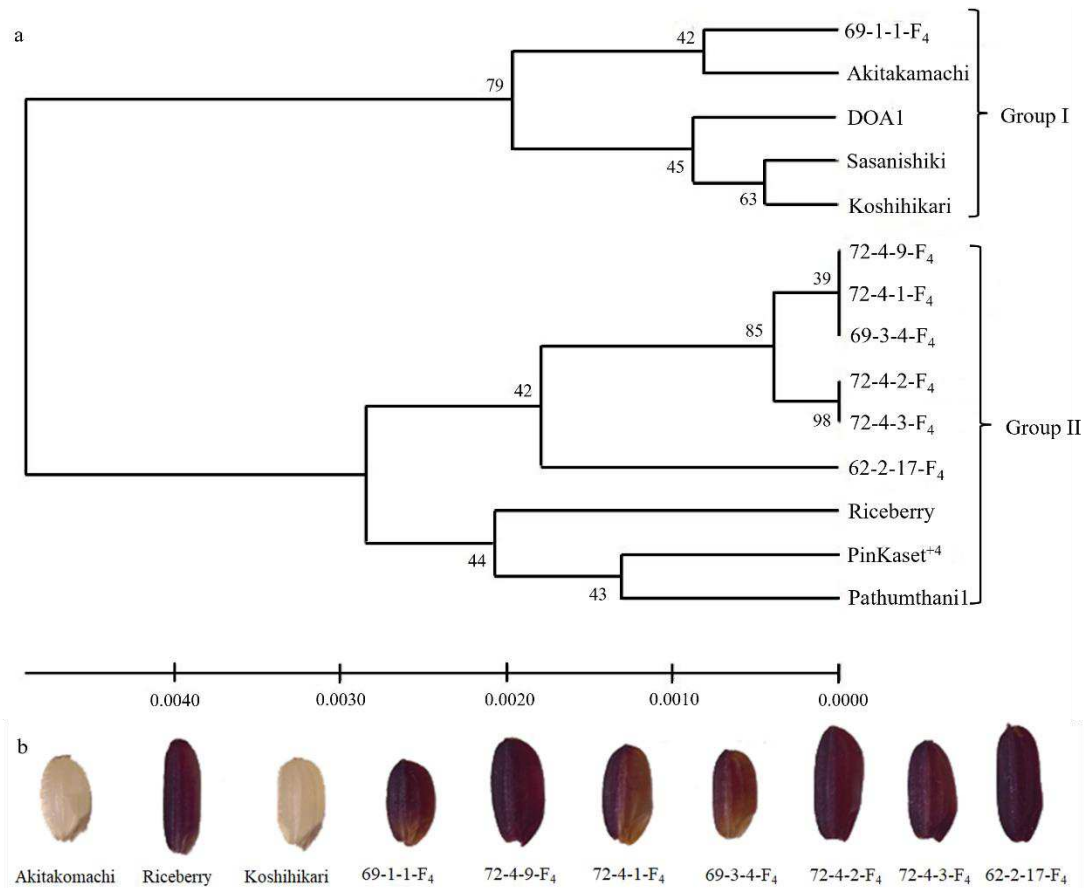


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676 Figure 2 F₂ segregation of (a) grain color, (b) grain length and (c) grain width compared with

677 Akitakomachi and Riceberry.

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679

680 Figure 3 Phylogenetic tree (a) of the breeding lines in F₄ and controlled varieties based on genotyping
 681 by sequencing. The phylogenetic tree was classified into two groups, and the numbers at the
 682 nodes indicate the percentages obtained with 1000 bootstrap replicates. The unpolished
 683 grains (b) of seven selected lines in F₄ compared with their parents and Koshihikari.

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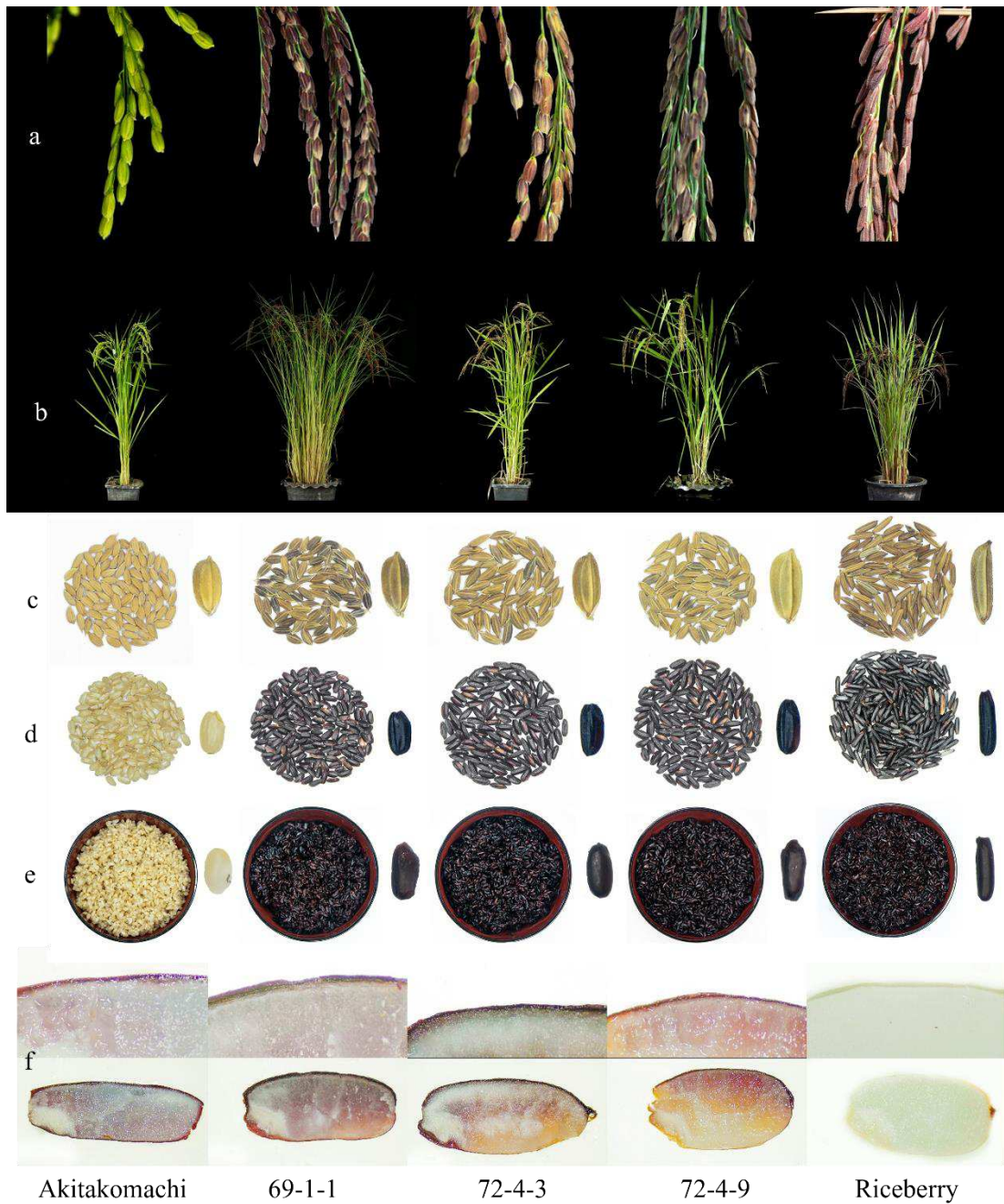
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692 Figure 4 Plant type (a and b), paddy grain (c), unpolished grain (d), cooked grain (e) and longitudinal

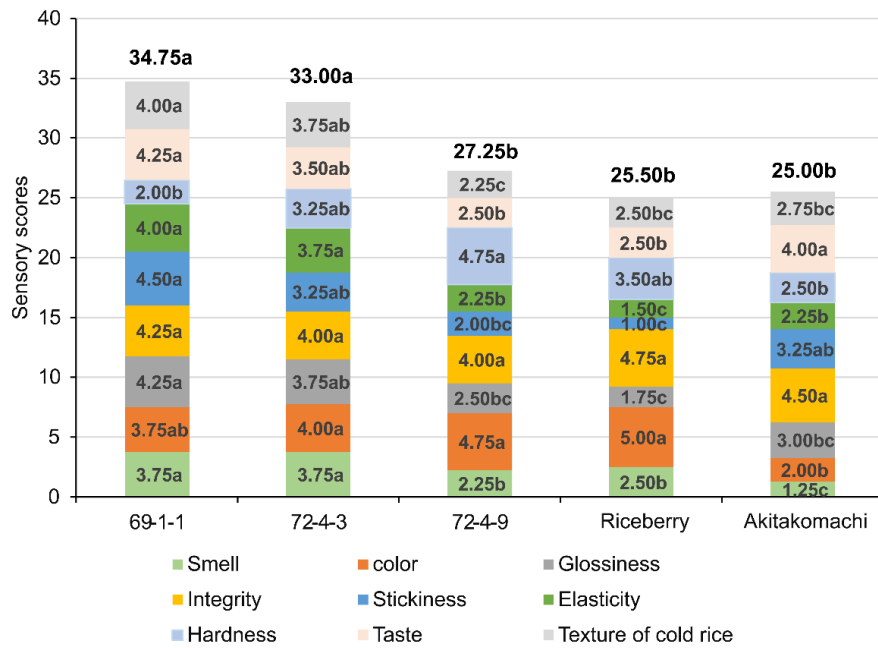
693 section of unpolished grain (f) of the three candidate lines in the F₇ generation compared

694 with their parents in WS20.

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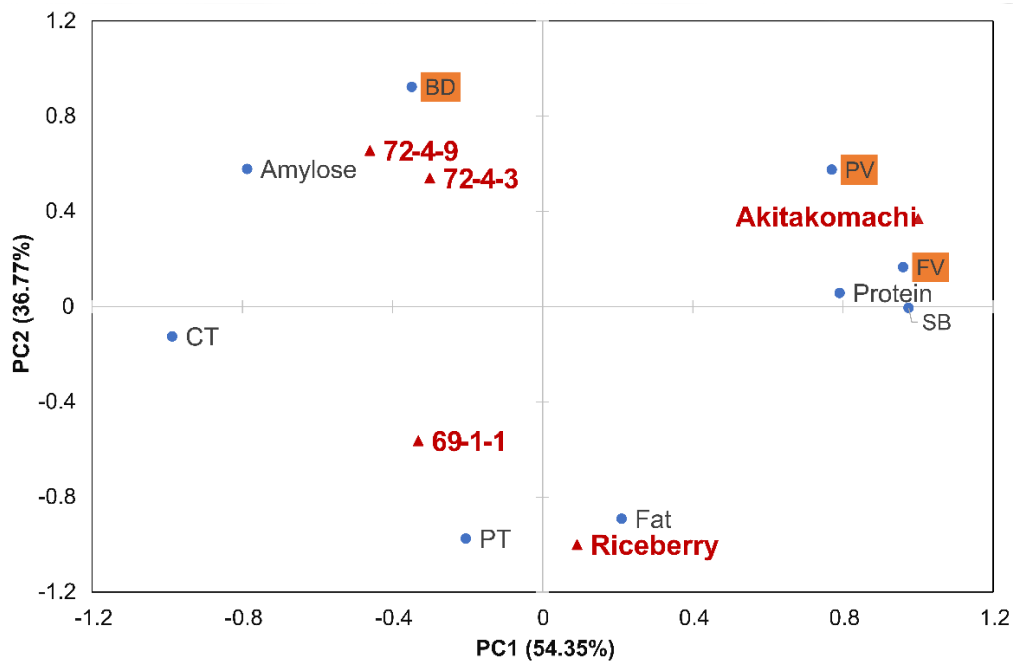
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699 Figure 5 Sensory test results of the unpolished grains of the three candidate lines in the F₇ generation

700 compared with their parents in WS20.



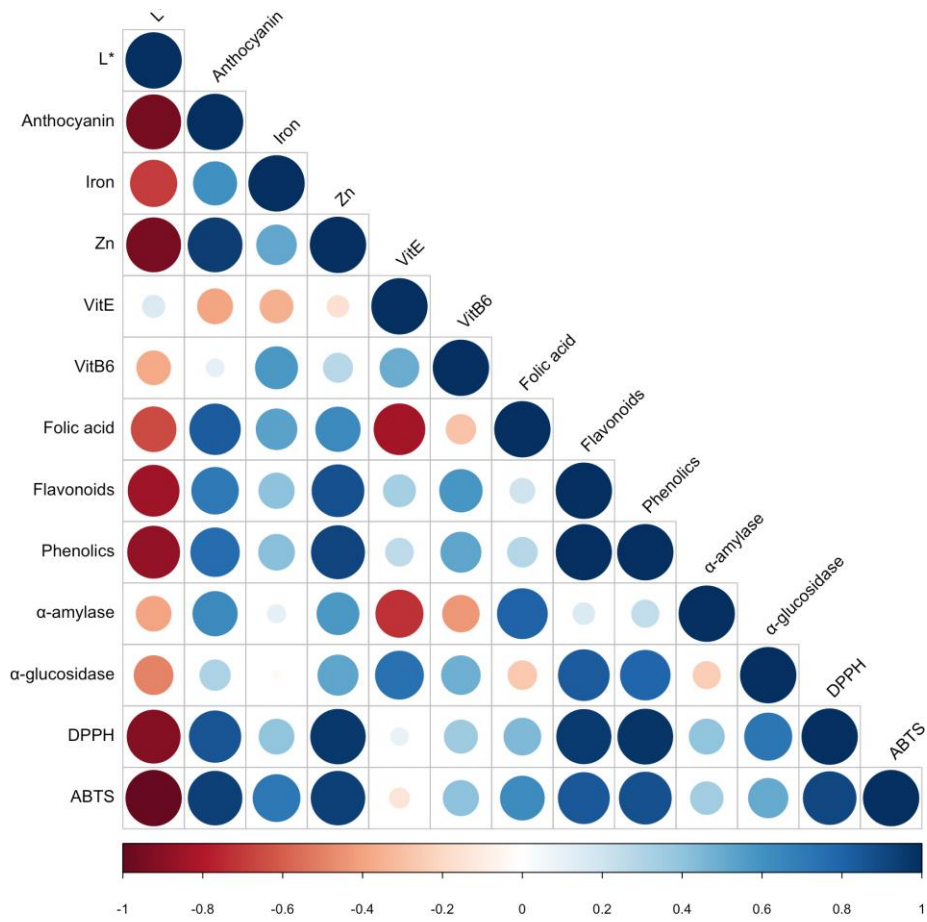
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702 Figure 6 Biplot graph of the PC1 score versus the PC2 score for the grain quality of the three

703 candidate lines and the control varieties. AK, Akitakomachi, RB, Riceberry, BD,

704 Breakdown, FV, Final viscosity, PV, peak viscosity, SB, Setback, CT, Cooking time, PT,

705 Pasting temperature.



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707 Figure 7 Correlation coefficients (r) among color intensity (L^*), nutrition, phytochemical, antioxidant
 708 and antidiabetic activity.

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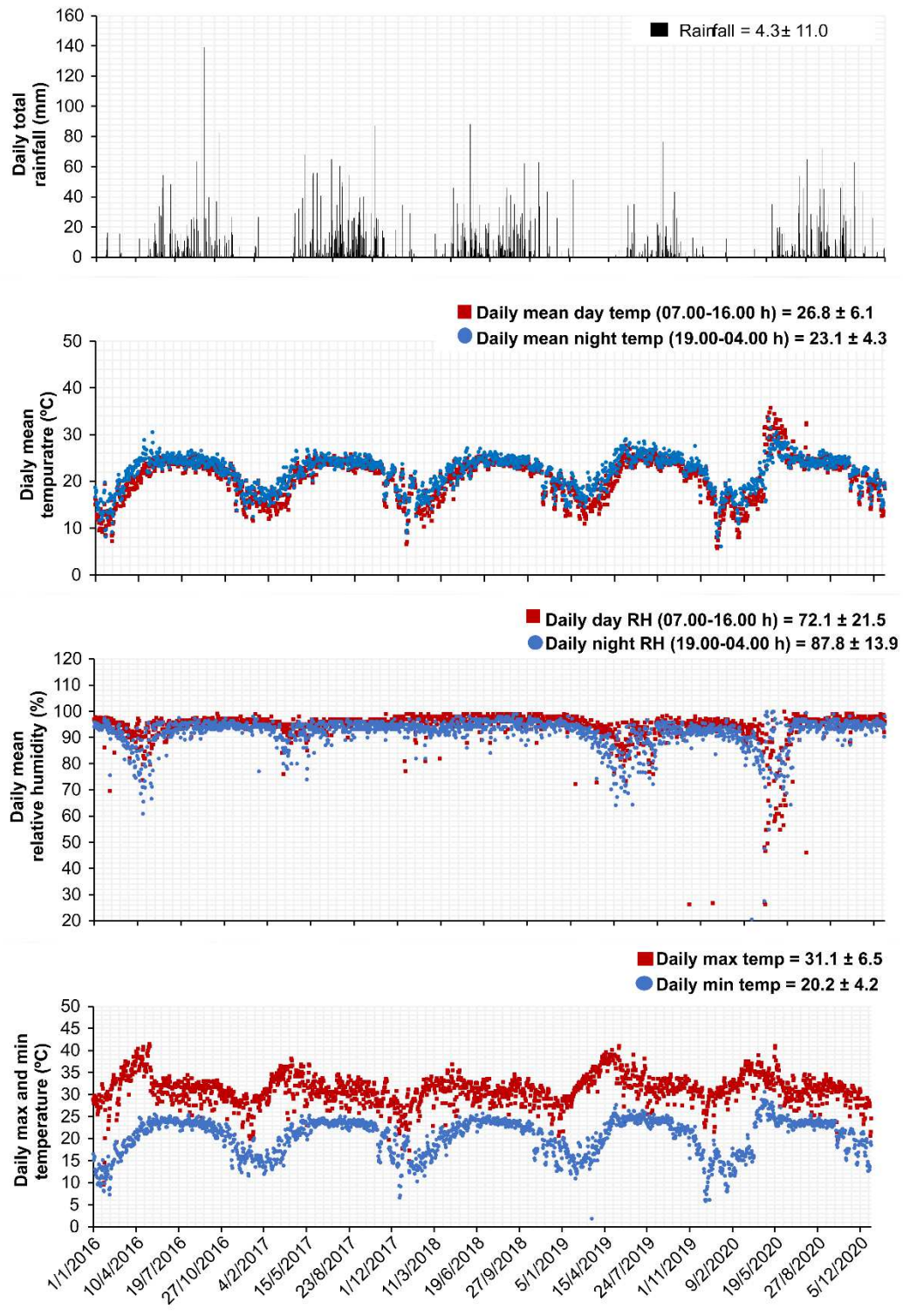
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718 Figure 1S Weather data from WS16-WS20 in the Phan District, Chiang Rai Province.

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726 Fig. 2S F₂ segregation of grain color derived from Riceberry x Akitakomachi.