Application of γ-aminobutyric acid (GABA) and nitrogen regulates aroma biochemistry in fragrant rice

Wenjun Xie1 | Umair Ashraf1,2 | Dating Zhong1 | Rongbin Lin1 | Peiqi Xian1 | Tong Zhao1 | Huoyi Feng1 | Shuli Wang1,3 | Meiyang Duan1,3 | Xiangru Tang1,3 | Zhaowen Mo1,3

1College of Agriculture, South China Agricultural University, Guangzhou, China  
2Department of Botany, Division of Science and Technology, University of Education, Lahore, Punjab, Pakistan  
3Scientific Observing and Experimental Station of Crop Cultivation in South China, Ministry of Agriculture, P. R. China, Guangzhou, China

Correspondence  
Zhaowen Mo and Xiangru Tang, College of Agriculture, South China Agricultural University, Guangzhou, 510642, Guangdong, China.  
Emails: scaumozhw@126.com and tangxr@scau.edu.cn

Funding information  
National Natural Science Foundation for Young Scientists, Grant/Award Number: 31601244; Special Fund for National Natural Science Foundation of China, Grant/Award Number: 31271646

Abstract

The 2-acetyl-1-pyrroline (2AP) is a key aroma compound in fragrant rice. The present study assessed the γ-aminobutyric acid (GABA) and nitrogen (N) application induced regulations in the biochemical basis of rice aroma formation. Four N levels, that is, 0, 0.87, 1.75, and 2.61 g/pot, and two GABA treatments, that is, 0 mg/L (GABA0) and 250 mg/L (GABA250), were applied to three fragrant rice cultivars, that is, Yuxiangyouzhan, Yungengyou 14, and Basmati-385. Results showed that GABA250 increased 2AP, Na, Mn, Zn, and Fe contents by 8.44%, 10.95%, 25.70%, 11.14%, and 43.30%, respectively, under N treatments across cultivars. The GABA250 further enhanced the activities of proline dehydrogenase (PDH), ornithine aminotransferase (OAT) (both at 15 days after heading (d AH), and diamine oxidase (DAO) (at maturity) by 20.36%, 11.24%, and 17.71%, respectively. Significant interaction between GABA and N for Mn, Zn, and Fe contents in grains, proline content in leaves, GABA content in leaves at 15 d AH and maturity stage (MS), Δ1-pyrroline-5-carboxylic acid (P5C) contents in leaves at 15 d AH, and Δ1-pyrroline-5-carboxylate synthase (P5CS), PDH, and OAT activities in leaves at MS was noted. Moreover, the 2AP contents in grains at MS showed a significant and positive correlation with the proline contents in the leaves at 15d AH. In conclusion, GABA250 enhanced the 2AP, Na, Mn, Zn, and Fe contents, as well as the enzyme activities involved in 2AP biosynthesis. Exogenous GABA and N application improved the 2AP contents and nutrient uptake in fragrant rice.

KEYWORDS

2-acetyl-1-pyrroline, enzyme activity, fragrant rice, nutrient contents, γ-aminobutyric acid

1 | INTRODUCTION

Rice is an important cereal crop, which fulfills the dietary needs of billions of people globally (Mahajan, Sekhon, Singh, Kaur, & Sidhu, 2010), whereas fragrant rice is well praised and liked by the consumers due to its high grain quality, strong aroma, and good taste (Chutipaijit & Sutjaritvorakul, 2018; Sasmal & Pal, 2018). “Basmati” and “Jasmine” are the typical fragrant rice types, which are world-famous due to their long-grain and fragrant characters. Previous studies have revealed that hundreds of volatile compounds have been detected in aromatic rice, among them 2-acetyl-1-pyrroline (2AP) was recognized as a key aroma compound (Maraval et al., 2008; Shao et al., 2013).
The betaine-aldehyde dehydrogenase (BADH)-related gene was found to be involved in aroma formation in fragrant rice (Bradbury, Fitzgerald, Henry, Jin, & Waters, 2005; Bradbury, Gillies, Brushett, Waters, & Henry, 2008; Chen et al., 2008; Fitzgerald, Hamilton, Calingacion, Verhoeven, & Butardo, 2008; Kovach, Calingacion, Fitzgerald, & McCouch, 2009; Niu et al., 2008). Previous studies revealed the activities of proline dehydrogenase (PDH), Δ1-pyrroline-5-carboxylate synthase (P5CS), ornithine aminotransferase (OAT), and diamine oxidase (DAO), as well as the concentrations of proline and Δ1-pyrroline-5-carboxylic acid (P5C) contents, were involved in 2AP formation in fragrant rice (Ghosh & Roychoudhury, 2018; Huang et al., 2008; Li, Ashraf, et al., 2016; Mo, Fan, et al., 2016; Mo, Huang, et al., 2016; Poonlapdechea et al., 2012).

The biosynthesis and regulations of 2AP are largely affected by not only the genetic factors but also the external environmental and crop management factors (Bao et al., 2018; Hasanuzzaman et al., 2019). The genetic factors play an important part in determining the quality of aromatic rice (Bao et al., 2018; Bradbury et al., 2008), which, on the other hand, are much dependent on prevailing environmental conditions during the growth period and also by pre- and postharvest management techniques (Gay et al., 2010). For example, nitrogen fertilization has great impacts on 2AP formation and accumulation. Recently, Ren et al. (2017) showed that 60 kg/ha pre- and postharvest management techniques (Gay et al., 2010). For example, nitrogen fertilization has great impacts on 2AP formation and accumulation. Recently, Ren et al. (2017) showed that 60 kg/ha nitrogen with water deficit conditions at tillering stage could enhance the 2AP contents in grains significantly, whereas Mo et al. (2018) also demonstrated that different nitrogen application levels affected the 2AP contents in fragrant rice differently. Early studies revealed that BADH induced the biosynthesis of γ-aminobutyric acid (GABA) by γ-aminobutyraldehyde, while the inactivation of BADH in aromatic rice induced the accumulation of Δ1-pyrroline (Bradbury et al., 2008). Moreover, the 2AP and GABA contents in grain were increased under salt treatment and shading condition (Mo et al., 2015; Poonlapdechea et al., 2012) and there was a significant positive correlation between 2AP and GABA in the grains of “Yuxiangyouzhan” rice cultivar (Mo et al., 2015). Hence, there might be a possibility for exogenous GABA to regulate the 2AP biosynthesis.

GABA is well known as a plant signaling molecule and is involved in various physio-biochemical processes in plants (Fait, Fromm, Walter, Gall, & Fernie, 2008; Routray & Rayaguru, 2018). GABA has been reported to increase plant resistance ability against several environmental stresses (Yang, Shewfelt, Lee, & Kays, 2008). Regulatory effects of GABA in plant physio-biochemical metabolism have been previously reported in Arabidopsis, tobacco, maize, barley, and rice (Batsushansky et al., 2014; Li, Guo, Yang, Meng, & Wei, 2016; Nayar, Kaur, Kaur, & Singh, 2014; Song, Xu, Wang, Wang, & Tao, 2010; Yu & Sun, 2007), whereas exogenous GABA applications could improve salt stress resistance in wheat seedlings (Li, Guo, et al., 2016) and enhance photosynthesis capacity in maize seedling (Li, Liu, et al., 2016). In sum, GABA plays a crucial role in plant growth to respond to the external environmental conditions (Fait et al., 2008; Li, Guo, et al., 2016); nevertheless, endogenous GABA levels in plants are very low, but it can be accelerated under stress conditions (Kinnersley & Turano, 2000). Hence, GABA may influence the physiological metabolism of fragrant rice by affecting either aroma biosynthesis or plant growth. Reports are available for GABA-induced regulations in stress tolerance in various crops; however, effects of exogenous GABA application coupled with various N levels on aroma biosynthesis are rarely investigated. Therefore, the present study was aimed to evaluate the effects of GABA under various levels of N on physio-biochemical attributes, aroma biosynthesis, and enzyme involved in 2AP formation in fragrant rice.

2 | MATERIALS AND METHODS

2.1 | Plant material and growing conditions

The seeds of three aromatic rice cultivars, that is, Yuxiangyouzhan (YX), Yugengyou14 (YG), and Basmati-385 (BS), were collected from the College of Agriculture, South China Agricultural University, Guangzhou, China. These aromatic rice cultivars are newly developed and famous countrywide due to their aroma. A pot experiment was performed in the greenhouse of the Experimental Research Farm, College of Agriculture, South China Agricultural University, Guangzhou, China, during the early growing season of 2017. Rice seedlings (30 days old) were transplanted into the soil-containing pots (31cm in diameter and 29 cm in height) on 4 April. The experimental soil was sandy loam containing soil organic matter 18.65 g/kg, total nitrogen 1.17 g/kg, available phosphorus 32.69 mg/kg, available potassium 185.28 mg/kg, and pH 6.44. The yearly average temperature of the region lies between 21 and 29°C and is characterized by a subtropical monsoonal type of climate.

2.2 | Treatments and plant sampling

The treatment without GABA application was denoted as GABA0 (double-distilled water sprinkler application), and the GABA at 250mg/L (GABA250) was applied at panicle initiation stage, whereas four nitrogen rates, that is, 0, 0.87, 1.75, and 2.61 g/pot denoted as ZN, LN, MN, and HN, were applied as basal dose. Flag leaves at the heading stage (HS), 15 days after heading stage (15d AH), and maturity stage (MS), whereas mature panicles at MS were collected and stored at -80°C for physio-biochemical assays.

2.3 | Determination of 2AP contents in grains

The 2AP concentration was determined by synchronization distillation and extraction method (SDE) combined with GCMS-QP 2010 Plus (Shimadzu Corporation) according to Huang et al. (2012) and the contents of 2AP were expressed as μg/g.

2.4 | Determination of Na, Mn, Zn, and Fe contents in grains

The Na, Mn, Zn, and Fe contents in grains were determined according to the method of Wu et al. (2014) with some modification. In brief, the grain sample was oven-dried at 80°C to constant weight and ground in powder, and then weighed 0.25 g of the sample. The
weighted samples were dry-digested in a muffle furnace at 500°C for 6 hr, and then added to 40 ml HNO₃·H₂O (1:1). The contents of Na, Mn, Zn, and Fe were determined using a flame atomic absorption spectrometry (SHIMADZU AA-6300C AA spectrometer).

2.5 | Measurement of protein contents in grains

After sun drying, about 1.5 kg grains from each treatment were taken to estimate the protein contents by using an Infratec 1241 grain analyzer (FOSS-TECATOR; Mo et al., 2015).

2.6 | Determination of proline, GABA, and pyrroline-5-carboxylic acid (P5C) contents in leaves

The proline contents in fresh leaf samples were determined by Bates, Waldren, and Teare (1973) by using ninhydrin, and the absorbance was read at 520 nm. The final proline contents were expressed as µg/g fresh weight (FW) of leaves. The GABA contents were measured according to the methods described by Zhao et al. (2009), and the GABA contents were expressed as µg/g. The P5C concentration was estimated by following the methods of Wu, Chou, Wu, Chen, and Huang (2009). The reaction mixture contained 0.2 ml supernatant of enzyme extract, 0.5 ml of 10% trichloroacetic acid (TCA), and 0.2 ml of 40 mM 2-aminobenzaldehyde. The absorbance was read at 440 nm, and the contents were expressed as µmol/g.

2.7 | Determination of the activities of proline dehydrogenase (PDH), Δ1-pyrroline-5-carboxylic acid synthetase (P5CS), ornithine aminotransferase (OAT), and diamine oxidase (DAO)

The PDH activity was assayed by following the methods of Tateishi, Nakagawa, and Esaka (2005) and Ncube, Finnie, and Van (2013). The absorbance after reaction was read, and the reaction mixture contained L-proline (15 mM), cytochrome c (0.01 mM), phosphate buffer (100 mM, pH 7.4), 0.5% (v/v) Triton X-100, and the enzyme extract (0.1 ml) in a total volume of 0.5 ml. The reaction mixture was incubated at 37°C for 30 min, and the reaction was terminated by adding 0.5 ml of 10% trichloroacetic acid (TCA). After adding 0.5 ml of 0.5% 2-aminobenzaldehyde in 95% ethanol, the mixture was further incubated at 37°C for 10 min and centrifuged at 8,000 rpm for 10 min, and the absorbance of the supernatant was read at 440 nm, the absorbance change of 0.1 in one minute was defined as one unit of enzyme activity, and the activity was expressed as U/g FW.

The DAO activity was assayed by using the methods described by Su, An, Zhang, and Liu (2005). The reaction solutions (3.0 ml) contained 2.5 ml 0.1 M sodium phosphate buffer (pH 6.5), 0.1 ml crude enzyme extracts, 0.1 ml peroxidase (250 U/ml), and 0.2 ml 4-aminoantipyrine/N, N-dimethylamine reaction solutions. The reaction was initiated by the addition of 0.1 ml 20 mM Put. A 0.01 value of the changes in absorbance at 440 nm was regarded as one activity unit of the enzyme, and the activity was expressed as U/g FW.

2.8 | Statistical analyses

The pots were arranged in a completely randomized design (CRD), and the data were analyzed by using Statistix version 8 (Analytical Software). Relationships among the indexes were evaluated using correlation analyses by Statistix version 8 (Analytical Software). Means among treatments were compared based on the least significant difference test (LSD) at the 0.05 probability level.

3 | RESULTS

3.1 | 2AP contents

The 2AP contents in grains were substantially affected by GABA treatment (T), variety (V), nitrogen (N), and T × V. Compared with GABA0, the GABA250 significantly increased the 2AP contents by 8.44% under nitrogen treatments across cultivars. With the increase in the application dose of nitrogen fertilizer, the 2AP contents in grains of all rice cultivars were increased. Moreover, the highest 2AP in the application dose of nitrogen fertilizer, the 2AP contents in grains of aromatic rice cultivars. Compared with GABA0, GABA250 led to a decrease in the protein content by 3.76% under

The OAT activity was measured according to the methods of Chen, Chen, Lin, and Kao (2001) and Umair, Leung, Bland, and Simpson (2011). The reaction medium contained 100 mM potassium phosphate buffer pH 8.0, 50 mM ornithine, 20 mM α-ketoglutarate, 1 mM pyridoxal 5-phosphate, and the enzyme extract (0.1 ml)—the final total volume was 1 ml. The reaction medium was incubated at 37°C for 30 min. The reaction was stopped by adding 0.5 ml trichloroacetic acid (10%), and the color was developed by incubating the reaction mixture with 0.5 ml α-amino benzaldehyde (0.25%) in ethanol (95%) for 1 hr. After centrifugation at 8,000 rpm for 10 min, the clear supernatant fraction was taken to measure the absorbance at 440 nm. The absorbance change of 0.1 in one minute was defined as one unit of enzyme activity; the activity was expressed as U/g FW.

The DAO activity was assayed by using the methods described by Su, An, Zhang, and Liu (2005). The reaction solutions (3.0 ml) contained 2.5 ml 0.1 M sodium phosphate buffer (pH 6.5), 0.1 ml crude enzyme extracts, 0.1 ml peroxidase (250 U/ml), and 0.2 ml 4-aminoantipyrine/N, N-dimethylamine reaction solutions. The reaction was initiated by the addition of 0.1 ml 20 mM Put. A 0.01 value of the changes in absorbance at 440 nm was regarded as one activity unit of the enzyme, and the activity was expressed as U/g FW.

2.8 | Statistical analyses

The pots were arranged in a completely randomized design (CRD), and the data were analyzed by using Statistix version 8 (Analytical Software). Relationships among the indexes were evaluated using correlation analyses by Statistix version 8 (Analytical Software). Means among treatments were compared based on the least significant difference test (LSD) at the 0.05 probability level.

3 | RESULTS

3.1 | 2AP contents

The 2AP contents in grains were substantially affected by GABA treatment (T), variety (V), nitrogen (N), and T × V. Compared with GABA0, the GABA250 significantly increased the 2AP contents by 8.44% under nitrogen treatments across cultivars. With the increase in the application dose of nitrogen fertilizer, the 2AP contents in grains of all rice cultivars were increased. Moreover, the highest 2AP in the application dose of nitrogen fertilizer, the 2AP contents in grains of aromatic rice cultivars. Compared with GABA0, GABA250 led to a decrease in the protein content by 3.76% under

3.2 | Protein contents

The T, V, N, T × V, and V × N significantly affected the protein contents in grains of aromatic rice cultivars. Compared with GABA0, GABA250 led to a decrease in the protein content by 3.76% under
N treatments across cultivars. With the increased application dose of N fertilizer, the mean protein contents in grains were increased. BS showed higher mean protein content than YX and YG (Figure 2).

3.3 | Na, Mn, Zn, and Fe contents in grains

Compared with GABA0, Na content in grains under GABA250 was significantly higher, which was 5.33%. Similar trends were also recorded for Mn, Zn, and Fe contents. Compared with GABA0, the Mn, Zn, and Fe contents under nitrogen treatments across cultivars were increased by 25.72%, 11.14%, and 43.30%, respectively, under GABA250. Moreover, GABA and nitrogen interaction affected Na, Mn, Zn, and Fe content in grains significantly. For YX, higher Na, Mn, Zn, and Fe content in grains were found under LN (124.48 mg/kg), ZN (132.33 mg/kg), HN (11.64 mg/kg), and ZN (63.78 mg/kg) under GABA0 treatment, respectively, while higher Na, Mn, Zn, and Fe content in grains were found under HN (188.95 mg/kg), ZN (180.85 mg/kg), ZN (35.89 mg/kg), and ZN (102.38 mg/kg), respectively. For YG, higher Na, Mn, Zn, and Fe content in grains were found under LN, ZN, and ZN, respectively, and HN and MN treatment accumulated highest Zn content under GABA0 and GABA250 treatment, respectively. For BS, highest Mn, Zn, and Fe contents in grains were found under ZN treatment, and HN and LN treatment accumulated highest Zn content under GABA0 and GABA250 treatment, respectively (Table 1).

3.4 | Proline contents in leaves

Proline content in leaves at HS, 15 d AH, and MS were significantly affected by V, N, T × N, V × N, and T × V×N. The T × V significantly affected the proline content in leaves at HS and 15 d AH. At GABA250, the proline contents in leaves were substantially reduced MS. For YX, under GABA0 treatment, proline content in MN, HN, and LN treatment was the highest at HS, 15 d AH, and MS, respectively, while under GABA250 treatment, proline content in HN, MN, and ZN treatment was the highest at HS, 15 d AH, and MS, respectively. For YG, ZN treatment resulted in the highest proline contents in leaves at HS under GABA0 treatment, while HN treatment produced the highest proline content in leaves at 15 d AH and...
The proline contents in leaves under HN treatment remained the highest under GABA250 treatment where the proline contents were 14.404, 15.096, and 5.690 µg/g FW for HS, 15 d AH, and MS, respectively. For BS, MN produced the highest proline content in leaves at HS; ZN and HN resulted in the highest proline in leaves at 15 d AH under GABA0 and GABA250, respectively, and HN and ZN had the highest proline in leaves at MS under GABA0 and GABA250, respectively (Table 2).

3.5 | P5C concentration

GABA treatment (T) did not affect the P5C contents in leaves, whereas the cultivars were found statistically significant regarding P5C contents in leaves. Significant differences were observed for T × V, N, T × N, V × N, and T × V×N at 15 d AH. Furthermore, YG produced lower P5C content in leaves as compared to YX and BS, while P5C contents substantially differed among different growth stages (Table 3).
### TABLE 2  Effect of γ-aminobutyric acid (GABA) and nitrogen on proline accumulation in leaves in different fragrant rice genotypes (μg/g fresh weight (FW))

| Treatment | HS | 15d AH | MS  |
|-----------|----|--------|-----|
| GABA0     |    |        |     |
| YX        |    |        |     |
| ZN        | 21.129 | 6.988  | 9.056 |
| LN        | 19.546 | 7.207  | 7.522 |
| MN        | 27.310 | 6.300  | 6.571 |
| HN        | 15.956 | 8.245  | 9.444 |
| YG        |    |        |     |
| ZN        | 21.341 | 10.250 | 4.837 |
| LN        | 14.544 | 11.332 | 5.257 |
| MN        | 13.959 | 13.342 | 5.109 |
| HN        | 9.199  | 14.543 | 7.708 |
| BS        |    |        |     |
| ZN        | 12.915 | 9.331  | 7.691 |
| LN        | 13.901 | 8.496  | 8.685 |
| MN        | 16.443 | 6.810  | 8.622 |
| HN        | 16.300 | 8.253  | 11.624 |
| Mean      | 16.879 | 9.258  | 7.677a |

| GABA250   |    |        |     |
| YX        |    |        |     |
| ZN        | 22.918 | 8.671  | 8.357 |
| LN        | 17.828 | 6.249  | 7.058 |
| MN        | 20.753 | 8.247  | 6.814 |
| HN        | 23.858 | 9.287  | 11.624 |
| Mean      | 16.145 | 9.122  | 6.663b |

**ANOVA**

- **T**: 1.29ns 0.27ns 48.11**
- **V**: 76.52** 122.05** 238.41**
- **T × V**: 3.94* 17.0** 3.36ns
- **N**: 5.61** 19.44** 43.24**
- **T × N**: 15.14** 4.09* 16.37**
- **V × N**: 6.42** 10.53** 7.69**
- **T × V × N**: 9.88** 2.66* 4.28**

Note: Means with different lowercase letter(s) differ statistically at p < .05.

Abbreviations: BS, Basmati-385; d AH, days after heading; HS, heading stage; MS, maturity stage; YG, Yugengyou14; YX, Yuxiangyouzhan.

*Significant at p < .05; **Significant at p < .01; ns: nonsignificant.

### TABLE 3  Effect of γ-aminobutyric acid (GABA) and nitrogen on P5C accumulation in leaves in different fragrant rice genotypes (µmol/g FW)

| Treatment | HS | 15d AH | MS  |
|-----------|----|--------|-----|
| GABA0     |    |        |     |
| YX        |    |        |     |
| ZN        | 0.509  | 1.085  | 1.010 |
| LN        | 0.428  | 0.857  | 1.243 |
| MN        | 0.642  | 1.015  | 0.979 |
| HN        | 0.614  | 1.355  | 1.265 |
| YG        |    |        |     |
| ZN        | 0.391  | 0.945  | 0.534 |
| LN        | 0.484  | 0.636  | 0.511 |
| MN        | 0.378  | 0.648  | 0.639 |
| HN        | 0.352  | 0.289  | 0.367 |
| BS        |    |        |     |
| ZN        | 1.228  | 0.819  | 0.864 |
| LN        | 2.148  | 0.709  | 1.194 |
| MN        | 0.673  | 0.774  | 0.994 |
| HN        | 0.657  | 0.628  | 0.910 |
| Mean      | 0.709  | 0.813  | 0.876 |

| GABA250   |    |        |     |
| YX        |    |        |     |
| ZN        | 0.545  | 0.895  | 0.907 |
| LN        | 0.494  | 1.274  | 0.921 |
| MN        | 0.426  | 1.395  | 1.090 |
| HN        | 0.548  | 1.180  | 1.296 |
| YG        |    |        |     |
| ZN        | 0.722  | 0.488  | 0.465 |
| LN        | 0.437  | 0.547  | 0.832 |
| MN        | 0.414  | 0.560  | 0.461 |
| HN        | 0.487  | 0.456  | 0.628 |
| BS        |    |        |     |
| ZN        | 0.936  | 1.005  | 0.746 |
| LN        | 0.628  | 0.860  | 0.781 |
| MN        | 0.764  | 1.328  | 1.130 |
| HN        | 1.313  | 0.714  | 0.815 |
| Mean      | 0.643  | 0.892  | 0.839 |

**ANOVA**

- **T**: 0.56ns 6.65ns 0.65ns
- **V**: 12.89** 81.69** 31.39**
- **T × V**: 1.15ns 8.47** 1.1ns
- **N**: 0.96ns 3.55* 2.23ns
- **T × N**: 2.60ns 4.98** 1.02ns
- **V × N**: 1.01ns 4.32** 1.78ns
- **T × V × N**: 2.67* 2.88* 2.19ns

Note: Means with different lowercase letter(s) differ statistically at p < .05.

Abbreviations: BS, Basmati-385; d AH, days after heading; HS, heading stage; MS, maturity stage; YG, Yugengyou14; YX, Yuxiangyouzhan.

*Significant at p < .05; **Significant at p < .01; ns: nonsignificant.
3.6 | GABA contents

Cultivar significantly differed regarding the accumulation of GABA contents in leaves; however, GABA treatment (T), nitrogen (N), T × V, V × N, and T × V×N did not affect GABA content in leaves significantly. On the other hand, significant differences were observed in GABA content for T × N at 15 d AH and MS, whereas YG produced lower GABA content in leaves as compared to YX and BS, while GABA contents substantially differed (p < .05) among different growth stages (Table 4).

3.7 | Activities of P5CS, PDH, OAT, and DAO

No significant difference was found regarding P5CS activity in leaves under GABA treatment (T). Variety (V) and nitrogen (N) substantially affected the P5CS activity in leaves of all rice cultivars. Significant effect of T × V on P5CS activity in leaves at HS and 15 d AH was observed. Significant interactive effect of V × N and T × V×N on P5CS activity in leaves at 15 d AH and MS and T × N at MS was noted. Lower P5CS activity in leaves was investigated in YG as compared to YX and BS. Under GABA250 treatment, lower nitrogen application improved P5CS activities in the leaves of all aromatic rice cultivars (Table 5).

GABA treatment (T) showed a significant difference in PDH activity in leaves at 15d AH. Variety (V) and T × V significantly affected PDH activity in leaves at HS and 15 d AH. Significant effect of nitrogen (N) on PDH activity in leaves at HS and MS was observed. Remarkable T × N and T × V×N effect on PDH activity in leaves at MS and HS was detected, respectively. Higher PDH activity in leaves was observed under GABA250, and a significant increase in PDH activity in leaves at 15 d AH by 20.364% under GABA250 compared to GABA0 was investigated. Under GABA250 treatment, higher PDH activity in leaves at MS was found for lower N application treatment (Table 6).

Significant effects of GABA treatment (T) were noted on OAT activity in the leaves of aromatic rice cultivars at 15d AH. Varieties (V) were also differed statistically (p < .05) regarding OAT activity, while nitrogen (N) did not affect OAT activity in leaves significantly. Moreover, significant effects of T × V, T × N, and T × V×N on OAT activity in leaves at MS were observed, whereas the T × N effect on OAT activity in leaves at 15 d AH and MS has also remained significant. The OAT activity in the leaves at 15 d AH was increased by 11.242% under GABA250 than GABA0. Under GABA250 treatment, higher OAT activity in leaves was found in ZN for YG and LN and MN for YX (Table 7).

GABA treatment (T) significantly affected DAO activity in leaves at MS, whereas variety (V) showed differential responses regarding DAO activity. Nitrogen (N) significantly affected DAO activity in all rice cultivars at HS and MS. The interactive effects of V × N and T × V×N on DAO activity in leaves at MS and 15 d AH were found significant, respectively. The DAO activity in the leaves at MS was increased by 17.712% under GABA250 compared to GABA0 (Table 8).

3.8 | Correlation analyses

The associations of micronutrients, that is, Na, Mn, Fe, and protein contents in grains, have remained nonsignificant (P > 0.05) with grain 2AP contents. The grain 2AP contents showed a significant and positive correlation with proline contents in leaves at 15d AH under both GABA0 and GABA250. Moreover, under GABA0, significant negative correlations were found between grain 2AP and P5C contents at 15d AH and MS, PDH activity at 15d AH, OAT activity at 15 d AH, P5CS activity at HS, DAO activity at 15 d AH, DAO activity at MS, GABA content at HS, GABA content at 15 d AH, and GABA content at MS. Under GABA250, significant negative correlation relationship exists between 2AP content in grains and proline content at HS and MS, P5C content at 15d AH and MS, PDH activity at HS and 15d AH, OAT activity at 15 d AH and MS, P5CS activity at HS, 15d AH and MS, and GABA content at HS, 15 d AH, and MS (Table 9).

4 | DISCUSSION

Influences of the GABA and N application regulations in the 2AP and the activities of the enzymes involved in its biosynthesis were assessed in this study. The 2AP is recognized as the key compound for the fragrance of aromatic rice in many previous reports (Buttery, Ling, Juliano, & Turnbaugh, 1983; Magnus, Juliano, & Peter, 2002; Poonlapdhecha et al., 2012). In this study, the 2AP content in grain increased with the improvement of N level and the 2AP content was significantly increased under GABA250 treatment (Figure 1). Previous studies have indicated that salt and shading treatment increased the 2AP and GABA contents in grain (Mo et al., 2015; Poonlapdhecha et al., 2012) and a significant positive correlation between 2AP and GABA in the grains was also observed for the “Yuxiangyouzhan” (Mo et al., 2015). Additionally, nitrogen fertilization has great impacts on 2AP formation and accumulation (Mo et al., 2018; Ren et al., 2017). Therefore, the main reasons that GABA and N regulate the aroma formation are as follows: (a) GABA is directly related to 2AP formation; and (b) the GABA and N regulate the physio-chemical parameters of 2AP formation.

Moreover, compared with GABA0, Na, Mn, Zn, and Fe contents in grains under GABA250 were substantially increased. This was supported by the study of Kinnersley and Lin (2000), who have reported that the Lemna minor plants treated with 10 mM GABA yielded higher levels of Mn and Zn than the untreated plants. The increase in the content of micronutrients was mainly caused by the enhancement of assimilation capacity, which was induced by GABA application. The interaction of GABA and N affected the proline contents in grains significantly (Table 1).

Proline contents in leaves at HS, 15 d AH, and MS were significantly affected by V, N, T × N, and V × N; however, the proline contents in leaves under GABA250 treatment were substantially reduced at MS (Table 2), which suggested the transportation of proline from leaves to grains during grain filling and lead to higher 2AP
TABLE 4  Effect of γ-aminobutyric acid (GABA) and nitrogen on GABA content in leaves in different fragrant rice genotypes (µg/g FW)

| Treatment | HS | 15dAH | MS   |
|-----------|----|-------|------|
| GABA0     |    |       |      |
| YX        |    |       |      |
| ZN        | 107.71 | 86.64 | 113.94 |
| LN        | 128.58 | 94.32 | 91.73  |
| MN        | 107.98 | 94.29 | 84.59  |
| HN        | 119.58 | 91.01 | 116.99 |
|          |    |       |      |
| GABA250   |    |       |      |
| YX        |    |       |      |
| ZN        | 23.34  | 11.17 | 7.21   |
| LN        | 30.99  | 21.23 | 7.26   |
| MN        | 32.81  | 27.22 | 7.22   |
| HN        | 43.60  | 19.38 | 9.19   |
|          |    |       |      |
| BS        |    |       |      |
| ZN        | 112.70 | 94.30 | 88.34  |
| LN        | 118.78 | 116.35| 106.45 |
| MN        | 110.16 | 129.76| 79.58  |
| HN        | 110.25 | 101.03| 107.46 |
| Mean      | 87.21 | 73.89 | 68.33  |
| Analysis of variance |
| T         | 0.22ns | 0.91ns | 0.18ns |
| V         | 745.94**| 127.65**| 274.74**|
| T × V     | 2.87ns | 0.15ns | 2.15ns |
| N         | 0.27ns | 0.87ns | 2.23ns |
| T × N     | 0.92ns | 3.45* | 2.92* |
| V × N     | 1.61ns | 0.50ns | 1.18ns |
| T × V × N | 0.86ns | 1.50ns | 2.19ns |

Note: Means with different lowercase letter(s) differ statistically at p < .05.
Abbreviations: BS, Basmati-385; d AH: days after heading; HS, heading stage; MS: maturity stage; YG: Yugengyou14, YX, Yuxiangyouzhan.
*Significant at p < .05;
**Significant at p < .01; ns: non-significant.

TABLE 5  Effect of γ-aminobutyric acid (GABA) and nitrogen on P5CS activity in leaves in different fragrant rice genotypes (U/g FW)

| Treatment | HS | 15dAH | MS   |
|-----------|----|-------|------|
| GABA0     |    |       |      |
| YX        |    |       |      |
| ZN        | 9.859  | 10.585| 14.537 |
| LN        | 10.166 | 10.493| 13.369 |
| MN        | 7.455  | 14.659| 14.055 |
| HN        | 7.670  | 9.901 | 10.987 |
|          |    |       |      |
| GABA250   |    |       |      |
| YX        |    |       |      |
| ZN        | 6.103  | 6.362 | 7.148  |
| LN        | 6.840  | 6.054 | 6.953  |
| MN        | 5.151  | 4.491 | 7.628  |
| HN        | 6.556  | 6.712 | 6.137  |
| Mean      | 8.637  | 17.195| 16.956 |
| Analysis of variance |
| T         | 0.12ns | 4.10ns | 1.90ns |
| V         | 53.54**| 9.06**| 37.81**|
| T × V     | 4.13*  | 4.24* | 2.29ns |
| N         | 17.75**| 3.64* | 16.09**|
| T × N     | 1.79ns | 2.68ns| 16.42**|
| V × N     | 1.17ns | 3.33* | 11.76**|
| T × V × N | 0.89ns | 3.65**| 19.55**|

Note: Means with different lowercase letter(s) differ statistically at p < .05.
Abbreviations: BS, Basmati-385; d AH: days after heading; HS, heading stage; MS: maturity stage; YG: Yugengyou14; YX, Yuxiangyouzhan.
*Significant at p < .05;
**Significant at p < .01; ns: nonsignificant.
TABLE 6  Effect of γ-aminobutyric acid (GABA) and nitrogen on PDH activity in leaves in different fragrant rice genotypes (U/g FW)

| Treatment | HS  | 15d AH | MS  |
|-----------|-----|--------|-----|
| GABA0     |     |        |     |
| YX        |     |        |     |
| ZN        | 10.199 | 11.685 | 8.765 |
| LN        | 11.310 | 9.203  | 8.204 |
| MN        | 9.132  | 12.794 | 12.783 |
| HN        | 8.364  | 14.100 | 16.545 |
| YG        |     |        |     |
| ZN        | 12.046 | 8.838  | 8.450 |
| LN        | 10.854 | 9.262  | 7.812 |
| MN        | 9.890  | 8.341  | 9.441 |
| HN        | 7.841  | 4.548  | 8.603 |
| BS        |     |        |     |
| ZN        | 12.216 | 12.417 | 8.933 |
| LN        | 18.202 | 12.989 | 7.712 |
| MN        | 11.141 | 15.156 | 9.650 |
| HN        | 16.070 | 11.131 | 8.168 |
| Mean      | 11.278 | 10.872 | 9.589 |

GABA250

| Treatment | HS  | 15d AH | MS  |
|-----------|-----|--------|-----|
| YX        |     |        |     |
| ZN        | 14.581 | 11.497 | 13.909 |
| LN        | 13.485 | 14.091 | 17.909 |
| MN        | 11.184 | 12.768 | 9.088 |
| HN        | 13.766 | 12.085 | 8.281 |
| YG        |     |        |     |
| ZN        | 8.140  | 10.907 | 13.768 |
| LN        | 10.978 | 8.476  | 11.693 |
| MN        | 10.439 | 9.232  | 11.152 |
| HN        | 11.815 | 9.101  | 7.455 |
| BS        |     |        |     |
| ZN        | 15.938 | 18.453 | 17.408 |
| LN        | 12.453 | 15.353 | 22.175 |
| MN        | 13.292 | 16.744 | 9.976 |
| HN        | 13.742 | 18.318 | 9.430 |
| Mean      | 12.484 | 13.086 | 12.687 |

ANOVA

| Factor | Value |
|--------|-------|
| T      | 4.92ns |
| V      | 52.90** |
| T × V  | 15.72** |
| N      | 3.06*  |
| V × N  | 2.47ns |
| T × V × N | 4.43** |

Note: Means with different lowercase letter(s) differ statistically at p < .05.
Abbreviations: BS, Basmati-385; d AH, days after heading; HS, heading stage; MS, maturity stage; YG, Yugengyou14; YX, Yuxiangyouzhan.
*Significant at p < .05;
**Significant at p < .01; ns: nonsignificant.

TABLE 7  Effect of γ-aminobutyric acid (GABA) and nitrogen on OAT activity in leaves in different fragrant rice genotypes (U/g FW)

| Treatment | HS  | 15d AH | MS  |
|-----------|-----|--------|-----|
| GABA0     |     |        |     |
| YX        |     |        |     |
| ZN        | 50.436 | 118.519 | 175.335 |
| LN        | 53.765 | 131.904 | 180.132 |
| MN        | 39.305 | 167.937 | 173.495 |
| HN        | 58.381 | 154.551 | 174.264 |
| YG        |     |        |     |
| ZN        | 80.354 | 77.057  | 88.004 |
| LN        | 77.795 | 85.609  | 98.689 |
| MN        | 72.791 | 76.401  | 176.657 |
| HN        | 68.798 | 65.836  | 187.467 |
| BS        |     |        |     |
| ZN        | 137.599 | 154.256 | 196.877 |
| LN        | 157.291 | 155.757 | 178.268 |
| MN        | 135.698 | 177.890 | 193.295 |
| HN        | 143.348 | 135.035 | 174.143 |
| Mean      | 89.630 | 125.063 | 166.385 |

GABA250

| Treatment | HS  | 15d AH | MS  |
|-----------|-----|--------|-----|
| YX        |     |        |     |
| ZN        | 49.478 | 142.124 | 174.679 |
| LN        | 56.952 | 162.462 | 203.454 |
| MN        | 57.504 | 182.127 | 182.437 |
| HN        | 49.556 | 157.765 | 175.051 |
| YG        |     |        |     |
| ZN        | 93.848 | 121.396 | 112.825 |
| LN        | 85.471 | 77.531  | 108.065 |
| MN        | 85.621 | 76.574  | 93.593 |
| HN        | 87.751 | 78.630  | 90.222 |
| BS        |     |        |     |
| ZN        | 154.083 | 160.150 | 196.942 |
| LN        | 148.063 | 163.815 | 227.015 |
| MN        | 156.887 | 171.943 | 209.212 |
| HN        | 134.361 | 174.965 | 197.217 |
| Mean      | 96.631 | 139.123 | 164.226 |

ANOVA

| Factor | Value |
|--------|-------|
| T      | 6.01ns |
| V      | 376.67** |
| T × V  | 1.23ns |
| N      | 1.26ns |
| T × N  | 2.62ns |
| V × N  | 0.89ns |
| T × V × N | 1.39ns |

Note: Means with different lowercase letter(s) differ statistically at p < .05.
Abbreviations: BS, Basmati-385; d AH, days after heading; HS, heading stage; MS, maturity stage; YG, Yugengyou14; YX, Yuxiangyouzhan.
*Significant at p < .05;
**Significant at p < .01; ns: nonsignificant.
accumulation in grain. So, correlation analysis revealed that the 2AP content in fragrant rice was significantly and positively associated with proline content in leaves at 15 d AH (Table 9). Our results confirm the previous reports in which proline is reported as the precursor of 2AP in fragrant rice, and its higher level often results in more 2AP contents (Huang et al., 2008; Poonlaphdecha et al., 2012). Generally, genetic factors largely determine the aroma formation in fragrant rice (Bradbury et al., 2008; Fitzgerald et al., 2008); nevertheless, many environmental factors and the cultivation practices may have a significant influence on aroma volatiles in rice (Li, Ashraf, et al., 2016; Mo et al., 2015; Yang et al., 2012). Nitrogen fertilizer is also one of the important factors that could lead to the improvement of 2AP accumulation significantly (Ren et al., 2017; Sikdar, Rahman, Islam, Yeasmin, & Akhter, 2008), even there are still different arguments on whether or not N could improve the 2AP content in fragrant rice (Itani, Tamaki, Hayata, Fushimi, & Hashizume, 2004; Li et al., 2014; Yoshihashi, 2005). Moreover, GABA and nitrogen application did not

**TABLE 8** Effect of γ-aminobutyric acid (GABA) and nitrogen on DAO activity in leaves in different fragrant rice genotypes (U/g FW)

| Treatment | HS | 15d AH | MS |
|-----------|----|--------|----|
| GABA0     |    |        |    |
| YX        |    |        |    |
| ZN        | 5.393 | 5.457 | 6.350 |
| LN        | 5.499 | 5.651 | 6.929 |
| MN        | 3.518 | 6.613 | 7.199 |
| HN        | 4.352 | 5.682 | 6.114 |
| YG        |    |        |    |
| ZN        | 3.569 | 2.761 | 5.394 |
| LN        | 3.232 | 3.772 | 5.020 |
| MN        | 3.518 | 3.186 | 4.227 |
| HN        | 3.069 | 2.772 | 3.752 |
| BS        |    |        |    |
| ZN        | 2.843 | 4.553 | 3.985 |
| LN        | 4.376 | 4.386 | 5.866 |
| MN        | 3.169 | 4.062 | 5.775 |
| HN        | 2.555 | 2.981 | 5.039 |
| Mean      | 3.758 | 4.323 | 5.471 b |
| GABA250   |    |        |    |
| YX        |    |        |    |
| ZN        | 4.435 | 5.766 | 6.158 |
| LN        | 3.968 | 7.288 | 9.200 |
| MN        | 3.733 | 6.146 | 8.288 |
| HN        | 3.623 | 6.238 | 7.803 |
| YG        |    |        |    |
| ZN        | 3.897 | 5.059 | 6.453 |
| LN        | 3.858 | 3.656 | 5.150 |
| MN        | 2.446 | 4.375 | 5.009 |
| HN        | 3.491 | 3.039 | 4.780 |
| BS        |    |        |    |
| ZN        | 3.996 | 4.622 | 5.376 |
| LN        | 3.508 | 4.661 | 7.609 |
| MN        | 4.147 | 4.777 | 6.379 |
| HN        | 2.877 | 5.525 | 5.119 |
| Mean      | 3.665 | 5.096 | 6.444 a |
| ANOVA     |    |        |    |
| T         | 0.24ns | 9.78ns | 15.95* |
| V         | 8.59** | 46.90** | 16.82** |
| T × V     | 2.74ns | 0.38ns | 0.17ns |
| N         | 4.92** | 1.32ns | 4.23** |
| T × N     | 0.93ns | 0.49ns | 0.29ns |
| V × N     | 1.58ns | 0.81ns | 2.86* |
| T × V × N | 2.17ns | 2.80* | 0.99ns |

Note: Means with different lowercase letter(s) differ statistically at p < .05.

**TABLE 9** Correlation analyses between 2AP content in grains and the investigated indices under GABA0 and GABA250

| Investigated parameters | GABA0 | GABA250 |
|-------------------------|-------|---------|
| Na content in grains    | −.4746ns | −.2631ns |
| Mn content in grains    | .3313ns | −.0151ns |
| Zn content in grains    | −.4603ns | −.5893* |
| Fe content in grains    | .3417ns | .0134ns |
| Protein content in grains | −.4748ns | −.3527ns |
| Proline content at HS   | −.4481ns | −.6470* |
| Proline content at 15d AH | .9274** | .6353* |
| Proline content at MS   | −.5366ns | −.7836** |
| P5C content at HS       | −.4523ns | −.3187ns |
| P5C content at 15d AH   | −.6096* | −.8059** |
| P5C content at MS       | −.8328** | −.6759* |
| GABA content at HS      | −.8910** | −.9388** |
| GABA content at 15 d AH | −.8627** | −.9112** |
| GABA content at MS      | −.8924** | −.9406** |
| PDH activity at HS      | −.3483ns | −.6681* |
| PDH activity at 15d AH  | −.8127** | −.7318** |
| PDH activity at MS      | −.2231ns | −.3915ns |
| OAT activity at HS      | −.2374ns | −.1241ns |
| OAT activity at 15 d AH | −.8752** | −.9272** |
| OAT activity at MS      | −.3232ns | −.9449** |
| P5CS activity at HS     | −.7635** | −.8134** |
| P5CS activity at 15 d AH | −.4447ns | −.9008** |
| P5CS activity at MS     | −.3725ns | −.6652* |
| DAO activity at HS      | −.3407ns | −.4584ns |
| DAO activity at 15 d AH | −.7044* | −.7094** |
| DAO activity at MS      | −.6377* | −.6116* |

Abbreviation: ns, non-significant.

*Significant at p < .05; **Significant at p < .01.

acc
affect the GABA contents in leaves significantly (Table 4), whereas significant but negative correlations between 2AP content in grain and GABA in leaves were detected (Table 9). Differences in opinions exist among scientists regarding relationships of GABA with 2AP. For example, Poonlaphdecha et al. (2012) revealed that 2AP was correlated with proline content but not with the GABA content in leaves, whereas the positive correlation between 2AP and GABA contents in fragrant rice grains was detected by Mo et al. (2015). Furthermore, GABA treatment did not affect the PSC contents, whereas the cultivars were found statistically significant regarding PSC contents in leaves of all rice cultivars. Significant differences in PSC contents were observed for T × V, N, T × N, V × N, and T × V×N at 15 d AH (Table 3), whereas GABA treatment enhanced the activities of PDH, OAT at 15 d AH, and DAO at maturity. The significant interaction between GABA and N for P5CS, PDH, and OAT activities in leaves at MS was observed. In addition, GABA treatment enhanced the activities of PDH, OAT at 15 d AH, and DAO at maturity. The significant interaction between GABA and N for P5CS, PDH, and OAT activities in leaves at MS was observed (Tables 5–8). Moreover, the 2AP contents in grains at maturity showed significant negative associations with some of the investigated enzyme activity in leaves at some growth stage (Table 9). The difference between this study and other previous study is mainly due to the difference in cultivars, application treatments, the correlation analysis that was between 2AP and the biochemistry parameters in different plant parts. Previous studies reported that grain 2AP contents in fragrant rice are directly associated with the activities of PDH, P5CS, OAT, and DAO (Li, Ashraf, et al., 2016), whereas Ghosh and Roychoudhury (2018) reported that aromatic rice types have higher activities of PDH, P5CS, and OAT and P5C contents than nonaromatic rice types. Furthermore, Mo et al. (2017) also demonstrated that PDH activity and proline content were connected to the 2AP formation and accumulation. Differences in the concentration of 2AP in different plant parts showed their differential abilities to accumulate the 2AP contents (Buttery et al., 1983; Maraval et al., 2010). Micronutrients such as Mn, Zn, and Fe also play important roles in modulation of 2AP contents (Hu, Xu, & Huang, 2001; Huang et al., 2008; Huang, Xiao, & Tang, 2010; Li, Ashraf, et al., 2016; Tang & Wu, 2006). Present study indicated that exogenous GABA application improved the uptake of Na, Mn, Zn, and Fe in all rice cultivars. Overall, GABA application in interaction with N substantially modulated the 2AP contents in grains by affecting the enzyme activities involved in the 2AP formation; however, further studies are needed to better understand the involvement of and/or mechanism of GABA and N to regulate the 2AP biosynthesis and the enzymes involved in the whole process.

5 | CONCLUSION

GABA250 can increase 2AP, Na, Mn, Zn, and Fe contents, but decrease in protein content in grains as compared to GABA0. The GABA250 treatment further enhanced the activities of PDH and OAT at 15 d AH and DAO activity at maturity but reduced the proline contents at maturity. Significant interactive effect of GABA and nitrogen was observed for Mn, Zn, and Fe content in grains, proline content in leaves, P5C content in leaves at 15 d AH, GABA content in leaves at 15 d AH and MS, and P5CS, PDH, and OAT activities in leaves at MS. Overall, GABA treatment improved the 2AP content and nutrient uptake in all rice cultivars, whereas GABA and nitrogen revealed significant interaction effect on nutrient content in grains and some physiological parameters in leaves that involved in 2AP formation.

ACKNOWLEDGMENTS

We acknowledge the funding provided by the National Natural Science Foundation for Young Scientists (31601244) and the Special Fund for National Natural Science Foundation of China (31271646).

CONFLICT OF INTEREST

The authors declare that they have no conflict of interest.

ETHICAL APPROVAL

The study did not involve any human or animal testing.

INFORMED CONSENT

Written informed consent was obtained from all study participants.

ORCID

Umair Ashraf https://orcid.org/0000-0003-0041-7706
Zhaowen Mo https://orcid.org/0000-0002-1887-9389

REFERENCES

Bao, G., Ashraf, U., Wang, C., He, L., Wei, X., Zheng, A., ..., Tang, X. (2018). Molecular basis for increased 2-acetyl-1-pyrroline contents under alternate wetting and drying (AWD) conditions in fragrant rice. Plant Physiology and Biochemistry, 133, 149–157. https://doi.org/10.1016/j.plaphy.2018.10.032
Bates, L. S., Waldren, R. P., & Teare, I. D. (1973). Rapid determination of free proline for water-stress studies. Plant and Soil, 39, 205–207. https://doi.org/10.1007/BF00018060
Batushansky, A., Kirma, M., Grillich, N., Toubiana, D., Pham, P. A., Balbo, I., ..., Fait, A. (2014). Combined transcriptomics and metabolomics of arabidopsis thaliana seedlings exposed to exogenous GABA suggest its role in plants is predominantly metabolic. Molecular Plant, 7, 1065–1068. https://doi.org/10.1093/mp/ssu017
Bradbury, L. M. T., Fitzgerald, T. L., Henry, R. J., Jin, Q. S., & Waters, D. L. E. (2005). The gene for fragrance in rice. Plant Biotechnology Journal, 3, 363–370. https://doi.org/10.1111/j.1467-7652.2005.00131.x
Bradbury, L. M. T., Gillies, S. A., Brushett, D. J., Waters, D. L. E., & Henry, R. J. (2008). Inactivation of an aminoaldehyde dehydrogenase is responsible for fragrance in rice. Plant Molecular Biology, 68, 439–449. https://doi.org/10.1007/s11103-008-9381-x
