Utilization

Biochemical Constituents Related to Odor Generation in Some ICRISAT Pearl Millet Materials

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Development of mousy odor in the meal shortly after grain milling is an important constraint to the wider acceptability and utilization of pearl millet (Pennisetum glaucum (L.) R. Br.). The hydrolytic breakdown of meal lipids (Kaced et al. 1984; Kadlag et al, 1995) and enzymatic degradation of meal phenolics, C-glycosylflavones (Reddy et al. 1986), have been speculated to cause odor generation in the stored meal. Recently, Bangar (1998) has shown that phenolics and peroxidase activity (POD), mainly from the germ fraction of the seed, are responsible for odor generation in pearl millet meal. To study genetic variation and identify low POD types, 29 pearl millet genotypes from the International Crops Research Institute for the Semi-Arid Tropics (ICRISAT) (including several parents of mapping populations) with green, brown, or white seed colors were analyzed for crude fat, fat acidity, total phenolics (AOAC 1990), polyphenol oxidase (PPO), and peroxidase (POD) activities (Kumar and Khan 1982).

The genotypes studied showed a wide variation in these constituents (Table 1). The total lipids ranged between 3.3 to 7.6%, the fat acidity (a measure of lipolytic fat degradation) in 30-day stored meal varied from 206 to 680 mg KOH (100 g meal)⁻¹, and the total phenolics from 228 to 486 mg (100 g)⁻¹. Seed color was not related with total lipids, fat acidity, or content of total phenolics except that IP 18293-which has a purple pericarphad the highest total phenolics content and lowest POD activity. Among the enzymes, POD activity varied markedly (54-332 units g⁻¹ min⁻¹) in different genotypes. The low POD activity genotypes identified are IP 18293, Banner Pop, 863B, and,ICMP 451-.P8. Since the POD activity is related to odor generation, the genotypes with low POD activities may be advantageous to develop new cultivars with improved meal shelf life. As three of the low POD activity lines identified in this study (IP 18293, 863B,

and ICMP 451-P8) have been used as parents of pearl millet mapping populations (Hash and Witcombe 1994; Devos et al. 1995), it should be possible to rapidly map quantitative and quantitative trait loci associated with this trait at relatively low cost. Further, the substantial difference detected between near-isogenic lines ICMP 85410 (E_1E_1) and ICMR 94410 (e_1e_1) (Bidinger et al. in press) suggest that some of the genetic factors controlling this trait may be linked with alleles conferring early flowering.

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| | | Crude | | Total | | |
|-------------|-------------|---------|----------------------|-----------------------------|------------------|-------------------------|
| 0.111 | Seed | fat | Fat | polyphenols | | DDO ³ |
| Cuitivar | color | (%) | acidity ¹ | [mg (100 g) ⁻¹] | POD ² | PPO ³ |
| Raj 171 | Green | 6.12 | 451 | 282 | 136 | 66.4 |
| ICMB 89111 | Green | 5.04 | 680 | 324 | 248 | 52.2 |
| PT 732B | Green | 3.30 | 343 | 330 | 240 | 67.4 |
| CZ-IC 923 | Green | 6.26 | 603 | 312 | 160 | 80.0 |
| GICV 93191 | Green | 5.62 | 451 | 396 | 200 | 77.1 |
| PCB-IC 148 | Green | 5.62 | 343 | 360 | 168 | 83.8 |
| CZ-IC416 | Green | 7.36 | 461 | 258 | 164 | 88.6 |
| 843B | Light green | 4.30 | 559 | 390 | 112 | 69.6 |
| ICMR 94410 | Light green | 4.02 | 304 | 288 | 184 | 88.6 |
| 1CMP 85410 | Light green | 6.01 | 284 | 270 | 332 | 69.6 |
| LGD-1-B-10 | Light green | 6.14 | 412 | 282 | 192 | 91.2 |
| Tlft 238D1 | Light green | 3.54 | 372 | 288 | 112 | 66.4 |
| 863B | Light green | 4.16 | 255 | 300 | 64 | 83.8 |
| ICMV 91773 | Light green | 5.02 | 412 | 270 | 188 | 60.8 |
| CMB 88004 | Dark green | 5.02 | 647 | 246 | 148 | 52.0 |
| AIMP 92901 | Dark green | 5.54 | 363 | 300 | 138 | 60.8 |
| RCB-IC 911 | Dark green | 5.16 | 265 | 228 | 136 | 60.5 |
| ICMS 7703 | Brown | 6.60 | 539 | 228 | 120 | 66.4 |
| CMB 90111 | Brown | 7.60 | 274 | 312 | 116 | 75.2 |
| CMR 356 | Brown | 6.54 | 285 | 396 | 128 | 94.4 |
| CMP 451-P8 | Brown | 5.04 | 412 | 366 | 108 | 90.4 |
| IP 18292 | Brown | 5.78 | 490 | 390 | _ | _ |
| PRLT2/89-33 | Brown | 6.32 | 461 | 396 | 216 | 100.8 |
| CZ-IC 618 | Brown | 5.77 | 274 | 258 | 124 | 89.6 |
| Nokha Local | Brown | 5.51 | 421 | 234 | 120 | 110.4 |
| Barmer Pop | Brown | 5.60 | 401 | 228 | 92 | 100.8 |
| Balu Local | Brown | 5.91 | 578 | 246 | - | — |
| P 18293 | Dark brown | 4.32 | 343 | 486 | 54 | 66.4 |
| CMB 88006 | Light brown | 4.88 | 206 | 312 | 184 | 60.5 |
| Ranges | | 3.3-7.6 | 206-680 | 228-486 | 54-332 | 60.5-110 |

Table 1. Crude fat, fat acidity, total polyphenols, peroxidase (POD), and polyphenol oxidase (PPO) activity in whole-grain meal produced from some pearl millet genotypes with varying seed colors.

1. Mg KOH (100 g meal)⁻¹.

2. Units g⁻¹ min⁻¹.

3. Units g⁻¹ h⁻¹.