

A study of puroindoline b gene involvement in the milling behavior of hard-type common wheats

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1 A study of puroindoline b involvement in the milling behavior of hard type common wheats 2 3 4 P. Lasme^{1,4}, F.-X. Oury², C. Michelet³, J. Abecassis¹, F. Mabille¹, C. Bar 5 6 L'Helgouac'h⁴, V. Lullien-Pellerin¹ 7 8 9 10 1. INRA, UMR 1208, Ingénierie des Agropolymères et Technologies Emergentes, 2 Place 11 Viala, F-34060 Montpellier Cedex 01, France 12 2. INRA, UMR 1095 Génétique Diversité et Ecophysiologie des Céréales, 234 avenue du 13 Brézet, F-63039 Clermont-Ferrand, Cedex 2, France 3. UFS, 17 rue du Louvre, F-75001 Paris, France 14 15 4. ARVALIS-Institut du végétal, Station Expérimentale, F-91720 Boigneville, France 16 17 Corresponding author: Lullien-Pellerin V., UMR Ingénierie des Agropolymères et 18 19 Technologies Emergentes, 2 Place Viala, F-34060 Montpellier Cedex 01, France. Tél: 33 20 (0)4 99 61 31 05, Fax : 33 (0)4 99 61 30 76, e-mail : lullien@supagro.inra.fr. 21 22 List of Abbreviations 23 24 cv., cultivar 25 DArT, Diversity Array Technology 26 EBA, Endosperm Breakage Ability 27 HI, Hardness Index 28 FAR, Farina 29 NIL, Near-Isogenic Line

NIRS, Near Infrared Spectroscopy

- 31 PSI, Particle Size Index
- 32 SKCS, Single Kernel Characterization System
- 33 TFY, Total Flour Yield
- 34 TW, Test Weight
- 35 TKW, Thousand Kernel Weight

- 37 Keywords
- Hardness, near-isogenic lines, vitreousness, endosperm reduction, environmental conditions

ABSTRACT

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Differences in the milling behavior among hard type common wheat (T. aestivum) cultivars are well known by the millers. Among them, the French Soissons cultivar which contains the *Pinb-D1d* allelic form of puroindoline b gene is particularly distinguished for its high milling value. Near-isogenic lines differing by the allelic forms of puroindoline b gene, Pinb-D1d or Pinb-D1b (one of the most frequent alleles found in the European wheat population), were constructed. Grain characteristics obtained after wheat cultivation in distinct environmental conditions were compared between near-isogenic lines and Soissons, as well as their fractionation behavior. Results showed that near-isogenic lines containing the *Pinb-D1d* allele displayed lower values of grain hardness and vitreousness than the corresponding lines, containing the *Pinb-D1b* allelic form, in the same cultivation conditions. Both the genetic background and the environmental conditions appeared to affect the grain texture. Measured SKCS hardness index values of the samples under study were found to be correlated with the vitreousness values. Studies of the milling behavior helped to point out that grain vitreousness is an important factor acting on endosperm breakage ability whatever the genetic background of the wheat. Our results also demonstrated that, at similar levels of vitreousness, the endosperm of Soissons could me more easily reduced than other wheat lines.

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Introduction

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Hardness and vitreousness are both characteristics of the endosperm from common wheat grains that are recognized to affect their mechanical properties (Haddad et al 1999, 2001) and thus play a role in the grain milling behavior (Greffeuille et al 2006, 2007). Indeed, mechanical properties of the endosperm are known to affect the flour particle size distribution, the degree of starch damage and the water absorption capacity of flours. A number of methods have been developed in order to evaluate the mechanical resistance of the endosperm (Turnbull and Rahman 2002). These methods are either based on the size of the particles after grinding (i.e. particle size index (PSI)) or more related to the required crushing force to break the grains as determined with Single Kernel Characterization System (SKCS) developed by Martin et al (1993). PSI is a direct measurement of the percentage of particles under a specific size (75 µm) which can be generated after grain grinding and is used to classify wheats in the two classes soft or hard depending on the obtained PSI values (Williams and Sobering 1986). Near-infrared reflectance spectroscopy (NIRS) is often used instead of PSI (Saurer 1978) to classify wheats based on the distinct scattering at specific wavelengths of ground grains which differ by hardness and is calibrated with grains distinguished by PSI. However, SKCS allows to classify wheats according to the required force for grain breaking and defines four ranges based on the kernel texture characteristics (Gaines et al 1996). A hardness index (HI) is calculated to express the average crushing resistance of the grain sample. The endosperm hardness is found to be mainly genetically controlled by the Hardness (Ha) locus on the short arm of Chromosome 5D (Chantret et al 2005) and particularly linked to the expression of wild-type or variant allelic forms of two genes encoding specific proteins called puroindolines A and B (PinA and PinB) which are found to be located at the starch granule and protein network interface (Feiz et al 2009). These proteins are mainly found in the starchy

endosperm (Wiley et al 2007), but some authors have also found one of them (PinB) in the aleurone layer (Caparelli et al 2005; Dubreil et al 1998). However, Lesage et al (2011) studying their expression during grain development confirmed their endosperm location. Indeed expression of the wild alleles of these genes (Pina-D1a, Pinb-D1a) leads to a soft mechanical behavior of the grain endosperm whereas absence or mutations in one of the genes results in a hard phenotype (Giroux and Morris 1998; Lillemo and Morris 2000; Beecher et al 2002; Wanjugi et al 2007). Several possible mutations in the puroindoline genes have been identified and recently reviewed by Morris and Bhave (2008). But the most frequent in Europe corresponds to a point mutation within the gene encoding PinB (Pinb-D1b) resulting to a Gly to Ser substitution in position 46 of the corresponding protein sequence (Huang and Röder 2005; Lillemo and Morris 2000). Nevertheless other QTL for hardness have also been pointed out as reviewed in Turnbull and Rahman (2002) and Bhave and Morris (2008). It is also recognized that some variation of the hardness level and thus of the milling performance could be due to environmental factors. Indeed temperature and light intensity during grain filling and the rate of grain drying (Parish and Halse 1968) were pointed out to affect other characteristics of the endosperm texture known as vitreousness. Differences in vitreousness could be determined visually by the level of translucence or using new developed methods, i.e. a machine vision (Symons et al 2003) or dual energy X-ray imaging (Neethirajan et al 2007), as demonstrated with durum wheats. Relationships between the Ha locus and vitreousness have been suggested previously (Nelson et al 1995) but were not confirmed by other authors (Weightman et al 2008). A number of mutations in puroindoline genes were found to affect the milling behavior as well as the flour properties (Chen et al 2007; Greffeuille et al 2006; Hogg et al 2005; Ma et al 2009; Martin et al 2001), none of them examined the respective contribution of the genetic versus the environmental factors on wheat grain fractionation behavior.

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In this paper, near-isogenic lines (NILs) containing either the most frequent allele of puroindoline b gene in cultivated French wheats, *Pinb-D1b*, (found for example in Caphorn cultivar (cv.)), or containing the type d allele of puroindoline b gene, *Pinb-D1d*, (found in Soissons cv. which is well known for its high flour producing level), were grown in different environments in order to study the respective effect of puroindoline genotype, site and nitrogen fertilization on the endosperm ability for size reduction in relation to the endosperm texture.

Materials and Methods

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Wheat material and field conditions

Three pairs of near-isogenic lines (NIL1 to NIL3) produced by UFS (Fr.) and differing by the allelic forms (b or d) of *Pinb-D1* gene were used in this study. The near-isogenic lines (NILs) were obtained from two different crosses and after selection of the two allelic forms at the F6:F7 step (F7 siblings drawn from the same F6 parent plant). Genetic similarity within NILs was confirmed using diversity array technology (DArT) markers (Akbari et al 2006) generated by Triticarte Pty.Ltd (www.triticarte.com.au). The wheat cultivar Soissons was used as a reference sample. All the genotypes were grown in France in four different locations: Cappelle (50°29' N/3°10' E), Clermont-Ferrand (45°47' N/3°8' E), Maule (48°54' N/1°51' E) and Orgeval (48°55' N/1°58' E) and harvested in summer 2008. In each location, two types of nitrogen fertilization were used, the first one (N condition) corresponded to intensive farming with nitrogen fertilization around 300 kg/ha adjusted to high yield objectives (around 9t/ha depending on the location), and taking into account the residual nitrogen remaining in soil at early spring. In the second one (N+ condition), an additional nitrogen dose (50 kg/ha) was added at flowering. Grain from these different samples (48 samples from near-isogenic lines and 8 samples from Soissons) were collected and cleaned to remove broken kernels or impurities and stored at 4°C before milling.

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Measurement of physical and biochemical characteristics of the grain

- 141 Test weight (TW) and thousand-kernel weight (TKW) were measured according to the AACC
- 142 Method 55-10 (AACC 2000) and the method NF EN ISO 520 (Janv. 2011,

- 143 www.boutique.afnor.org), and expressed in kg/hl or in g respectively. NIRS Hardness was
- evaluated using a Percon NIRS apparatus (AACC method 39-70A, AACC 2000)
- Grain vitreousness was assessed by analysis of kernel cross sections obtained using a Pohl
- grain cutter (Versuchs and Lehranstalt, Brauerei, Berlin, Germany). Depending on the
- percentage of vitreous surface in the analyzed grains, they were classed into 5 groups (i.e.
- grains which displayed less than 25 % of vitreousness were classed in the first group, those
- with around 25 % of vitreousness constituted the second group, those with around 50%, 75 %
- or 100 % of vitreousnesswere classed in the third, four and fifth group respectively). The
- number of grains in each class (N1, number of grains in class 1 for example) was multiplied
- by the corresponding factor of vitreousness in the considered class and the percentage of
- vitreousness in the grain sample was calculated using the following equation.
- 154 Percentage of vitreousness = (N1*0.00 + N2*0.25 + N3*0.50 + N4*0.75 +
- N5*1)/(N1+N2+N3+N4+N5), where the sum of the analyzed grains in each class, N1 to N5,
- 156 was around 500.
- 157 A Perten single-kernel characterization system (SKCS) 4100 (Perten Instruments North
- America INC, Springfield, IL) was used to collect data on 100 kernels and to calculate a mean
- hardness index (HI) using AACC method 55-31 (AACC 2000). This method also allows to
- determine grain weight which was found to be highly correlated with TKW measured using
- 161 the ISO method (r=0.84).
- 162 Grain moisture and grain protein content were determined according to AACC methods 44-16
- and 39-35, respectively and expressed in percentage (AACC 2000).

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Milling behavior characterization

- Five hundred grams of cleaned grain was conditioned to 16 % moisture and milled after 24 h
- at room temperature using a patented prototype (French Patent Chopin, INRA, ARVALIS, N°

0905572, 02/20/2009) developed in order to evaluate the wheat milling value. This milling device is equipped with two break rolls, one sizing and one reduction roll. This milling process thus allows the production of two break flours, one sizing and one reduction (in two steps) flours as well as a coarse bran, a fine bran and a short fraction. Sieves of 200 µm mesh were used to isolate the break and sizing flours whereas a 160 µm mesh was used to isolate the reduction flour. Each collected fraction was weighed at each step and expressed either as a percentage of the total grain weight or percentage of the charged product weight at each step. Incomplete milling was also undertaken to analyze more specifically behavior of the inner endosperm part. In this case, coarse endosperm particles, also named farina as defined in International Association of Cereal Science and Technology ICC dictionary (http://old.icc.or.at/cgi-bin/dic/anetdic.cgi?lang=fr) obtained after the first break rolls (FAR1) were recovered and directly reduced at the sizing step, skipping the second break roll. A mass ratio between flour produced at the sizing step from coarse endosperm particles obtained at the first break roll (FAR1 in incomplete milling) and FAR1 was determined (= EBA1) to more specifically study the endosperm behavior in size reduction. Similarly the ratio between sizing flour produced during milling from coarse endosperm particles obtained after the two breaking steps (FAR2) and FAR2 was determined to study the endosperm ability to break (=EBA2).

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Particle size distribution analysis

- Particle size distribution of farina or resulting flour was analyzed with a laser beam particle
- size analyzer (Coulter Co, Miami, FL) using ethanol as a solvent and the sonication mode.
- 190 Distribution was expressed as the percentage in volume according to the particle size.

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Statistical analyses

Analysis of variance, comparisons of means using Tukey's test (confidence interval 90%) and simple Pearson's correlations were performed using XLStat software (2008 version 7.03, Addinsoft, Paris, Fr.).

Results and discussion

NILs construction and grain characterization

In order to test the impact of two of the most frequent alleles (*b* and *d*) of *Pinb-D1* found in French cultivated wheats and located on chromosome 5DS, near-isogenic lines were produced from two different crosses. Three pairs of lines (NIL1 to NIL3) were retained and analyzed for genetic similarity using diversity array technology (DArT) markers. As expected, similarity appeared higher between NIL2 and NIL3 which were derived from the same cross (14.1% differences revealed using the DArT markers), whereas NIL1, coming from a different cross appeared more distant (26.1 and 27.3% differences with NIL2 and NIL3, respectively according to DArT markers). Within NIL1, NIL2 and NIL3, differences between the 2 genotypes, which either present the *Pinb-D1b* allelic form where Gly46 is replaced by Ser (Lillemo and Morris 2000) or the *Pinb-D1d* allelic form where Trp44 is replaced by Arg (Day et al 2006), were equal to 1.4, 0.7 and 0.9%, respectively, according to DArT markers, and thus lower than the theoretical level (=3.12%) of residual heterozygosity at this selection step.

Table 1

All of the samples were grown in four distinct locations and with two distinct levels of nitrogen fertilization in order to increase the variability of conditions under which grains were produced, thus increasing the probability of obtaining distinct grain texture. Mean values and standard errors of main grain characteristics are summarized in Table 1 for each NIL and were compared with cv. Soissons grown in similar conditions. TKW and TW mean values were found to be around 45 g and 79.5 kg/hl for NIL1 and NIL2, respectively whereas NIL3

exhibited higher values for TKW (49.5 g) and slightly higher values (80 kg/hl) for TW. No differences were observed taking into account both of the allelic forms. But very low variability (less than 10%) was observed between samples for these two characteristics (especially for TW), as also found for Soissons, which displays lower TKW but TW values similar to those of NIL1 and NIL2. However, NIRS and SKCS hardness, as well as vitreousness, were found to display high variability. Therefore, the experimental plan appeared to be relevant to study the effect of grain texture on wheat grain milling behavior. As expected due to the puroindoline genotype of the considered samples, NIRS hardness values correspond to hard wheat type. Hardness and vitreousness of NIL samples were generally found to display higher values compared to Soissons samples grown in similar conditions. Furthermore, NIL1 appeared to display the highest SKCS hardness as well as the highest level of vitreousness in comparison with the other near-isogenic lines. Mean values for hardness and vitreousness appeared to be significantly higher in genotypes displaying *Pinb-D1b* compared with those with *Pinb-D1d* (except for NIL3), despite similar protein content in grains. Thus, the effect on grain characteristics of the background genotype, or *Pinb-D1* allele (b or

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Table 2

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TKW was found to be mainly affected by the genetic background whereas TW was mostly affected by the growth location even if both genetics and the environment were found to act on these grain characteristics. Interestingly NIRS hardness was found to depend mainly on *Pinb-D1* allele and was also affected by the combination of location and nitrogen amount. Indeed NIRS hardness was already known to depend on the *Pinb-D1* allele when lines

d), as well as of the growth location and nitrogen level, were further analyzed (Table 2).

displaying a wild type form (Pinb-D1a), corresponding to soft phenotype, were compared to lines containing a mutated form corresponding to hard phenotype (Morris and Massa 2003). But, the effect of the *Pinb-D1* allele among the hard wheat types is unexpected as well as the variability of its effect according to the growth conditions. In comparison with NIRS hardness, the measured SKCS hardness index was found to be mainly influenced by the growth location and genetic background. Similar influence of growth location and genetic background was also observed for vitreousness. Therefore for these set of genotypes, the SKCS hardness index seems to take into account the vitreousness differences between samples better than NIRS hardness measurements. This could be due to the differences between NIRS and SKCS measurements. Indeed SKCS HI value determines the required force for grain breaking and helps to define four ranges based on the kernel texture characteristics (Gaines et al 1996), which depend both on the genetic background and the growth conditions, whereas NIRS hardness is more related to the size of the products obtained after grinding. Therefore relationships between these grain characteristics appeared important to analyze. Furthermore, protein content of the samples under study did not appear to be correlated with the puroindoline allele but to depend on the NIL type. As expected, protein content was also found to depend on both the growth location and the nitrogen level. Indeed, Daniel and Triboi (2000) have already demonstrated the effect on nitrogen addition and temperature conditions on grain protein content. The relationships between grain characteristics were therefore studied using Pearson's test. The highest correlation between the analyzed grain characteristics was found between SKCS hardness index and vitreousness (r = 0.92), as well as with the protein content (r = 0.84), whereas NIRS hardness was significantly correlated with these grain parameters but at a lower level (r around 0.4). Morris and Massa (2003) also compared different texture measurements (i.e. NIR hardness, SKCS and PSI) and already pointed out the lower

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relationship between NIR hardness determination and SKCS HI within a hardness class in comparison with the relationship established when cultivars from both hard and soft classes were taken into account. As SKCS measurement is more related to compression tests and produces a four-class frequency distribution of hardness data for each sample, this hardness index value more clearly reflect the intrinsic sample variability as already pointed out by Morris et al (2008) similarly to the vitreousness measurement. The narrower genetic background of the studied population could also explain these correlations. However relationships between SKCS HI and vitreousness were already reported in common wheat (r=0.7; Orucevic et al 2008) or durum wheat $(R^2 = 0.99; Sissons et al 1999)$. In barley, relationships between SKCS and light transflectance measurement as well as visual assessment of mealiness were also pointed out (Gamlath et al 2005; Psota et al 2008). TW was also found to be significantly and positively correlated with vitreousness and hardness either measured with NIRS or SKCS, as well as with the protein content. By contrast, TKW was negatively correlated with vitreousness and SKCS HI and protein content. Indeed an increase in the protein level appears to be correlated with an increase of vitreousness which affects the grain density, as already reported in durum wheat (Samson et al 2005), but probably occurs inversely to the grain size. Similar negative correlation between TKW and SKCS hardness index was also pointed out by Gazza et al (2008) studying wheat grains grown in distinct locations.

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Endosperm breakage ability (EBA) and grain milling behavior

All of the grain samples were milled in order to study their behavior depending on the grain nature and characteristics. The effect of the endosperm structure on size reduction by grinding was particularly examined. Therefore, the main values measured were: percentage of coarse endosperm particles, known as farina, produced at the first break roll (FAR1) or after two

breaking steps (FAR2), the ratio between flour produced from farina and corresponding farina obtained either at the first break roll (EBA1) or after two break rolls (EBA2), and the total flour yield (TFY). A variance analysis (Table 3) was performed in order to study the effect of the genetic background, *Pinb-D1* allele, and environmental conditions on the retained parameters.

Table 3

Production of farina as well as its size reduction was found to be mainly related to the wheat growth location and genetic background. The effect of the *Pinb-D1* allele was also found to be significant but at a lower level than the other parameters. Indeed endosperm from wheat grains carrying the *Pin-b-D1d* allele showed a slightly higher ability to break compared with grains carrying the *Pinb-D1b* allele. The effect of nitrogen addition in soil was also found to be significantly related to the farina production and to the endosperm ability to produce flour at the first break (EBA1). TFY was also related to the wheat growth location and the genetic background. But in this case the genetic background appears as the predominant factor.

parameters and the grain characteristics were analyzed (Table 4).

Table 4

Clear positive relationships between the ability of coarse endosperm particles to produce flour at the sizing step and TFY were found. Indeed EBA1 could help to explain 50 % of the TFY which is in accordance with previous studies (Greffeuille et al 2007). As expected however, production of farina which is inversely proportional to the flour production at breaking stage

appears to be negatively correlated to the TFY. Analysis of relationships between milling behavior and grain characteristics clearly pinpointed the influence of vitreousness on the production and size reduction of farina. Indeed an increase of grain vitreousness appears to increase the farina yield and negatively affects its breakage ability, leading therefore to flour yield reduction. Therefore as location and genetic background were found as the main factors contributing to vitreousness (Table 2), they were also found to be significantly related to the size reduction ability of farina (Table 3). As SKCS hardness index and the protein content were found to be highly related to vitreousness, it was not surprising that similar correlations could be observed between the milling analyzed parameters and these grain characteristics. Results also showed more significant correlations at the first breaking step where farina originated mainly from the inner part of the endosperm (EBA1). If Pinb-D1 allele was found significantly related to the milling grain behavior as measured by the farina and flour yields and the farina breakage ability, its influence appears less than the site of cultivation and the genetic background. As this influence of puroindoline allele could simply be due to differences in vitreousness (Tables 1 and 2) between grains carrying either Pinb-D1b or Pinb-D1d, deeper analysis between EBA1 and vitreousness was carried out on the overall set of samples and compared with the cv. Soissons (carrying Pinb-D1d) grown in the same conditions (Fig. 1).

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Figure 1

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Results confirmed the existence of a negative linear relationship between the endosperm ability for size reduction and grain vitreousness whatever the sample analyzed. It also pointed out the major effect of the genetic background differences in comparison with the carried puroindoline allele. A Fisher test enabling the pair-wise comparison of the linear regressions

(Tomassone et al. 1983) showed that they were significantly different (p<0.001), except the regressions for NIL2 and NIL3, which indeed derived from the same parental cross. Furthermore, coarse endosperm particles (farina) from the first breaking step (FAR1) from cv. Soissons grains were found to display a higher ability for size reduction (expressed by EBA1) than the near-isogenic lines even if grains of similar vitreousness were compared. Similar conclusions were also obtained if vitreousness values were replaced by the grain protein content or SKCS hardness index (data not shown).

Study of the particle size before and after farina reduction

In order to further study the differences in farina ability for size reduction among the examined wheat grain samples, size distribution profiles of farina from the first breaking step (FAR1) were analyzed before and after sizing reduction. For this study, only samples grown under a standard nitrogen amount on the sites leading to the most contrasted grain vitreousness were retained. These locations were Cappelle, where samples displayed vitreousness between 20 and 45%, and Maule where grain vitreousness was between 50 and 70%. Typical curves of particle size distribution from farina before and after reduction are shown in Fig. 2.

Figure 2

Before reduction, size of the farina particles were between 300 and 2000 µm and centered around 800 µm as illustrated in Fig. 2A with NIL1-*Pinb-D1b*. However comparison of the mean size of farina particles clearly pointed out differences between the two locations (Table 5). Indeed, the location which was shown to lead to the most vitreous samples (Maule) was found to correspond to the largest farina particles. Significant differences between mean

particle size of farina from the distinct near-isogenic lines was also observed, NIL3 displaying the lowest mean particle size. No distinction between lines with the different alleles of Pinb-D1 was pointed out even if potential interactions between genetic background and Pinb-D1 was revealed using variance analysis (p<0.05).

Table 5

- After farina breakage, distribution of the particle size was enlarged and showed a complex profile between 5 and 2000 µm (Fig. 2A). Particles from the most vitreous grains (grown in Maule location) appeared less reduced compared to the other grain samples whatever the genotype, however, main differences between curves were difficult to pinpoint. Therefore, a mathematical model was further used in order to fit the data of particle size distribution and determined their main characteristics for each sample. The experimental curve between volume according to the size of the particles displayed four discrete peaks, the model was thus considered as the sum of four Gaussian-like shaped functions, $F(x) = \sum_{i=1}^{i=4} F_i(x)$ where F_1 , F_2 F_3 , F_4 correspond to large, medium-size, small and finest particles, respectively. Each function was expressed with the following equation:
- $F_i(x) = D_i \cdot \exp^{-\frac{1}{2} \left(\frac{\log(x) \log(\overline{m}_i)}{\log(\sigma_i)} \right)^2},$

where D_i corresponds to the peak height, x to the particle size, \overline{m}_i to the mean particle size of the sub-population *i.e.* the particle size at the peak of each Gaussian-like curve, σ_i the deviation around the mean particle size. Values for D_i , \overline{m}_i , were first estimated from the experimental curve and σ_i first fixed to 0.5. Then, these parameters were adjusted by minimization of the differences between the area under the experimental data curve $\Phi(x)$ and the mathematical model using the solver function of Excel (Microsoft Office 2007). The

following function, $\min \left[\sum_{x \in R} |F(x) - \Phi(x)| \right]$, where R was the set of observed particle size values, was used to minimize the differences between data and the model curve for large and medium-size particle peaks whereas $\min \left[\sum_{x \in P} |F(e_x) - \Phi(e_x)| \right]$ was used to minimize differences due to the small and the finest particles as it takes into account the number of the particles instead of the volume and was thus more adapted to these type of particles. This model was found to fit correctly the data curves as illustrated in Fig. 2B, and to help the determination of \overline{m} , σ and D for each type of particle. Indeed, root mean squared errors were found between 0.09 and 0.14 for curves corresponding to particles of farina before breakage and between 0.02 and 0.08 for curves corresponding to particles after farina breakage by the sizing step. Table 5 summarizes the results obtained for mean particle size of the farina particles in each peak obtained after farina breakage. It also reports the percentage of the total particle volume in each peak. Analyses of the volume of particles distributed in the four peaks deduced from the curve of each sample after farina size reduction revealed that around 20 to 28% of the material was not reduced with the equipment used. As no differences were observed between samples, this could correspond to the grinding limit of the reduction cylinder. Sixty to seventy percent of the volume of particles were recovered in the peak of medium-size particles after the sizing step. The other peaks, corresponding to small and finest particles, represented 6-8 % and 1.5-3 %, respectively. Comparison of the mean size of particles in the major peak (medium-size particles) from distinct samples revealed a larger size of particles obtained from grains grown in the location (Maule) where grains displayed a higher vitreousness. Moreover, significant differences between NIL were also pointed out. Medium-size particles of NIL 2 were found to display the lower mean value (182.5 μm) compared to NIL 3 (190.8 μm) and NIL1 (222.3 µm) which was in accordance with the greatest ability of NIL 2 to be reduced, as expressed by the endosperm breakage ability (EBA1) values (%) shown in Fig. 1. Indeed in

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the prototype used for milling, separation of flour particles from farina occurred on a sieve of 200 µm mesh, therefore differences of the EBA1 values could be due to differences of the mean medium-particle size obtained after size reduction of farina from the distinct NIL or from samples grown at distinct locations. Thus, if differences in location leading to distinct level of vitreousness mainly impact on endosperm breakage ability as measured by EBA1, the effect of genetic background also appeared as an important factor to consider. Indeed comparison of the mean medium-particle size values obtained from NILs with those obtained after farina reduction from Soissons cv. clearly pointed out the lower particle size value from this cultivar (mean size = 167.3) which is in accordance with the highest ability of the endosperm from these grains to be reduced as measured by the ratio of flour to the farina production (EBA1 in Fig. 1).

Conclusions

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434 The growing location was found to affect the grain vitreousness level and differences in 435 vitreousness were found depending on the Pinb-D1 allele of the near-isogenic line (grains 436 from lines containing the *Pinb-D1d* allele were found to be less vitreous). 437 The increase in vitreousness of samples with a similar genetic background was found to 438 deeply affect the endosperm ability for size reduction and thus the efficiency of flour 439 production from farina. Therefore in order to select the sample displaying the best ability for 440 flour production among a set of grain samples from the same hard wheat cultivar, millers have 441 to choose the one with the lowest vitreous grains. The potential of SKCS hardness index 442 measurement to evaluate the wheat grain vitreousness within hard wheats has also been 443 highlighted. Furthermore, differences in endosperm breakage ability appeared to be mainly 444 due to the genetic background rather than to the Pinb-D1 allele and the higher endosperm 445 ability for size reduction from Soissons grains was pointed out. The size of the medium-size 446 particles obtained from farina size reduction appears as a critical parameter to relate with this 447 better endosperm size reduction ability and higher flour yield. 448

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		TKW	TW	NIRS	SKCS-	Vitreousness	Protein
NILs/Cultivar		(g)	(kg/hl)	hardness	HI	(%)	content (%)
NIL1							
Pina-D1a/Pinb-D1b	mean	45.3 a	79.9 ab	82.4 b	61.7 d	62.0 d	13.4 c
	SD	1.8	0.8	3.8	10.1	15.5	1.1
Pina-D1a/Pinb-D1d	mean	44.9 a	79.4 a	73.3 a	55.6 c	54.3 c	13.0 c
	SD	2.7	1.1	7.2	10.4	15.8	0.9
NIL2							
Pina-D1a/Pinb-D1b	mean	45.8 a	79.6 ab	80.3 ab	47.7 ab	44.4 b	11.6 a
	SD	2.3	1.1	7.7	7.4	16.1	0.7
Pina-D1a/Pinb-D1d	mean	45.4 a	79.4 a	73.1 a	45.4 a	40.6 a	11.4 a
	SD	3.0	0.7	10.9	10.0	16.8	1.0
NIL3							
Pina-D1a/Pinb-D1b	mean	49.9 b	80.4 b	83.8 b	49.9 b	53.3 с	12.1 b
	SD	2.3	1.1	9.8	12.6	20.5	1.3
Pina-D1a/Pinb-D1d	mean	49.3 b	80.1 ab	77.1 ab	50.0 b	51.5 c	12.1 b
	SD	3.9	1.1	11.8	12.2	20.5	1.5
Soissons							
Pina-D1a/Pinb-D1d	mean	42.9	79.3	64.9	41.4	42.1	12.4
	SD	2.9	1.3	6.0	9.1	14.4	1.2

Table 1: Mean values and standard deviations (SD) of main grain characteristics from three pairs of near isogenic lines (NIL1 to NIL3) differing by *Pinb-D1* allele and the French cultivar *Soissons*. Each distinct sample was grown in four locations in 2008, with two levels of nitrogen fertilization. Each line corresponds therefore to eight distinct grain samples. Identical letters indicate homogeneous groups provided by Tukey's test. TKW = thousand-kernel weight; TW = test weight; NIRS = near-infrared reflectance spectroscopy; and SKCS HI = single-kernel characterization system hardness index.

		TKW	TW	NIRS	SKCS-	Vitreousness	Protein
Studied factor	df			hardness	HI		content
Genetic background							
(NIL)	2	96.4***	2.7**	ns	623.1***	999.3***	12.0***
Pinb-D1 allele	1	ns	ns	697.7***	92.7**	233***	ns
Location	3	65.3***	9.1***	295.0***	1320.8***	3962.1***	10.4***
Nitrogen	1	ns	ns	275.6**	73.8**	413.7***	8.3***
NIL* Pinb	2	ns	ns	ns	38.0*	35.4**	ns
NIL* Location	6	8.1**	ns	ns	54.0***	77.0***	0.9***
Location* Nitrogen	3	ns	ns	334.4***	27.9*	60.5***	0.7**
Model fitting R ²		0.92	0.83	0.84	0.97	0.99	0.96

Table 2: Effect of genotype, *Pinb-D1* allele (*b* or *d*), growth location and nitrogen level on grain characteristics of the three pairs of near-isogenic lines. *, **, *** indicate significance at the 0.05, 0.01, 0.001 probability levels, respectively, ns nonsignificant effect. Values are mean squares. Only significant interactions are pointed out.

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Studied factor	df	FAR1	FAR2	EBA1	EBA2	TFY
Genetic background (NIL)	2	8.4***		211.6***	98.1***	7.2***
Pinb-D1 allele	1	2.4*	4.2*	35.9***	20.9*	0.9*
Location	3			229.8***	/	3.2***
Nitrogen NIL*Pinb	I	20.5***	43.4***	9.1*** 2.8*	ns ns	ns 0.7**
NIL* Location	2	1.8***	5.1*	4.0***	ns	0.5**
Pinb* Location	6	1.3*	ns	1.7*	ns	ns
NIL * Nitrogen	2	ns	3.4*	ns	ns	ns
Location* Nitrogen	3	2.8***	ns	ns	ns	ns
Model fitting R ²		0.96	0.92	0.99	0.90	0.92

Table 3: Effect of genotype, *Pinb-D1* allele (*b* or *d*), growth location and nitrogen level on grain milling characteristics of the three near-isogenic lines. FAR1 = percentage of coarse endosperm particles (farina) obtained after the first break roll; FAR2 = percentage of farina obtained after two breaking steps; EBA1 = endosperm breakage ability (the ratio between flour produced from farina and the corresponding source farina) of FAR1; EBA2 = endosperm breakage ability of FAR2; TFY = total flour yield; NIL = near-isogenic line; df = degree of freedom; *, **, and *** indicate significance at the 0.05, 0.01, and 0.001 probability levels, respectively; and ns = nonsignificant effect. Values are mean squares. Only significant interactions are pointed out.

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	FAR1	FAR2	EBA1	EBA2	TFY
Variables					
TKW (g)	-0.33	-0.19	0.34	0.20	0.60
TW (kg/hl)	0.60	0.64	-0.55	-0.57	-0.03
NIRS hardness	0.50	0.47	-0.43	-0.41	-0.14
SKCS-HI	0.80	0.79	-0.92	-0.82	-0.73
Vitreousness (%)	0.90	0.86	-0.91	-0.79	-0.59
Protein content					
(%)	0.88	0.78	-0.85	-0.68	-0.63
FAR1					
(%)					
FAR2					
(%)	0.85				
EBA1	-0.79	-0.76			
EBA2	-0.61	-0.69	0.86		
TFY					
(%)	-0.49	-0.42	0.71	0.59	

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Table 4: Relationships between grain characteristics and milling behaviour assessed with Pearson's correlations. Significant correlations with $P < 0.0\overline{1}$ are marked in bold. Correlations are only presented once in the table. Reported values are r. FAR1 = percentage of coarse endosperm particles (farina) obtained after the first break roll; FAR2 = percentage of farina obtained after two breaking steps; EBA1 = endosperm breakage ability (the ratio between flour produced from farina and the corresponding source farina) of FAR1; EBA2 = endosperm breakage ability of FAR2; TFY = total flour yield; TKW = thousand-kernel weight; TW = test weight; NIRS = near-infrared reflectance spectroscopy; and SKCS

HI = single-kernel characterization system hardness.

			FAR1	hefore	Particles from FAR1 reduction after sizing rolls							
			FAR1 before reduction		Large p	Large particles Medium- size particles		Small particles		Finest particles		
		Grain	Particle	Particle	Particle	Particle	Particle	Particle	Particle	Particle	Particle	Particle
		vitreousness	size	volume	size	volume	size	volume	size	volume	size	volume
			(µm)	(%)	(µm)	(%)	(µm)	(%)	(µm)	(%)	(µm)	(%)
Genetic												
background												
NIL1 (n=4)	Mean	51.9 ^b	836.8 ^b	98.8 ^a	909.9 ^a	20.7 ^a	222.3 ^c	70.1 ^a	30.2 ^a	8.0 ^b	4.7 ^a	1.5 ^a
	SD	20.7	17.5	1.7	43.6	3.9	29.7	3.8	5.9	1.6	0.4	0.2
NIL2 (n=4)	Mean	43.9 ^a	859.9 ^c	97.8 ^a	942.6 ^a	23.1 ^a	182.5 ^a	69.3 ^a	26.2 ^a	6.0 ^a	5.2 ^a	2.3 ^b
	SD	22.4	16.9	1.6	13.4	3.3	37.7	2.8	5.4	1.3	0.3	0.5
NIL3 (n=4)	Mean	49.2 ^b	812.7 ^a	96.6 ^a	870.4 ^a	26.4 ^a	190.8 ^b	64.8 ^a	25.8 ^a	6.7 ^{ab}	4.8 ^a	2.1 ^{ab}
	SD	30.7	32.5	1.0	36.2	5.2	33.6	5.0	5.0	2.0	0.3	0.4
Soissons (n=2)	Mean	41.8	809.6	96.5	906.0	28.1	167.3	62.1	24.0	6.8	5.2	3.0
	SD	24.9	24.1	0.9	80.8	4.9	30.4	4.7	3.0	0.9	0.7	0.7
Location												
Cappelle (n=6)	Mean	27.1 ^A	821.7 ^A	96.9 ^A	921.4 ^A	26.1 ^A	166.6 ^A	66.2 ^A	23.2 ^A	5.9 ^A	4.8 ^A	2.2 ^B
20 % < v <45 %	SD	5.9	31.5	1.6	23.2	4.1	22.3	4.0	2.4	1.3	0.3	0.6
Maule (n=6)	Mean	69.5 ^B	851.2 ^B	98.6 ^A	893.8 ^A	20.7 ^A	227.4 ^B	69.9 ^A	31.6 ^B	7.8 ^B	5.0 ^A	1.7 ^A
50 % < v <70 %	SD	6.2	19.2	1.2	55.9	3.4	16.2	4.2	3.9	1.6	0.4	0.2

Table 5: Mean values and standard deviations (SD) of particle size (μm) and percent volume of particles from farina (FAR1) before and after reduction obtained after milling of the different grain samples (NIL1 to NIL3) differing by *Pinb-D1* allele and the French cultivar *Soissons* grown in two distinct locations leading to contrasted vitreousness (v). Identical letters indicate homogeneous group provided by Tukey's test applied separately for the two factors, NIL = near isogenic line.

658 Figure Captions

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- Figure 1: Correlation between grain vitreousness and endosperm size reduction ability at first
- break (EBA1). Data are from Soissons which carries Pinb-D1d (*); NIL1 carrying either
- 662 Pinb-D1b (□) or Pinb-D1d (■); NIL2 carrying either Pinb-D1b (○), or Pinb-D1d (●); NIL3
- carrying either *Pinb-D1b* (\triangle), or *Pinb-D1d* (\triangle). Equations of regression lines for the
- different genotypes are, respectively:
- Soissons: y=-0.20x + 49.4; $r^2=0.90$
- NIL1: Pinb-D1b: y=-0.21x + 41.1; r^2 =0.96; Pinb-D1d: y=-0.24x + 43.6; r^2 =0.95
- NIL2: Pinb-D1b: y=-0.19x + 44.6; r^2 =0.83; Pinb-D1d: y=-0.26x + 47.7; r^2 =0.76
- NIL3: Pinb-D1b: y=-0.22x + 44.7; $r^2=0.94$; Pinb-D1d: y=-0.21x + 45.0; $r^2=0.90$.

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- 670 Figure 2: Percentage of volumes of farina particles according to their size before (solid
- 671 symbols) and after sizing step (open symbols). Experimental data (A) from NIL1-Pinb-D1b
- grown in two distinct locations (without nitrogen addition) which results in vitreous (squares)
- or non vitreous endosperm (diamond-shaped). Example of modeling curves (B) for the
- of Vitreous sample of NIL1-Pinb-D1b (squares). Squares are the experimental data, four peaks
- are generated by mathematical treatment of the data and correspond either to large particles
- 676 (dashed line), medium-size particles (unbroken line), small particles (dotted line) and to the
- finest particles (not shown at this representation scale as it only corresponds to less than 1.3%
- of the particle total volume).

680 Figure 1

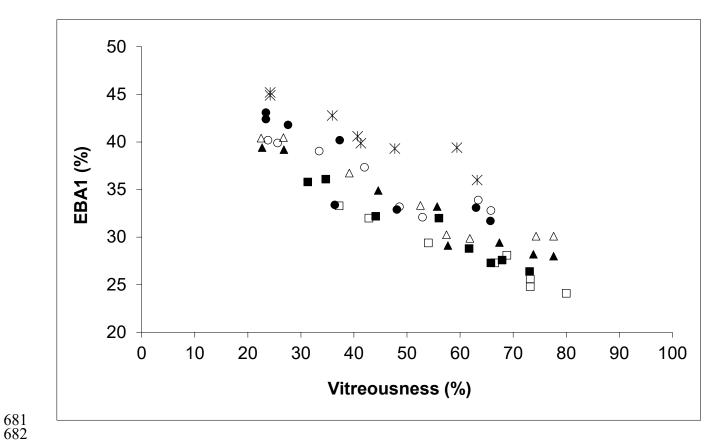


Figure 2

