

# Effect of wheat puroindoline alleles on functional properties of starch

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**Abstract** Puroindoline *a* and *b* (*Pina*, *Pinb*) form the molecular basis of bread wheat grain hardness. Varieties with a softer endosperm and a wild genotype, in which both *Pina* and *Pinb* were present, seemed to produce less damaged starch flour than hard varieties, where *Pin* mutations occurred and changed the starch rheological properties. The functional property of starch samples extracted from wheat varieties with different *Pin* alleles was evaluated. Starch morphology was characterized by scanning electron microscopy and laser light scattering. Thermal properties were evaluated by differential scanning calorimetry. Amylose content, starch damage and rapid visco-analyser (RVA) parameters were also determined. Significant variations ( $P < 0.05$ ) were identified between different *Pin* variants for the distribution pattern of starch granule volume, amylose content, starch damage, RVA viscosity breakdown and retrogradation, gelatinisation transition temperatures and enthalpies. Hard genotypes presented higher medium diameter granules and lower enthalpic values. However, the differences detected are more evident among varieties that present both *Pina* and *Pinb*, than among those presenting only one of the two (*a* or *b*).

**Keywords** Wheat starch · Puroindolines · Hardness · RVA · DSC · SEM

## Introduction

Endosperm texture is a useful characteristic to evaluate the technological ability of wheat varieties [1] affecting milling yield, flour particle size, shape and degree of starch damage. Grain texture is controlled by the hardness locus (*Ha*) on the distal end of the short arm chromosome 5D [2]. The genes of three polypeptides (puroindoline-*a*, puroindoline-*b*, and grain softness protein—GPS), which are components of friabilin [3, 4], have been identified as being closely linked to the *Ha* locus. Puroindolines (*Pina* or *pina-D1a* and *Pinb* or *pinb-D1a*) are thus generally considered to be genetic markers of endosperm texture [5, 6]. Soft wheats (wild type of *Triticum aestivum* L.) present both *Pina* and *Pinb*, whilst in hard wheats *Pina* is absent (null mutation, gene *pina-D1a* is not expressed), or several mutations are found to occur in *Pinb* [7–9]. In previous studies, commercial Portuguese bread wheat varieties were characterised in terms of their technological end-use, and different allelic puroindolines variants were identified [10, 11].

Puroindolines, which are found at the surface of starch granules, are two major isoforms (*Pina* and *Pinb*) and basic cysteine-rich proteins [12] with a molecular mass of about 13 kDa. Both may be extracted in Triton X-114 and possess a unique tryptophan-rich hydrophobic domain that is responsible for the strong affinity of puroindolines to lipids [13]. It has been suggested [4] that these lipid/protein interfaces play a major role in the texture of wheat endosperm by controlling the interactions between the starch granules and the protein matrix. Puroindoline lipid binding affinity affects flour properties, by producing stable foams on dough

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[14], and dough properties, by interfering in the formation and expansion of the gas cell [15].

Starch is the primary component of wheat flour and plays an important role in the quality of the end product. Most of the functional attributes of starch can be related to temperature-dependent interactions with water during gelatinisation, pasting and gelation (retrogradation) processes. The thermal behaviour depends on the physicochemical composition of starch, where the influence of the amylose/amylopectin ratio, amylose–lipid complex, branch chain length or degree of polymerisation of amylopectin has been demonstrated [16–18]. Physical treatments that occurred during the milling process changed the molecular arrangement, resulting in starch damage, which influenced starch functionality [19]. Differences were also detected in the gelatinisation behaviour of granule size classes [20–23].

Inasmuch as puroindolines are closely associated with starch granules, and given their effects on grain texture and starch's influence on the quality of wheat-based end products, the aim of the present study is to contribute to the knowledge of the influence of allelic puroindoline variability on the functional properties of bread wheat starch.

## Materials and methods

### Materials

Twelve varieties of bread wheat (Table 1) presenting different allelic puroindoline compositions and technological properties (such as grain hardness, flour water absorption, gluten strength and dough tenacity) were selected on the basis of the results of previous studies [10, 11]. Puroindoline allelic composition was obtained by DNA isolation [24] and PCR amplification of *Pina* and *Pinb* coding regions with specific primers that identify loss of *Pina* [12] or point mutation in *Pinb* [7, 25]. A waxy wheat variety presenting quite different properties was also analysed.

### Methods

Starch was extracted by a modified version of the method described by Hayakawa et al. [26]. The dough obtained from mixing purified water with flour was washed until the water came out clear. The recovered slurry was centrifuged thrice at 3,000g, the supernatant was discarded after each centrifugation and the starch was resuspended. The final starch slurry was then lyophilised. The apparent starch amylose content (% as is basis) and that of starch damage were determined in duplicate samples by a colorimetric method [27] and enzymatic assay kit (Megazyme, Wicklow, Ireland), respectively, following the AACC method 76–31 [28].

**Table 1** Puroindoline allelic composition and technological parameters of selected bread wheat varieties

Allelic composition of puroindolines	Hardness	W ( $10^{-4}$ J)	P (mm)	Abs (%)
<i>Amazonas</i> , <i>Centauro</i> , <i>Doirado</i>	Soft	149–215	53–72	49.5–53.6
<i>Argueil</i> , <i>Búfalo</i> , <i>Golia</i> , <i>Trapio</i>	Hard	144–301	85.3–135	59–64
<i>Idra</i> , <i>Lâncer</i> , <i>Pinzon</i> , <i>Regain</i>	Hard	108–436	52.3–125	52.1–58.9
<i>Armie</i>	Hard	189	71	55.5

Genotype that expresses wild type Puroindoline-a absence  
Mutation in puroindoline-b, substitution of Gly to Ser in position 46  
Mutation in puroindoline-b, substitution of Trp to Arg in position 44

Hardness NIR grain endosperm hardness, W ( $10^{-4}$  J) alveograph gluten strength, P (mm) alveograph dough tenacity, Abs (%) farinograph water absorption capacity

Starch granule morphology and surface were observed by scanning electron microscopy (SEM). Samples were spread on aluminium stubs, previously covered with a double-sided sticky tape. The surfaces were sputter-coated with gold. Images from starch granules were collected at 1500× amplification.

Starch granule size distribution was evaluated by laser light scattering (Mastersizer X, Malvern, UK). A polydisperse mode of analysis and a 300 mm lens were used. Starch powder was dispersed in isopropyl alcohol in the equipment circulation unit to attain an obscuration of 15–20%. Before measuring, the sample was circulated in the equipment with mechanical agitation and ultrasound for 1 min to dissolve the starch clots. Measurements were taken at 2 min intervals. Size distribution was determined in four replications of duplicate samples ( $n = 8$ ), and the results were expressed in terms of volume (%) occupied by starch type A (diameter  $> 9.48 \mu\text{m}$ ), type B ( $5.24 < \emptyset \leq 9.48 \mu\text{m}$ ) and type C (diameter  $\leq 5.24 \mu\text{m}$ ) granules. In addition, a medium sphere diameter ( $D_{(4,3)}$ ) was calculated based on the medium volume of the three granules fractions.

The apparent viscosity of starch suspensions (2.85 mg starch/25 ml purified water) was analysed using a rapid visco-analyser (RVA; Newport Scientific Pvt. Ltd.) in duplicate samples. The applied trial programme was based on the method used by Batey and Curtin [29]. The evaluated parameters were: maximum viscosity and the time (min) needed to reach it; minimum (or trough) and final viscosities; breakdown (BREAK; maximum viscosity minus minimum viscosity); setback 1 (final viscosity minus maximum viscosity); and setback 2 (STB.2; final viscosity minus minimum viscosity). All the results are expressed in RVA units (1 RVAU  $\approx 12$  cP).

Thermal properties were evaluated by DSC analysis, performed in a Shimadzu DSC 50 equipped with a TA 50 SI thermal analyser. Helium (99.95% purity) was the purge gas and flowed at approximately  $20 \text{ mL min}^{-1}$ . The calorimeter was calibrated according to a standard procedure laid down in the manufacturer's user manual. The DSC instrument was calibrated using indium (mp  $156.6 \text{ }^\circ\text{C}$ ,  $\Delta H_f = 28.45 \text{ J g}^{-1}$ ) and deionised water (mp  $0 \text{ }^\circ\text{C}$ ,  $\Delta H_m = 333 \text{ J g}^{-1}$ ). Triplicate samples of about  $5.0 \pm 0.5 \text{ mg}$  were weighted directly in aluminium pans (inner volume  $40 \mu\text{L}$ ) at a water/starch ratio of 70:30 (w/w), covered and hermetically sealed into place, and allowed to equilibrate at room temperature before scanning. An empty, hermetically sealed aluminium pan was used as reference. Samples were subjected to the following heating programme: from 30 up to  $100 \text{ }^\circ\text{C}$  at a heating rate of  $20 \text{ }^\circ\text{C/min}$ . Thermal transitions were defined in terms of onset ( $T_0$ ), peak ( $T_p$ ), and endset ( $T_c$ ) transition temperature. Heat of transition or enthalpy  $\Delta H$  ( $\text{J g}^{-1}$ ) was evaluated from peak areas and the

results were expressed per weight (g) of sample. The manufacturer's software programme was used to analyse and plot the thermal data.

Results were subjected to variance analysis (general linear model procedure,  $F$ -test on type III sum of squares; [30]), which was conducted in such a way as to detect significant differences ( $P < 0.05$ ) between puroindoline protein alleles. Significantly ( $P < 0.05$ ) different puroindoline protein allele means were classified by Duncan's test. Linear relationships between previously determined starch and flour technological quality parameters [10] were examined by generating Pearson correlation coefficients.

## Results and discussion

As observed by SEM (Fig. 1), wheat starch granules from all varieties showed typical morphological characteristics: lenticular large granules, as well as spherical small ones. No marked morphological differences were observed between the starch granules of different wheat varieties. However, the surface of starch granules from "hard" cultivars (Fig. 1a, d) seems to present more grooves than that of those from soft and waxy varieties (Fig. 1b, c, respectively).

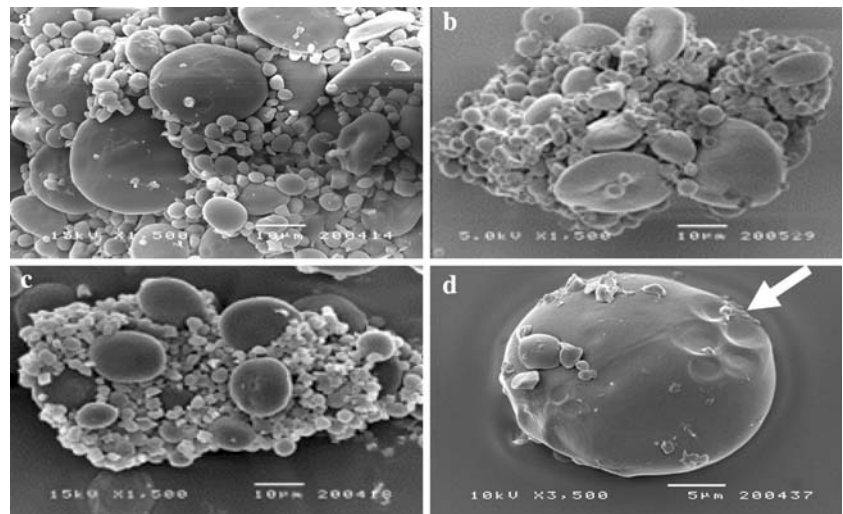
All varieties showed a population with three typical starch granule sizes: small granules (type C); midsize granules (type B); and large granules (type A). Highly significant differences were detected between puroindoline alleles as regards mean starch granule size distribution and medium diameter ( $D_{(4,3)}$ ; Fig. 2, Table 2), ranging from 19.9 to  $22.8 \mu\text{m}$ .

The histograms presented in Fig. 2a show that type A (granule with diameter  $> 9.48 \mu\text{m}$ ) represents more than 75% of the volume in all the studied varieties. Figures ranging from 57.9 to 76.9 % for granule type A population were obtained [31] in 12 soft wheat cultivars, but the cut-off points that delineate the class is  $9.9 \mu\text{m}$  instead of  $9.48 \mu\text{m}$ . Soft varieties, with both puroindolines-*a* and -*b* [pin (*a, a*)], showed a significantly higher percentage of volume occupied by small (type C) and medium (type B) size granules, and a significantly lower percentage occupied by large (type A) size granules than was the case in hard [pin (*a, b*), pin (*b, a*) and pin (*a, d*)] wheat bread varieties.

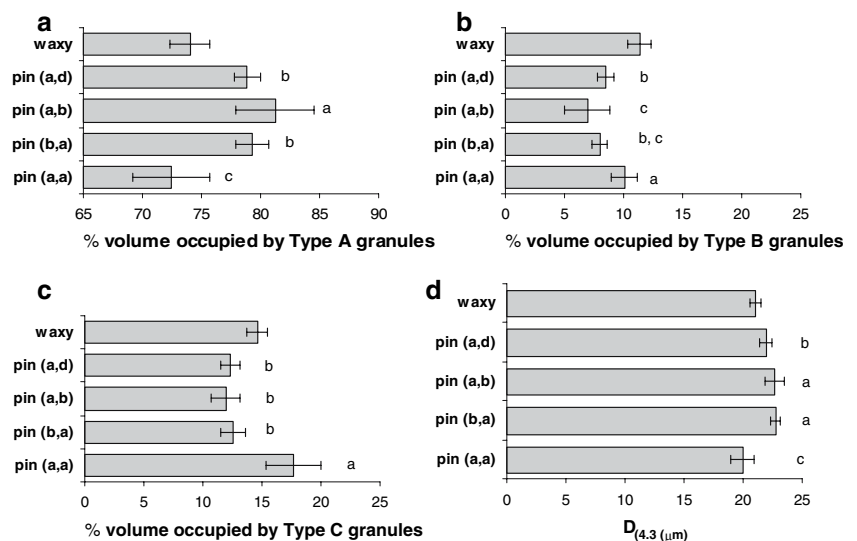
The results also revealed a smaller medium diameter (Fig. 2d) in soft varieties (pin (*a, a*)) than in the hard ones. Waxy starch presented an intermediate size.

The relationship between the size distribution of starch granules and wheat texture (hardness) has not yet been clearly established by the results of other authors [32, 35]. Differences observed in the range of A, B and C sizes, classes and hardness values for the analysed cultivars may be the primary reason for the difference encountered. It should

**Fig. 1** Images of starch granules from different wheat varieties obtained by SEM. **a** Hard wheat *Trapio*; **b** soft wheat *Centrauro*; **c** waxy wheat at 1,500 $\times$  magnification; **d** hard wheat *Trapio* at 3,500 $\times$  magnification. Arrow indicates the grooves in the starch granule



**Fig. 2** Histograms representing the mean and standard deviations of % volume occupied by type A, B and C granules and medium diameter type D starch granules<sup>(4,3)</sup> by allelic puroindoline and in waxy wheat. Histograms with the same letters (a, b or c) correspond to an absence of significantly different means at  $P < 0.05$ . Error bar signifies variability among four replications of duplicate samples ( $n = 8$ ) from the same genotype



**Table 2** Analysis of variance (mean squares) for starch granule size distribution of 12 bread wheats grouped in four puroindoline alleles ( $n = 85$ )

Source	df	Volume, Type A (%)	Volume, Type B (%)	Volume, Type C (%)	$D_{(4,3)}$ ( $\mu\text{m}$ )
Puroindolines	3	375.36*	36.56*	171.60*	42.92*
Replicates	7	9.26	2.25	3.14	0.80
Error	85	6.81	2.59	2.09	0.54

\* Significant at  $P = 0.001$

be noted that the varieties selected in the present study show a large gap in hardness values, contrary to the majority of values presented by other authors. While in the present study NIR hardness ranges from 34.2 to 105.7, in other studies the variation is lower [34]. Despite a large gap in hardness (16–100), Igrejas et al. [35] detected a higher number of small starch granules in hard varieties than in

soft ones, in a population derived from a synthetic amphihexaploid wheat. Discrepant results can be attributed to the different nature of the germplasm, and the analytical methodologies employed are not the same. In addition, different techniques for size determination (light microscopy, Coulter Counter, laser light scattering) and comparisons using different terms of expression (volume, number, or superficial area distribution) may also significantly affect results.

Significant differences between puroindoline alleles were also found for amylose content, starch damage, RVA breakdown, RVA setback 2 and thermal DSC properties (onset and peak temperature, enthalpy or heat required to gelatinise), as shown in Table 3.

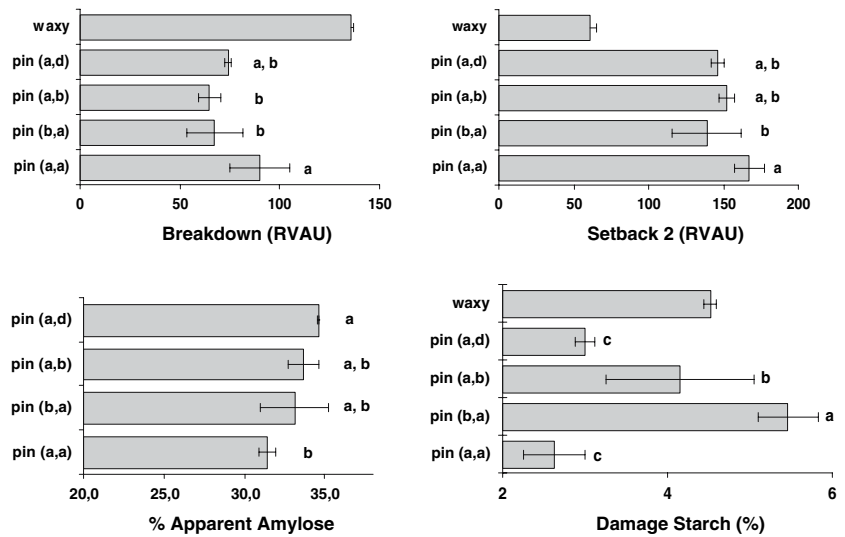
The apparent starch amylose contents of hard genotypes was found to be higher than those of the soft ones (average of 31.4%), but only the difference for the hard allele pin (a, d), with an average of 34.6%, is significant (Fig. 3). As expected, significant differences between the hard and soft

**Table 3** Analysis of variance (mean squares) for amylose, starch damage, RVA breakdown and setback, DSC onset and peak temperature and enthalpy of 12 wheats grouped in four puroindoline alleles

Source	df	Amylose (%)	Starch damage (%)	RVA break.	RVA STB. 2	df	DSC $T_0$ (°C)	DSC $T_p$ (°C)	DSC $\Delta H$ (J/g)
Puroindolines	3	7.78*	10.15***	833.56**	865.12*	3	56.22**	33.34*	4376.46***
Repetitions	1	0.01	0.05	53.52	5.79	2	1.97	0.49	2.94
Error	19	2.32	0.38	139.35	219.65	30	11.03	9.39	118.92

\*, \*\*, \*\*\* Significant at  $P = 0.05$ ,  $P = 0.01$  and  $P = 0.001$ , respectively

**Fig. 3** Histograms representing the mean and standard deviations for RVA setback 2, breakdown, damage starch and apparent amylose content, by allelic puroindoline and in waxy wheat. Histograms with the same letters (a, b or c) correspond to an absence of significantly different means at  $P < 0.05$ . Error bar signifies variability among duplicate samples from the same genotype



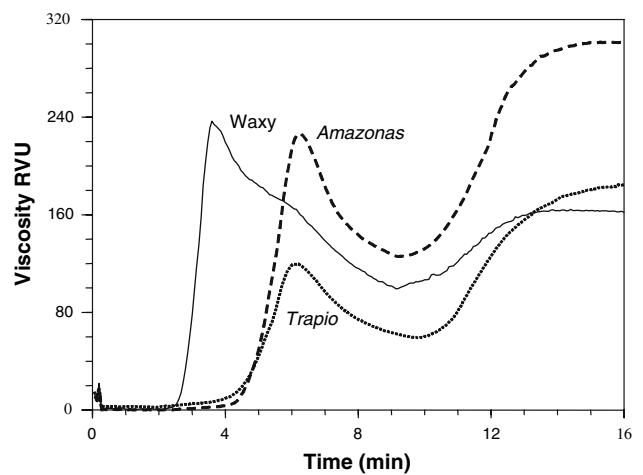
genotypes were obtained for starch damage: higher values were found in harder genotypes and the differences were more pronounced between pin (a, a) and pin (b, a) alleles.

The pasting viscosity profile exhibits significant differences between the soft genotype and the hard alleles pin (b, a) and pin (a, b); when *Pina* is absent, the breakdown and setback are significantly lower than in the wild genotype pin (a, a); the absence of *Pinb* has the same effect on breakdown.

Apparent amylose content is related with the maximum and minimum viscosities, thereby demonstrating a highly significantly negative correlation (result not shown) between apparent amylose and RVA breakdown.

In the present study, due to its different gelatinisation profile (as is also shown by the RVA results, Figs. 3, 4) waxy starch was used only as a reference. The expected pasting profile was obtained and reached the highest value for peak viscosity earlier (3.6 min.), with higher breakdown (135 RVU) and lower setback values (60 RVU) than the other genotypes. This is in agreement with previous findings [36, 40], where the rapid swelling was attributed to a high amylopectin content.

The DSC results (Fig. 5) lend further insight into the understanding of the different behaviour of hard and soft

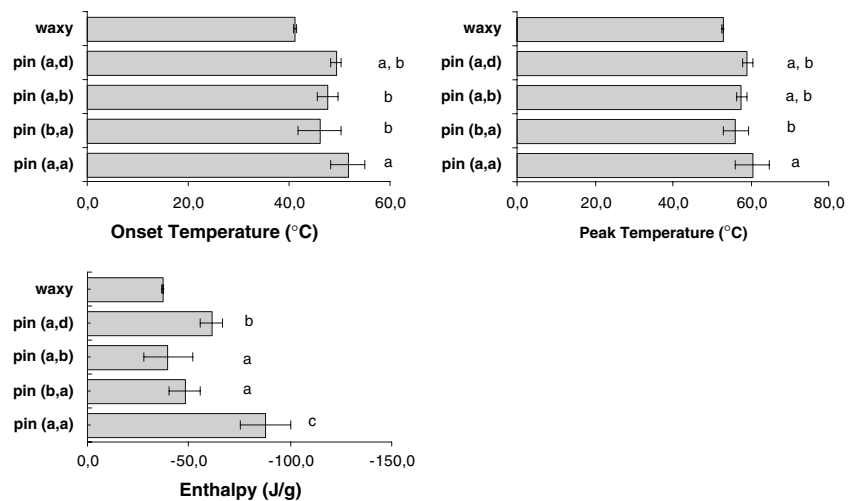


**Fig. 4** RVA pasting profile of starch from waxy, soft Amazonas and hard Trapio genotypes

genotypes, which is clear from the onset gelatinisation temperatures and in which the enthalpic values constitute the most discriminating parameter.

Overall, the onset temperature varied between 41.2 and 55.4 °C, and  $T_p$  from 51.1 to 65.9 °C, with significant variation between genotypes. No significant difference was

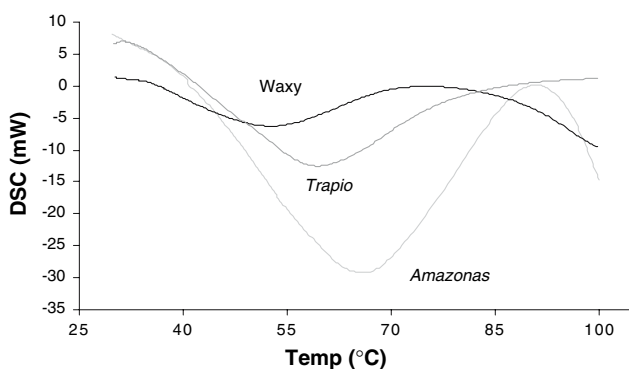
**Fig. 5** Histograms representing the mean and standard deviations of onset and peak temperatures and DSC enthalpy, by allelic puroindoline composition and for waxy wheat. Histograms with the same letters (a, b or c) correspond to an absence of significantly different means at  $P < 0.05$ . Error bar signifies variability among triplicate samples from the same genotype



detected in the conclusion/final temperatures. In fact, when both puroindolines are present, the gelatinisation onset and peak temperatures are higher than when either one of them is missing. This aspect is more evident when looking for energy values.

Starch from soft wheat varieties required higher temperatures to start gelatinisation (onset temperature) and also more energy to complete gelatinisation than hard wheat equivalents, as evidenced by thermograms (Fig. 6) for *Amazonas*-pin (a, a) and *Trapio*-pin (b, a) starch.

In native starch granules, gelatinisation corresponds to granule hydration and swelling, followed by the thermal disordering of crystalline structures. It was assumed that starch granule size might influence granule hydration and swelling. The small-granule starches presented a higher ratio of surface area per unit weight of starch, and thus hydrated and swelled more efficiently than the large-granule starches. In this respect, some studies [20, 41] have reported higher gelatinisation enthalpies for A-type starch granules than for B-type ones, although another work [42] reports larger gelatinisation temperature ranges for B-type compared with A-type granules. In the present study, it was observed that starch



**Fig. 6** DSC profile obtained for waxy *Trapio*-pin (b, a) and *Amazonas*-pin (a, a) starch

from soft wheat varieties presented a lower starch granule medium diameter, the values for this parameter being negatively correlated to gelatinisation energy values (data not shown). In other words, the former needed higher energy values to complete the gelatinisation process than did the hard varieties. A higher peak gelatinisation temperature for small granules than for the larger fractions was also obtained [20, 23], a difference that was attributed to the accumulation of defects during starch biosynthesis. Large granule fractions accumulated more crystallite defects than small granules during starch deposition—a tendency that was enhanced in hard genotypes. The defects could be related to other factors like starch granule composition and degree of crystallisation [43]. The gelatinisation endotherm may also be attributed to a disordering of amylopectin crystallites [16]. However, the role of crystallinity was secondary [23], and the different thermal behaviour may well be due to the different starch amylose/amylopectin composition. Despite the influence of the amylose content, no significant difference in gelatinisation temperatures was detected between normal and waxy starch [40]. Waxy alleles of hard *Búfalo*, *Golia* and *Pinzon* and soft *Amazonas* and *Centauro* have been characterised [44, 45] and exhibit wild genotype (*Wx-A1-a*, *Wx-B1-a*, *Wx-C1-a*). Moreover, the apparent amylose content in the starch of hard genotypes [pin (a, b), pin (b, a)] ranging from 33.1 to 33.7% does not significantly differ from that of soft genotypes (31.4%), and so the differences encountered cannot be laid at the door of this parameter. Hardness proved to be negatively correlated to the surface lipid content [46]. In softer wheats containing greater lipid-complexed amylose [34], the different behaviour detected can thus be attributed to the magnitude of the lipid-amylose complex or amylopectin—crystalline/amorphous. These characteristics affect granule starch chain organisation and the degree of crystalline organisation within the granules and inter-chain associations. On the other hand, the physical damage to starch granules during milling converts crystalline

amylopectin into amorphous amylopectin, with the formation of some low molecular weight fragments [47]. A large proportion of very short amylopectin chains ( $DP < 8$ ) in the large granules are related to crystallite defects [23]. This phenomenon is more pronounced in hard genotypes with greater starch damage. According to Matsuki et al. [48], this is probably the reason for the lower gelatinisation temperatures and enthalpy of these genotypes. That conclusion is also supported by León et al. [19], who detected a similar influence of starch damage on the decrease in enthalpy.

Waxy starch, which has almost no amylose, displays lower transition temperatures and enthalpy than the other genotypes, and so it gelatinises easily compared to the study varieties. This probably means that in these cases, the higher amylopectin increases the disordering phenomenon, although Fujita et al. [43] obtained a higher gelatinisation temperature for a waxy genotype compared to a non-waxy type.

Differences in thermic properties between the three hard allelic variants pin (*a*, *d*), pin (*a*, *b*) and pin (*b*, *a*) were only detected for gelatinisation enthalpy values with the higher value for allele with the Trp/Arg mutation.

Correlation coefficients between previously determined physicochemical grain and flour traits [10] and starch functionality were calculated (Table 4) in order to draw conclusions about the influence of starch on the rheological behaviour of wheat flours.

As expected, the protein content was not correlated with the starch parameters, and hardness was significantly correlated with the same parameters as those affected by puroindoline composition. Grain hardness evaluated by NIR correlated significantly ( $P < 0.05$ ) with starch damage ( $r = 0.83$ ), medium diameter ( $r = 0.66$ ), RVA setback ( $r = -0.62$ ) and DSC transition temperatures  $T_0$  ( $r = -0.64$ ) and  $T_c$  ( $r = -0.61$ ). This is corroborated by Noda et al. [49], who obtained positive correlation between the starch gelatinisation temperature

and the softness of wheat. Flour sedimentation volumes evaluated by sodium dodecyl sulphate test correlated significantly ( $P < 0.05$ ) with RVA pasting parameters (except setback).

Flour water absorption is significantly ( $r = 0.92$ ,  $P < 0.001$ ) correlated with starch damage and negatively correlated ( $r = -0.66$ ) with setback 2. Dough tenacity, as evaluated by alveograph ( $P$ ), is significantly negatively correlated with starch damage and DSC gelatinisation temperatures  $T_0$ ,  $T_p$  and  $T_c$ . The strong relation between grain hardness, flour water absorption, starch damage and dough tenacity was extensively reported [1, 10, 19, 20]. Gluten proteins are the main compounds responsible for dough tenacity. In addition, different thermal behaviours were found [50] in gluten fractions that were isolated from one soft variety (*Amazonas*) and one hard variety (*Sorraia*). The soft variety required more energy to onset and to develop the transition than the hard variety. However, despite the importance of  $P$  contribution to dough strength (as expressed by  $W$  values), no relationship was found between this parameter and starch properties.

## Conclusions

The analysis of medium granule diameter in 13 bread wheat variety starches clearly revealed lower values for soft genotypes than for hard genotypes.

Starch granule size distribution, starch damage, RVA breakdown, RVA setback and DSC  $T_0$ ,  $T_p$  and  $\Delta H$  parameters were affected by the presence of both puroindolines *a* and *b*; however, more research is needed with other germplasm as isogenic lines to surpass these results in bread wheat breeding programmes. The analytical methodologies used to assess the gelatinisation phenomenon (RVA breakdown/DSC  $T_0$ ; RVA Setback 2/DSC  $T_p$ ) led to similar results as regards discrimination of the puroindoline alle-

**Table 4** Correlation and level of significance between some of the most relevant parameters of starch and flours analysed

Flour starch	Protein (%)	Hard.	SDS (mm)	Abs (%)	$P$ (mm)	$W$ ( $10^{-4}J$ )
Damage (%)	-0.0594 ns	0.8327***	-0.2533 ns	0.9255***	0.8508***	0.4019 ns
Peak. (RVA)	0.2511 ns	-0.3114 ns	0.5894*	-0.3998 ns	0.0101 ns	0.3338 ns
Trough. (RVA)	0.2249 ns	-0.2344 ns	0.605 *	-0.3863 ns	0.1113 ns	0.3539 ns
End. (RVA)	0.2984 ns	-0.451 ns	0.5934*	-0.562 ns	-0.1125 ns	0.2138 ns
STB. 2 (RVA)	0.3269 ns	-0.6202*	0.4614 ns	-0.6601	-0.3574 ns	0.0039 ns
$T_0$ (°C)	-0.1524 ns	-0.635 *	-0.08 ns	-0.5586 ns	-0.7709 **	-0.4649 ns
$T_p$ (°C)	-0.1063 ns	-0.5626 ns	0.0533 ns	-0.4771 ns	-0.6506 *	-0.2609 ns
$T_c$ (°C)	-0.0805 ns	-0.6095 *	0.1791 n.s	-0.5481 ns	-0.6516 *	-0.182 ns
$\Delta H$ (J/g)	-0.4559 ns	-0.5382 ns	-0.2925 ns	0.4598 ns	0.4425 ns	0.1689 ns
$D_{(4,3)}$ ( $\mu m$ )	-0.4378 ns	0.6606*	-0.3465 ns	0.5156 ns	0.5096 ns	0.2188 ns

\*, \*\*, \*\*\* Significant at  $P = 0.05$ ,  $P = 0.01$  and  $P = 0.001$ , respectively

les—the  $\Delta H$  parameter being the most determinant in this respect.

The higher values for the transition temperatures and enthalpy of soft genotypes may reflect a more ordered/stable granular organisation when both puroindolines *a* and *b* are present in the endosperm than when one of them is absent. This is also supported by a significant negative correlation between starch thermal properties and grain hardness.

It was concluded that hard varieties require a low level of energy to onset and to develop the transition during the gelatinisation process. This aspect has important practical and useful implications for technological wheat operations, most of which—milling and the extrusion processes, for example—are energy dependent.

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