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Chemometric Approach to Characterization of Flour Mill Streams: Chemical and Rheological Properties

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Abstract The aim of the present study was to utilize chemometric methods (the principal component analysis and hierarchical cluster analysis) for monitoring the certain aspects of flour mill streams quality and their interrelation to selected rheological properties. Thirty-seven flour mill streams were separated from industrial mill of 300 t/day capacity. All flour streams were analyzed for ash, protein, wet gluten, and damaged starch content and rheological properties as determined by Brabender Farinograph, Extensograph, and Amylograph. The obtained results indicated that break, sizing, and reduction flour streams exhibited different rheological behavior in relation to a change in protein, wet gluten, ash, and mechanically damaged starch content within the milling passages. Rheological properties of dough during mixing and kneading as well as during extension were different with regard to the technological phase of milling from which they were extracted. The obtained results could be utilized for selection of certain flour streams in production of special-purpose flours.

Keywords Flour mill streams · Chemical properties · Rheological properties · Principal component analysis

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Introduction

The main objective of industrial flour milling is to extract the maximum quantity of flour of desired quality. It is accomplished by the successive grinding and sieving operations during which the gradual fragmentation of wheat kernel occurs. Industrial wheat milling process comprises three main stages: grinding within roller mills, sieving, and purifying. The grinding stage comprises several phases—passages where certain amount of flour is produced, removed from the process and combined to provide final flour (Pomeranz 1988; Kent and Evers 1994; Posner and Hibbs 1997; Owens 2001; Sugden and Osborne 2001). A comprehensive knowledge about the distribution of certain chemical, biochemical, and rheological properties between mill streams is an important issue for optimization of final product quality (Ramseyer et al. 2011; Liu et al. 2011). Therefore, the determination of different aspects of quality of flour mill streams has been the focus of attention of numerous authors—the content and distribution of total protein and ash (Prabhasankar et al. 2000; Loza-Garay and Flores 2003; Okrajková et al. 2007), wet gluten (Pojić et al. 2004), lipids and fatty acids (Prabhasankar and Haridas Rao 1999; Prabhasankar et al. 2000), enzymes (Rani et al. 2001; Gebruers et al. 2002; Every et al. 2006a), damaged starch (Sutton and Simmons 2006; Banu et al. 2010; Pojić et al. 2012), pentosans (Wang et al. 2006a; Ramseyer et al. 2011), and antioxidants (Engelsen and Hansen 2009). Protein composition of different flour mill streams has also been investigated (Menkovska et al. 2002; Sutton and Simmons 2006; Wang et al. 2006b; Okrajková et al. 2007; Liu et al. 2011) as well as distribution of sulfur and free amino acids (Liu et al. 2011). The diversity in protein composition of

63 flour mill streams as well as the presence of variable amount of
 64 certain kernel constituents influences the different rheological
 65 properties of flour mill streams (Pojić et al. 2004; Banu et al.
 66 2010; Liu et al. 2011). The data on the distribution of
 67 certain constituents and functional properties between flour
 68 mill streams could be utilized for the monitoring of the
 69 efficiency of milling procedure, mill settings, and blending
 70 of flour mill streams to obtain certain end-use quality that
 71 meet specific customer demands (Prabhasankar et al. 2000;
 72 Spasojević et al. 2000; Liu et al. 2011).

73 The implementation of more efficient tools for monitor-
 74 ing, optimization, characterization, handling of raw mate-
 75 rials, intermediate and final products, as well as for predic-
 76 tion of quality throughout the whole production chain has
 77 become a must for the modern food industry if it strives for
 78 competitive position in the market. Throughout the produc-
 79 tion process, a number of different measurements are
 80 performed resulting in a large amount of data, used for one
 81 specific purpose (Bro et al. 2002). However, by combining
 82 all available information, the extraction of even more rele-
 83 vant information from the collected data is possible by
 84 application of multivariate statistics, mathematical model-
 85 ing, and computing—all of them combined into a highly
 86 multidisciplinary research discipline—chemometrics (Bro et
 87 al. 2002; Gemperline 2006). It has become an irreplaceable
 88 tool for exploratory analysis of large datasets, multivariate
 89 quality assurance and quality control, detection of adultera-
 90 tion, estimation of chemical, physical, and sensory proper-
 91 ties of foods (Gemperline 2006). Despite the wide applica-
 92 tion in food science, relevant literature is lacking in exam-
 93 ples related to the use of chemometrics in studying the
 94 quality characteristics of flour mill streams, with the excep-
 95 tion of the study by Dornez et al. (2006) who utilized the
 96 principal component analysis to study the distribution of
 97 arabinoxylans, endoxylanases, and endoxylanase inhibitors
 98 in industrial wheat roller mill streams. The application ex-
 99 amples of chemometrics in the quality control of processed
 100 food with spectroscopic data are numerous, and beyond the
 101 scope of this study. Another example on the application of
 102 chemometrics in the quality control of processed food in-
 103 volved the utilization of chemometrics to study the influence
 104 of commercial bread improvers on dough rheology and
 105 hearth bread properties (Aamodt et al. 2003), the monitoring
 106 of authenticity of Brazilian UHT milk (Souza et al. 2011),
 107 the characterization of Brazilian lager and brown ale beers
 108 (Granato et al. 2011), and the discrimination between low-
 109 and full-fat yogurts (Cruz et al. 2013).

110 The objective of this study was to characterize the
 111 flour mill streams in terms of their chemical and rheo-
 112 logical properties and to assess the interdependence be-
 113 tween them by using two chemometric techniques—
 114 principal component analysis (PCA) and hierarchical
 115 cluster analysis (HCA).

Materials and Methods

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Samples

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118 Thirty-seven wheat flour mill streams were obtained from a
 119 commercial wheat flour mill (capacity 300 t/day): 8 break
 120 mill streams (1B-I₁, 1B-I₂, 2B-I₁, 2B-I₂, 3B-I₂, 3B-I₂, 4B-I₁,
 121 and 4B-I₂), 11 sizing mill streams (1R-I₁, 1R-I₂, 1R-II₁, 1R-
 122 II₂, 1R-III₁, 1R-III₂, 2R-I₁, 2R-I₂, 2R-II₁, 2R-II₂, and 2R-
 123 III), 17 reduction mill streams (1M-I, 1M-II, 2M-I, 2M-II,
 124 3M-I, 3M-II, 3M-III, 4M-I, 4M-II, 5M-I, 5M-II, 6M, 7M,
 125 8M, 9M-I, 9M-II, and 9M-III) and one bran duster flour
 126 stream (Fs–Vs). The flour stream samples were collected in
 127 accordance with the milling diagram during a fixed time
 128 interval taking into account the succession of the milling
 129 passages (Fig. 1).

Methods

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131 The selected physicochemical properties of flour mill
 132 streams were determined as follows: moisture content
 133 according to ICC 109/1, protein content according to ICC
 134 105/2, ash content according to ICC 104/1, and wet gluten
 135 content according to ICC 106/2 (ICC 1996) and damaged
 136 starch content according to AACC method 76–33.01
 137 (AACC 2000). Selected rheological properties included
 138 Brabender Farinograph water absorption and softening de-
 139 gree determined according to ICC 115/1; Brabender
 140 Extensograph energy, dough resistance, and extensibility
 141 according to ICC 114/1; and Brabender Amylograph peak
 142 viscosity ICC 126/1. All selected properties of flour streams
 143 were analyzed in duplicate.

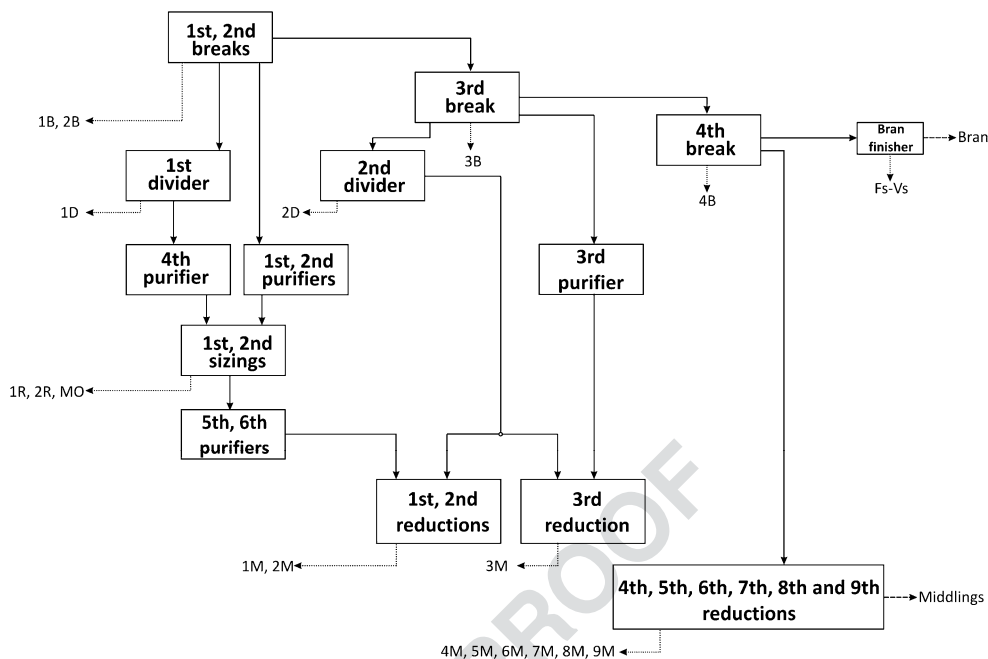
Data Analysis

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145 A one-way analysis of variance (ANOVA) was used to test
 146 the significant differences in chemical and rheological qual-
 147 ity between flour mill streams. ANOVA was followed by
 148 Fisher’s least significant difference test, where the differ-
 149 ences between means at the 5 % level ($p < 0.05$) were con-
 150 sidered significant.

151 The subsequent data analysis included two pattern rec-
 152 ognition methods, the PCA and HCA applied to the data for
 153 each technological phase of milling—break, sizing, and
 154 reduction. Therefore, three different matrices were formed:
 155 8 rows (samples) × 11 columns (quality indicators) for break
 156 phase, 11 rows (samples) × 11 columns (quality indicators)
 157 for sizing phase, and 18 rows (samples) × 11 columns (qual-
 158 ity indicators) for reduction phase (Cruz et al. 2013). PCA
 159 and Pearson correlation coefficients (r) were used for find-
 160 ing the relationships between selected quality properties of
 161 flour mill streams. PCA was performed following the mean
 162 centering, performed as a preprocessing method to equalize

Fig. 1 A simplified mill flow diagram



163 the statistical importance of all the variables, with cross-
 164 validation (Beebe et al. 1998; Aamodt et al. 2003; Granato
 165 et al. 2011). HCA was performed following autoscaling as a
 166 preprocessing step. Sample similarities were calculated on
 167 the basis of the squared Euclidean distance, while the Ward
 168 hierarchical agglomerative method was used to establish
 169 clusters (Granato et al. 2011; Cruz et al. 2013). Statistical
 170 analysis was carried out using the software XLSTAT, ver-
 171 sion 2012.2.02 (Addinsoft, NY, USA).

172 **Results and Discussion**

173 Physicochemical Properties of Flour Mill Streams

174 *Break Flour Mill Streams*

175 The role of break system is to open the wheat kernel and to
 176 gradually scrape the endosperm from the bran throughout
 177 four break passages (1B, 2B, 3B, and 4B), which result in
 178 stock with maximally possible large middlings and minimal
 179 quantity of flour with minimal bran contamination. Certain
 180 amount of flour is produced within the break stage either
 181 due to releasing particles from fractured endosperm or par-
 182 ticle attrition (Kent and Evers 1994; Posner and Hibbs 1997;
 183 Catterall and Cauvain 2007). The ground stock from the first
 184 two break passages are additionally classified on the two
 185 grading systems (D1 and D2), where additional amount of
 186 flour is separated (Fig. 1). This flour is not a result of
 187 grinding, but extended sieving of stock from the preceding
 188 breaking passages. Physicochemical properties of break
 189 flour streams indicated a wide variation in their quality

(Table 1). Protein content (P) of break flours gradually
 increased over the successive break passages from 11.7 to
 16.1 % dry matter (dm) being in accordance with the previ-
 ously reported results (Prabhasankar et al. 2000; Rani et al.
 2001; Dornez et al. 2006; Wang et al. 2007; Banu et al.
 2010; Sakhare et al. 2012). The increase in protein content
 of break flours is affected by the increase in the presence of
 peripheral endosperm and bran particles rich in protein
 (Wang et al. 2007). Wet gluten content followed the similar
 trend as protein content and gradually increased over the
 successive break passages from 31.3 % for 1B to 39.7 % for
 3B. Ash content (A) of break flours varied between 0.44 and
 0.75 % dm with noticeable increase from 2B to 4B due to
 gradual roll gap decrease which resulted in release of aleu-
 rone layer, fine bran, and germ particles along with the
 endosperm particles (Sakhare and Inamdar 2011). Although
 1B flour often has the lowest ash content among break flours,
 1B flour was characterized by higher ash content in relation
 to 2B flour stream, probably due to the release of accumu-
 lated mineral dust from kernel crease (Sakhare and Inamdar
 2011). Damaged starch content showed an increasing trend
 from 12.7 to 20.2 % (U) Chopin Dubois (UCD) as the grind-
 ing progressed from 1B to 4B showing the opposite ten-
 dency observed by Banu et al. (2010).

Rheological properties of dough as determined by empiri-
 cal rheological tests such as Brabender Farinograph, and
 Extensograph reflect the processing quality of flours and
 are commonly used as a part of routine quality control within
 the baking industry (Stojceska et al. 2007; Dapčević
 Hadnađev et al. 2011). The presence of different anatomic
 parts of wheat kernel in break flours caused their different
 rheological behavior (Hayta and Schofield 2004).

Table 1 Selected physicochemical and rheological properties of break flour streams

Passage	P % dm	WG %	A % dm	WA %	SD BU	E cm ²	R BU	Ex mm	PV BU	DS UCD
t1.4	11.8±0.06 n	31.7±0.43 e,f	0.48±0.00 k	50.0±0.25 s	82±2.9 e,f,g	44±1.5 o,p	172±2.9 l,m	163±2.1 f,g	173±2.9 k	12.7±0.14 q,r
t1.5	11.7±0.05 n,o	31.3±0.20 f,g	0.49±0.01 j,k	50.5±0.15 r,s	73±2.9 f,g,h	48±1.5 o	174±5.1 l,m	169±2.5 e	180±5.0 k	13.2±0.21 q
t1.6	12.8±0.01 k,l	32.2±0.10 e	0.45±0.00 l	52.1±0.40 o,p,q	48±2.9 k,l,m	103±4.6 c,d	285±5.0 e	178±3.8 c,d	222±5.8 h,i	14.4±0.28 p
t1.7	12.7±0.02 l	31.9±0.38 e	0.44±0.01 l	52.2±0.47 o,p,q	52±2.9 j,k,l	99±4.0 d,e	292±2.9 d,e	182±3.8 c	237±2.9 h	16.0±0.21 n,o
t1.8	15.1±0.03 e	39.4±0.10 a,b	0.44±0.00 l	54.0±0.30 m,n	43±2.9 l,m,n	122±4.2 a	230±3.0 i,j	231±4.6 b	220±5.0 i	17.6±0.14 j,k
t1.9	15.3±0.09 e	39.7±0.76 a	0.57±0.01 i	54.2±0.45 m,n	50±5.0 k,l,m	126±5.0 a	232±3.6 i,j	235±2.5 b	217±2.9 i	17.2±0.14 k,l
t1.10	16.3±0.10 d	39.0±0.75 b,c	0.73±0.03 g	56.9±0.57 k,l	55±5.0 j,k,l	106±3.3 c	190±4.2 k	248±4.2 a	220±5.0 i	20.2±0.14 h,i
t1.11	16.1±0.02 d	38.6±0.23 c	0.75±0.01 g	56.6±0.35 l	60±2.9 i,j,k	112±5.5 b	193±4.7 k	251±2.1 a	225±5.0 h,j	20.2±0.14 h,i

Mean value±standard deviation. Figures followed by the same letter within the same column are not significantly different ($p < 0.05$)

P protein content, *WG* wet gluten content, *A* ash content, *WA* Farinograph water absorption, *SD* Farinograph softening degree, *E* Extensograph energy, *R* Extensograph resistance, *Ex* Extensograph extensibility, *PV* peak viscosity, *DS* damaged starch

PCA was performed for all analyzed chemical and rheological characteristics in order to observe the quality variation among the flour mill streams and their interrelationships (Martens and Martens 2001).

This method was chosen due to its suitability for identifying the patterns in data and data display which provides evident observation of their similarities and differences indicating the correlation between variables. The original variables are compressed into a small number of new variables—principal components (PCs) describing independent variation structures in the data. PCs are used as new axes in a plot of samples (score plot) and a corresponding plot of variables (loading plot). Hence, the properties that are related and those that are most important in distinguishing between samples are easily visualized. Two variables that appear close to each other in the plot indicate their positive correlation, whereas variables appearing on opposite sides are negatively correlated. Variables located in perpendicular directions along the PC axis are independent of each other (Aamodt et al. 2003; Domez et al. 2006; Shin et al. 2010).

Figure 2a shows the PCA score plot of the first two principal components (PC1 vs. PC2) derived from selected physicochemical properties of break flours, revealing the similarities and differences between them. A clear separation of the flour samples from different passages as well as grouping of the flour samples originated from the same passage is noticeable (Fig. 2a). PCA loading plot revealed that 93.39 % of the variability in the data was explained by the first two PCs, where 67.40 % of the variability was explained by PC1 and 25.99 % by PC2 (Fig. 2b). By examination of PCA loading plot, it was noticeable that increase in protein, wet gluten, and mechanically damaged starch content affected the increase in the Farinograph water absorption ($r=0.97$, $r=0.86$, and $r=0.93$, respectively) which varied from 50.0 to 56.9 % as the grinding progressed from 1B to 4B (Fig. 2b, Table 1). The tail-end break flours appeared to have more acceptable water absorption values for bread making than those from head-end passages. The intensity of mechanical damage of starch influences the quality of flour in terms of water absorption, rheological behavior, and fermentation of the leavened products thus determining bread crumb firmness and crust color (Di Stasio et al. 2007). Higher water binding capacity is also attributed to the presence of arabinoxylans, mainly located in the aleurone and bran (Pomeranz 1988; Domez et al. 2006; Wang et al. 2006a; Fustier et al. 2009a, b; Banu et al. 2010; Ramseyer et al. 2011). Ramseyer et al. (2011) reported strong correlation of total and water-unextractable arabinoxylan content with ash content, which reflected the presence of bran in flour streams suggesting the differences in flour functionality. Protein and wet gluten content were positively correlated with the Extensograph extensibility ($r=0.99$ and $r=0.96$, respectively) and energy ($r=0.79$ and $r=0.77$, respectively). The Extensograph

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275 extensibility exhibited the gradual increase from 163 mm for 1B
 276 to 251 mm for 4B, while the Extensograph energy gradually
 277 increased from 44 cm² for 1B to 126 cm² for 3B (Table 1). The
 278 positive relationship between protein content and dough exten-
 279 sibility along the first PC for patent flour was previously
 280 reported by Aamodt et al. (2003). Fustier et al. (2009a, b)
 281 reported the positive relation between protein content and the
 282 Alveograph deformation energy, being the measure of exten-
 283 sional work as Extensograph energy is. Dobraszczyk and
 284 Salmanowicz (2008) reported close relation between protein
 285 content and Kieffer maximum extensibility and area under
 286 Kieffer force–distance curve. Kieffer dough extensibility rig
 287 measures large deformation of dough in uniaxial extension
 288 providing results that correspond to Extensograph results
 289 (Dunnwind et al. 2004). Resistance to extension of break flour
 290 streams was not related to measured chemical properties of
 291 break flours and manifested no particular trend over the break
 292 passages, being the lowest for 1B (170 BU) and highest for 2B
 293 (290 BU; Table 1). Although Aamodt et al. (2003) reported
 294 that resistance to extension was related to flour protein con-
 295 tent along the first PC, it must be noted that the distribution
 296 of the various protein groups occurs during milling process,
 297 which is reflected in a corresponding variation of the rheo-
 298 logical properties of flour streams (Wang et al. 2007). The
 299 obtained results were also consistent with the results of
 300 Dobraszczyk and Salmanowicz (2008) who reported no rela-
 301 tion between protein content and Kieffer maximum force that
 302 corresponds to the Extensograph resistance to extension. The
 303 rheological properties of flour mill streams are also dependent
 304 on the presence of glutathione, being impaired with the
 305 increasing content of glutathione typical for the tail-end pas-
 306 sages (Every et al. 2006b).

307 The peak viscosity of break flours as being located in
 308 perpendicular directions along the PC1 axis appeared to be
 309 independent of selected chemical parameters (Fig. 2b). The
 310 peak viscosity values varied within narrow range 175–
 311 225 BU, indicating that the viscosity was not a consequence
 312 of mechanical damage of starch but of flour origin.

313 HCA was performed, primarily due to its ability to
 314 reveal a natural groupings (or clusters) within a given data
 315 set, in the form of a special graph—dendrogram. Thus, the
 316 visualization of clusters and correlations among samples or
 317 variables is enabled (Souza et al. 2011). Similar to PCA,
 318 HCA revealed three distinct clusters of break flour mill
 319 streams (Fig. 2c). Cluster 1 contained the initial break
 320 flours (1B-I1 and 1B-I2), being in accordance with the
 321 sample grouping in the PCA score plot. The loading plot
 322 indicated that their position was related to the highest
 323 Farinograph softening degree. Cluster 2 contained flour
 324 streams from second break passage (2B-I1 and 2B-I2),
 325 remained separated from other samples indicating that their
 326 Extensograph resistance significantly differed from the
 327 other break flours. Cluster 3 contained the terminal break

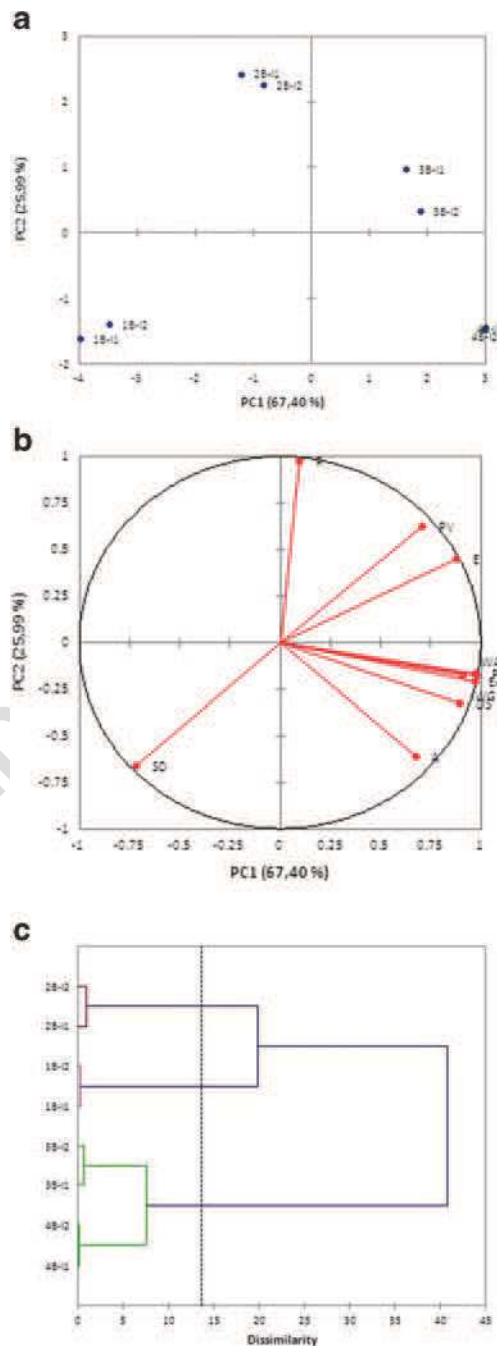


Fig. 2 Break flour mill streams: PCA score plot (PC1 vs PC2) (a). PCA loading plot (PC1 vs PC2) for physicochemical and rheological properties (b). HCA dendrogram (c). *P* Protein content, *A* ash content, *WG* wet gluten content, *DS* damaged starch content, *WA* Farinograph water absorption, *SD* Farinograph softening degree, *E* Extensograph energy, *Ex* Extensograph extensibility, *R* Extensograph resistance to extension, *PV* amylograph peak viscosity

flour streams (3B-I1, 3B-I2, 4B-I1, and 4B-I2) which
 appeared separated from each other in PCA score plot
 due to significantly different values of the Extensograph
 energy and extensibility, ash, and damaged starch content

333 *Sizing Flour Mill Streams*

334 The main objective of the sizing system is to separate bran and
 335 endosperm particles that remained together after breaking
 336 passages, to reduce the size of large endosperm particles and
 337 to enhance the particle size uniformity. The selected physico-
 338 chemical and functional properties of sizing flour mill streams
 339 are shown in Table 2. Due to the absence of bran and peripheral
 340 endosperm particles, sizing flours were of lower ash
 341 content compared to that of break flours. Ash content of sizing
 342 flours varied within narrow range of 0.30–0.34 % dm, indicating
 343 that flour originates from the central parts of endo-
 344 sperm. Moreover, the sizing flour streams were characterized
 345 by lower protein and wet gluten content in comparison to
 346 those of breaking passages, varying within the range of
 347 10.4–11.8 and 23.0–33.0 % dm, respectively. Damaged starch
 348 content of sizing flours varied within broad range from 0.2 to
 349 23.1 UCD, with no regular trend over the sizing passages.

350 Although no discernable pattern in variation of selected
 351 chemical and rheological properties of mill streams within
 352 the sizing system could be noticed, the PCA score plot of the
 353 first two principal components (PC1 vs. PC2) revealed clear
 354 separation of flours originated from different sizing pas-
 355 sages. The PCA loading plot revealed that 73.48 % of the
 356 variability in the data was explained by the first two PCs,
 357 where 45.82 % of the variability was explained by PC1 and
 358 27.66 % by PC2 (Fig. 3a). Farinograph water absorption
 359 increased with the increase in protein ($r=0.75$), ash ($r=0.80$),
 360 and damaged starch content ($r=0.98$; Fig. 3b). Moreover, in-
 361 crease in wet gluten content caused the decrease in Farinograph
 362 softening degree ($r=-0.74$), while Extensograph energy ($r=$
 363 0.62) and extensibility ($r=0.67$) increased with the increase in
 364 protein content (Fig. 3b). Increase in mechanically damaged
 365 starch granule content caused the decrease in Amylograph peak
 366 viscosity ($r=-0.94$) due to increased susceptibility of mechan-
 367 ically damaged starch granule to α -amylase hydrolysis.

368 Similar to PCA, HCA revealed three distinct clusters of
 369 sizing flour mill streams (Fig. 3c). Cluster 1 contained the first
 370 flours from the first two passages of sizing (1R-II, 1R-I2, 2R-
 371 II, and 2R-I2), which in PCA score plot appeared separated
 372 from each other due to significantly different values of
 373 Farinograph water absorption, protein, and damaged starch
 374 content (Fig. 3a and b, Table 2). Cluster 2 comprised the
 375 second and third flours from the initial passage of sizing
 376 (1R-III1, 1R-II2, 1R-III1, and 1R-III2), while cluster 3 com-
 377 prised the remaining flours from the second sizing passage
 378 (2R-III1, 2R-II2, and 2R-III), which corresponded to the sam-
 379 ple grouping as appeared in PCA score plot (Fig. 3a).

380 *Reduction Flour Mill Streams*

381 The role of the reduction system is to reduce the size of
 382 endosperm particles into as much high-quality flour as

Table 2 Selected physicochemical and rheological properties of sizing flour streams

Passage	P % dm	WG %	A % dm	WA %	SD BU	E cm ²	R BU	Ex mm	PV BU	DS UCD
1R-I ₁	11.8±0.01 n	29.6±0.10 i	0.33±0.01 n,o,p	56.2±0.57 l	23±2.9 p,q	101±3.2 c,d,e	327±2.9 c	156±4.7 h,i,j	288±2.9 e,f	23.1±0.14 f
1R-I ₂	11.8±0.02 n	29.0±0.10 i	0.34±0.00 n,o	56.7±0.61 k,l	35±5.0 n,o,p	97±4.7 e	350±5.0 b	150±2.3 k,l	283±2.9 f	22.4±0.14 f,g
1R-II ₁	11.5±0.06 o,p	31.1±0.30 g	0.32±0.00 o,p,q	53.5±0.25 m,n,o	23±0.0 p,q	104±5.1 cd	360±5.0 a,b	154±3.2 i,j,k	380±5.0 b	13.2±0.07 q
1R-II ₂	11.3±0.04 p,q	29.1±0.35 i,j	0.32±0.00 o,p,q	53.0±0.41 n,o,p	18±2.9 q	90±3.2 f	365±5.0 a	139±3.2 m	380±5.0 b	12.2±0.28 r
1R-III ₁	11.3±0.01 p,q	27.4±0.30 k,l,m	0.31±0.01 p,q	51.2±0.30 q,r,s	23±2.9 p,q	100±6.2 d,e	322±2.9 c	164±4.0 e,f	425±0.0 a	0.2±0.07 x
1R-III ₂	11.0±0.02 r,s,t	33.3±0.20 d	0.30±0.01 q	51.1±0.55 q,r,s	23±2.9 p,q	84±2.5 g	302±2.9 d	150±1.5 k,l	428±2.9 a	2.0±0.07 w
2R-I ₁	11.1±0.01 q,r,s	27.5±0.04 k,l,m	0.31±0.00 p,q	54.6±0.47 m	52±2.9 k,l,m	62±3.0 k,l,m	222±2.9 j	155±2.1 h,i,j,k	288±2.9 e,f	16.2±0.21 m,n
2R-I ₂	11.0±0.04 r,s,t	27.0±0.31 m,n	0.32±0.00 o,p,q	54.4±0.35 m,n	45±0.0 l,m,n	70±2.0 i,j	253±2.9 g,h	156±2.5 h,i,j	302±2.9 d,e	18.3±0.14 j
2R-II ₁	10.7±0.05 u,v,w	26.6±0.20 n,o	0.30±0.01 q	51.7±0.38 p,q,r	28±2.9 o,p,q	75±3.5 h,i	363±5.8 a,b	126±3.0 o,p	393±5.8 b	4.4±0.07 v
2R-II ₂	10.6±0.02 v,w,x	27.1±0.33 l,m,n	0.31±0.00 p,q	51.7±0.36 p,q,r	42±2.9 m,n,o	85±2.3 f,g	367±15.3 a	139±1.5 m	390±5.0 b	6.4±0.21 t
2R-III	10.4±0.01 x	23.0±0.39 s,t	0.32±0.01 o,p,q	51.5±0.55 q,r	65±0.0 h,i,j	80±2.5 g,h	370±5.8 a	131±3.2 n,o	385±5.0 b	5.5±0.07 u

Mean value±standard deviation. Figures followed by the same letter within the same column are not significantly different ($p<0.05$)

P protein content, WG wet gluten content, A ash content, WA Farinograph water absorption, SD Farinograph softening degree, E Extensograph energy, R Extensograph resistance, Ex Extensograph extensibility, PV peak viscosity, DS damaged starch

383 possible. At this stage, bran and germ particles are flattened
 384 which enable their separation during sieving. The reduction
 385 flour mill streams were characterized by general gradual
 386 increase in protein (from 10.5 to 17.8 % dm) and ash (from
 387 0.35 to 2.52 % dm) content being the highest in the tail-end
 388 reduction flours due to the increasing contamination of
 389 millstreams with aleurone, bran, and germ particles
 390 (Table 3). Prabhasankar et al. (2000) and Dornez et al.
 391 (2006) observed no regular trend in protein content increase
 392 from head-end to tail-end reduction passages, whereas
 393 Gebruers et al. (2002) indicated a regular trend in protein
 394 content increase for both break and reduction rolls. Dornez
 395 et al. (2006) indicated that the increase in protein content
 396 over reduction stage is not always obvious due to smaller
 397 differences in protein content of reduction passages in relation
 398 to that of break passages. Wet gluten content gradually
 399 increased from 18.0 % for 1 M to 30.5 % for 3 M. The
 400 content of mechanically damaged starch increased from 9.7
 401 to 29.6 UDC being the highest in the tail-end reduction
 402 flours due to more severe grinding at tail-end reduction
 403 passages (Table 3). By studying the microstructure of tail-
 404 end flour streams from roller mill, Gangadharappa et al.
 405 (2008) indicated the presence of deformed A-type (lenticular
 406 shaped) and intact B-type (spherical shaped) starch granules
 407 in the very tail-end flour stream, whereas preceding
 408 flour stream contained slightly visible structural deforma-
 409 tions of A type starch granules.

410 The qualitative differences within the reduction flours
 411 could be visualized in the PCA score plot, where tail-end
 412 reduction flours (6M–9M) were located to the left and
 413 separated from head-end reduction flours (1M–5M) to the
 414 right in the score plot (Fig. 4a). Bran duster flour stream
 415 (Fs–Vs) appeared to be entirely distinct from tail-end
 416 reduction flours. In the corresponding loading plot (Fig. 4b),
 417 79.55 % of the variability in the data was explained by the
 418 first two PCs, where 62.51 % of the variability was
 419 explained by PC1 and 17.05 % by PC2. The increasing
 420 content of peripheral endosperm in flour resulted in increas-
 421 ing water absorption throughout the reduction system due to
 422 their higher water-binding capacity (Banu et al. 2010). In
 423 particular, the increase in water absorption was affected by
 424 the increase in protein ($r=0.63$), ash ($r=0.58$), and damaged
 425 starch content ($r=0.49$). The increase in Farinograph soft-
 426 ening degree was affected by increase in protein ($r=0.70$),
 427 ash ($r=0.66$), and damaged starch content ($r=0.54$), show-
 428 ing the opposite trend in comparison to the break mill-
 429 streams. Moreover, the increased presence of peripheral
 430 kernel particles rich in protein and minerals affected the
 431 decrease in Extensograph energy ($r=-0.75$ and $r=-0.76$,
 432 respectively) and resistance to extension ($r=-0.84$ and $r=-$
 433 0.74 , respectively) being in contradiction with break flour
 434 mill streams. The opposite functional behavior between
 435 break and reduction streams in terms of total flour protein

436 content was also observed by Wang et al. (2007) who
 437 reported the positive correlation between loaf volume and
 438 protein content of break streams, and negative correlation
 439 between loaf volume and protein content of reduction
 440 streams. The reason for diverse relation of certain

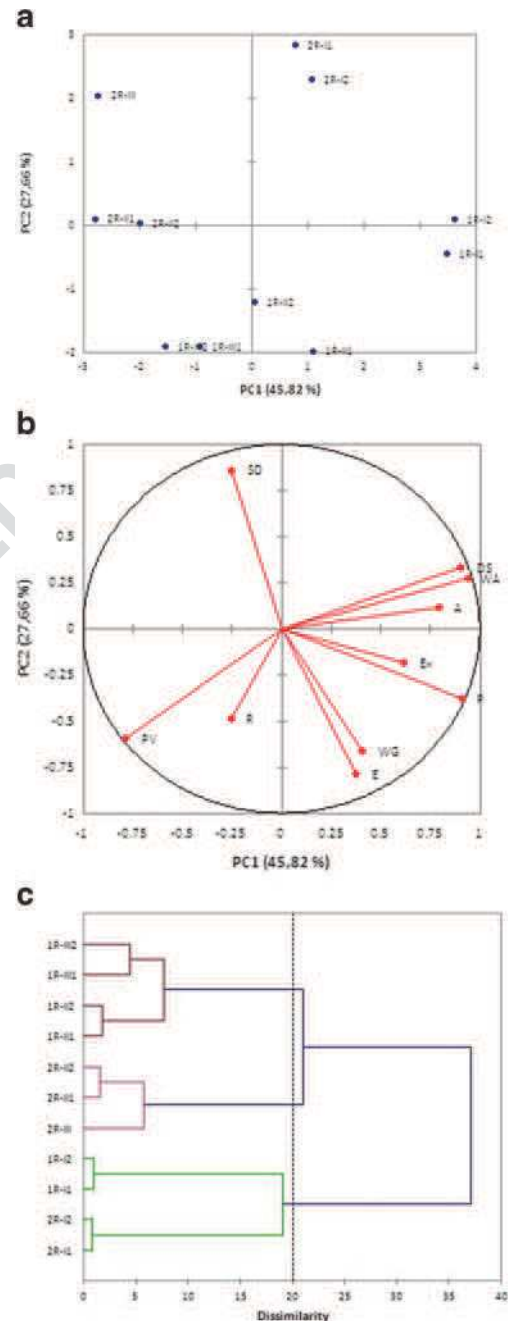


Fig. 3 Sizing flour mill streams: PCA score plot (PC1 vs PC2) (a). PCA loading plot (PC1 vs PC2) for physicochemical and rheological (b). HCA dendrogram (c). *P* Protein content, *A* ash content, *WG* wet gluten content, *DS* damaged starch content, *WA* Farinograph water absorption, *SD* Farinograph softening degree, *E* Extensograph energy, *Ex* Extensograph extensibility, *R* Extensograph resistance to extension, *PV* amylograph peak viscosity

t3.1 **Table 3** Selected physicochemical and rheological properties of reduction flour streams

t3.2 t3.3	Passage	P % dm	WG %	A % dm	WA %	SD BU	E cm ²	R BU	Ex mm
t3.4	1M-I	10.8±0.12 t,u,v	23.1±0.34 s,t	0.35±0.01 n	58.1±0.15 j,k	63±2.9 h,i,j	66±3.2 j,k	302±2.9 d	131±2.0
t3.5	1M-II	10.5±0.01 w,x	22.7±0.02 t,u	0.38±0.02 m	58.5±0.21 j	78±2.9 e,f,g	57±3.5 m,n	303±2.9 d	124±2.5
t3.6	2M-I	11.2±0.03 q,r	25.4±0.32 p	0.38±0.00 m	60.7±0.41 h,i	90±0.0 d,e	54±0.6 n	273±15.3 f	130±3.8
t3.7	2M-II	10.9±0.03 s,t,u	24.1±0.24 q	0.38±0.00 m	59.0±0.35 j	75±5.0 f,g,h	59±0.2 l,m,n	290±10.0 d,e	127±2.1
t3.8	3M-I	11.8±0.06 n	27.6±0.12 k,l	0.40±0.00 m	62.5±0.32 f,g	70±0.0 g,h,i	71±0.5 i,j	252±10.4 g,h	152±1.5
t3.9	3M-II	11.5±0.05 o,p	28.9±0.16 j	0.45±0.01 l	62.7±0.25 f,g	62±2.9 i,j,k	69±0.9 j	333±11.5 c	130±3.0
t3.10	3M-III	11.0±0.08 l	22.2±0.19 u	0.50±0.01 j,k	65.0±0.32 d,e	78±2.9 e,f,g	44±1.5 o,p	257±10.4 g	118±2.5
t3.11	4M-I	13.3±0.02 r,s,t	26.3±0.23 o	0.51±0.01 j	62.0±0.23 g,h	68±2.9 g,h,i	63±2.1 k,l	220±5.0 j	160±2.5
t3.12	4M-II	13.1±0.11 i,j	26.4±0.42 o	0.49±0.01 j,k	59.5±0.41 i,j	73±2.9 f,g,h	66±1.2 j,k	240±5.0 h,i	158±2.5
t3.13	5M-I	12.2±0.05 m	23.5±0.18 r,s	0.48±0.02 k	63.6±0.31 e,f	85±5.0 e,f	56±3.0 n	252±5.8 g,h	134±3.6
t3.14	5M-II	13.0±0.07 j,k	23.5±0.10 r,s	0.63±0.01 h	65.6±0.15 d	100±5.0 d	39±1.5 p	217±15.3 j	122±2.3
t3.15	6M	14.3±0.04 g	23.9±0.32 q,r	0.89±0.01 e	69.0±0.35 c	118±2.9 c	26±2.1 q	163±11.5 m	117±2.1
t3.16	7M	16.1±0.03 d	27.5±0.26 k,l,m	1.35±0.01 d	68.6±0.15 c	123±2.9 b,c	23±3.0 q	112±2.9 n	148±2.1
t3.17	8 M	14.8±0.04 f	27.8±0.52 k	1.39±0.01 c	70.5±0.25 b	132±2.9 a,b,c	16±0.6 r	123±11.5 n	90±1.5 r
t3.18	9 M-I	17.8±0.05 a	18.0±0.50 w	2.52±0.02 a	72.0±0.29 a	130±2.9 a,b,c	n/a	n/a	n/a
t3.19	9 M-II	17.3±0.05 b	19.1±0.60 v	2.09±0.01 b	71.5±0.25 a,b	140±5.0 a	n/a	n/a	n/a
t3.20	9 M-III	16.7±0.12 c	18.5±0.62 w	2.07±0.01 b	58.8±0.25 j	80±0.0 e,f,g	n/a	n/a	n/a
t3.21	Fs-Vs	13.4±0.03 h	30.5±0.30 h	0.79±0.01 f	71.0±0.49 a,b	135±5.0 a,b	56±1.5 n	183±5.8 k,l	175±2.6

Mean value±standard deviation. Figures followed by the same letter within the same column are not significantly different ($p<0.05$)

P protein content, *WG* wet gluten content, *A* ash content, *WA* Farinograph water absorption, *SD* Farinograph softening degree, *E* Extensograph energy, *R* extensibility, *PV* peak viscosity, *DS* damaged starch

441 rheological properties of dough of break and reduction
 442 millstreams might be associated with different distribution
 443 of protein fractions among them (Figs. 2b and 4b; Liu et al.
 444 2011). Namely, the different rheological behavior of break
 445 and reduction flour streams could be explained by higher
 446 ratio of polymeric to monomeric proteins of break than of
 447 reduction streams (Wang et al. 2007). Break passages espe-
 448 cially head-end passages release relatively pure endosperm
 449 particles, while tail-end milling passages tend to scrape off
 450 residual endosperm particles from peripheral layer of kernel,
 451 together with fine bran and germ particles (Sakhare et al.
 452 2012). Resistance to extension was negatively correlated
 453 with damaged starch content. Jovanovich et al. (2003) indi-
 454 cated that the positive relation between Alveograph dough
 455 tenacity and damaged starch content regardless of the dif-
 456 ferences in wheat and milling fractions. When performing
 457 Alveograph test, which is conducted with constant moisture
 458 content of dough, flour containing higher damaged starch
 459 content would not be completely hydrated resulting in
 460 higher dough tenacity. The increased presence of peripheral
 461 kernel particles rich in protein, minerals, and amylolytic
 462 enzymes affected the decrease in Amylograph peak viscosi-
 463 ty ($r=-0.79$ and $r=-0.78$, respectively). The fact that α -
 464 amylase is mostly located in the peripheral parts of a wheat
 465 kernel indicated similar distribution of α -amylase and ash
 466 content among flour mill streams, where tail-end flours were
 467 characterized by higher α -amylase activity (Rani et al. 2001;
 468 Every et al. 2002; Dornez et al. 2006). More intense shear
 469 stresses imposed on starch granules during reduction stage
 470 induced the stronger mechanical damage of starch granules
 471 than during break stage. Since the recorded peak viscosity is
 472 an indirect measure of the present α -amylase status along
 473 with the starch granule quality, the higher α -amylase activi-
 474 ty and higher quantity of mechanically damaged starch in
 475 the tail-end reduction passages resulted in lower peak vis-
 476 cosity (Fustier et al. 2009a, b).

477 HCA revealed three distinct clusters of reduction flour
 478 mill streams (Fig. 4c). Cluster 1 contained the flour streams
 479 from head-end reduction passages (1M-I, 1M-II, 2M-I, 2M-
 480 II, 3M-I, 3M-II, 3M-III, 4M-I, 4M-II, and 5M-I) located to
 481 the left side in the corresponding PCA score plot due to the
 482 differences in Alveograph peak viscosity, Extensograph resis-
 483 tance and energy between this group and the remaining
 484 group of reduction flours. Cluster 2 included 5M-II, 6M,
 485 7M, 8M, and Fs–Vs flours, while cluster 3 comprised the
 486 terminal reduction flour streams (9M-I, 9M-II, and 9M-III).
 487 The corresponding loading plot indicated that the position of
 488 these terminal reduction flours was related to their high
 489 protein and ash content (Fig. 4a and b).

490 The noticeable differences in the rheological behavior of
 491 break, sizing, and reduction flours could be generally attrib-
 492 uted to the presence of different proportion of various ana-
 493 tomic part of a wheat kernel in them (Wang et al. 2007; Fustier

et al. 2009a, b). In particular, flour streams differ in protein 494
 concentration and its molecular composition—amount and 495
 size distribution of polymeric and monomeric proteins, 496
 amount of free thiol groups, and amount of disulfide linkings 497
 (Menkovska et al. 2002; Sutton and Simmons 2006; 498
 Okrajková et al. 2007; Wang et al. 2007). Moreover, the 499

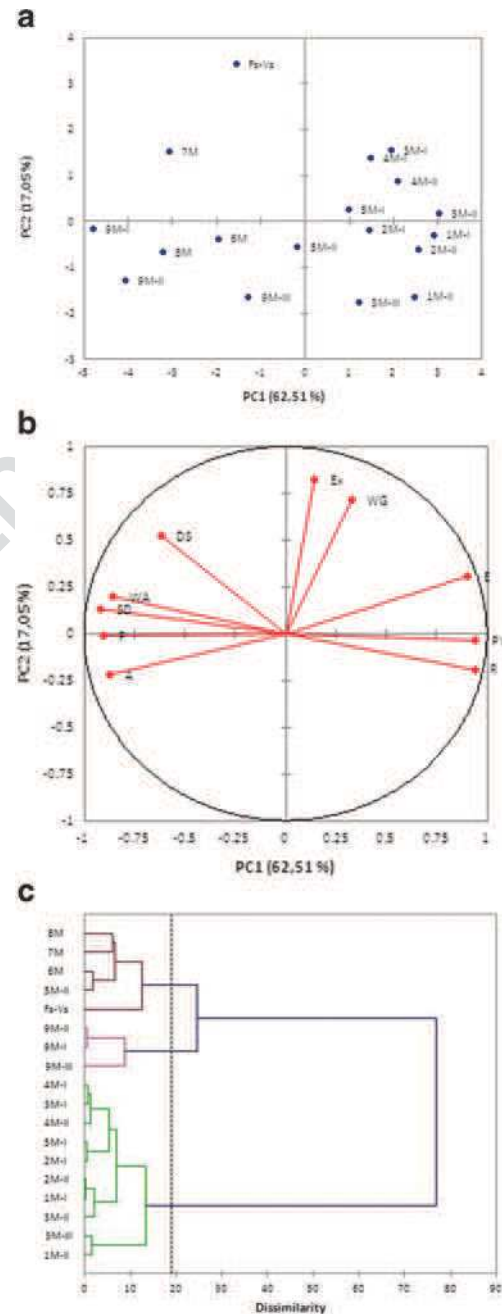


Fig. 4 Reduction flour mill streams: PCA score plot (PC1 vs PC2) (a). PCA loading plot (PC1 vs PC2) for physicochemical and rheological properties (b). HCA dendrogram (c). *P* Protein content, *A* ash content, *WG* wet gluten content, *DS* damaged starch content, *WA* Farinograph water absorption, *SD* Farinograph softening degree, *E* Extensograph energy, *Ex* Extensograph extensibility, *R* Extensograph resistance to extension, *PV* amylograph peak viscosity

500 variable distribution of lipids, fatty acids, enzymes (α -amy-
 501 lase, protease, lipoxygenase, polyphenol oxidase, peroxidase),
 502 and mechanically damaged starch granules within the flour
 503 mill streams also contribute to the variable rheological behav-
 504 ior of flour mill streams (Prabhasankar and Rao 1999;
 505 Prabhasankar et al. 2000; Rani et al. 2001; Di Stasio et al.
 506 2007). The different rheological behavior of flour streams
 507 could be associated to different sulfur content between break
 508 and reduction flour streams, being the higher for break flours
 509 as indicated by Liu et al. (2011). Higher sulfur content is
 510 associated with the increase in the proportion of LMW sub-
 511 units of glutenin and increased dough extensibility and de-
 512 creased dough elasticity thereof (Zhao et al. 1999; Liu et al.
 513 2011). Liu et al. (2011) indicated that sulfur content of flour
 514 mill streams supplemented with ash content appeared to be
 515 useful to evaluate their quality due to significant correlations
 516 with bread loaf volume, Farinograph peak time, and
 517 Extensograph parameters. Better rheological properties and
 518 higher gluten strength of break flours in comparison to those
 519 of reduction flours could be explained by lower degree of
 520 disulfide cross-linkings and higher amount of albumins and
 521 globulins mainly located in the peripheral parts of wheat
 522 kernel (Okrajková et al. 2007; Wang et al. 2007). Every et
 523 al. (2006a) reported higher concentration of enzyme peroxi-
 524 dase in the break flours, which oxidizes and cross-links ferulic
 525 acid residues on arabinoxylan and produces a secondary pen-
 526 tosan network through the gluten network that improves rhe-
 527 ological properties of dough made from break flours. More
 528 apparent variability in rheological properties of break flours in
 529 comparison to those of reduction flour could be attributed to
 530 the amount of gliadin, sodium dodecyl sulfate (SDS)-soluble
 531 glutenin, and SDS-insoluble glutenin that changed markedly
 532 within the breaking stage in contrast to reduction stage (Sutton
 533 and Simmons 2006). Sutton and Simmons (2006) also found
 534 that tailing passages exposed to more intensive grinding and
 535 consequently more intensive molecular disruption, character-
 536 ized by higher thiol content (Sutton and Simmons 2006) and
 537 higher damaged starch content (Jovanovich et al. 2003;
 538 Gangadharappa et al. 2008) affecting certain rheological and
 539 processing quality properties.

540 **Conclusion**

541 Flour mill streams from consecutive milling passages largely
 542 differ in composition, rheological properties, and hence in
 543 overall technological functionality. The selected chemometric
 544 techniques, the principal component analysis, and the hierar-
 545 chical cluster analysis could be effectively used for the visu-
 546 alizations of the performance of milling procedure by mon-
 547 itoring the different aspects of flour mill streams quality and
 548 their interdependency. Break, sizing, and reduction flour
 549 streams exhibited different rheological behavior in relation

550 to a change in protein, wet gluten, ash, and mechanically
 551 damaged starch content within the milling process.
 552 Rheological properties of dough made of flour streams
 553 recorded during mixing and kneading were different with
 554 regard to the technological phase of milling from which
 555 they were extracted. Softening degree of break flours was
 556 not related to measured chemical properties, while that of
 557 sizing and reduction flours was negatively correlated with
 558 wet gluten content and positively correlated with protein, ash,
 559 and damaged starch content, respectively. Moreover, the ex-
 560 tensional properties of dough made of flour streams were
 561 dependent on the phase of milling from which they were
 562 extracted so that extensibility and energy was positively cor-
 563 related with protein and wet gluten content of break and sizing
 564 flours. Energy and resistance to extension of dough made
 565 from reduction flours was negatively correlated with protein
 566 and ash content. Pasting properties of sizing and reduction
 567 flour streams were negatively correlated with ash and mechan-
 568 ically damaged starch content, while such dependence was not
 569 observed for break flours.

570 Although the results in this study were obtained by
 571 milling certain wheat within one milling diagram, it might
 572 be expected that the interrelation between certain aspects of
 573 flour millstream quality could be also applicable in the
 574 processing of wheat of different quality and/or using differ-
 575 ent milling diagram. Hence, the selection of certain flour
 576 streams for production of special-purpose flours, as well as
 577 the selection of appropriate flours for specific bakery prod-
 578 ucts could be enabled.

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 583

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