

Adaptability and stability of white oat cultivars in relation to chemical composition of the caryopsis

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Abstract – The objective of this work was to characterize the chemical properties of white oat (*Avena sativa*) caryopsis and to determine the adaptability and stability of cultivars recommended for cultivation in the state of Rio Grande do Sul, Brazil. The trials were carried out in the 2007, 2008 and 2009 crop seasons, in three municipalities: Augusto Pestana, Capão do Leão, and Passo Fundo. Fifteen cultivars were evaluated in a randomized block design, with four replicates. The contents of protein, lipid, and nitrogen-free extract were evaluated in the caryopsis. Cultivar performances for the measured characters varied according to location and year of cultivation. The cultivar URS Guapa showed high content of nitrogen-free extract and low contents of protein and lipid in the caryopsis. 'FAPA Louise' showed high content of lipid, whereas 'Albasul', 'UPF 15', and 'UPF 18' showed high content of protein and low content of nitrogen-free extract. There is no evidence of an ideal biotype for the evaluated characters, which could simultaneously show high average performance, adaptability to favorable and unfavorable environments, and stability.

Index terms: *Avena sativa*, bissegmented linear regression, lipid, nitrogen-free extract, nutritional quality, protein.

Adaptabilidade e estabilidade de cultivares de aveia-branca quanto à composição química da cariopse

Resumo – O objetivo deste trabalho foi caracterizar as propriedades químicas da cariopse de aveia-branca (*Avena sativa*) e determinar a adaptabilidade e estabilidade de cultivares recomendadas para cultivo no Rio Grande do Sul. Os ensaios foram realizados nas safras de 2007, 2008 e 2009, em três municípios: Augusto Pestana, Capão do Leão e Passo Fundo. Foram avaliadas 15 cultivares, em delineamento de blocos ao acaso, com quatro repetições. Foram determinados os teores de proteína, lipídeos e extrativos não nitrogenados na cariopse. O desempenho das cultivares quanto aos caracteres avaliados variou conforme o local e ano de cultivo. A cultivar URS Guapa apresentou elevado conteúdo de extrativos não nitrogenados na cariopse, e baixos de proteína e lipídeo. 'FAPA Louise' apresentou elevado conteúdo de lipídeo, enquanto que 'Albasul', 'UPF 15' e 'UPF 18' apresentaram elevado conteúdo de proteína e baixo teor de extrativos não nitrogenados. Não há evidência de um biótipo ideal para os caracteres avaliados, que apresente, simultaneamente, elevado desempenho médio, adaptabilidade a ambientes favoráveis e desfavoráveis, e estabilidade.

Termos para indexação: *Avena sativa*, regressão linear bissegmentada, lipídeo, extrativos não nitrogenados, qualidade nutricional, proteína.

Introduction

The environment tends to affect the contents of protein, lipid, and nitrogen-free extract in white oat (*Avena sativa* L.) caryopses (Beber et al., 2002; Zhu

et al., 2004; Gatto, 2005; Peterson et al., 2005; Martinez et al., 2010). Due to the characteristically polygenic behavior detected in these characters (Zhu et al., 2004; Orr & Molnar, 2007; Crestani et al., 2012), the genotype x environment (G x E) interaction does not

provide a deep understanding of genotype performance in different environmental conditions. The analysis of stability and adaptability allows the identification of genotypes responsive to environmental variations and with predict performance (Cruz et al., 2004).

Among available methods to identify the adaptability and stability of genotypes, the one proposed by Cruz et al. (1989) is usually adopted for evaluation of annual species, tested in several environments (more than eight). This method has been adopted to evaluate grain yield of white oat genotypes in different environments and cultivation schemes (Lorencetti et al., 2002, 2004). According to this model, adaptability and stability are established based on bissegmented linear regressions, in which the dependent variable is expressed as an index that measures the quality of the evaluated environments. According to Cruz et al. (1989), the ideal genotype should have high average performance, adaptability to favorable and unfavorable environments, and regression deviations equal to zero (stability).

Considering the chemical attributes of oat caryopsis, genotypes with grains suitable for human diet should have low calory, low contents of lipid and nitrogen-free extract, and high contents of protein and dietary fiber (Holland, 1997; Chernyshova et al., 2007). For feedstuffs, however, genotypes should have high calory, characterized by high contents of lipid, protein, and nitrogen-free extract (Holland, 1997; Marinissen et al., 2004). The fraction of nitrogen-free extract corresponds to nonstructural carbohydrates in the caryopsis, mainly starch, sucrose, and pectin. The understanding of the G x E interaction for caryopsis chemical quality of white oat cultivars is very important for their proper recommendation, aiming at the production of high-quality grains, destined for specific market niches. Moreover, the estimates of environment effects on cultivar phenotypic expression is essential to direct future selection strategies of superior white oat genotypes for chemical quality of grains.

The objective of this work was to characterize the chemical properties of white oat (*Avena sativa*) caryopsis, in relation to protein, lipid and nitrogen-free extract contents, and to determine the adaptability and stability of cultivars recommended for cultivation in the state of Rio Grande do Sul, Brazil, regarding these characteres.

Materials and Methods

Fifteen white oat cultivars, recommended for cultivation in the state of Rio Grande do Sul, Brazil, were evaluated during the crop seasons of 2007, 2008, and 2009. The trials were carried out in the municipalities Augusto Pestana (28°27'S, 53°54'W, 328-m altitude), in a Rhodic Hapludox (Latossolo Vermelho distrófico, according to Santos et al., 2006); in Capão do Leão (31°45'S, 52°29'W, 13-m altitude), in a Typic Hapludox (Latossolo Vermelho-Amarelo); and in Passo Fundo (28°15'S, 52°24'W, altitude 687m), in a Rhodic Hapludox. All trials were performed in a randomized block design, with four replicates. A density of 250 viable seeds per square meter was used. Replicates were formed by plots with five 5.0-m rows, spaced in 0.20 m, with measurements made in the three core lines. Soil preparation, liming, and fertilization were made following the recommendations for white oat in Brazil (Comissão Brasileira de Pesquisa de Aveia, 2006). The results of soil chemical analysis for each year are shown in Table 1. Tebuconazole fungicide was applied for the control of shoot disease, during the development cycle.

Chemical composition of white oat grains was evaluated as to the contents of protein, lipid, and nitrogen-free extract in the caryopsis. For these analyses, samples of 300 manually dehulled grains (length larger than 2 mm) were used. These grains were collected with a bulk harvest in each plot. Caryopses were ground in a Willey type mill with 0.25 mm sieve opening. Milled material was analyzed by a near-infrared reflectance spectrophotometer (NIRS) model 5000 (Perstorp Analytical Co., Maryland, USA). Calibration curves for the determination of chemical quality traits was obtained using the software NIRSystem (Infrasoft International, 1996), considering the analysis of 100 samples, following the recommendations of Cunniff (1995) and American Association of Cereal Chemists (1996). The protein value was obtained from the nitrogen content in the samples, using the correction factor of 6.25. NIRS readings were performed in triplicate and the results were expressed in dry weight basis (g per 100 g).

Analysis of variance was performed to identify the effects of cultivar, year, location, and of the interaction between these factors. Adaptability and stability parameters were estimated according to

Cruz et al. (1989) model, based on the bissegmented linear regression. This procedure has the average (β_{oi}) and the linear performance in unfavorable (β_{1i}) and favorable environments ($\beta_{1i} + \beta_{2i}$) as adaptability parameters. The genotype stability is evaluated by the regression deviation ($\sigma^2_{\delta_{ij}}$) regarding environmental variations, according to the model,

$$Y_{ij} = \beta_{oi} + \beta_{1i}I_j + \beta_{2i}T(I_j) + \delta_{ij} + \bar{\varepsilon}_{ij},$$

in which: Y_{ij} corresponds to the average of genotype i in the j environment; β_{oi} is the general average of genotype i over all tested environments; β_{1i} is the linear regression coefficient which evaluates the response of i^{th} genotype to the variation of unfavorable environments; I_j is the coded environmental index; β_{2i} is the linear regression coefficient which evaluates the i^{th} genotype response to the variation of favorable environments; $T(I_j) = 0$, if $I_j > 0$, or $T(I_j) = I_j + I_+$, if $I_j < 0$, being I_+ the average of I_j positive indexes; δ_{ij} is the regression deviations of genotype i in the environment j ; and $\bar{\varepsilon}_{ij}$ is the average experimental error. The means were compared with the Scott-Knott test, at 5% probability, in order to analyze the performance of genotypes in different crop seasons and cultivation sites. Additionally, the broad sense heritability (h_a^2), evidenced by the means of the characters in each environment tested, was determined according to Carvalho et al. (2001). All statistical procedures were performed with the aid of Genes software (Cruz, 2006).

Results and Discussion

Triple interaction between genotypes (G), cultivation year (Y), and locations (L) was significant

for all evaluated traits, except lipid contents (Table 2). No significant interaction between genotypes and locations was detected for the contents of protein and nitrogen-free extract (NFE). Further interactions between traits were all significant, as well as the effects of the major factors (G, Y, and L). According to these results, cultivars varied as to their contents of protein, lipid and NFE in the caryopsis, and the average performance of these traits is affected differently by the tested years and cultivation sites.

The highest mean square for lipid content was found for the cultivar effect (Table 2), indicating that genotype has a major role on the phenotypic expression of this trait, which agrees with Peterson et al. (2005). However, for the contents of protein and NFE, the highest mean squares were determined for cultivation year, followed by Y x L interaction. Beber et al. (2002) reported average values of 16.80 and 6.84 g per 100 g for the contents protein and lipid, respectively, in Brazilian oat cultivars. This is slightly lower than those observed in the present work, which were 18.71 and 7.88 g per 100 g.

Cultivar performances varied over the years and the cultivation sites (Figure 1 and 2). Lipid contents did not differ significantly over the years in many cultivars, in each location. This can be clearly observed in Capão do Leão and Passo Fundo. However, the contents of protein and NFE varied considerably between cultivars over the crop seasons in these locations.

In general, white oat stands out for its high level of proteins of excellent nutritional value, with balanced amino acid profile and high digestibility (Beber et al., 2002; Zhu et al., 2004). The highest mean performance for protein content was observed

Table 1. Soil chemical properties and meteorological conditions in three cultivation sites in the state of Rio Grande do Sul, Brazil.

Crop season	Location	Clay -----(%) -----	OM -----	SMP index	P ----- (mg dm ⁻³) -----	K	Temperature (°C) ⁽¹⁾			Precipitation (mm)
							Maximum	Mininum	Mean	
2007	Augusto Pestana	50.0	3.2	6.6	31.0	385.0	23.3	11.4	17.3	863.0
	Capão do Leão	24.0	2.2	6.8	29.1	143.0	19.5	10.8	15.7	798.8
	Passo Fundo	56.0	5.5	5.4	15.0	183.8	21.3	11.3	16.3	1,263.5
2008	Augusto Pestana	54.0	2.9	6.2	26.0	292.0	23.1	11.3	17.2	963.6
	Capão do Leão	16.0	1.5	6.8	8.6	64.0	20.4	11.7	16.0	575.3
	Passo Fundo	46.0	3.3	6.0	5.0	204.0	20.7	11.5	16.1	1,118.8
2009	Augusto Pestana	56.0	2.8	6.4	25.0	215.0	22.8	9.7	16.2	1,407.4
	Capão do Leão	21.0	2.1	6.5	19.7	80.0	20.0	11.0	15.0	952.0
	Passo Fundo	57.4	5.4	5.4	15.1	186.0	20.5	11.3	15.4	1,425.7

⁽¹⁾Data concerning the period from June to November of each year. OM, soil organic matter.

Table 2. Analysis of variance for the contents of protein, lipid, and nitrogen-free extract (NFE) in the caryopsis, measured in 15 white oat cultivars grown in three municipalities of the state of Rio Grande do Sul, Brazil, in the crop seasons of 2007, 2008, and 2009.

Source of variation	Df	Mean square		
		Protein	Lipid	NFE
Genotype (G)	14	13.309**	7.703**	43.496**
Year (Y)	2	153.029**	4.799**	817.798**
Location (L)	2	22.663**	3.713**	19.735**
G x Y	28	2.177**	0.545**	6.391**
G x L	28	0.817 ^{ns}	0.539**	2.899 ^{ns}
Y x L	4	94.192**	5.750**	101.253**
G x Y x L	46	2.507**	0.321 ^{ns}	5.459**
Block	3	2.309	0.045	6.057
Error	402	0.917	0.257	2.634
Average (g per 100 g)	-	18.709	7.877	67.563
Standard deviation	-	0.958	0.507	1.623
CV (%)	-	5.118	6.437	2.402

^{ns}Nonsignificant. ** and *Significant at 1 and 5% probability, respectively, by the F test. CV, coefficient of variation.

in the cultivars UPF 18 (19.80 g per 100 g), UFRGS 14 (19.58 g per 100 g), UPF 15 (19.44 g per 100 g), Albasul (19.25 g per 100 g), and UPFA 22 (19.14 g per 100 g) (Table 3). These cultivars can represent important sources of positive genes to increase grain protein in breeding programs for improving grain chemical quality. However, when it comes to cultivar recommendations, it is important to emphasize that these genotypes had variable adaptability and stability for this trait. Cultivars UPF 18, Albasul and UPFA 22 did not show adaptability to both unfavorable and favorable environments, since their β_{1i} values were equal or higher than 1.0, and their $\beta_{1i} + \beta_{2i}$ values were equal or smaller than 1.0. According to Cruz et al. (1989), the model $\beta_{1i} < 1.0$ represents adaptability to unfavorable environments, and $\beta_{1i} + \beta_{2i} > 1.0$ corresponds to adaptability to favorable

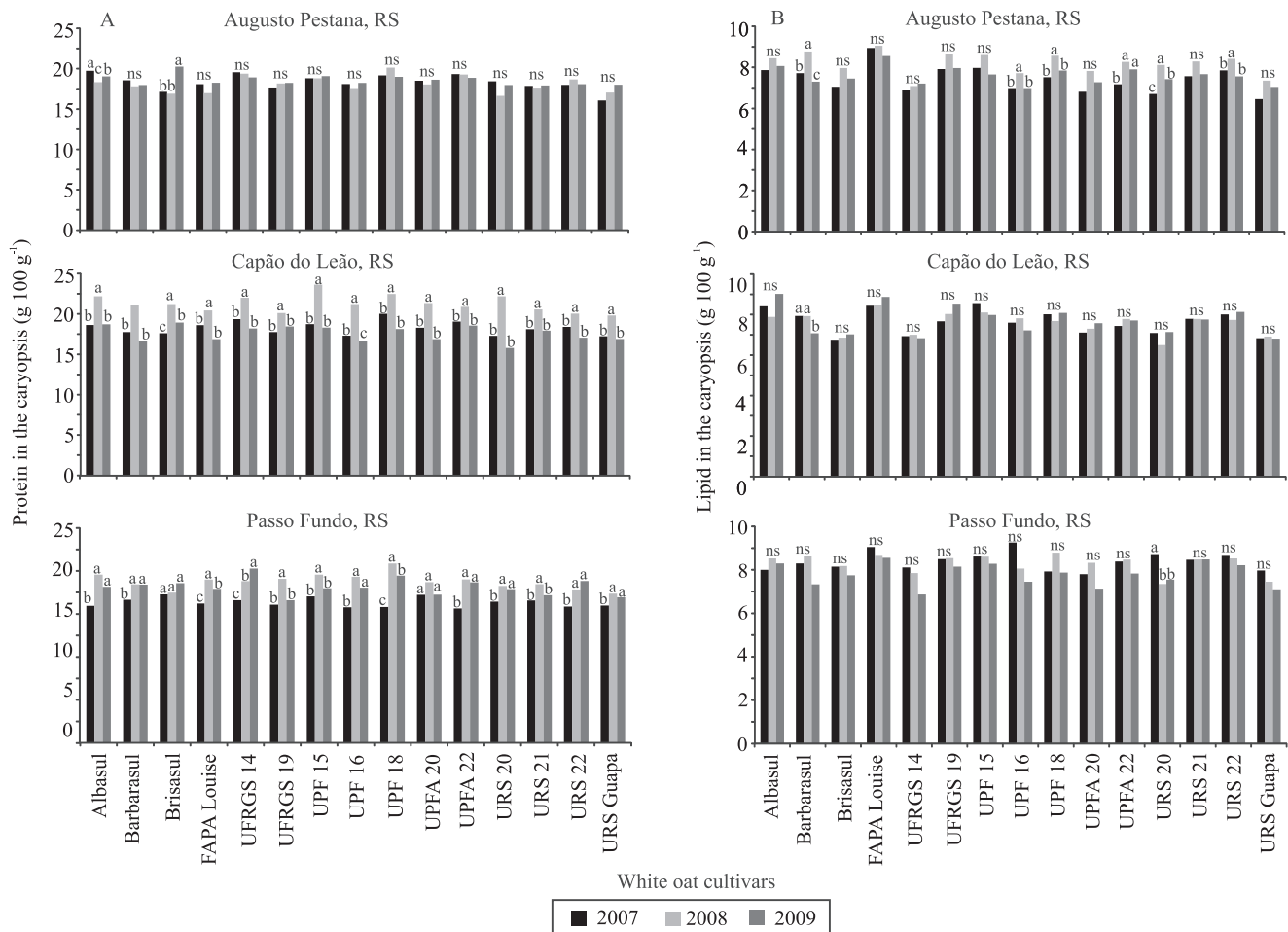


Figure 1. Performance of white oat cultivars in relation to the contents of protein (A), and lipid (B) in the caryopsis (g per 100 g). Means followed by equal letters do not differ by Scott-Knott test, at 5% probability. ^{ns}Nonsignificant.

environments. The cultivars URS 20 and UPFA 15 were adapted to favorable environments, and Brisasul, UFRGS 19, and URS 21, to unfavorable environments. However, the cultivars Brisasul and UFRGS 19, together with UPF 18 and URS 20, were unstable, i.e. they showed significant regression deviations ($\sigma^2_{\delta_i} \neq 0$), which indicate unpredictable behavior across the environments. The lowest mean protein content was shown by 'URS Guapa' (17.60 g per 100 g), which was stable in the tested environments.

The lowest environment index (I_j) for protein content (-1.44) was detected in 2007, in Passo Fundo, where the lowest average performance (17.27 g per 100 g) occurred (Table 4). The highest mean was

obtained in Capão do Leão, in 2008 ($I_j = 2.55$; 21.26 g per 100 g).

According to the model of Cruz et al. (1989), an ideal genotype would be a cultivar which concomitantly shows high protein content (β_{oi}), adaptability to unfavorable ($\beta_{li} < 1.0$) and favorable environments ($\beta_{li} + \beta_{2i} > 1.0$), and stability ($\sigma^2_{\delta_i} = 0$). No ideal genotype was found among the analyzed cultivars, which met these criteria.

As to lipid content in the caryopsis, cultivars URS Guapa and UFRGS 14 showed the lowest performance (7.12 and 7.22 g per 100 g, respectively) while the highest average were found in FAPA Louise (8.77 g per 100 g) (Table 3). Only 'URS 20' was adapted to favorable environments. However, this cultivar had an unstable behavior, along with Albasul, Barbarasul and UPF 16. The lowest environment index for genotype performance as to lipid content was detected in 2007, in Augusto Pestana ($I_j = -0.45$), where the lowest average performance (7.43 g per 100 g) was observed (Table 4). The best favorable environment was Augusto Pestana, in 2008 ($I_j = 0.33$, average of 8.21 g per 100 g).

Nitrogen-free extract had an overall average performance of 67.56 g per 100 g, ranging from 65.54 to 69.75 g per 100 g (Tables 2 and 3). Gatto (2005) observed a range of NFE contents of 69.8 to 75.7 g per 100 g, in the caryopses of 22 Brazilian white oat genotypes grown in Southern Brazil. 'Albasul' and 'UPF 18' were adapted only to favorable environments. However, these cultivars showed low performance (66.04 and 65.54 g per 100 g, respectively) and low stability. Higher performance was shown by 'URS Guapa' (69.75 g per 100 g) and 'URS 20' (68.93 g per 100 g); however, they showed low adaptability, and the cultivars URS 20, Brisasul and UFRGS 19 had low stability as well. The lowest environment index for nitrogen-free extract was observed in 2008, in Capão do Leão ($I_j = -3.64$; 63.93 g per 100 g), and the best favorable environment was Augusto Pestana ($I_j = 2.77$; 70.33 g per 100 g) in 2007 (Table 4).

Alternative models of stability and adaptability analysis adopting nonparametric methods have been used for genotype evaluations of autogamous annual species (Backes et al., 2005; Amorin et al., 2006; Elias et al., 2007). These methods can represent effective alternatives to evaluate the stability and

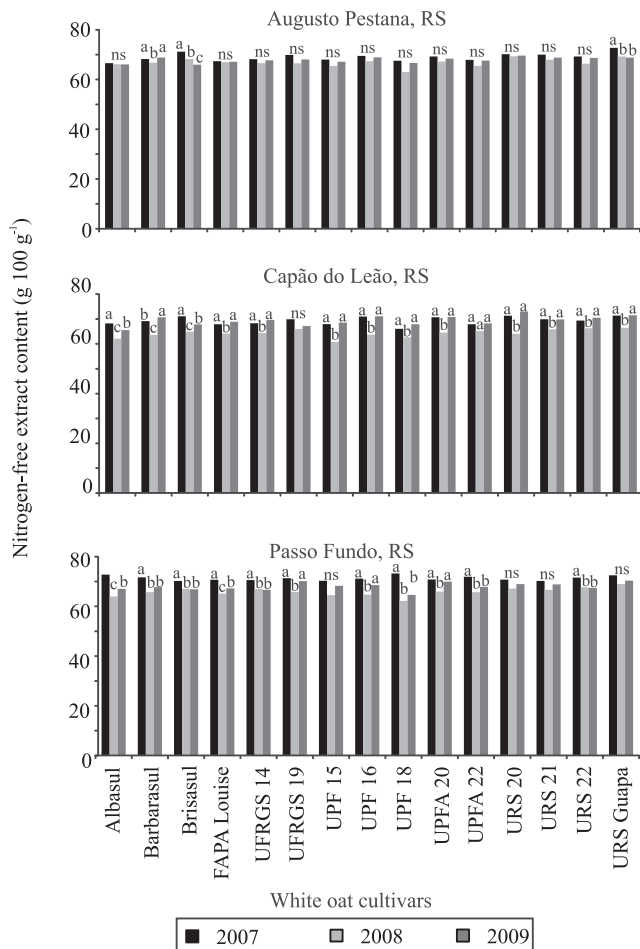


Figure 2. Performance of white oat cultivars as to the nitrogen-free extract contents in the caryopsis (g per 100 g). Means followed by equal letters do not differ by Scott-Knott test, at 5% probability. ^{ns}Nonsignificant.

Table 3. General averages (b_0 ; g per 100 g) of protein, lipid, and nitrogen-free extract content in the caryopsis and parameters of adaptability (b_1 , ($b_1 + b_2$)) and stability ($\sigma_{\delta_i}^2$) according to Cruz et al. (1989) model, in white oat cultivars grown in three municipalities of the state of Rio Grande do Sul, Brazil, in the crop seasons of 2007, 2008, and 2009⁽¹⁾.

Cultivar	b_0	Average (U)	Average (F)	b_1	$b_1 + b_2$	$\sigma_{\delta_i}^2$	R ²
Protein content in the caryopsis							
Albasul	19.25a	18.54	20.68	1.09 ^{ns}	1.38 ^{ns}	1.50 ^{ns}	87.68
Barbarasul	18.48b	17.71	20.01	1.07 ^{ns}	0.84 ^{ns}	1.01 ^{ns}	89.54
Brisasul	18.71b	18.18	19.77	0.69*	0.93 ^{ns}	6.70**	39.81
FAPA Louise	18.37b	17.65	19.81	1.05 ^{ns}	0.64 ^{ns}	1.01 ^{ns}	88.57
UFRGS 14	19.58a	18.82	21.10	1.13 ^{ns}	0.45 ^{ns}	1.91 ^{ns}	82.09
UFRGS 19	18.34b	17.87	19.27	0.64*	1.03 ^{ns}	2.17*	66.70
UPF 15	19.44a	18.62	21.10	1.14 ^{ns}	2.19**	0.16 ^{ns}	99.03
UPF 16	18.36b	17.42	20.23	1.34*	0.95 ^{ns}	0.44 ^{ns}	96.81
UPF 18	19.80a	18.85	21.69	1.45**	0.81 ^{ns}	2.76**	84.26
UPFA 20	18.64b	18.08	19.77	0.80 ^{ns}	1.44 ^{ns}	1.01 ^{ns}	87.97
UPFA 22	19.14a	18.60	20.23	0.90 ^{ns}	0.53 ^{ns}	1.89 ^{ns}	75.46
URS 20	18.20b	17.24	20.12	1.34*	1.64*	2.36*	87.07
URS 21	18.35b	17.82	19.40	0.70*	1.08 ^{ns}	0.41 ^{ns}	92.31
URS 22	18.41b	17.82	19.59	0.91 ^{ns}	0.17**	1.23 ^{ns}	81.58
URS Guapa	17.60c	17.02	18.68	0.75 ^{ns}	0.92 ^{ns}	1.34 ^{ns}	78.64
Lipid content in the caryopsis							
Albasul	8.32b	7.97	8.49	0.94 ^{ns}	1.09 ^{ns}	1.18**	25.42
Barbarasul	7.92c	7.36	8.20	1.43 ^{ns}	0.52 ^{ns}	0.90**	49.03
Brisasul	7.48d	7.32	7.56	0.46 ^{ns}	2.27 ^{ns}	0.40 ^{ns}	36.19
FAPA Louise	8.77a	8.58	8.86	0.46 ^{ns}	1.95 ^{ns}	0.43 ^{ns}	30.50
UFRGS 14	7.22e	6.91	7.38	0.88 ^{ns}	1.19 ^{ns}	0.29 ^{ns}	55.14
UFRGS 19	8.25b	7.91	8.42	0.91 ^{ns}	1.98 ^{ns}	0.29 ^{ns}	60.40
UPF 15	8.30b	7.87	8.51	1.09 ^{ns}	0.04 ^{ns}	0.34 ^{ns}	59.72
UPF 16	7.70d	7.05	8.03	1.72*	1.04 ^{ns}	0.81**	61.11
UPF 18	8.06c	7.65	8.27	1.15 ^{ns}	0.98 ^{ns}	0.42 ^{ns}	58.00
UPFA 20	7.49d	6.98	7.74	1.40 ^{ns}	0.38 ^{ns}	0.20 ^{ns}	80.34
UPFA 22	7.91c	7.54	8.10	1.08 ^{ns}	0.60 ^{ns}	0.18 ^{ns}	74.03
URS 20	7.42d	7.13	7.56	0.84 ^{ns}	5.24**	0.62*	62.29
URS 21	8.06c	7.80	8.18	0.69 ^{ns}	0.38 ^{ns}	0.14 ^{ns}	58.34
URS 22	8.15b	7.77	8.34	0.97 ^{ns}	1.20 ^{ns}	0.13 ^{ns}	76.78
URS Guapa	7.12e	6.78	7.30	0.99 ^{ns}	0.69 ^{ns}	0.19 ^{ns}	69.31
Nitrogen-free extract content in the caryopsis							
Albasul	66.04d	64.26	67.47	1.07 ^{ns}	2.41**	7.29*	83.03
Barbarasul	67.60b	65.39	69.37	1.18 ^{ns}	0.78 ^{ns}	1.92 ^{ns}	94.08
Brisasul	67.71b	66.21	68.91	0.79 ^{ns}	1.26 ^{ns}	12.09**	57.44
FAPA Louise	66.82c	65.32	68.03	0.85 ^{ns}	1.07 ^{ns}	1.22 ^{ns}	93.49
UFRGS 14	67.22c	65.60	68.52	0.80 ^{ns}	0.77 ^{ns}	2.14 ^{ns}	87.40
UFRGS 19	67.87b	66.55	68.92	0.76 ^{ns}	0.97 ^{ns}	6.90*	67.02
UPF 15	66.33d	64.20	68.04	1.31*	0.91 ^{ns}	3.29 ^{ns}	92.01
UPF 16	68.00b	65.54	69.97	1.36*	0.47 ^{ns}	1.55 ^{ns}	96.19
UPF 18	65.54d	62.57	67.92	1.37*	2.43**	10.36**	83.14
UPFA 20	68.17b	66.34	69.64	1.07 ^{ns}	0.52 ^{ns}	2.56 ^{ns}	90.60
UPFA 22	67.08c	65.49	68.35	0.79 ^{ns}	1.43 ^{ns}	2.51 ^{ns}	87.19
URS 20	68.93a	66.80	70.63	1.23 ^{ns}	0.01*	5.72*	84.76
URS 21	68.24b	66.78	69.41	0.80 ^{ns}	0.16 ^{ns}	0.64 ^{ns}	95.51
URS 22	68.13b	66.38	69.52	0.76 ^{ns}	0.77 ^{ns}	3.15 ^{ns}	80.90
URS Guapa	69.75a	68.17	71.02	0.88 ^{ns}	1.06 ^{ns}	3.54 ^{ns}	83.95

⁽¹⁾Means followed by equal letters do not differ by Scott-Knott test, at 5% probability. ^{ns}Nonsignificant. ** and *Significant at 1 and 5% probability, respectively, by t test. (U), unfavorable environment; (F), favorable environment.

adaptability of genotypes and show advantages over parametric models: reduction of the tendentiousness caused by points way out of the fitted regression; no assumption requirement about the characteristics of the phenotypic value distribution; parameters are easy to use and interpret; and low sensibility to the addition or removal of one or few genotypes from the analysis (Cruz, 2006).

There is little information about environmental effects that can interfere in the NFE contents of oat grains. Much of this fraction is formed by starch, which correspond to nearly 60% dry matter of the oat grain (Liu et al., 2010). Rhymer et al. (2005) observed that the environment was the dominant factor contributing to the total variation in starch contents of oat genotypes. Many reports indicate that environmental effects may interfere in the protein and oil contents in the oat caryopsis (Beber et al., 2002; Zhu et al., 2004; Peterson et al., 2005; Martinez et al., 2010). Humphreys et al. (1994) and by Kolchinski & Schuch (2003, 2004) reported that application of nitrogen fertilizer increases protein content in the grains, and decreases oil content. Moreover, temperature variation during oat development may have great influence on lipid synthesis in the grain. According to Martinez et al. (2010), the increase of total lipid content is favored by lower temperatures and, in these conditions, the synthesis of unsaturated lipids and the level of unsaturation are increased.

Despite the amplitude in the performance observed in the chemical composition of white oat caryopsis (Figure 1 and 2), a general behavior can be noticed along favorable and unfavorable environments (Table 4). In general, cultivars characterized by a higher or a lower average performance for a specific character tended to have this behavior in both environmental conditions. The highest amplitudes were observed for protein contents, in which the average performance of cultivars was 7.7% lower than the overall average, in unfavorable environment, and 13.6% higher than the overall average, in the most favorable environment. For the contents of lipid and NFE, variation was close to 5% on the overall average. Passo Fundo in 2007 and Augusto Pestana in 2009 represented the environments with the most effect on the genotypes performance, which was evidenced by the lowest heritabilities (<50%) observed for all evaluated characters. However, in general, the heritabilities found were of high magnitude, higher than 70% in seven of the nine tested environments, suggesting that the genetic variance played a major role in the phenotypic variance.

There are evidences in the literature that these traits have polygenic inheritance, with strong contribution of additive effects, and that the environmental contribution for the phenotypic definition vary (Zhu et al., 2004; Orr & Molnar, 2007; Crestani et al., 2012).

Table 4. General averages (b_0 ; g per 100 g) of protein, lipid, and nitrogen-free extract content in the caryopsis, environment index (I_j) according to Cruz et al. (1989) model, and broad sense heritability (h_a^2) in white oat cultivars grown in three municipalities of the state of Rio Grande do Sul, Brazil, in the crop seasons of 2007, 2008, and 2009.

Season	Augusto Pestana			Capão do Leão			Passo Fundo		
	Average	I_j	h_a^2 (%)	Average	I_j	h_a^2 (%)	Average	I_j	h_a^2 (%)
Protein content in the caryopsis									
2007	18.33	0.38 (U)	79.85	18.26	-0.45 (U)	74.34	17.27	-1.44 (U)	31.13
2008	18.10	-0.61 (U)	91.97	21.26	2.55 (F)	80.27	19.85	1.14 (F)	81.52
2009	18.56	-0.15 (U)	40.91	17.58	-1.13 (U)	83.32	19.17	0.47 (F)	90.74
Lipid content in the caryopsis									
2007	7.43	-0.45 (U)	91.93	8.02	0.14 (F)	89.93	8.08	0.21 (F)	30.13
2008	8.21	0.33 (F)	94.91	7.96	0.09 (F)	93.03	7.99	0.12 (F)	72.77
2009	7.59	-0.28 (U)	48.34	8.10	0.23 (F)	92.73	7.50	-0.38 (U)	77.49
Nitrogen-free extract content in the caryopsis									
2007	69.06	1.49 (F)	84.41	68.94	1.38 (F)	85.29	70.33	2.77 (F)	35.59
2008	66.84	-0.72 (U)	89.33	63.93	-3.64 (U)	75.90	64.98	-2.59 (U)	79.66
2009	67.90	0.34 (F)	24.58	69.01	1.45 (F)	87.17	67.09	-0.48 (U)	71.52

(U), unfavorable environment; (F), favorable environment.

Conclusions

1. White oat performance as to grain chemical composition is affected by cultivation site, crop season, and genotypes.

2. There is no ideal genotype, among the evaluated cultivars, with high average performance, high adaptability, and high stability, simultaneously.

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Received on April 14, 2012 and accepted on January 7, 2013