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Jieyu Zhang , Xiaolong Li , [Junzhen Wang](#) , Yang Li , Qiaohui Yang , [Dabing Xiang](#) , Yan Wan , Eviatar Nevo , Jun Yan , Fan Yu , [Liang Zou](#) *

Posted Date: 14 September 2023

doi: 10.20944/preprints202309.0934.v1

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Article

Wild Oats Offer New Possibilities for Forage Because of the Higher Nutrition Content and Feed Value

Zhang Jieyu ¹, Li Xiaolong ¹, Wang Junzhen ², Li Yang ¹, Yang Qiaohui ¹, Xiang Dabing ¹, Wan Yan ¹, Nevo Eviatar ³, Yan Jun ¹, Fan Yu ^{1,*} and Zou Liang ^{1,*}

¹ Key Laboratory of Coarse Cereal Processing, Ministry of Agriculture and Rural Affairs, Sichuan Engineering & Technology Research Center of Coarse Cereal Industrialization, College of Food and Biological engineering, Chengdu University, Chengdu 610106, People's Republic of China

² Sichuan Liangshan Yi Autonomous Prefecture Agricultural Science Research Institute, Liangshan Prefecture 615000, P.R. China

³ Institute of Evolution, University of Haifa, Haifa 31905, Israel

* Correspondence: Zou Liang Chengdu University, Longquanyi District, Chengdu City, Sichuan Province, P.R. China Tel.: 028-84616653; Email: zouliang@cdu.edu.cn; Yu Fan Chengdu University, Longquanyi District, Chengdu City, Sichuan Province, P.R. China Tel.: 028-84616653; Email: fandavi@163.com

Abstract: To explore the forage quality of wild oat populations under different growth environments, the basic nutrient composition and forage quality of five wild oat populations in Israel were compared. Ash, crude protein, and soluble sugar content of oats were significantly ($P < 0.05$) related to some forage quality and possible environment of population origin. Among them, the environmental factors of origin had significant effects on the indicators of ash, crude protein, and soluble sugars of oat straw, but less on the contents of crude fat, total phosphorus, and total potassium. Meanwhile, the annual rainfall and the number of days of rainfall in the country of origin greatly influenced the cellulose content. Finally, we investigated the similarity of feeding values of different populations using PCA analysis. The results showed that, in conclusion, the higher variability of wild oat populations may be a new possibility brought by feeding oats.

Keywords: wild oats; feeding value; Germplasm resource; environmental factor; origin

1. Introduction

Oats are important annual forage and food crop of the genus oats in the family Gramineae, divided into two types: naked oats (*Avena nuda*) and leather oats (*A. Sativa*), which are grown all over the world. Oat cultivation in China is mainly distributed in northeast, north, and southwest China (Masood et al., 2008), and is recognized as one of the world's eight major food crops and an excellent forage crops (Masood et al., 2008). Oat seeds are very rich in protein, fat and other nutrients and can be an excellent dietary product, with a protein content of more than 15.0% and a fat content of 8.8% (Meen et al., 2017; Masood et al., 2008). Oats do not have harsh growing conditions, and can be grown in poor and arid areas, but also because of its advantages high yield is often planted in pasture or high Beit-Orenude areas as livestock feed (Li et al., 2013; Li et al., 2011; Xiao et al., 2017). It is also often grown as livestock feed in pastoral areas or at high Beit-Orenudes for its high yield. About 73% of the world's oat production is forage, mainly for processing and eating only 12%, so oats are an important forage quality raw material in the world (Biel et al., 2020; Suttle et al., 2004.; Varma et al., 2016; Wang et al., 2002).

The nutritional content of crude protein, crude fat, starch, and β -glucan varied somewhat between oat varieties, and the starch pasting characteristics and thermodynamic properties were influenced by different source environments (Chen et al., 2020; John, 1995; Francis W & Associates, 2011). The forage quality of oats is influenced by different factors, and the available studies show that: oats require a large amount of water during the growing period, and the water supply status has an important influence on the growth and development, yield, and quality formation of oats (Sojka et al., 1998; Sui et al., 2018). The water requirements of oats are different in

different fertility stages. Fixed drip irrigation can effectively increase the crude protein, crude fat, and glucan content of oats (Enciso et al., 2007; Ngouajio et al., 2007; Oktem et al., 2003; Wu et al., 2009). It has also been shown that the selection of the appropriate sowing stage has a positive impact on the quality of oats. It has also been shown that the selection of suitable sowing and mowing periods can improve the yield and nutritional quality of oats (Li et al., 1996; Li et al., 2015; Ma et al., 2010a, 2010b; Yue & Wang, 2019). The increase in cumulative temperature during the growing period had a significant effect on the growth of oat forage (Akhtar et al., 2006; Costa et al., 2014; Du et al., 2008), and the increase in cumulative temperature from nodulation to tiller was not conducive to the formation of oat forage yield and quality (Robertso et al., 2004), but the increase in cumulative temperature between seedling and tiller could improve the content of crude fat and acidic detergent fiber (Li et al., 2001; Zhang et al., 2002; Zhou et al., 2021). However, the increase in temperature between seedling and tillering increased crude fat and acidic detergent fiber content. Israel is one of the origins of wheat crops (including oats), and its wild oats have rich genetic diversity and are excellent genetic resources for improving forage oat varieties. Previous studies have shown that oats from Israel vary greatly in nutritional composition between groups, presumably due to their genetics and differences in the growth environment (Loskutov et al., 2017; Welch et al., 1997; Xu et al., 2020; Zong et al., 2017). However, further research is needed on the differences in nutrients such as ash, crude fat, etc. between groups.

Currently, there have been many studies on the nutritional quality of oat forage, but there are fewer studies on the nutritional composition and forage quality of different oat populations in their places of origin. In this study, we analyzed the nutritional composition and forage quality of 18 genotypes of oats from six oat populations and explored the potential relationship between them based on the environmental factors of different populations' origins to provide a reference for oat forage quality improvement and breeding.

2. Materials and methods

2.1. Test materials

A total of six oat populations were used in this experiment. Five wild oat populations were collected from Israel by Prof. Yan Jun of Chengdu University, and one cultivated oat population was provided by Prof. Hu Yinguang of Northwest Agriculture and Forestry University of Science and Technology, respectively. The ecogeographic data of different populations are shown in Table 2.

2.2. Test methodology

The experiment was conducted on 2018.10-2019.03, 2019.10-2020.03, 2020.10-2021.03 with 60 seeds of uniform size and intact shape from each of six cultivated oat varieties and 18 wild oat genotypes. Seeds were sown in strips using a randomized partitioned design with 20 seeds per row, a row length of 1.5 m, row spacing of 50 cm, setting three replications, and making protected rows. Oats were mowed in January 2019-January 2021 during the tasseling period, harvested and killed at 105°C for 15 min, dried at 75°C to constant weight, crushed with a plant grinder, passed through a 40 mesh sieve, and stored in self-sealing bags at room temperature, protected from light.

2.3. Measurement indicators and methods

Crude protein (CP), crude fat (Ether extract, EE), ash content (Ash) (Li et al., 2016; Yuan et al., 2021), neutral detergent fiber (NDF) (Rohwede & Barnes, 1978), acid detergent fiber (ADF) (Rohweder & Barnes, 1978) of oats were The content of crude fat (EE) (Li et al., 2016), ash content (Ash), neutral detergent fiber (NDF), and acid detergent fiber (ADF) were determined using a FOSS NIR quality analyzer.

Water soluble carbohydrates (WSC) was extracted by anthrone colorimetry and then measured by UV spectrophotometer; Total phosphorus (TP) was measured by ammonium molybdate spectrophotometry; Total calcium (TK) was measured by atomic absorption spectrometry. Total phosphorus (TP) was determined by ammonium molybdate spectrophotometry; Total potassium (TK)

was determined by atomic absorptionspectrometry(Guo et al., 2003); Inorganic phosphorus (IP) was measured spectrophotometrically(Zheng et al., 2017).

Relative feeding quality is calculated using the equation, see equation (2.1) and equation (2.2)(Zhu et al., 2019).

Relative Feeding Quality (RFQ)

$$RFQ = TDN(\%DW) \times DMI(\%BW)/1.23 \text{ (2.1)}$$
The prediction model equation for total digestible nutrients (TDN) is.

$$TDN (\%DM) = 82.38 - (0.7515 \times ADF) \text{ (2.2)}$$

2.4. Data analysis

Data were collated using Excel, Turkey-Kramer HSD (P<0.05) using JMP Pro to detect the significance of differences, and Spielman rank correlation coefficient and ANOVA were used to analyze the correlation between oat nutrient content, foragequality, and place of origin.

Table 1. Oat material and its origin.

Community	Type	Place of Origin	Community	Type	Place of Origin
Tabigha	wild oats	Israeli	Evolution Canyon	wild oats	Israeli
Nahef	wild oats	Israeli	Beit-Oren	wild oats	Israeli
Sede Boqer	wild oats	Israeli	Hu	Cultivated oats	Hebei, China

Table 2. Ecogeographic data on the origin of the six oat origin.

Population	Ln/°	Lt/°	Al/m	Tm/°C	Ta/°C	Tj/°C	Td/°C	Tdd/°C	Ev/m	Rn/m	Rd/d	Hu14/%	Huan/%
Tabigha	35.5	32.9	0.0	23.6	31.9	14.3	17.4	10.3	164.0	437.0	49.0	43.8	58.2
Nahef	35,3	32.9	275.0	15.3	23.5	8.3	15.7	8.9	156.4	662.0	53.0	50.0	62.1
Sede Boqer	34,9	32.5	10.0	20.1	25.9	13.4	13.5	8.9	132.0	539.0	45.0	66.2	72.3
Evolution Canyon	34.6	32.4	90.0	22.7	27.2	14.1	13.2	9.2	144.0	602.0	50.2	65.9	65.4
Beit-Oren	35.0	32.7	50.0	20.2	26.2	12.7	13.0	9.0	133.0	507.0	48.2	65.5	72.5
Hu	114.5	40.5	1300.0	9.2	23.3	-8.2	31.2	14.2	18.7	31.9	6.4	13.0	15.2

Note: Ln: longitude; Lt: latitude; Al: Beit-Orenude; Tm:three-year annual mean temperature; Ta: three-year August mean temperature; Tj: three-year January mean temperature; Td: three-year seasonal temperature difference; Tdd: three-year diurnal temperature difference; Ev: three-year annual evaporation; Rn: three-year annual rainfall; Rd: three-year annual number of days of rainfall; Hu14%: three-year 14:00 mean fitness; Huan: three-year annual mean humidity, same below.

3. Results and analysis

3.1. Nutritional content analysis of different oat populations

In this study, six oat populations with 18 genotypes sown in fall 2018, fall 2019, and fall 2021 were tested for nutrients such as ash, crude fat, crude protein, soluble sugars, total phosphorus, and total potassium (Table 3). The results showed that the ash content of the Sede Boqer and Evolution Canyon oat populations was significantly higher than the other populations, reaching 9.25% and 9.04% (average data of 3 years), respectively. the crude fat content of the Beit-Oren (9.82%) was significantly lower than the other five wild populations, differing by 8.24% from the Tabigh, which

had the highest crude fat content. The crude protein content of Hu, Nahef was lower, differing by 11.96% and 18.96% from the Evolution Canyon population, which had the highest content. In terms of soluble sugars, the Tabigha population had the highest content of 21.39%; the Hu had the lowest content of 10.51%, with a difference of 50.87% and a significant difference between populations. In terms of total phosphorus, the Nahef population had the lowest total phosphorus content of 0.31%. Total potassium and total phosphorus contents did not differ significantly among the six populations. The breeding components of evolved canyon oat populations differed significantly among the six populations with the highest content of each indicator, which could be potentially useful.

To investigate the differences in nutrient composition among populations in more depth, the genotypic nutrient composition among populations was studied. Eighteen genotypes from six populations were analyzed for nutrient content, and data were analyzed for the five genotypes with the highest mean and the five genotypes with the lower mean (Figure 1). The results showed that the five genotypes with higher ash content, Ecg13, Ecg12, Ecg1, Se13, and Se16, were all around 10%, about 1.07% higher than the mean. The ash content of wild oats Evolution Canyon and Sede Boqer was higher. There was no significant difference in the crude fat content of oats, suggesting that the effect of stock origin on crude fat may be minimal. Nah1, which had the lowest crude protein content, was 7.39% lower than Ecg12. The crude protein content between Be1 and Ecg1 was close, at 11.32% and 11.44%, respectively. The XO-1-3 genotype of oats had nearly 40% lower soluble sugar content than the highest Se6. Nah10 had 0.89% total phosphorus content, much higher than the average value of 0.43%. XO-1-60 had a lower total potassium content of 2.4%, while Nah2 had the highest total potassium content of 4.15%.

Table 3. Analysis of nutritional content of different oat populations.

Year	Population		Ash/%	EE/%	CP/%	WSC/%	TP/%	TK/%
2018.10-2019.01	Tabigha	Mean	8.33±1.21	8.25±0.58	11.52±1.39	26.63±4.20 a	0.40±0.08	3.37±0.39
			cd	a	b		b	a
		Range	6.67~9.42	7.57~9.04	9.67~12.92	27.76~31.48	0.30~0.48	2.84~3.80
	Nahef	CV/%	13.71%	6.63%	11.37%	14.86%	18.28%	10.97%
		Mean	7.35±2.27	7.89±0.14	10.80±1.99	13.91±4.59 b	0.52±0.51	3.52±0.51
			d	bc	b		a	a
	Sede Boqer	Range	4.43~9.96	7.26~8.18	8.19~12.68	9.12~19.68	0.27~1.86	2.99~4.17
		CV/%	29.15%	3.82%	17.41%	31.14%	16.58%	13.63%
	Evolution Canyon	Mean	9.84±0.26	7.96±0.48	12.68±0.71	25.67±3.22 a	0.41±0.046	3.78±1.13
			a	a	ab		b	a
		Range	9.47~10.06	7.34~8.38	12.04~13.68	21.37~28.36	0.35~0.45	0.35~4.08
	Evolution Canyon	CV/%	2.48%	5.73%	5.32%	11.84%	10.54%	7.18%
		Mean	9.23±0.10	7.98±0.27	14.01±1.95	26.26±6.21 a	0.41±0.04	3.21±0.46
			a	a	a		b	a

2019.1 0- 2020.0 1	Beit-Oren	Rang e	9.81~10.08	7.03~9.08	11.39~15.7 9	17.94~30.97	0.27~0.46	2.76~3.48
		CV/ %	0.92%	10.21%	13.14%	22.30%	8.74%	13.49%
		Mea n	8.75±0.49 bc	7.46±0.18 d	12.01±0.79 ab	20.94±13.32 abc	0.43±0.04 b	3.20±0.21 ac
		Rang e	7.99~9.21	7.19~7.67	11.24~13.1 0	4.11~34.35	0.38~0.49	2.67~3.48
		CV/ %	5.25%	2.34%	6.22%	59.99%	7.61%	6.15%
		Mea n	9.39±0.23 ab	8.00±0.49 ab	10.35±0.63 b	14.28±12.52 b	0.40±0.06 b	3.12±0.54 a
	Hu	Rang e	9.07~9.76	7.36~8.67	9.51~11.32	0.08~29.97	0.34~0.48	3.12±0.54 ab
		CV/ %	2.34%	5.68%	5.21%	82.66%	13.65%	16.33%
		Mea n	8.28±0.71 cd	8.27±0.28 a	6.12±0.53 b	19.76±1.18 b	0.33±0.13 ab	2.55±0.24 b
		Rang e	6.64~9.52	7.45~9.01	5.48.75	17.78~21.56	0.19~0.49	2.28~2.85
		CV/ %	13.71%	6.55%	8.15%	9.18%	36.71%	9.02%
		Mea n	7.35±2.27 d	7.91±0.32 ab	5.82±0.78 b	11.37±2.69 b	0.27±0.08 b	2.91±0.39 a
	Nahef	Rang e	4.43~9.96	7.53~8.32	4.77~6.44	8.18~14.51	0.20~0.37	2.61~3.44
		CV/ %	29.15%	3.79%	12.57%	22.29%	26.58%	12.54%
		Mea n	9.87±0.44 a	7.93±0.28 ab	9.02±2.87 a	11.59±5.82 b	0.34±0.10 ab	2.92±0.41 a
		Rang e	9.38~10.11	7.31~8.44	5.55~12.23	6.71~19.30	0.20~0.41	2.45~3.40
		CV/ %	2.48%	5.73%	29.97%	47.35%	27.35%	13.25%
		Mea n	9.93±0.10 a	7.86±0.16 a	8.51±2.88 a	13.53±2.27 b	0.27±0.01 ab	2.44±0.25 b
	Sede Boqer	Rang e	9.81~10.08	7.03~9.08	5.79~12.23	11.31~16.47	0.26~0.29	2.10~2.69
		CV/ %	0.92%	10.21%	31.88%	15.83%	4.01%	9.73%
	Evolutio n Canyon	Mea n	9.93±0.10 a	7.86±0.16 a	8.51±2.88 a	13.53±2.27 b	0.27±0.01 ab	2.44±0.25 b
		Rang e	9.81~10.08	7.03~9.08	5.79~12.23	11.31~16.47	0.26~0.29	2.10~2.69
		CV/ %	0.92%	10.21%	31.88%	15.83%	4.01%	9.73%
		Mea n	9.93±0.10 a	7.86±0.16 a	8.51±2.88 a	13.53±2.27 b	0.27±0.01 ab	2.44±0.25 b

2020.10-2021.01	Beit-Oren	Mea	8.77±0.23	7.47±0.12	6.78±0.80 b	12.07±1.17 b	0.33±0.05	2.64±0.14
		n	bc	b			ab	ab
		Rang	7.99~9.21	7.08~7.87	5.97~7.80	11.04~13.74	0.326~0.38	2.47~2.84
		e						
	Hu	CV/	5.25%	2.34%	11.11%	9.66%	15.51%	4.89%
		%						
		Mea	9.49±0.32	8.07±0.49	7.44±0.70	9.50±4.25 a	0.35±0.05	2.57±0.30
		n	ab	a	ab		a	b
	Tabigha	Rang	9.18~9.77	7.37~8.66	6.72~8.33	3.00~14.67	0.28~0.40	2.17~2.82
		e						
		CV/	2.27%	5.73%	8.81%	44.42%	12.89%	11.13%
		%						
	Nahef	Mea	7.51±0.55	8.36±0.50	11.38±0.42	17.78±1.37 a	0.20±0.00	2.63±0.09
		n	b	a	a		a	a
		Rang	7.00~8.32	7.66~8.88	10.84~11.8	8.07~12.47	0.19~0.20	2.53~2.76
		e			4			
	Sede Boqer	CV/	6.95%	5.62%	3.52%	11.98%	2.35%	3.27%
		%						
		Mea	7.85±0.85	7.98±0.89	9.54±0.56 c	18.08±5.31 a	0.14±0.03	2.28±0.27
		n	ab	ab			bc	b
	Evolution Canyon	Rang	6.71~8.58	6.89~8.98	8.77~10.01	13.10~26.72	0.10~0.16	1.94~2.56
		e						
		CV/	10.16%	10.47%	5.57%	27.69%	18.16%	11.00%
		%						
	Beit-Oren	Mea	8.07±0.38	7.88±0.23	9.36±0.22 c	17.82±2.51 a	0.12±0.01	2.06±0.08
		n	a	ab			c	c
		Rang	7.63~8.63	7.56~8.10	9.03~9.59	14.24~22.03	0.11~0.13	1.96~2.16
		e						
	Beit-Oren	CV/	4.44%	2.75%	2.25%	13.30%	6.37%	3.81%
		%						
		Mea	7.95±0.52	7.21±1.47	9.78±0.31 c	16.00±4.19	0.14±0.01	2.28±0.09
		n	ab	b		ab	bc	b
	Beit-Oren	Rang	7.24~8.74	6.05~9.21	9.35~10.08	7.66~20.79	0.13~0.15	2.16~2.38
		e						
		CV/	6.20%	19.19%	2.95%	24.71%	5.96%	3.67%
		%						
	Beit-Oren	Mea	7.74±0.38	7.91±0.90a	7.64±0.08 d	14.79±1.48 b	0.07±0.01	1.85±0.16
		n	ab	b			d	d
		Rang	7.21~8.29	6.64~8.84	7.51~7.77	13.01~16.80	0.05~0.10	1.65~2.03
		e						

Hu	CV/ %	4.63%	10.70%	0.99%	9.43%	24.13%	8.06%
	Mea n	7.99±0.40 ab	8.39±0.91 a	10.66±1.53 b	10.51±1.77 c	0.17±0.08 ab	2.23±0.30 bc
	Rang e	7.65~8.71	7.17~9.23	8.66~12.18	9.09~13.56	0.07~0.26	1.85~2.56
	CV/ %	4.64%	10.18%	13.50%	15.87%	42.06%	12.75%

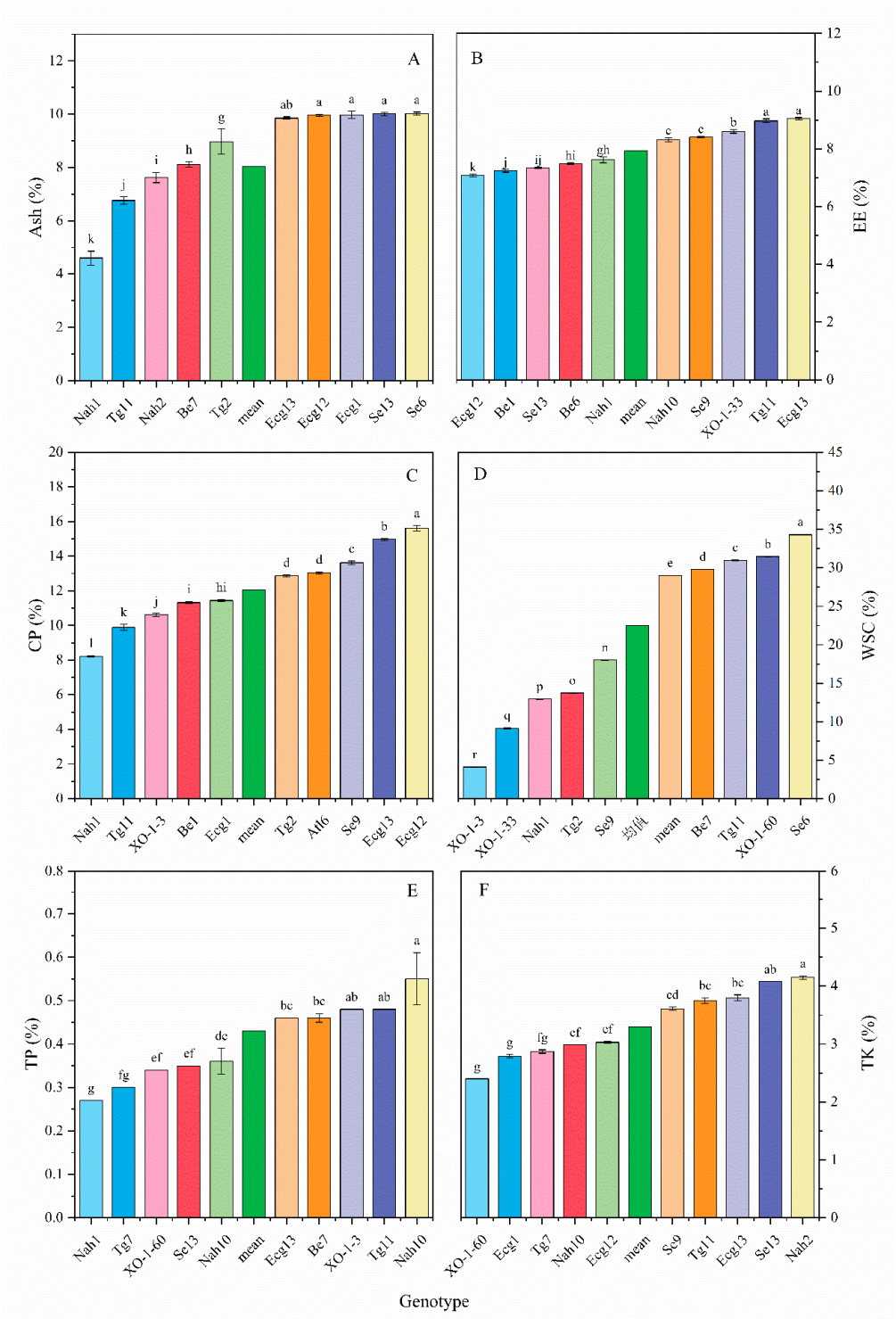


Figure 1. Average of Nutrient content in 3 years of different genotypes of oat.

3.2. Feed value analysis of different oat populations

Fresh-to-dry ratio, neutral detergent fiber, acid detergent fiber, lignin, and cellulose are five important indicators of oat forage quality. The analysis of forage quality data from six populations of oats (Table 4) revealed that. In terms of the first fresh-to-dry ratio, the Tabigha population had a significantly higher fresh-to-dry ratio than the other populations at 4.76% (average data of three years). The difference data of Hu(20.7%) which is the lowest neutral detergent fiber content and the Tabigha(26.61%) with the highest neutral detergent fiber content is 5.91%. The acidic detergent fiber of the artificially cultivated oat population Hu was significantly higher than other wild oat populations in all three years, reaching 5.99%, 29.93%, and 27.39%, respectively. In terms of lignin content, the Evolution Canyon population had the same amount as the Tabigha population at 1.15% in 2018, and the Hu had the highest amount in three years at 3.31% (average data). In terms of cellulose content, the Hu had significantly higher content than the other five populations at 71.88% (average data of three years), which was about eight times higher than the lowest content Sede Boque. It is note worth that there was no variability among the Tabigha, Nahef, Sede Boquer, and Evolution Canyon in lignin and cellulose in 2018 and 2019.

The forage quality of the 18 genotypes was analyzed (Figure 2) and the results showed that, in terms of moisture, the Se13 genotype had the highest content of 75.28%, while the XO-1-3 genotype had the lowest content of 59.05%. The fresh-to-dry ratio of the Be1 genotype of oats was 1.52%, which was significantly lower than the other 17 genotypes, while the Tg11 genotype had the highest content of 3.74%. In terms of lignin content, the XO-1-3 genotype had the highest content of 5.34%, while the Be1 genotype had the lowest content, with a difference of 4.49% between the two genotypes. XO-1-3 genotype had the highest cellulose content of 89.15% in all, and its cellulose content was 4.6 times higher than the mean value; Se 9, XO-1-3 and XO-1-60 genotypes of lignin content were close to each other, roughly 2.75%. This shows that the oat origin has a greater effect on lignin and cellulose and a smaller effect on other forage quality indicators. The six targets of XO which is planted by a human is smallest than other population. The forage value of cultivated oats is the lowest, and wild oats may have great forage potential.

Table 4. Analysis of feed value of different oat populations.

Year	Population		DFR/%	NDF/%	ADF/%	Lignin/%	Cellulose/%
2018.10	Tabigha	Mean	3.68±0.04	27.29±2.37	5.09±0.38 b	1.15±0.11 c	8.55±0.37 b
-			ab	ab			
		Range					
2019.03			3.64~3.75	25.56~30.55	4.68~5.58	1.02~1.32	7.99~9.10
		CV/%	1.15%	8.20%	7.06%	8.67%	4.10%
	Nahef	Mean	3.54±0.02 a	26.74±0.43 a	4.87±0.14 bcd	1.21±0.26 c	9.10±2.08 b
		Range					
			3.50~3.60	26.30~27.61	4.63~5.02	1.06~1.43	7.07~10.96
		CV/%	1.10%	2.23%	3.05%	10.80%	15.29%
	Sede Boquer	Mean	3.13±0.02 a	25.36±0.43 a	4.74±0.14 cd	2.42±0.26 b	7.98±2.08 b
		Range					
			3.10~3.16	24.77~25.84	4.56~4.94	2.11~2.79	5.99~10.86
		CV/%	0.53%	1.60%	2.68%	10.27%	24.63%

2019.10 - 2020.03	Evolution Canyon	Mean	3.24±0.04 a	25.08±1.24 a	4.90±0.24 bc	1.11±0.15 c	9.71±1.73 b
		Range	3.18~3.28	23.49~26.34	4.56~5.13	0.89~1.26	7.78~12.16
		CV/%	1.11%	4.66%	4.62%	12.56%	16.78%
	Beit-Oren	Mean	2.43±0.69 b	25.12±0.49 b	4.58±0.29 d	1.15±0.23 c	9.01±0.87 b
		Range	1.50~9.92	24.62~25.81	4.15~4.86	0.81~1.35	7.56~9.90
		CV/%	26.60%	1.82%	5.94%	18.66%	9.15%
	Hu	Mean	2.88±0.43 c	24.97±1.29 c	5.99±0.55 a	3.61±1.29 a	72.11±13.16 a
		Range	2.35~3.37	23.23~26.20	5.24~6.43	2.68~5.48	58.71~91.64
		CV/%	14.23%	4.87%	8.61%	33.75%	17.21%
	Tabigha	Mean	6.48±0.76 a	30.51±1.82 a	11.20±0.79 c	1.16±0.07 d	10.07±0.63 bc
		Range	5.50~7.62	28.15~32.81	9.82~12.36	1.04~1.28	8.92~10.81
		CV/%	11.77%	6.51%	6.65%	5.82%	5.89%
	Nahef	Mean	4.45±0.40 b	18.85±1.38 c	10.66±1.03 cd	1.45±0.21 c	9.45±1.09 bc
		Range	4.06~4.96	16.73~21.26	9.18~12.37	1.14~1.66	7.98~11.23
		CV/%	8.40%	6.89%	9.12%	13.31%	10.88%
	Sede Boqer	Mean	4.64±0.54 b	19.37±1.21 c	10.677±1.34 cd	1.17±0.49 d	9.23±0.51 bc
		Range	4.20~5.37	18.13~21.56	9.37~13.02	0.76~1.92	8.50~10.24
		CV/%	11.64%	5.26%	9.66%	24.31%	4.68%
	Evolution Canyon	Mean	4.24±0.20 b	19.12±1.23 c	9.89±0.85 d	1.23±0.30 cd	8.78±0.51 c
		Range	3.92~4.57	17.63~21.05	8.83~11.10	0.99~1.66	8.14~9.58
		CV/%	4.44%	6.05%	8.07%	23.15%	5.43%
	Beit-Oren	Mean	2.93±0.72 c	23.05±1.28 b	14.36±0.67 b	1.95±0.32 b	11.73±0.38 b
		Range	2.29~4.21	21.07~24.94	13.45~15.40	1.54~2.35	11.07~12.33
		CV/%	23.32%	5.24%	4.39%	15.50%	3.06%
	Hu	Mean	4.02±2.07 b	18.64±4.53 d	26.93±2.09 a	2.34±0.10 a	74.84±7.03 a
		Range	2.09~6.66	14.53~22.99	24.48~30.66	2.20~2.49	67.28~86.81
		CV/%	51.449%	24.30%	7.33%	3.99%	8.86%

2020.10	Tabigha	Mean	4.12±0.12 a	22.02±1.32 a	13.44±0.52 b	1.97±0.44 b	10.67±0.36 b
-							
2021.03		Rang e	4.01~4.28	20.20~24.59	12.37~14.13	1.43~2.53	10.27~11.39
		CV/%	2.67%	5.66%	3.62%	20.81%	3.22%
	Nahef	Mean	3.76±0.18 bc	19.93±0.78 cd	11.33±0.71 cd	1.64±0.08 c	9.98±0.56 bc
		Rang e	3.50~3.95	18.96~21.72	10.31~12.36	1.51~1.80	9.23~10.92
		CV/%	4.75%	3.93%	6.23%	5.04%	5.57%
	Sede Boqer	Mean	3.56±0.08 cd	18.73±2.24 d	10.78±1.92 d	1.51±0.30 c	8.99±1.55 c
		Rang e	3.45~3.65	15.78~22.09	8.08~12.85	1.10~1.78	6.82~10.65
		CV/%	2.12%	11.27%	16.83%	18.45%	16.23%
	Evolution Canyon	Mean	3.78±0.12 b	20.73±1.42 bc	12.28±0.85c	1.61±0.16 c	10.44±0.30 b
		Rang e	3.64~3.94	19.31~23.07	11.13~13.38	1.39~1.880	9.67~11.33
		CV/%	3.05%	6.47%	6.54%	9.39%	4.76%
	Beit-Oren	Mean	3.19±0.13 e	20.25±1.13 c	11.63±0.75 cd	1.43±0.28 c	9.90±0.76 bc
		Rang e	3.05~3.40	18.43~21.69	10.69~12.77	1.07~1.82	8.89~11.00
		CV/%	4.14%	5.57%	6.45%	19.30%	7.66%
	Hu	Mean	3.51±0.44 d	18.49±2.01 d	27.39±1.29 a	3.97±0.33 a	68.70±2.44 a
		Rang e	3.13~4.09	16.92~20.36	25.21~29.32	3.59~4.50	65.28~72.91
		CV/%	11.78%	10.87%	4.45%	7.92%	3.35%

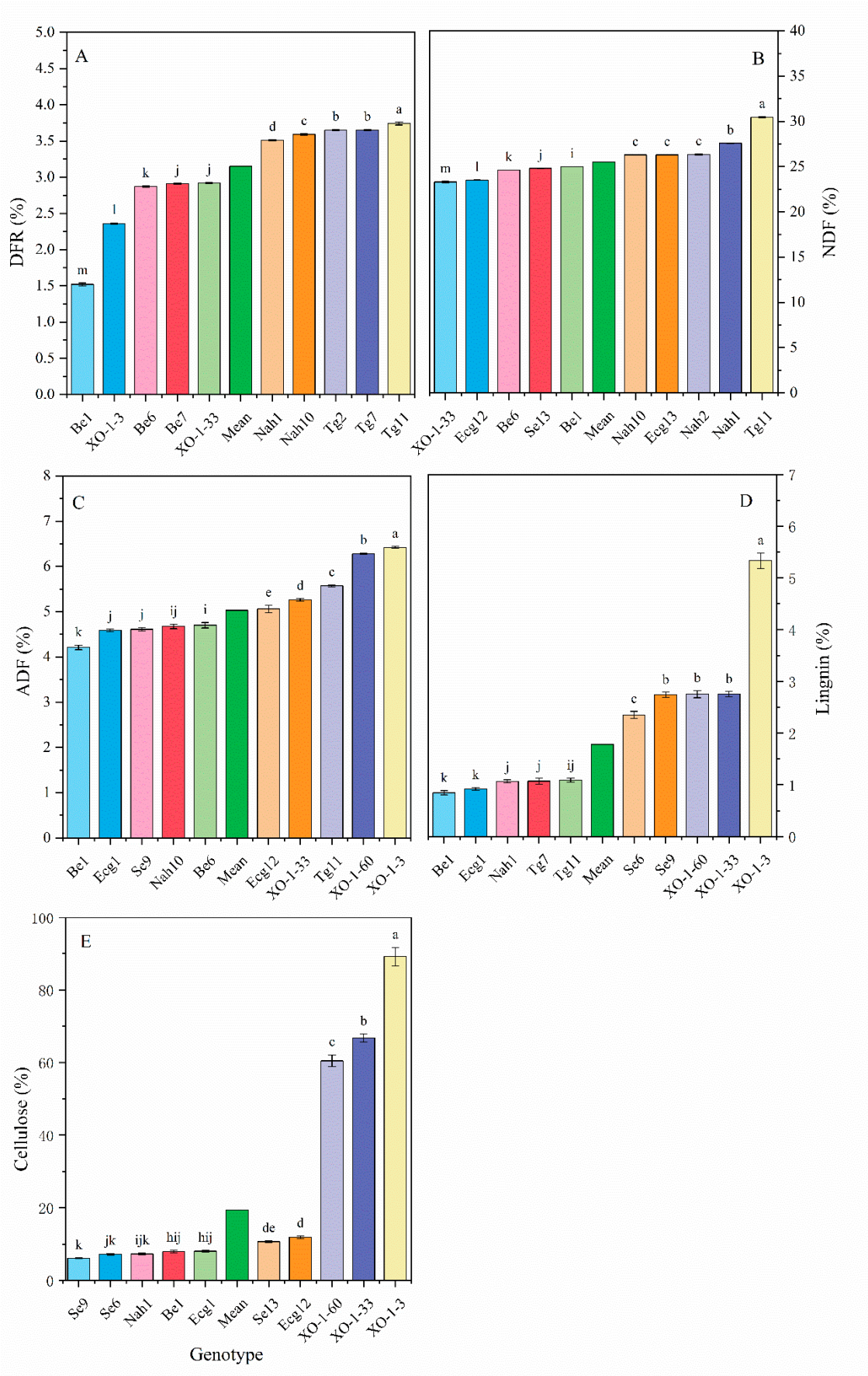


Figure 2. Average data of Feed value in 3 years of different oat genotypes.

3.3. The nutritional content of oat populations correlated with a place of origin

Analysis of oat population and ecogeographical factors revealed (as in Table 5) that nutrients such as crude protein, ash, and soluble sugars were significantly influenced by factors such as origin mile. Among them, crude protein was significantly positively correlated with Hu14, Huan, Tj, and Tm; while it was significantly negatively correlated with Ln, Td, and Tdd. Ash showed a significant positive correlation with Huan, while a significant negative correlation with Ev, Ln, Td, and Tdd. Soluble sugars showed significant positive correlations with Hu14, Ta, and Tm; and significant negative correlations with Al, Ln, and Lt. However, it is noteworthy that no significant correlations were observed for crude fat, total phosphorus, and total potassium in oats with factors such as ground mile of origin. The temperature at the origin of the oat population has a large effect on crude protein, ash, and soluble sugar, after which breeding can control seasonal temperature differences and diurnal temperature differences to enhance the nutrient content.

Table 5. Significant Spearman's rank correlation between nutritional content and ecogeographic factors in oat populations.

Nutrient content	Geographical environmental factors		Relevance	Prob> q
CP	Hu14		0.4965	0.0001
	Huan		0.3983	0.0029
	Ln		-0.5003	0.0001
	Td		-0.4465	0.0007
	Tdd		-0.3726	0.0055
	Tj		0.2741	0.0449
	Tm		0.2832	0.0380
Ash	Ev		-0.3417	0.0115
	Huan		0.3166	0.0197
	Ln		-0.5034	0.0001
	Td		-0.3449	0.0106
	Tdd		-0.3438	0.0109
WSC	Al		-0.4227	0.0015
	Hu14		0.2700	0.0483
	Ln		-0.2787	0.0413
	Lt		-0.3558	0.0083
	Ta		0.4893	0.0002
	Tm		0.4916	0.0002

3.4. Correlation between feed value and place of origin in oat populations

The analysis of oat populations and eco-geographic factors revealed (as in Table 6) that the indicators of lignin, cellulose, acidic detergent fiber, neutral detergent fiber, moisture, and the fresh-to-dry ratio of oat populations may be influenced by geographic environmental factors at the place of origin. Among them, lignin was significantly positively correlated with Ln, Lt, Al, and Td, while it was significantly negatively correlated with Tm, Tj, and Rn. Cellulose was negatively correlated with Tm, Tj, Ev, Rn, Ta, Rd, Hul14, and Huan; and positively correlated with Ln, Lt, Td, and Tdd. The acid detergent fiber was positively correlated with Lt and Al and negatively correlated with Rn and Rd; while neutral detergent fiber content was significantly influenced by only two geographic environmental factors, Rn and Rd, in a significant positive correlation, and was not significantly related to other influencing factors. In addition, we found that plant moisture and fresh-to-dry ratio could be influenced by Tm, Ta, Td, and other factors.

Table 6. Significant Spearman's rank correlation between feed value and ecogeographic factors in oat populations.

Feed indexes	Geographical environmental factors	Relevance	Prob> q
Lignin	Ln	0.2985	0.0283
	Lt	0.3172	0.0194
	Al	0.2996	0.0278
	Tm	-0.4948	0.0001
	Tj	-0.3205	0.0182
	Td	0.285	0.0367
	Rn	-0.382	0.0044
Cellulose	Ln	0.3705	0.0058
	Lt	0.3946	0.0032
	Tm	-0.4783	0.0003
	Tj	-0.3246	0.0166
	Td	0.4129	0.0019
	Tdd	0.4979	0.0001
	Ev	-0.3778	0.0049
	Rn	-0.3308	0.0146
	Ta	-0.4544	0.0006
	Rd	-0.2698	0.0485
	Hu14	-0.4159	0.0018

OM	Huan	-0.4675	0.0004
	Lt	-0.4783	0.0003
	Tm	0.3573	0.008
	Ta	0.3223	0.0175
	Td	-0.4406	0.0009
DFR	Ev	0.4488	0.0007
	Huan	0.3966	0.003
	Tm	0.4333	0.0011
	Ta	0.3994	0.0028
	Tj	0.2871	0.0353
ADF	Td	0.3672	0.0063
	Rd	0.4566	0.0005
	Huan	-0.402	0.0026
	Lt	0.501	0.0001
	Al	0.3863	0.0039
NDF	Rn	-0.3991	0.0028
	Rd	-0.2724	0.0463
	Rn	0.2807	0.0398
	Rd	0.4234	0.0014

3.5. Principle component analysis (PCA)

Principal component analysis (PCA) helps to study the analysis of nutritional content and forage quality among different populations, to select the best oat population, and to analyze the variability of oat population genes in nutritional content and forage quality in different origins. Figure 3 (A) shows the PCA of six oat populations in six nutritional content parameters studied, and Figure 3 (B) examines the populations in six forage quality parameters.

In the PCA analysis of feed value, the Hu population was significantly different from the other five populations, with the three genotypes XO-1-3, XO-1-33, and XO-1-60 of the Hu population positively correlated with variability in component 1 and the remaining five populations negatively correlated. In PC1 total variability was 55.50% and they were positively correlated with acidic detergent fiber (0.47), lignin (0.50 and cellulose (0.52) and negatively correlated with primary moisture (-0.48), fresh-to-dry ratio (-0.17) and neutral detergent fiber (-0.05).PC2 total variability was 25.1% and they were positively correlated with neutral detergent fiber, acidic detergent fiber, and fresh-to-dry ratio correlated,0.36, 0.62, and 0.68, respectively, and negatively correlated with primary moisture (-0.08), lignin (-0.07), and cellulose (-0.06).

From the analysis of nutritional content, there was no significant variability in forage quality among the five groups except for the Nahef group. the contribution of PC1 and PC2 was 33.75% and 61.63%, respectively, with a total coefficient of variation of 33.8% and PC2 of 27.9%. In PC1, the Evolution population genotype was positively correlated with variability, and the remaining five population genotypes were partly negatively and partly positively correlated. In PC1, there was a positive correlation with nutritional content in all six oat populations. PC2 was negatively correlated with Ash and CP, -0.42 and -0,29 respectively, but positively correlated with EE (0.52), WSC (0.02), TP (0.36), and TK (0.58).

The forage quality of cultivated oats Hu and the other five wild oats groups were far apart in Figure 4(A), and the five wild oats groups were more similar. Moreover, in the components of PC1, the variability of cultivated oats showed a positive correlation and the variability of wild oats showed a negative correlation. In terms of nutritional value Figure 4 (B), there is little convergence between cultivated and wild oats. the variability of Nah and Tab groups is greater, and Hu variability is less, probably the nutritional value of wild oats has more potential to be explored. The genetic diversity of the cultivated oat population decreased and was more homogeneous, however, the diversity and coefficient of variation were richer in the wild.

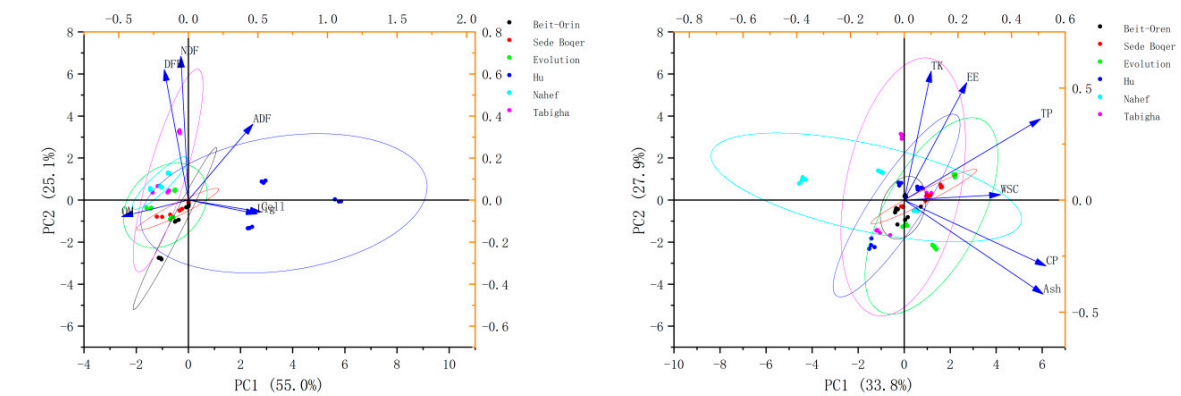


Figure 3. Principle component analysis (PCA).

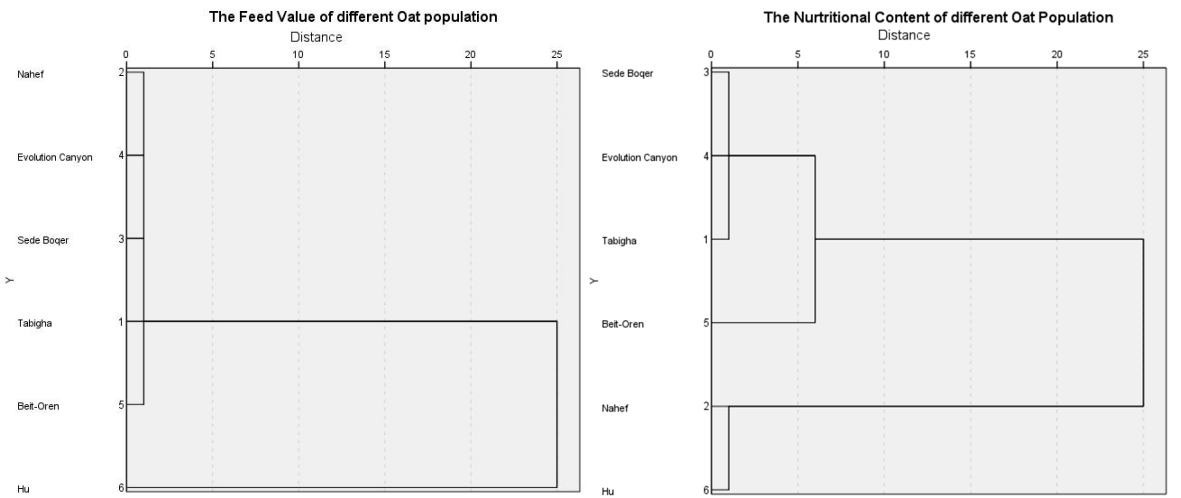


Figure 4. Cluster plotting of different communication.

3.6. Cluster plotting

The Wards method was used in performing the clustering analysis. From the feed value analysis, at a distance of 25%, the six oat populations were divided into two lineages, a Hu population from Zhangjiakou in Hebei, and five populations from Israel (Tabigha, Nahef, Sede Boqer, Evolution Canyon, and Beit-Oren). Within a distance of about 1.00%, they were further subdivided into five sub-clusters. The clustering results of the feeding value of the six groups are very clear, and wild oats have closer relatives and are completely independent of Chinese cultivated oats, which indicates that the feeding value of wild oats is unique.

In terms of nutritional content, there are two line segments at 25% dividing the six oat groups into two parts, one for Nahef and another part with four groups (Sede Boqer, Evolution Canyon, and Beit-Oren). The distance was further subdivided into two subgroups at 6.02%, except for Beit-Oren, where there was variability in the other three groups at very small distances. However, the Hu oat group from China was less differentiated from Nahef at a 1% distance. Therefore, it is speculated that the nutritional structure of the Naf oat population may be closer to that of cultivated oats, which is consistent with Figure 4.

4. Discussion

Wild oats are an important grass crop that is widely distributed in natural ecosystems around the world. Compared with cultivated oats, wild oats have higher nutritional quality and forage potential and therefore have received increasing attention (Nevo et al., 1984; Turpeinen et al., 2001). In the article, we will discuss the higher nutritional quality and forage potential of wild oats in protein content, crude fat, fiber content, and acid-detergent fiber.

Firstly, wild oats are rich in high-quality protein. A study found that the protein content and quality of wild oats were significantly better than that of cultivated oats. In a study, Australian scientists Harris and Mares found that the protein content of wild oats was higher than that of cultivated oats and that the amino acid composition of wild oats was more complete (Harris & Mares, 1974). Among them, wild oats have a higher content of essential amino acids, such as lysine, tryptophan, and isoleucine. These results indicate that wild oats have higher nutritional quality and can provide better nutritional security for humans. The experimental results of this study showed that the content indexes of various nutrients in the Evolution Canyon oat group from Israel were significantly higher than those in the other five oat groups, indicating that this wild group has a higher nutritional quality and may have potential value in forage utilization. The difference between the Tg11 genotype with the highest total potassium content and the Nah1 genotype with the lowest content was 0.21%, while the difference between the Nah2 genotype with the highest total phosphorus content and the XO-1-60 genotype with the lowest content was 1.77%. The present study further used Spearman's rank correlation analysis to conclude that the geographical environmental factors of the origin of the oat population had less influence on the nutrient composition of oats, and mainly had a more significant effect on ash, crude protein, and soluble sugars ($p < 0.05$). Crude protein was significantly positively correlated with Hu14, Huan, Tj, and Tm; while it was significantly negatively correlated with Ln, Td, and Tdd, which is consistent with Liu Wenting et al. study (Gao et al., 2015; Haitao et al., 2019; Peterson & David, 2004; Peltonen-Sainio et al., 2004). However, there was no significant correlation between total potassium, total phosphorus, and ecogeographic factors of the place of origin for oats, suggesting that these indicators may be less influenced by the environment.

Secondly, the cellulose content in wild oats is also relatively high. One study found that wild oats had a higher cellulose content than cultivated oats, with a relatively low content of neutral detergent fiber and a higher content of acid detergent fiber. This implies that wild oats are more digestible and can provide better feed value for animals (Wang et al., 2019; Zhang et al., 2019; Zheng et al., 2013). In addition, wild oats also contain higher levels of polyphenolic compounds, which are important for the prevention of intestinal diseases and the protection of cardiovascular health (Demirbas, 2005; Martinez et al., 2010). In this study, this study found that there were large differences between the forage quality of the oat groups, and the moisture, fresh-to-dry ratio, and neutral detergent fiber contents of the Hu group was significantly lower than those of the other five

wild groups, indicating that the five oat groups from Israel had higher forage value. The Hu cellulose content and lignin content were both higher than those of the other groups, at 71.88% and 3.11% (average data of three years), respectively. This suggests that the water use efficiency of wild oats in Israel may be higher. The genotypes with significantly higher cellulose content were all from the Hu population in Zhangjiakou, Hebei, suggesting that cultivated oats may be more “chewy” and less palatable. This study showed that the fiber content of different genotypes differed significantly, so the selection of forage oats for different groups should be carefully screened. Spearman’s rank significant correlation analysis revealed that several forage quality components were significantly influenced by geographic environmental factors of origin, and cellulose content was significantly influenced by 12 geographic environmental factors, which were negatively correlated with Tm, Tj, Ev, Rn, Ta, Rd, Hul14, and Huan; and positively correlated with Ln, Lt, Td, and Tdd, which is consistent with the study. (Zhang et al., 2016). In this study, the higher nutritional quality and forage potential of wild oats were investigated in terms of initial moisture, fresh-to-dry ratio, cellulose lignin content, neutral detergent fiber, and acid detergent fiber.

The forage potential of wild oats is also of interest. One study found that wild oats have a higher forage potential than cultivated oats. An Australian study found that wild oats grow faster and can produce high-quality forage in a short period. In addition, the biomass and nutritional value of wild oats were higher than that of cultivated oats (Robertson et al. 2004). Comparative studies have shown that the dry matter yield and forage dry matter yield of wild oats are about 30% higher than that of cultivated oats (Aanchaldeep et al., 2022; Buttar et al., 2018), while wild oats have higher crude protein, neutral detergent fiber, and acid detergent fiber content (Bhatta et al., 2003). In addition, wild oats have other physiological functions and nutritional values. For example, wild oats are rich in a variety of bioactive components, such as polyphenols, alkaloids, and flavonoids, which have various physiological functions such as antioxidant, antitumor, hypoglycemic, and hypolipidemic (Qin et al., 2019). These physiological functions and nutritional values make wild oats a natural resource with wide application value. In this experiment, we found that the differences between different genotypes of wild oats were greater than those of cultivated varieties, and the genetic diversity and population structure of wild oats were richer than those of cultivated ones (Montilla-Bascón et al., 2013; Kshitiz et al., 2022; Mohammad et al., 2022). This indicates that wild oats have greater potential for exploration.

In conclusion, wild oats have higher nutritional quality and forage potential compared to cultivated oats. Exploring protein content, amino acid composition, fiber content, and acid detergent fiber, wild oats have higher protein content, more complete amino acid composition, and higher fiber content, which can provide better feed value for animals. Wild oats also have a variety of physiological functions and nutritional values, so it has a wide range of application prospects in the field of food, medicine, and health products.

5. Conclusion

Oats have a very high nutritional quality and forage quality. The results of this experiment show that the protein content of wild oat populations (the average of five populations) in three-year cultivation can reach 9.35%, fat content of 7.86%. Higher than other wheat and corn, with high dietary fiber content, wild oats got a Cellulose content of only 9.57%, significantly lower than the cultivated population Hu (71.88%). Net milk energy production is also higher, and it is the most promising high-quality hay in the world for industrial production and a high-quality concentrate for various livestock. In this experiment, five oat populations and one cultivated population in Israel were studied, and the results showed that the ash, soluble sugar, and total potassium contents of the Sede Boquer population were significantly higher than those of the cultivated population. In terms of forage quality release, Hu showed significantly higher lignin and cellulose content than wild oat populations, indicating that cultivated oats are more resistant to stress, but also resulting in less palatability. Climatic analysis of Spearman rank correlations between populations and origin showed that several forage quality components were significantly influenced by geographic environmental factors of origin, and cellulose content was significantly influenced by 12 geographic environmental

factors, including negative correlations with Tm, Tj, Ev, Rn, Ta, Rd, Hul14 and Huan; and positive correlations with Ln, Lt, Td, and Tdd. Genetic variability of wild oats may be higher in the extreme arid weather and different latitudes and longitudes of origin in Israel. It has higher nutrient content than cultivated oats, offering greater possibilities for feed quality. In conclusion, this study provides important source material for fodder development and utilization of wild oats and good seed selection, and a better basis for further research on wild oats as an important fodder crop.

Acknowledgments: The authors thank Chengdu University for technical assistance and the Institute of Evolution, the University of Haifa for providing oat seed resources, and especially Liang Zou and Yu Fan for their role in facilitating this research. We thank , Li Xiaolong, Wang Junzhen, Li Yang, Yang Qiaohui, Xiang Dabing, Wan Yan, Eviatar Nevo, and Yan Jun for their work in the field and laboratory.

Funding information: National Natural Science Foundation of China (31860412), Talent Initiation Funding Project of Chengdu University (2081923007), Open Project Program of Irradiation Preservation Technology Key Laboratory of Sichuan Province, Sichuan Institute of Atomic Energy (FZBC206704), Ministry of Finance and Ministry of Agriculture and Rural Affairs: National Oat Buckwheat Industry Technology System (CARS-07-B-1).

Conflict of interest statement: The authors declare no conflicts of interest.

Data availability statement: Data are available from the corresponding author on reasonable request.

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