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Obtaining of Winter Rye (Secale Cereale L. ssp. Cereale) Haploid Embryos through Hybridization with Maize (Zea Mays L.)

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The aim of this study was to determine the effect of selected factors on rye (*Secale cereale* L.) haploid embryo production by the wide crossing method. The study was performed on fifteen winter rye genotypes. This is the first time for rye when besides the genotype, on the enlargement of ovaries and haploid embryo production, such factors as: type of auxin analogues 2,4-dichlorophenoxyacetic acid (2,4-D), 3,6-dichloro-2-methoxybenzoic acid (dicamba) and 4-amino-3,5,6-trichloropyridine-2-carboxylic acid (picloram), and the time between florets emasculation and pollination were investigated. All factors had a significant impact on rye ovary enlargement, however the haploid embryo formation depended only on rye genotype, not on kind of auxin and days between emasculation to pollination. In total, twenty one haploid embryos were formed by six genotypes of fifteen tested. On average, 13.86% (after 2,4-D treatment) to 20.05% (after dicamba treatment) enlarged ovaries per emasculated florets were obtained. Most of the ovaries enlarged when florets were pollinated 4 and 6 days after emasculation. The obtained haploid embryos did not germinate.

Keywords auxin, haploid embryo, maize, rye, wide crossing

Introduction

Rye (*Secale cereale* L.) is one of the most important cereals in Eastern and Northern Europe. Despite numerous beneficial characteristics of rye, its annual production decreases successively, which correlates with the lack of progress in its breeding compared with other cereals like wheat and barley. Unfortunately, the classical breeding of rye is very difficult because of its high self-incompatibility and inbred depression. On the other hand, rye is known as a recalcitrant plant species in regard to the *in vitro* culture response, thus the application of biotechnological methods (e.g., doubled haploid (DH) production, genetic transformation or selection of plants with beneficial changes of agronomic traits) is highly limited (Hromada-Judycka 2010). DH technology has become key tool for plant breeding. Using these techniques, the speed and efficiency of plant improvement pro-

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cesses can be significantly enhanced (Rubtsova 2013). Several methods of DH production have been investigated in cereals, including microspore and/or anther culture (androgenesis), ovary and ovule culture (gynogenesis) as well as wide hybridization through interspecific crosses. Major advantages of DH lines compared to pedigree lines include: maximum genetic variance among lines and testcross performance from the first generation, reduced length of breeding cycle, perfect fulfillment of distinctness, uniformity, stable varietal status, reduced costs in maintenance breeding as well as increased efficiency in marker-assisted selection, gene introgression and gene stacking (Sserumaga et al. 2015).

For cereals, androgenesis is the most commonly applied DH system. Anther or isolated microspore cultures have been more favorable than other methods. The production of DHs in rye has not been as efficient as for barley or wheat. Although high regeneration rates of green plants have been achieved with some rye lines or cultivars, rye can still be considered a recalcitrant species (Tenhola-Roininen 2006). Pioneering work on rye doubled haploid production was carried out by Wenzel and Thomas (1974) and Zenkteler and Misiura (1974) and concentrated solely on anther culture. Callus, embryos and green plants have been produced by microspore culture of F_1 winter hybrid types of S. cereale L. × S. vavilovii Grossh (Thomas and Wenzel 1975). In the nineties, the first studies with rye anther culture and success in high green plant regeneration rate were reported by Flehinghaus-Roux et al. (1995) and by Immonen and Anttila (1996). Since the first studies on anther or microspore cultures of rye, callus induction and embryo formation as well as albino and green plant rates vary depending on the genotype, physiological stage of donor plant, stress treatments, developmental stage of microspore as well as medium and culture conditions (Deimling and Flehinghaus-Roux 1997; Immonen 1999; Immonen and Anttila 1999, 2000). Rakoczy-Trojanowska et al. (1997) have suggested that various genotypes could have different requirements with regard to plant growth regulators and no universal medium exists for different rye genotypes. Compared to barley and wheat, the development of an efficient androgenic cell culture system for rye is less advanced. Problems associated with rye anther and microspore culture include poor embryogenic callus induction, limited green plant regeneration, a high proportion of albinos and severe genotype dependency. More effective methods are needed to produce doubled haploids from a wide range of genotypes (Altpeter and Korzun 2007).

Wide crossing with pollen from another cereal genus (e.g., *Hordeum bulbosum* or *Zea mays*) is an alternative method of producing haploid embryos. This method involves fertilization, but foreign chromosomes are eliminated during early embryo development. Advantages of this approach are that the methods involved (emasculation, pollination, embryo culture) are familiar to breeders, cross combinations can be manipulated to maximize haploid embryo production by eliminating genotypic dependency on the maternal side (Forster et al. 2007). In the study of Zenkteler and Nitsche (1984), where *Secale cereale* was pollinated with *Dactylis glomerata*, *Festuca glauca*, *Hordeum bulbosum* and *Zea mays*, globular hybrid embryo production was achieved. Haploid embryos from rye x maize crosses obtained by Laurie et al. (1990) died already at the early stages of development, while Deimling et al. (1994) and Altenhofer et al. (1997) observed conversion of rye embryos into plants. Ponitka and Ślusarkiewicz-Jarzina (2004), based on cleared ovule preparations, found irregularities in endosperm development. Usually the ovules contained both embryo and endosperm, but they sometimes contained embryos without endosperm or the endosperm alone. The developmental stage of such embryos for *in vitro* culture due to irregularities in endosperm development is the most common cause of dieback of haploid embryos obtained by the wide hybridization method. Similar results was observed for oat haploid embryos (Noga et al. 2016).

The aim of this study was to determine the effect of selected factors on rye (*Secale cereale* L.) haploid embryo production by the wide crossing method. The study was performed on fifteen Polish winter rye genotypes and investigated the importance of donor plant vernalization. Information that genotype, type of auxin and days between floret emasculation and pollination by maize on the efficiency of ovary enlargement and haploid embryo production could be helpful in optimization of rye doubled haploid production using this method.

Materials and Methods

Fifteen Polish F₁ winter rye (*Secale cereale* L.) genotypes: 2550, 94, 41, 31, 61, S1123/14, S1138/14, 127-2R, 186-R, 131-R, 135-2R, 143-3R, S1131/14, 2545 and 36 were provided by Danko Plant Breeding Ltd. Rye grains were treated with fungicide Funaben T (Synthos Agro Ltd., Poland) and placed in containers with perlite wetted with Hoagland medium (Hoagland and Arnon, 1938). Thus prepared seeds were vernalized for 7 and 9 weeks at 3 ± 1 °C with an 8 h photoperiod under 100 µmol m² s⁻¹ (PAR). After vernalization, five plants per genotype were planted individually to 3 dm³ pots filled with a mixture of soil with sand (3:1 v/v). Rye and maize plants were grown in a greenhouse in April and May, under controlled conditions, 800 µmol m⁻² s⁻¹ photosynthetic active radiation (PAR) and long day (16-h light, 8-h dark). Photoperiod and light intensity were maintained with the support of high pressure sodium lamps (400 W; Philips SON-T AGRO, Belgium) between 6–8 a.m., 6–10 p.m. and on cloudy days. Rye plants were grown at 21/17 °C day/night.

Maize (*Zea mays* L. var. *saccharata*) cv. Waza used as a pollen donor, was grown in a greenhouse at 25-28/17 °C day/night and the same light regime as rye. In order to synchronize the flowering of rye with pollen production by maize, maize seeds were sown for the first time about 3 weeks before the end of rye vernalization (7 or 9 weeks) and then 5 times at 7 day intervals.

All rye and maize plants were fertilized with a liquid medium once a week (Hoagland and Arnon 1938). Emasculation of rye florets was performed when the spikes remained prior to anthesis. Dependently of genotype 24 to 35 spikes were emasculated and covered with glassine bags. After 3–14 days spikes were pollinated with fresh maize pollen. Then, 1 day later rye pistils were treated with three different analogues of auxins: 2,4-dichloro-phenoxyacetic acid (2,4-D), 3,6-dichloro-2-methoxybenzoic acid (dicamba) or 4-amino-3,5,6-trichloropyridine-2-carboxylic acid (picloram) at a concentration of 100 mg dm⁻³. Two to 3 weeks later enlarged ovaries were isolated from the florets and surface-sterilized

in 70% v/v ethanol (1 min), 2.5% v/v calcium hypochlorite (8 min) and then washed four times with sterilized water. Embryos were isolated and placed on 190-2 medium (Zhuang and Xu 1983) with 9% (w/v) maltose solidified with 0.6% (w/v) agar. The pH of the medium was adjusted to 5.8 using 0.1 N NaOH before autoclaving at 121 °C for 20 min. Embryos were maintained under 100–150 μ mol m² s⁻¹ PAR at 21 °C and a 16 h photoperiod. All reagents used in the experiment were obtained from Sigma-Aldrich.

The analysis of rye ovaries enlargement and embryo formation after maize pollination was performed depending on the genotype, type of auxin used to stimulate the ovary enlargement and time between floret emasculation and pollination.

Statistical analysis

The results concerning the influence of the aforementioned factors were evaluated using two way ANOVA variance analysis and Duncan's test incorporated in the statistical package STATISTICA 12.0 (Stat-Soft Inc., USA). Significant differences between traits depending on source of variation are marked by asterisks (*, ***, significant at $p \le 0.05$, 0.001, respectively; ns – not significant). Mean values marked with the same letters do not differ significantly.

Results

To obtain the highest number of spikes per plant (even 11 spikes), vernalization should last over a period of 9 weeks (data not shown). Shorter vernalization (7 weeks) resulted in a lower number of heading spikes. In addition, to synchronize rye flowering with pollen production by maize, maize seeds should be sown for the first time approx. 3 weeks before the 9 weeks of vernalization and then 5 times at 7 day intervals. Rye needs approximately 60 days to reach appropriate stage of spike development for wide crossing (prior to anthesis). The analysis of variance showed significant differences in the efficiency of rye ovaries enlargement relative to the genotype, type of auxin and days between floret emasculation to pollination (Table 1). There were significant differences

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Trait	Source of variation	SS	df	MS	F	р
Enlarged ovaries	Genotype	4746.68	14	339.05	3.49	0.000***
	Type of auxin	545.82	2	272.91	2.62	0.049*
	Days between emasculation to pollination	4127.23	11	275.14	2.78	0.000***
	Genotype	5.43	14	0.38	2.62 2.78 2.02	0.015*
Embryos	Type of auxin	0.05	2	0.02	0.13	0.870 ^{ns}
	Days between emasculation to pollination	1.06	11	0.07	0.34	0.989 ^{ns}

Table 1. Analysis of variance showing the significance of genotype, type of auxin, days between florets emasculation to pollination on rye ovaries enlargement and haploid embryo production using the wide hybridization method

SS sum of squares; df degrees of freedom; MS mean squares; ns not significant; *, ***, significant at $p \le 0.05$, 0.001, respectively.

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among genotypes in the haploid embryo formation, in contrast to the type of auxin and days between floret emasculation to pollination. Overall, 435 rye spikes (17154 florets) from all genotypes were emasculated and pollinated with maize. Approximately two weeks after pollination, 2838 ovaries had been enlarged (Table 2).

Enlarged ovaries were more than twice bigger than the ovaries before pollination and non-enlarged ones (Fig. 1A, B and C). There was an average of 16.50% enlarged ovaries per pollinated floret. The highest number of enlarged ovaries per emasculated floret were observed in genotypes S1138/14, S1131/14, 2550 and 186-R (34.27%, 26.81%, 25.76%, 25.57%, respectively), whereas the lowest (3.82%) in genotype 61. In total, 21 haploid embryos (Fig. 1D) were formed by six genotypes of 15 tested. The highest number of haploid embryos, 12 (1.17% per emasculated floret), were obtained from genotype 2545, while genotype 2550 formed 4 haploid embryos (0.28% per emasculated floret), genotype 41, 2 embryos (0.16% per emasculated floret), and genotypes 127-2R, 31, 186-R only 1 embryo (0.11%, 0.09%, 0.07% per emasculated floret, respectively). The obtained haploid embryos did not germinate and convert into plants on the investigated 190-2 medium. The number of enlarged rye ovaries and haploid embryo formation depended on the time between emasculation of rye florets and pollination by maize. Ovary enlargement was observed in all the investigated time periods, however, most of them enlarged when florets were pollinated 4–11 days after emasculation (Fig. 2). In turn, haploid embryos

	Number of				Enlarged ovaries/	Haploid embryos/	
Genotype	Emasculated spikes	Pollinated florets	Enlarged ovaries	Haploid embryos	pollinated florets [%]	pollinated florets [%]	
2550	35	1435	360	4	25.76	0.28	
94	24	1008	88	0	8.73	0.00	
41	32	1248	209	2	16.75	0.16	
31	28	1120	143	1	12.77	0.09	
61	31	1178	45	0	3.82	0.00	
S1123/14	25	1050	87	0	8.29	0.00	
S1138/14	28	1036	355	0	34.27	0.00	
127-2R	26	936	173	1	18.48	0.11	
186-R	35	1365	349	1	25.57	0.07	
131-R	28	1148	103	0	8.97	0.00	
135-2R	26	988	178	0	18.02	0.00	
143-3R	28	1092	168	0	15.38	0.00	
S1131/14	29	1160	311	0	26.81	0.00	
2545	25	1025	197	12	19.22	1.17	
36	35	1365	72	0	5.27	0.00	
Total/Average	435	17154	2838	21	16.50	0.13	

Table 2. The effect of rye genotype on the efficiency of rye haploid embryos production

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Figure 1. Rye×maize wide crossing for haploid embryo production: **A** ovary before pollination; **B** enlarged ovaries after treatment with dicamba; **C** not enlarged ovary after treatment with dicamba; **D** haploid embryo

formed only when florets were pollinated 5–10 days after emasculation (Fig. 3). The highest percentage of enlarged ovaries per emasculated floret was 25.36% and 26.87%, respectively (Fig. 2).

The lowest ovary enlargement was observed when they were pollinated by maize 13 (5.26%) and 14 days after emasculation (2.56%). The highest percentage of enlarged ovaries not always resulted in the highest percentage of haploid embryo formation. When florets were pollinated 6 days after emasculation, 26.87% of enlarged ovaries formed 13 haploid embryos (0.33% per emasculated florets) (Fig. 3). Despite the 25.36% of enlarged ovaries when florets were pollinated 4 days after emasculation, haploid embryo formation was not observed. The number of enlarged ovaries was affected by the type of auxin and genotype (Table 3). The average number of enlarged ovaries per emasculated floret after dicamba treatment was 20.05%, while in the case of picloram and 2,4-D it was 16.01% and 13.86%, respectively.



Figure 2. The influence of the number of days between emasculation of rye florets and pollination with maize on the percent of enlarged ovaries per emasculated florets independently on genotype. Mean values marked with the same letters do not differ significantly according to Duncan test at $p \le 0.05$



Figure 3. The influence of the number of days between emasculation of rye florets and pollination with maize on the number of haploid embryos formation and the percent of haploid embryos per emasculated florets independently on genotype. Mean values marked with the same letters do not differ significantly (A–D – the number of haploid embryos; a–b – the percent of haploid embryos per emasculated florets) according to Duncan test at $p \le 0.05$

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Canatana	Enlarged ovaries per pollinated florets [%]			Haploid embryos per pollinated florets [%]			
Genotype	2,4-D	Dicamba	Picloram	2,4-D	Dicamba	Picloram	
2550	30.82	20.16	26.29	0.67	0.00	0.27	
94	2.38	18.03	7.74	0.00	0.00	0.00	
41	3.85	17.31	23.59	0.00	0.32	0.00	
31	15.42	16.88	4.69	0.00	0.00	0.31	
61	0.00	7.89	2.37	0.00	0.00	0.00	
S1123/14	5.95	8.33	13.81	0.00	0.00	0.00	
S1138/14	44.79	21.62	37.61	0.00	0.00	0.00	
127-2R	23.26	15.28	17.22	0.35	0.00	0.00	
186-R	10.43	40.51	33.33	0.17	0.00	0.00	
131-R	2.93	19.51	7.76	0.00	0.00	0.00	
135-2R	28.16	22.18	3.51	0.00	0.00	0.00	
143-3R	10.62	20.09	12.82	0.00	0.00	0.00	
S1131/14	19.25	30.91	30.63	0.00	0.00	0.00	
2545	5.23	37.67	11.65	0.00	1.90	1.36	
36	4.81	4.36	7.12	0.00	0.00	0.00	
Average	13.86	20.05	16.01	0.08	0.15	0.13	

Table 3. The effect of auxin on the number of enlarged ovaries and haploid embryos per pollinated florets [%] dependently on rye genotype.

Mean values marked with the same letters do not differ significantly according to Duncan test ($p \le 0.05$)

The highest number of enlarged ovaries per emasculated floret after 2,4-D treatment was obtained in two genotypes: S1138/14 and 2550 (44.79% and 30.82% respectively), after dicamba treatment in three genotypes: 186-R, 2545 and S1131/14 (40.51%, 37.67%) and 30.91% respectively), and after picloram in three genotypes S1138/14, 186-R and S1131/14 (37.61%, 33.33% and 30.63% respectively). Genotype 61 was the least responsive one, as after 2,4-D treatment the ovaries did not even enlarge. The highest frequency of haploid embryo production was recorded after dicamba treatment (0.15%), then after picloram and 2,4-D (0.13% and 0.08%, respectively). Dicamba caused the highest haploid embryo formation in genotype 2545 (1.19%). The same genotype formed 1.36% of haploid embryos after picloram. The frequency of haploid embryos in other genotypes ranged from 0.67% to 0.17% depending on auxin treatment.

Discussion

Tissue culture methods can effectively improve the breeding of rye. However, their application is highly limited by the absence of efficient procedures for plant regeneration in vitro, because rye is one of the most recalcitrant cereals with regard to the tissue culture

response, and successful regeneration is highly dependent on the genotype (Targońska et al. 2013). Alternatively, substantial changes to centromeric histone H3 (CENH3), such as replacing the hypervariable N-terminal tail of CENH3 with the tail of conventional histone H3 and fusing it to GFP (producing "tailswap-*cenh*3"), or complementing the cenh3.2-null mutant with homologs from the mustard family CENH3s creates haploid inducer lines in the model plant *Arabidopsis thaliana*. Haploidization occurred only when such a haploid inducer was crossed with a wild-type plant. The haploid inducer line proved to be stable upon selfing, suggesting that competition between modified and wild-type centromeres in the developing hybrid embryo results in the inactivation of the centromeres from the inducer parent. Consequently, chromosomes from the inducer parent are lost, and progeny can be recovered that retain only the haploid chromosome set of the wild-type parent (Karimi-Ashtiyani et al. 2015). CENH3 is almost universal in eukaryotes, so this method has the potential to produce haploids in any plant species.

Some *in vitro* techniques (ovary, anther, and microspore culture) appear to be more promising than traditional breeding techniques. Hybridization involving interspecific and intergeneric crosses encounters numerous obstacles, mainly concerning pre- and postzy-gotic barriers (Rybczyński 1990). Rye wide hybridization and the utilization of ovary or embryo culture for rye doubled haploid (DH) production have either never been investigated or their failures have not been published.

In contrast, anther and microspore cultures have been the subject of several studies over the past 40 years (Wenzel and Thomas 1974; Thomas and Wenzel 1975, Wenzel et al. 1977; Flechinghaus et al. 1991; Flechinghaus-Roux et al. 1995; Immonen 1999; Immonen and Anttila 1996, 1999, 2000; Rakoczy-Trojanowska et al. 1997; Guo and Pulli 2000; Ma et al. 2004; Mikołajczyk et al. 2012). Most of all, the authors identified basal media composition that enhanced haploid embryo induction and green plant regeneration. However, a problem in this technique is the formation of numerous chlorophyll-deficient plantlets and very low regeneration efficiency. Survival and response of the microspores have been maximized by growing the plants used as a source of androgenic cultures under optimal conditions (Hörlein 1991). Vernalization period of the donor plants is important for regeneration of androgenic plants. Best anther culture response has been observed after plants were vernalized at 2 °C for 8 to 14 weeks with an optimum at 10 weeks (Flehinghaus et al. 1991). Vernalization for less than 8 weeks resulted in donor plants with small number of spikes and did not develop. Different genotypes require distinct lengths of vernalization, but in general, longer duration treatments are advised (Deimling and Flehinghaus-Roux 1997). Also in our investigation, longer time of vernalization (9 weeks compared to 7 weeks) resulted in a higher spike production and plants were in a better condition.

Zenkteler and Nitsche (1984) reported that wide hybridization in cereals resulted in the globular hybrid embryo when *Secale cereale*, *Hordeum vulgare* and *Triticum aestivum* were used as female species in crosses with 15 different species of Gramineae. The embryos were present in embryo sacs of *S. cereale* pollinated with: *Dactylis glomerata*, *Festuca glauca*, *Hordeum bulbosum* and *Zea mays*. In all these cases, globular embryos were formed, however, they degenerated six to ten days after pollination. The endosperm

was very poorly, if at all, developed. In crosses of Secale cereale with Hordeum bulbosum or Zea mays, endosperm development often appeared normal without embryos. Nevertheless, endosperm development collapsed in all cases before the tenth day after pollination. Using interspecific pollination of rye with maize pollen, Laurie et al. (1990) observed that when the rye variety Petkus Spring was pollinated with Seneca 60, fertilization occurred in 18.7% of the florets studied. Most fertilized florets had both embryo and endosperm and preliminary studies indicated that the elimination of maize chromosomes occurred early in the development of the embryo and endosperm. Based on cleared ovule preparations, the development of the embryo and endosperm was observed at 48, 72 and 96 h after pollination. Embryos were formed in eight cross combinations (3.3% to 23.3% of embryos per ovules depending on rye genotype), but in five combinations the ovules had both embryo and endosperm nuclei. In a sample of 344 cleared ovules, 28 (8.1%) had both embryo and endosperm, 15 (4.4%) had only embryo, and 13 (3.8%) had only endosperm (Ponitka and Ślusarkiewicz-Jarzina 2004). In our study, haploid embryos were formed in six genotypes from fifteen tested and ovary enlargement varied from 34.27% to 3.82% depending on the genotype. Moreover, as in the cited works, ovules in the current study had no endosperm irrespective whether they formed embryos or not.

Zenkteler and Nitzsche (1984) observed that none of the investigated cereal crosses seemed to fulfill the conditions necessary for haploid production. Independently of the detected haploid cells, the frequency of haploid embryo formation was too low for practical application. On the other hand, Zenkteler and Nitzsche (1984) stated that the formation of hybrid embryos offered new possibilities for intergeneric hybridization and gene transfer. There are certainly problems with embryo culture techniques concerning extremely young embryos, but they should be overcome in the near future. Using the wide crossing technique, Deimling et al. (1994) obtained six embryos, from which two DH lines were obtained after pollination of 48,000 emasculated flowers. One was induced after pollination with pearl millet, the other with maize. The effort required to emasculate and pollinate such a large number of rye flowers to regenerate so few haploids is far too great for interspecific pollination to be a suitable haploidization technique in rye. In our study, from 17,154 pollinated florets 21 haploid embryos were formed, but they did not germinate and convert into plants. Tenhola-Roininen et al. (2006) reported that due to the low survival rate of green regenerants (36-61%) and low fertility (3-29%), only about 10-36% of all the regenerated plants would have been suitable for rye research or for breeding purposes.

Conclusions

The development of DHs is more problematic in out-crossing self-pollinating species. The above results clearly showed that wide hybridization between taxonomically diverse cereal crops could be achieved, but such observations would be of limited interest if plants could not be produced. In all of the crosses, it has proved very difficult to recover plants using conventional embryo rescue techniques. As mentioned by Zenkteler and Nitzsche (1984), the reasons for this are still not known, but it is probably caused by the

absence or poor development of the endosperm. Admittedly, some rye haploid embryos and few plants have been obtained by wide crossing, but the embryo rescue method is still necessary.

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