



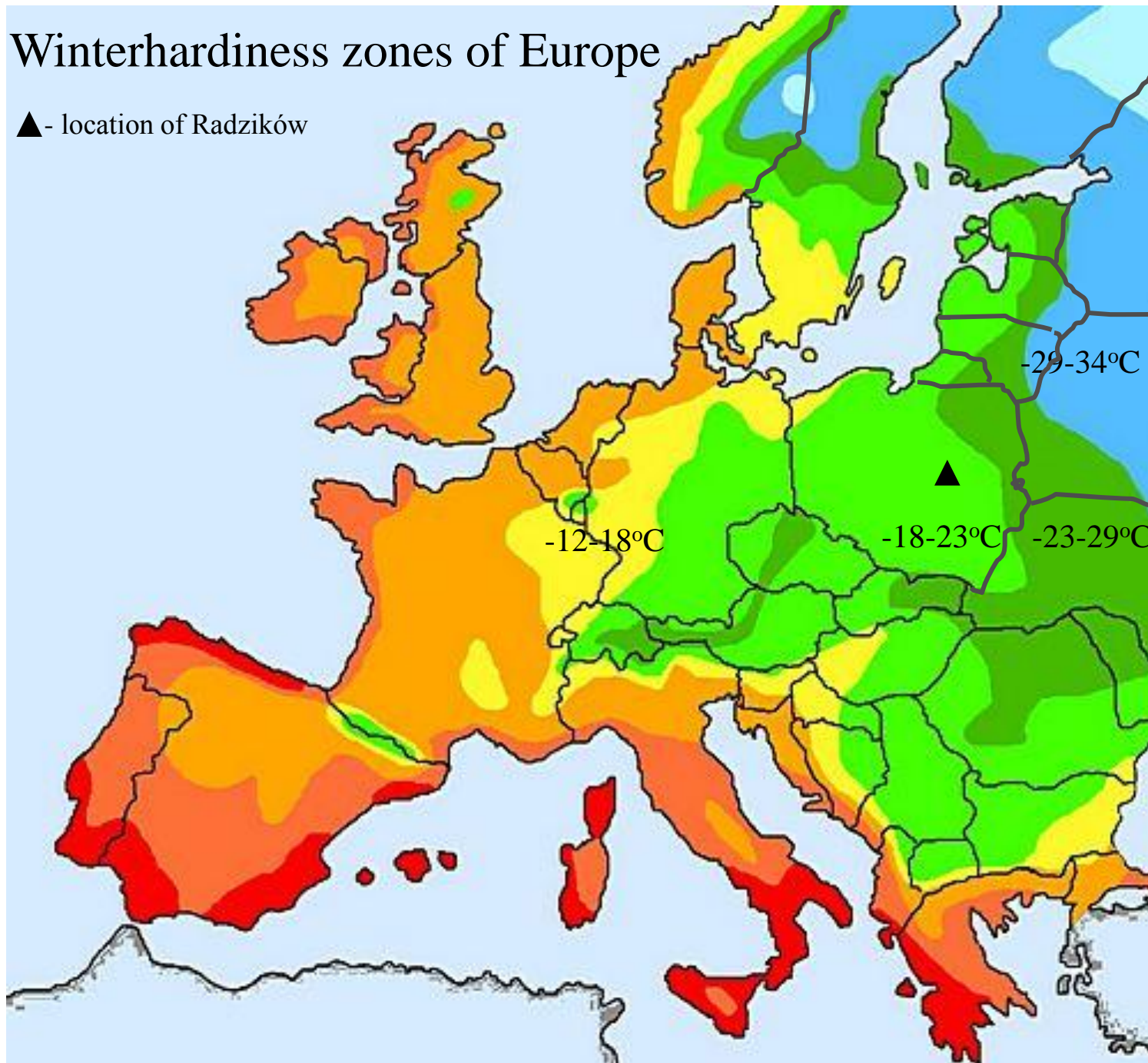
The ways of oat winterhardiness improvement in Poland

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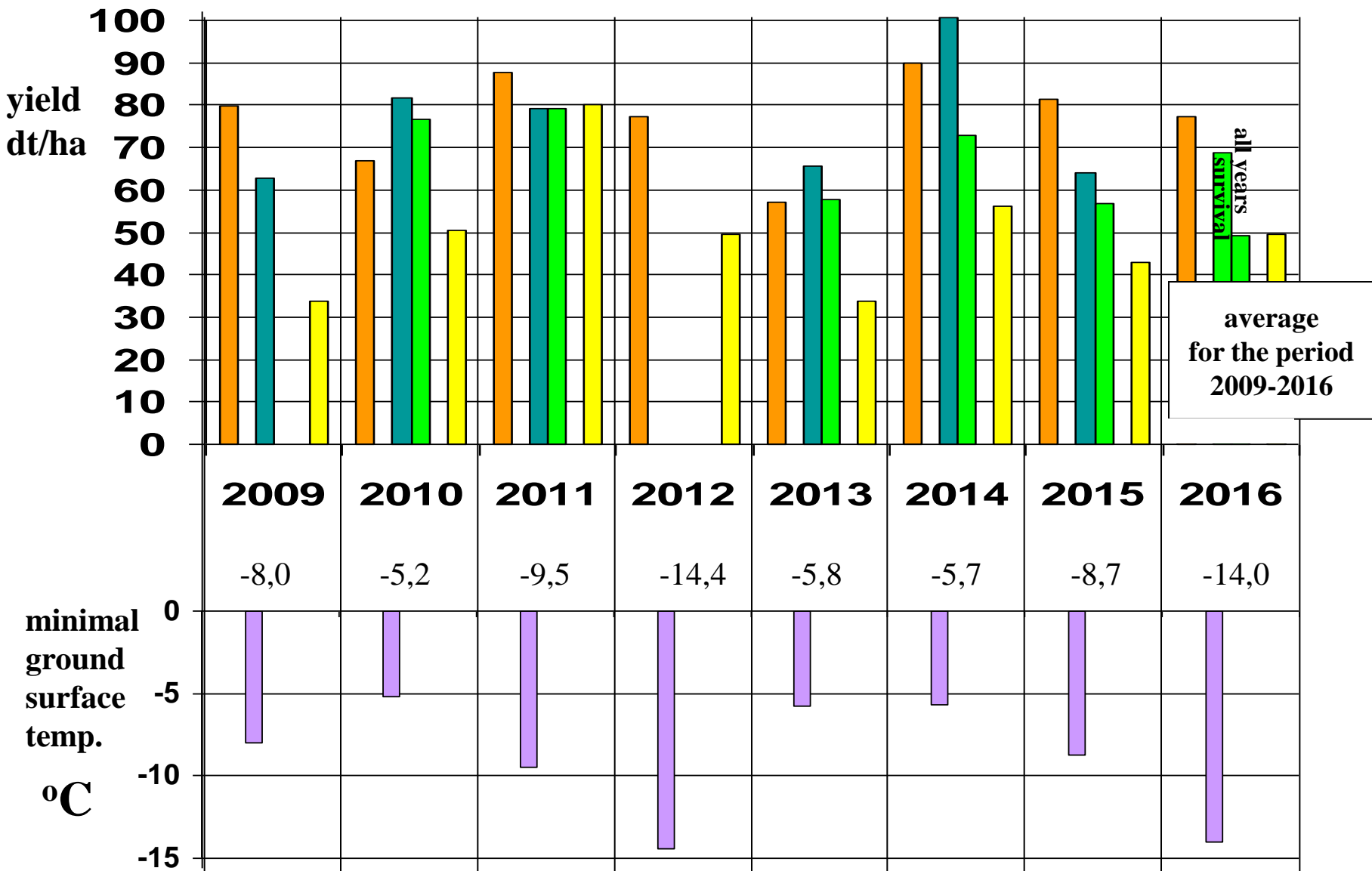
Winterhardiness zones of Europe

▲ - location of Radzików



Yields and minimal temperatures of ground surface. Radzików, 2009 - 2016

- 'Carola' – winter barley
- best yielding oat
- 'Krezus' – spring oat
- line 5Q5.2 (state trials)



Winter oat trial in Krzeczowice. February 2016

BARLEY



**Loss of chlorophyll in winter oat.
Radzików 2006**





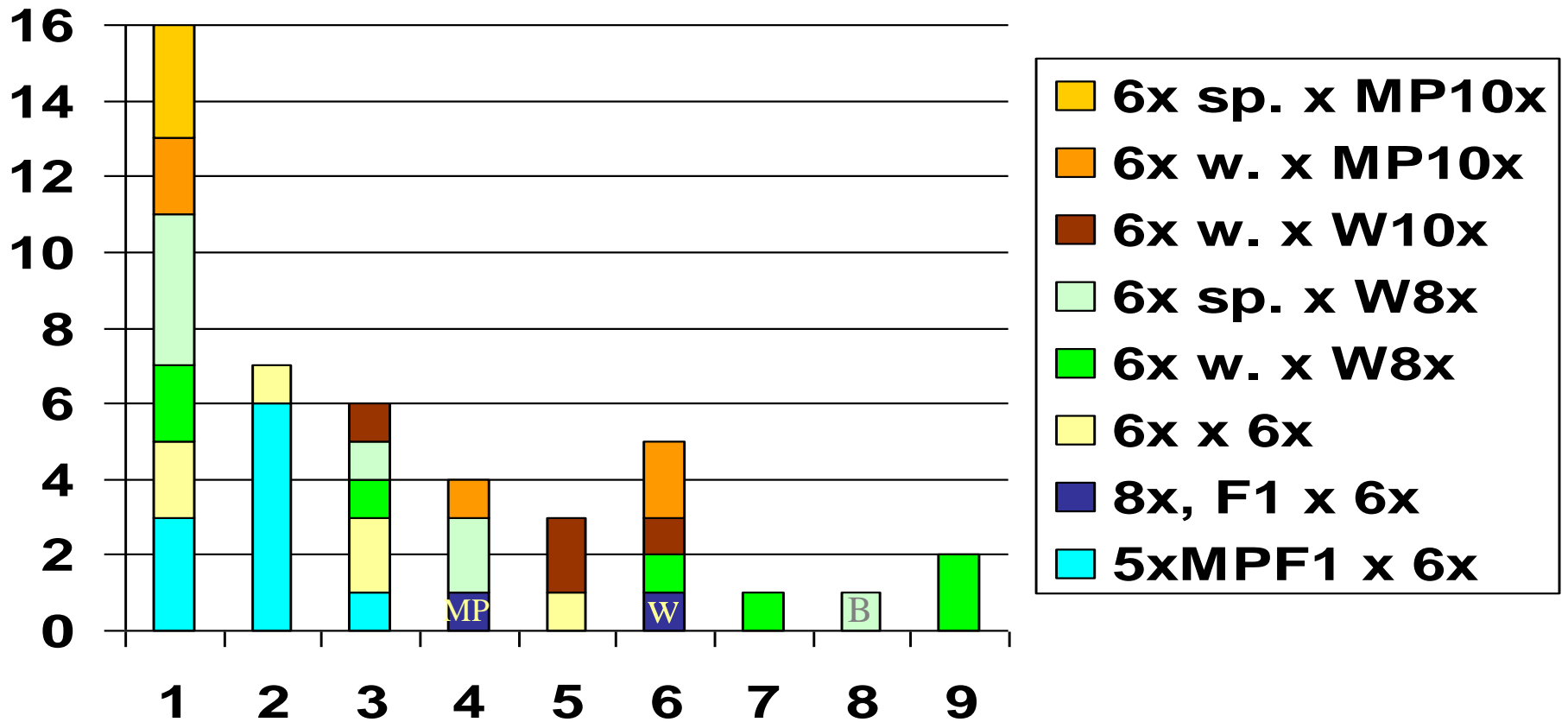
Avena macrostachya

wild autotetraploid
endemite from Atlas mountains
most distant to other *Avena* species,
outcrosser,
perennial,
winterhardy,
resistant to some diseases and pests



The octoploids and decaploids at winter sowing

Distribution of cold resistance scores in the best selections of 45 crosses between winter (w.) or spring (sp.) hexaploids (6x) of *A. sativa* and octoploid (8x) or decaploid (10x) hybrids (*A. sativa* x *A. macrostachya*) produced from the winter-hardy *A. sativa* cultivar ‘Wintok’ (W) or a non-winterhardy ‘Mirabel-Pendragon’ (MP) line.

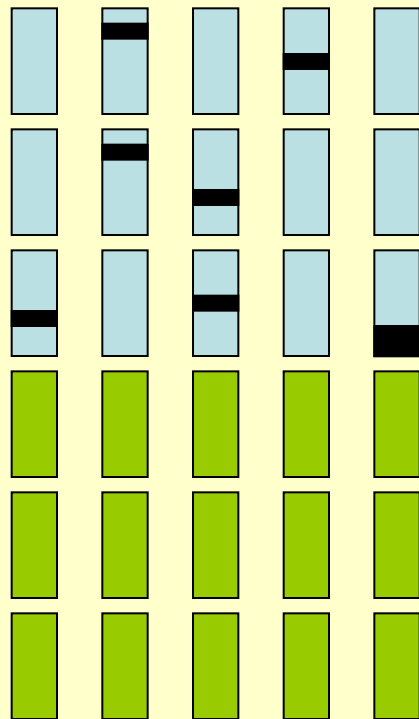


Breeding strategies after wide crossing:

	introgression	incorporation*	allopoloidization
trait genetic architecture:	simple	composed	composed
what transferred:	small fragment of alien DNA	numerous DNA fragments	whole alien genome
ploidy level	maintained	maintained	increased
breeding method:	back-crossing	intercrossing among recombinants	chromosome doubling and selfing

*) Simmonds N.W. 1993. Introgression and incorporation, strategies for the use of crop genetic resources. Biol. Rev. 68:539-562.

Which parent for crossing is better for re-building a genetic system decomposed by interspecific recombination?



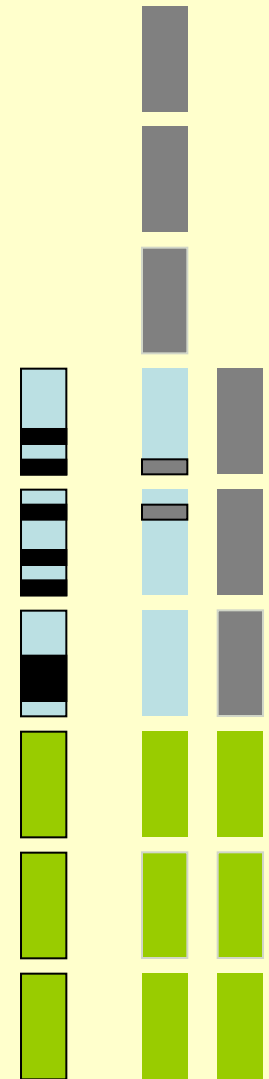
?

low
comple-
mentation
chance



?

meiotic
pairing
reduced



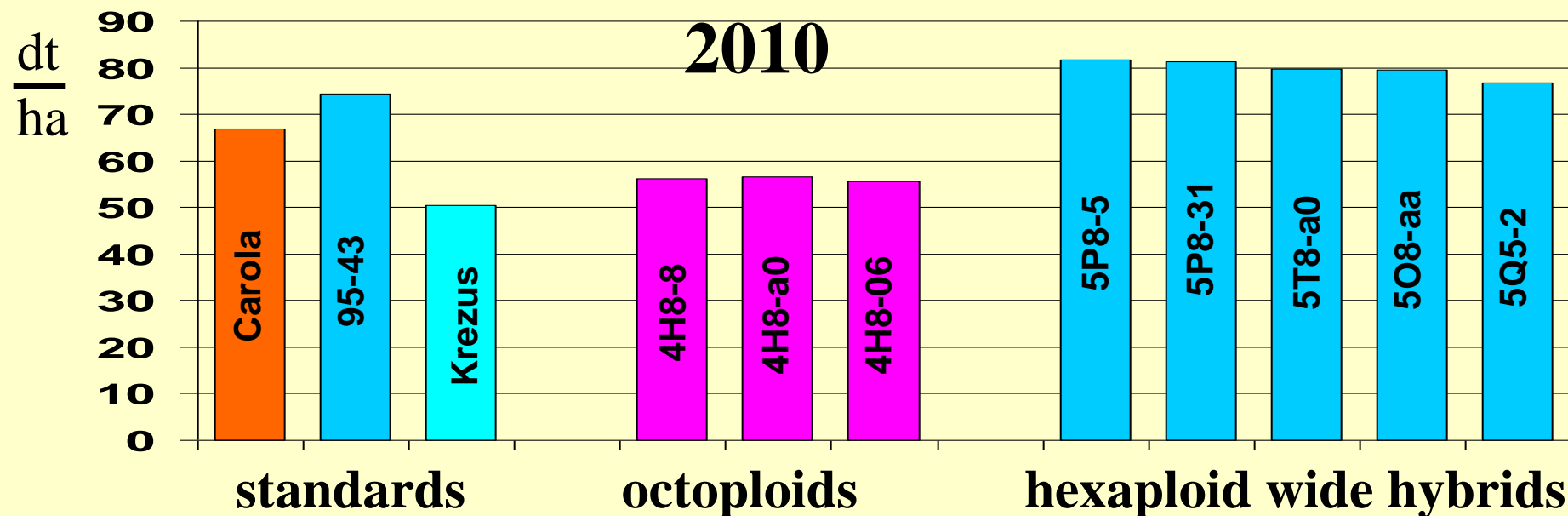
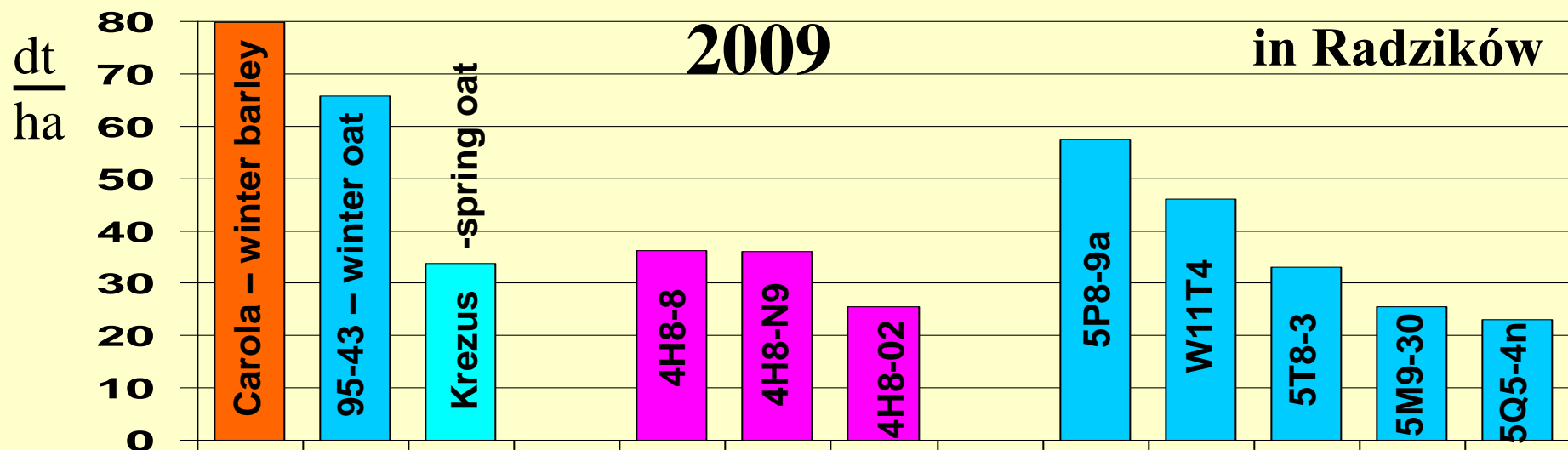


Octoploid lines, the 4H8 cross, 23.06.2008

Comparison of early allopoloids in triticale and oat

	8x or 6x triticale	8x oat A.s x A.m.
yield reduction in primary synthetics	~50%	~30%
time necessary to restore yield potential	~50 gns	?
winterhardiness	weak	high
time necessary to restore winterhardiness	~ 50 gns	none
disease resistance	high	variable
grain	large shrivelled	large plump
grain nutritious value	high	high

Yields of winter oat hybrids of *Avena sativa* with *A. macrostachya*





Krezus (TKW 25,7 g)



4H8.a (TKW 55,2 g)



6x




8x

Krezus (HC 26,2 %)

4H8.a (HC 25,0 %)

Conclusions

1. Crossing of common oat with 8x or 10x *sativa-macrostachya* allopolyploids is effective in transgressive improvement of winterhardiness.
2. Some spring oats are good cross components for winterhardiness improvement.
3. Relatively high performance of the *sativa-macrostachya* primary octopolyploids is a good prognostic for application of allopolyploidy in improvement of oat.
4. In spite of risk of winter killing, cultivation of the hexaploid oat wide hybrids in Poland is economically feasible on sufficiently fertile soils, in regions with stable snow cover.

A field of green oat plants with a yellow text box in the center. The text box contains the text "Thank you for the attention".

Thank you for the attention

Literature

Prospects for allopoloidization in oat – in relation to results in triticale

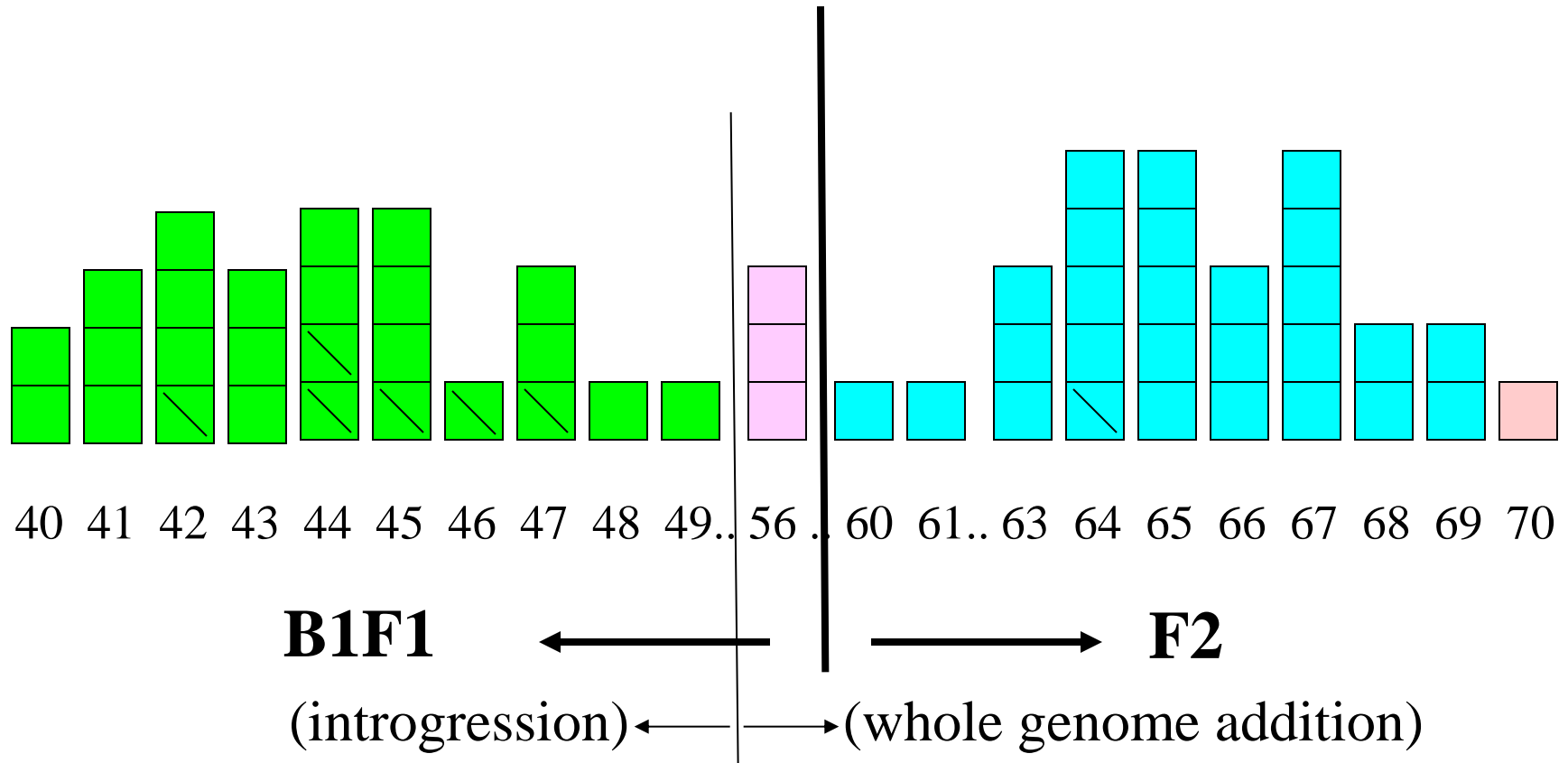
Hede A. 2001. A New Approach to Triticale Improvement. Paper presented at the Research Highlights of the CIMMYT Wheat Program, 1999–2000, Oaxaca, Mexico.

Loss of DNA

Ma X.F., Gustafson J.P. 2008. Allopolyploidization-accomodated Genomic Sequence Changes in Triticale. *Ann Bot.* 101(6): 825-832.

Chromosome numbers in B1F1 and F2 of the *Avena sativa* x *A. macrostachya* hybrids, 2004.

☐ - death after germination





8x

Winter oat nursery 23.04.2006



SIWEK (6x, cultivar)

TKW = 22,6 g



7.B/12 (6x x 10x, F₄ plant)

TKW = 29,1 g

Crossing and propagation of the hybrids:

No of cross combinations

No of pollinated flowers

No of F1 seeds obtained

embryo rescue

No of F1 plants recovered

greenhouse cloning

No of F1 plants after cloning

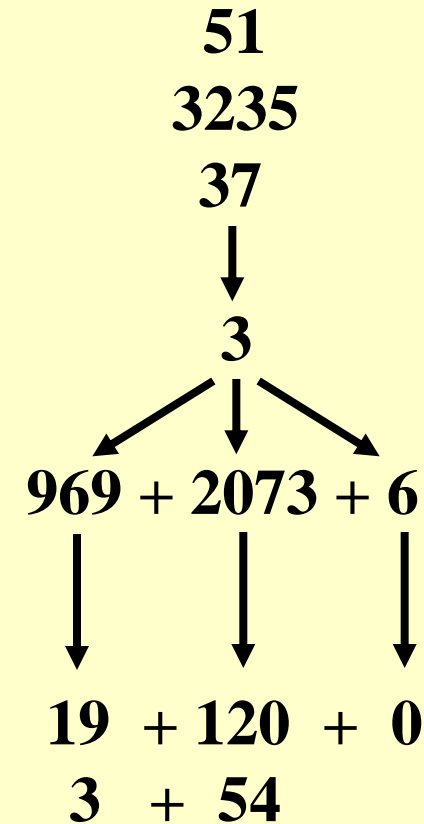
colchicine treatment

free winter oat pollination

No of F2+B1F1 seeds

No of F2+B1F1 seedlings

counting of chromosomes



Expected changes in genomic composition

