# Development of protein rich tetraploid oat- current state and prospects

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- Avena magna has the highest protein content (23%-27%) of all oat species
- The transfer of this protein content to the common oat is hindered by:
- <u>Difference in chromosome numbers</u> (2n=28 in *A. magna* and 2n=42 in *A. sativa*) resulting in hybrid self sterility but some seeds may be produced by pollination with parental pollen
- Protein content is a trait of quantitative nature

## New approach to exploit *A. magna* protein content

- <u>Domestication</u>: transferring the domestication syndrome of *A. sativa* to *A. magna*
- <u>Domestication syndrome</u>: A small number of traits that makes the difference between wild and domesticated state
- In oats, <u>fundamental character</u> of this syndrome is seed non-shattering at maturity, accompanied by glabrous and yellow lemmas and reduction of awns number and size and erect growth

### Genetics of domestication syndrome characters

- Each of these traits shows mono-genic inheritance
- Some of the characters are linked to each other
- As a result of this inheritance, the domestication syndrome can be transferred via the pentaploid bridge, in spite of the sever sterility of the hybrids

# The experimental procedure: first hybridization cycle

- 1. Crosses of A. sativa x A. magna
- 2. F1 x A.magna (wind pollination)

3. BC plants exhibit <u>variation in fertility</u>
(0-20%) and chromosome numbers (28-32).

 4. F2, BC selection of individuals for the next cycle

### Second hybridization cycle

- 1. A. sativa x Hybrid derivatives (2n=28)
- 2. F1 x Hybrid derivatives
- 3. BC, no wild phenotypes occur, selection for fertility
- 4. F2,BC, selection for fertility and individuals for initiating the third cycle
- The same procedure for the 4<sup>th</sup> and 5<sup>th</sup> cycles

#### Outcome of the hybridization cycles

2n=28 plants are morphologically indistinguishable from the common oat







#### THE PROTEIN

- 1. Protein content varied (16%-23%) among progeny of the 5<sup>th</sup> hybridization cycle
- 2. Individuals with the highest protein values were crossed again to wild *A. magna*
- 3. F1 hybrids were fertile
- 4. Individuals with wild phenotypes were eliminated in F2
- 5. Individuals with the highest protein contents were crossed with one another and the segregating generations have been reared by single seed descent(now at F4)

### Germplasm enhancement of the 2n=28 oat

 Number of traits are being introduced now to the domesticated 2n = 28 oat:

- Semi dwarf stature
- Day length insensitivity
- Naked seed

#### The future of domesticated 2n=28 oat

 1. Domesticated 2n=28 oat can be established as a crop if its protein yield would be superior to any common oat cultivar

• 2. Attaining the highest protein content per land unit can no longer be a one-man's project and requires collaborative efforts.

#### **Contact Information**

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