

# Next steps for perennial grain crops in Australia



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PROFITABLE PERENNIALS™ FOR AUSTRALIAN LANDSCAPES



# New studies

## Yet to be studied:

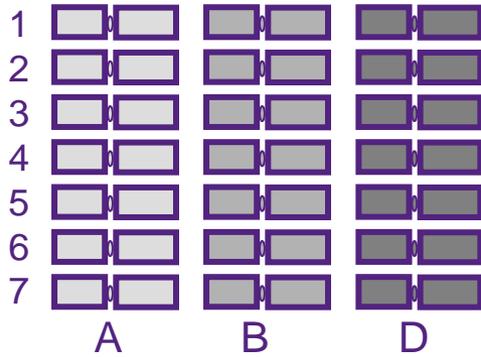
- \* Physiology of root systems – of perennial selections [PhD student]
  - \* Root mass and dynamics
  - \* water and nutrient dynamics and partitioning
- \* Forage management effects
- \* The physiology of summer survival
  - \* dormancy, dehydration tolerance
- \* Forage and grain quality
  - \* Grain quality assessment initiated (CSIRO)
- \* Other fungal diseases
  - \* yellow spot, nodorum, tritici, FHB
- \* Root and crown diseases
  - \* take-all, crown rot, common root rot, CCN, RLN



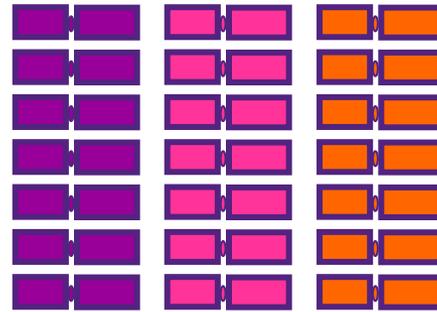
## Prospects for ongoing breeding of perennial wheat

- \* From what we have learned it seems sensible to breed deliberately at the amphiploid level for adequate perenniality
- \* It is greatly preferable to use a diploid perennial donor
- \* Comparison to triticale at 6x (ABR) or 8x (ABDR) level
- \* If the donor is a allopolyploid (mixed genomes) such as *intermedium* or *ponticum*, then the extra “genome” is synthetic and variable making subsequent interbreeding chaotic and inefficient ....

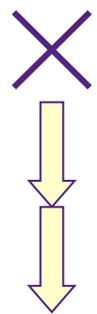
# Wheat



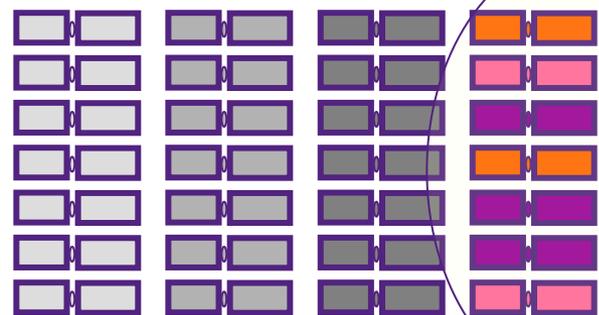
# *Thinopyrum intermedium*



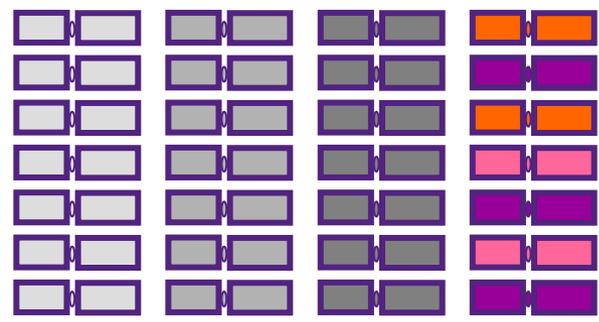
**Chromosomes**  
(showing one to represent a pair)

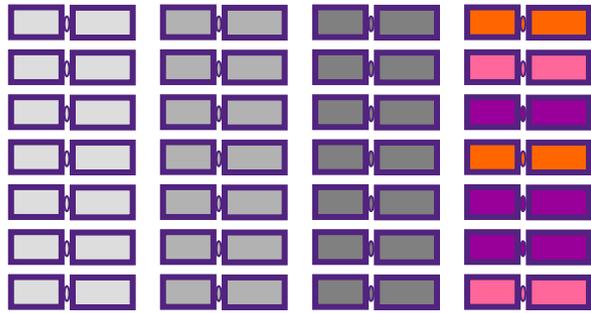


**Partial amphiploid**  
 $2n=56$

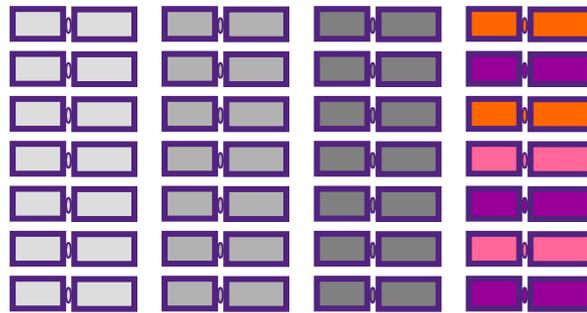


Variable composition of synthetic genome

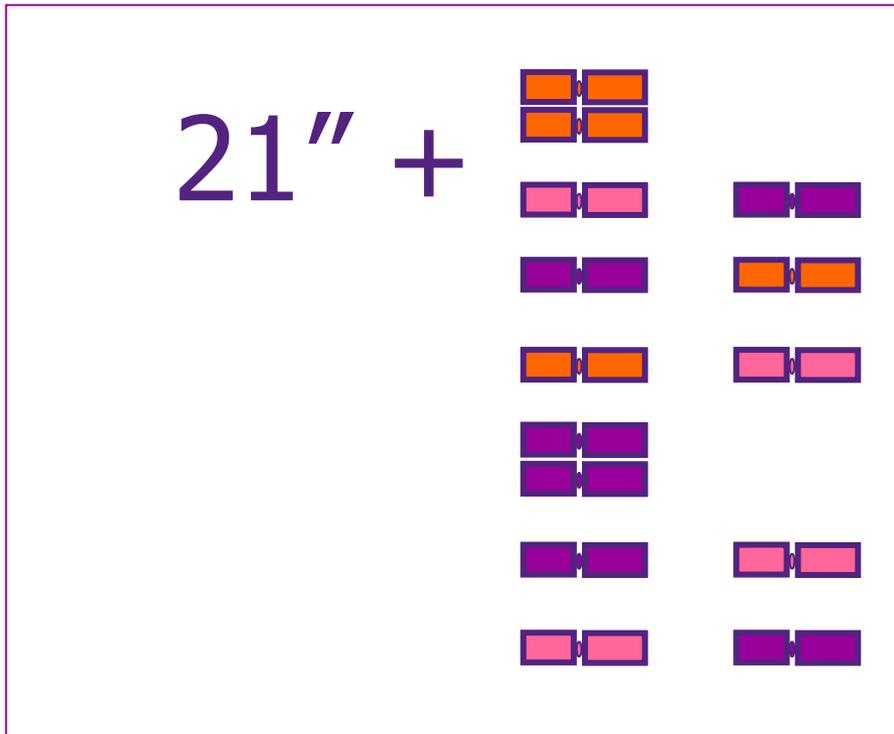




X



Showing one chromo to represent a pair



self



Variable chromo loss and instability



## Allo-polyploid donor makes subsequent breeding at the amphiploid level very difficult

- \* You make or collect a series of foundation lines, AABBDDXX, where X is a synthetic genome
  - \* Very difficult to recover stable  $2n=56$  progeny
  - \* Examples of such difficulties:
    - \* intercrossing Zhong lines (Banks et al 1993)
    - \* Ot38 x Zhong lines
    - \* OK72 x Ot38 (derivatives of *ponticum* and *intermedium*)
  
- \* Intercrossing synthetic octoploids makes a mess



## ***Elongatum* breeding**

- \* ***Elongatum* amphiploid breeding:**
  - \* The WSU program has demonstrated that the E genome is sufficient
  - \* Select adapted naturalised populations of *L. elongatum* ( $2n=14$ )
  - \* Cross to various 4x or 6x adapted wheats
  - \* Chromosome double F1 plants (colchicine)
  - \* Stabilise AABBEE and AABBDDDEE lines through selfing
    - \* Possibility that the genome balance ABE might be more robustly perennial than ABDE
  - \* Should be able to create many *foundation* breeding lines which are mutually compatible yet with genetic diversity to drive the breeding program



# *Elymus scaber* breeding



*Elymus scaber*, Common wheatgrass

- \* ***Elymus*** amphiploid breeding [Matt Newell postgrad project]
  - \* Assessment of five diverse accessions of *Elymus scaber*
  - \* An Australian endemic perennial Triticeae grass
  - \* Widely adapted
  - \* *E. scaber* appears to be a allo-hexaploid ( $2n=42$ ), genomes SYW
  - \* Some forms are  $2n=9x=63$
  - \* Interestingly it is ~95% apomictic
  
- \* *Elymus scaber* x wheat (4x or 6x, highly crossable genotypes)
- \* Chromosome double the F1 hybrids
  - \* Because 9x forms of scaber exist and are stable, possibility that ABSYW or ABDSYW might also be stable
  - \* Hybrids have been produced between wheat, barley and rye with other *Elymus* species



## Other breeding options

- \* Select increased grain yield and grain size within perennial grass
  - \* *Secale montanum*
  - \* Wheatgrass species
- \* Engineer grain size genes into perennial grass, e.g.
  - \* RNAi of wheat glucan water dikinase (GWD) gene (Ral et al 2010)
  - \* Overexpress wheat ExpA6 gene (Lizana et al 2010)
  - \* Overexpress rice GS3 gene (Li et al 2010)
  - \* Overexpress Brassica wri1 gene (Liu et al 2010)
  - \* Overexpress grain sucrose synthase gene
  - \* Apetala2 (lec 2 is an AP2)
  - \* Tomato Sw4.1 ABC transporter (Orsi and Tanksley, 2009)



## An international research possibility... sowing an idea you will hear more of on Thursday

### Molecular control of perenniality in grasses

- \* Use the power of the genomic tools of *Brachypodium* to find genes controlling perenniality in the Triticeae
- \* Iain Wilson (CSIRO) and John Vogel (USDA) showing interest in exploring the molecular genetics of perenniality with us
  - \* *B. distachyon* is an erect annual grass
  - \* There are also perennial species of the genus including *pinnatum*, *sylvaticum*, *glauco-virens*, *mexicanum*, *arbuscula*, *retusum*, *rupestre* and *phoenicoides*.
  - \* They can be intercrossed and hybrids were vigorous and long-lived
  - \* With some annual x perennial crosses there might be prospect of mapping perenniality
  - \* The perennial species can be sequenced and compared to the annual to find candidate genes for perenniality.