





International Adaptation Trial 2006

Ky L Mathews Tom Payne

Scott Chapman Jose Crossa

The Goal



- Better varieties to Australian growers, faster
- Evaluate Australian and CIMMYT germplasm in Australia and global wheat production regions
 - understand Australian germplasm on a global scale
 - link Australian production environments to global wheat production regions
- Probe and reference genotype sets
 - isolines, related lines
 - broadly adapted reference lines

Presentation Overview



- Results from the 2006 season
- Research overview
 - global adaptation of Australian and CIMMYT germplasm
 - 'window-on-wheat': a real-life example!
 - molecular data
 - protein analysis
- Planning for 2007
 - CIMMYT
 - sites
 - entries
 - research

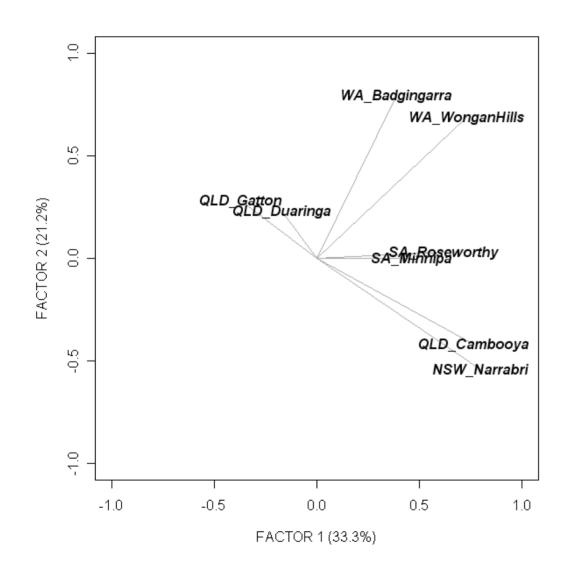
2006 locations





2006 Trials

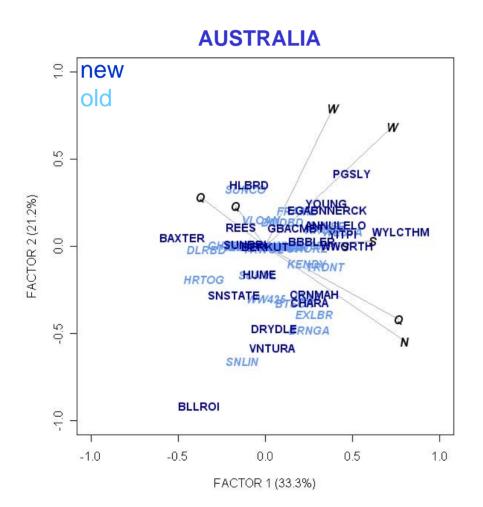


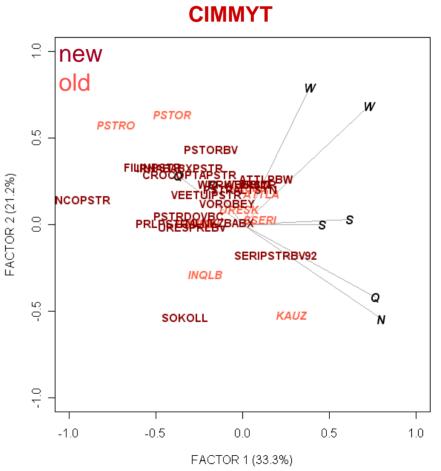


Trial	Yield (t/ha)	h _G ²
QLD_Gatton	4.90	0.864
QLD_Duaringa	3.37	0.907
WA_Badgingarra	1.16	0.706
WA_Wongan Hills	0.95	0.629
SA_Roseworthy	2.33	0.868
SA_Minnipa	0.40	0.777
QLD_Cambooya	4.37	0.939
NSW_Narrabri	3.22	0.949

And the genotypes...







2006 overall top performers



based on BLUPs from across site analysis which models between trial variance heterogeneity and correlation

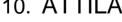


- WYALKATCHEM
- ANNUELLO
- WENTWORTH



- YITPI
- WESTONIA
- **SILVERSTAR**
- 9. KRICHAUFF
- 10. ATTILA

- 11. EGA BONNIE ROCK
- 12. BABBLER
- 13. GBA COMBAT
- 14. NESSER DWARF
- 15. KAUZ DWARF
- 16. ATTILA*2/PBW65
- 17. TRIDENT
- 18. SUPER SERI #1
- 19. WEEBILL1
- 20. FRAME



Yield Range: 2.6 – 3.7 t/ha



Global Adaptation

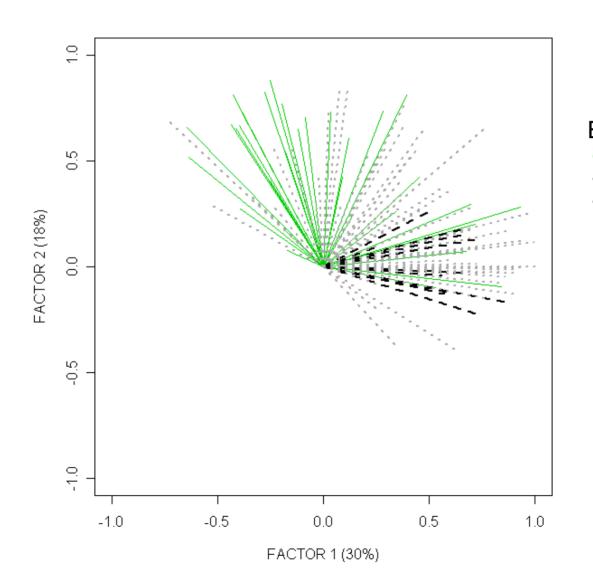


Mathew et al 2007 Global adaptation patterns of Australian and CIMMYT spring bread wheat

- investigate the GxE patterns for grain yield in the IAT
- pedigree relationship matrix used to partition the additive and non-additive genetic line effects Oakey et al (2006) TAG
- maturity used to explain the observed adaptive patterns

Global G x E





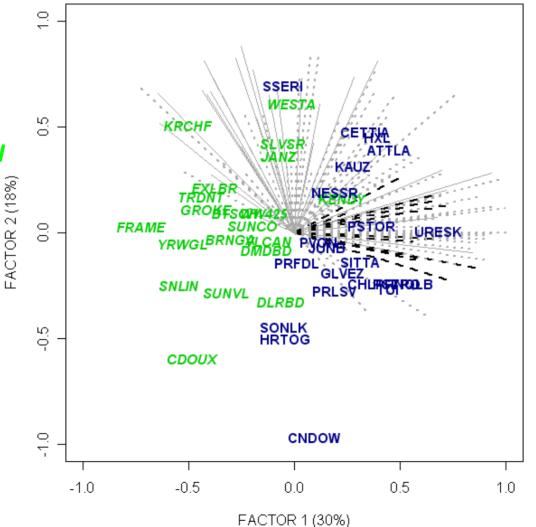
Environments



Global G x E







Environments

AustraliaCIANO

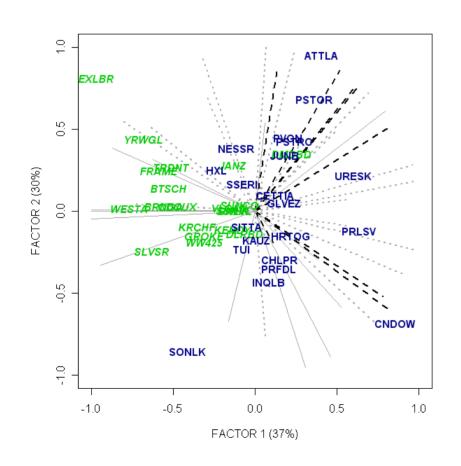
.....international

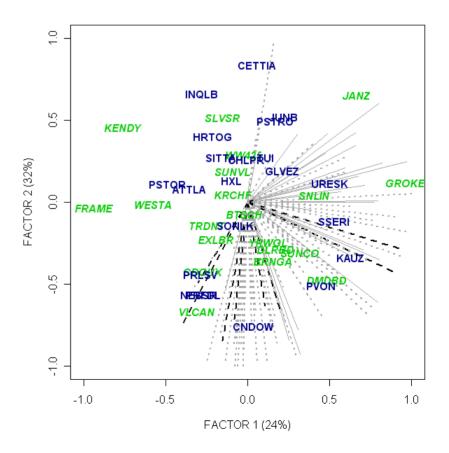
Additive and non-additive by environment



additive by environment

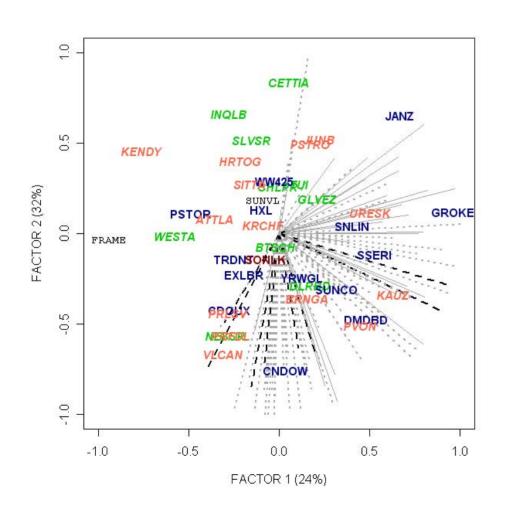
non-additive by environment

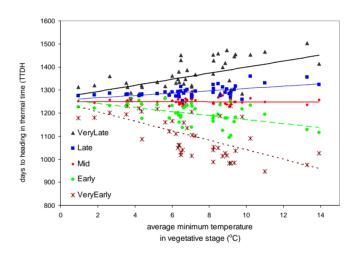




Maturity Class







VERY LATE
LATE
MID
EARLY
VERY EARLY

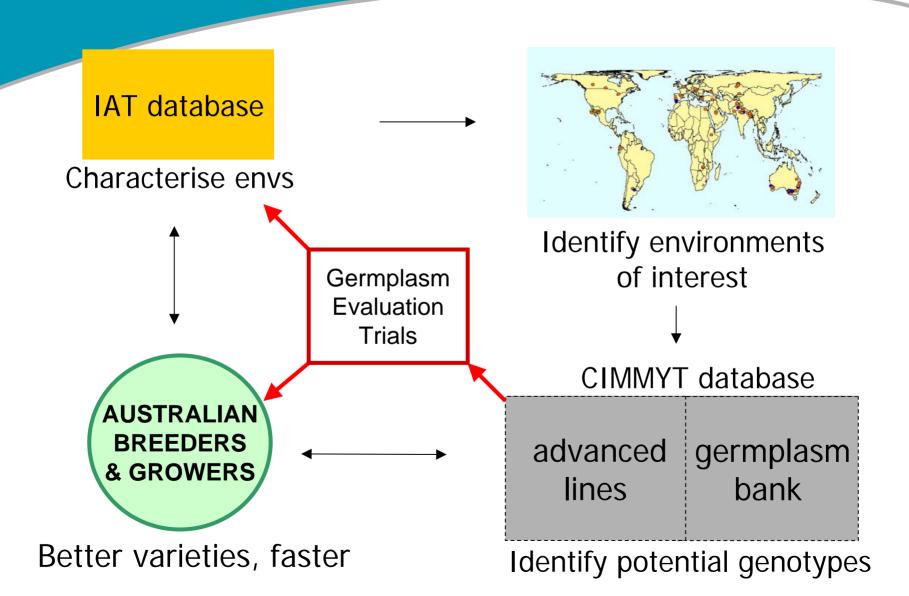
Key Points



- multiplicative mixed models be used in the analysis of multi-environment trials
- inclusion of pedigree information quantifies additive and non-additive effects
- Australian and CIMMYT germplasm are well adapted to their respective target environments and these environments are diverse
- maturity explains some of the adaptation differences between Australian and CIMMYT germplasm
- CIANO is a good source for northern Australian sites; southern sites may be better investigating later maturing lines suitable to the mega-env 9

Window-on-wheat





A real-life example! Selecting lines for high rainfall, irrigated trials



Identified three types of environments to select from

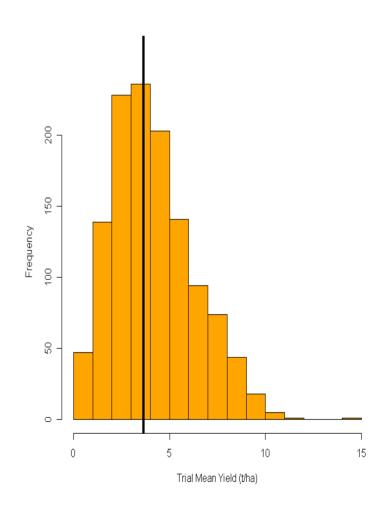
- 1. irrigated and high yield international trials, >4 t/ha
- international locations which correlate with southeastern Australia e.g. Wagga Wagga, Horsham, Robinvale, Moombooldool, Junee
- 3. international environments which correlate with irrigated IAT trials e.g. Tamworth.

International Nursery Data



Data used from IWIS (ICIS?)

- 1994 to 2004
- ESWYT, HRWYT, SAWYT
- traits: yield, days to heading, height
- 1231 trials
- 1253 genotypes
- 335 locations



The process of a real-life example! Selecting lines for high rainfall, irrigated trials



1231 trials; 1253 genotypes

- 1. irrigated and high yield international environments
 - > 4 t/ha and irrigated = yes
 - 341 trials with 1231 genotypes
- 2. international environments which correlate with south-eastern Australia
 - Wagga Wagga and Robinvale used
 - 199 trials with 1253 genotypes
- 3. international environments which correlate with irrigated IAT trials e.g. Tamworth
 - 241 trials with 1252 genotypes

Lessons from a real life example...



- Takes 2-3 'Ky-days'
- decreased the number of trials to investigate signficantly
- did not decrease the number of genotypes
- only single trial summaries provided
- so.... future developments
 - perform a GxE analysis on the selected trials and provide an across site BLUP to better identify the top performing lines, and hopefully decrease the quantity!!!

Other research activities



Molecular work

- DArTs have been run across 94 lines, not including durums
- data received 2 weeks ago
- Scott Chapman to outline future work

GxE quality

- seed supplied to Colin Cavanagh (CSIRO, Canberra) from 5 trials in 2005
 - Robinvale, Morowa, North Yuna, Wagga Wagga, Duaringa
 - Narrabri and Tamworth analysed in Brisbane
- analysis in progress...

Planning for 2007



CIMMYT

- Tom Payne providing entry list, some new lines from Ravi
 Singh and Yann Mannes
- 6 sets sent to southern latitude locations

Australia

- IAT grown with 5 Germplasm Evaluation Trials
- seed to be distributed by Agrisearch using DAFWA seed source
- entries see handout

research

- probe paper: crown rot and boron tolerance
- DArT
- protein

Acknowledgements



- Grains Research and Development Corporation
- International and Australian collaborators who grew the trial (and returned data!)
- Helena Oakey for assistance with pedigree analysis
- Andrew Milgate for requesting the window-on-wheat
- Chris Moore/Robin Wilson, DAFWA for seed source for 2007