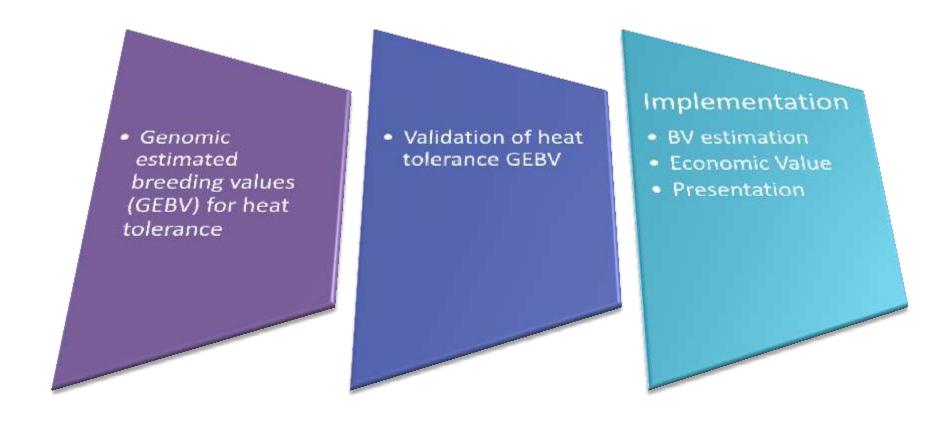


# Genomic selection for heat tolerance in Australian dairy cattle

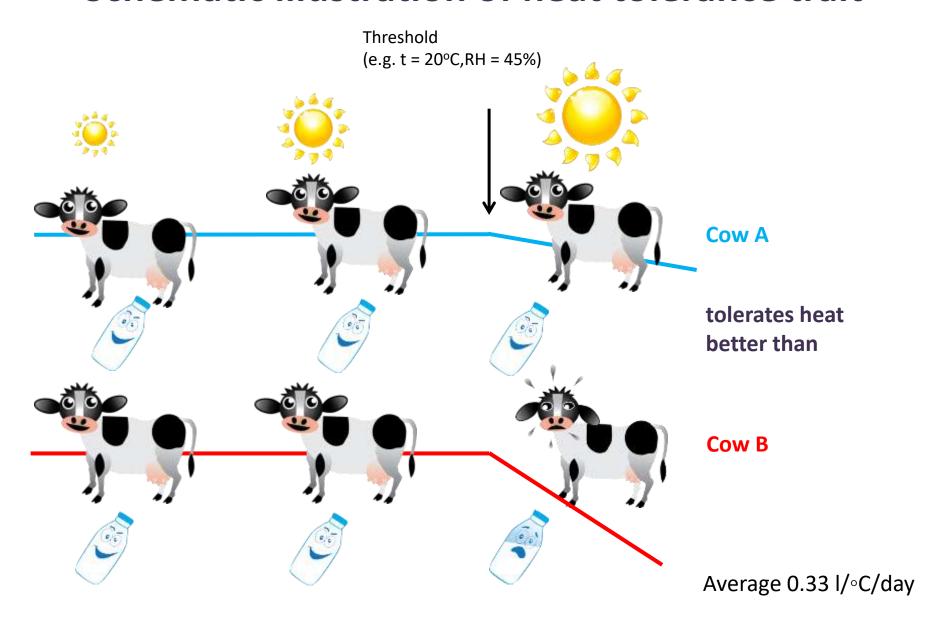
Thuy T.T. Nguyen, Josie Garner, M. Douglas, R.O. Williams, W.J. Wales, L.C. Marett, P.J. Bowman, C. Reich, M. Haile-Mariam, J.E. Pryce & B.J. Hayes



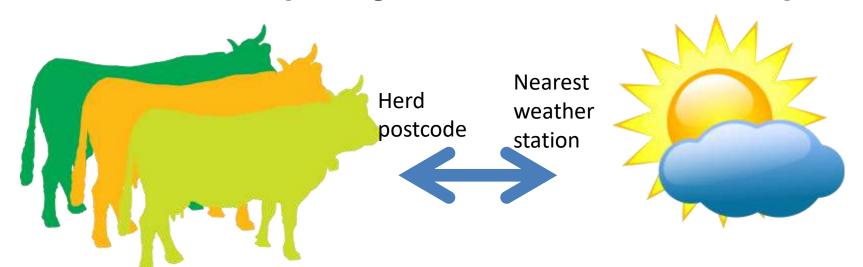
### **Outline**



#### Schematic illustration of heat tolerance trait



## Data (11 years, 2003-2013)

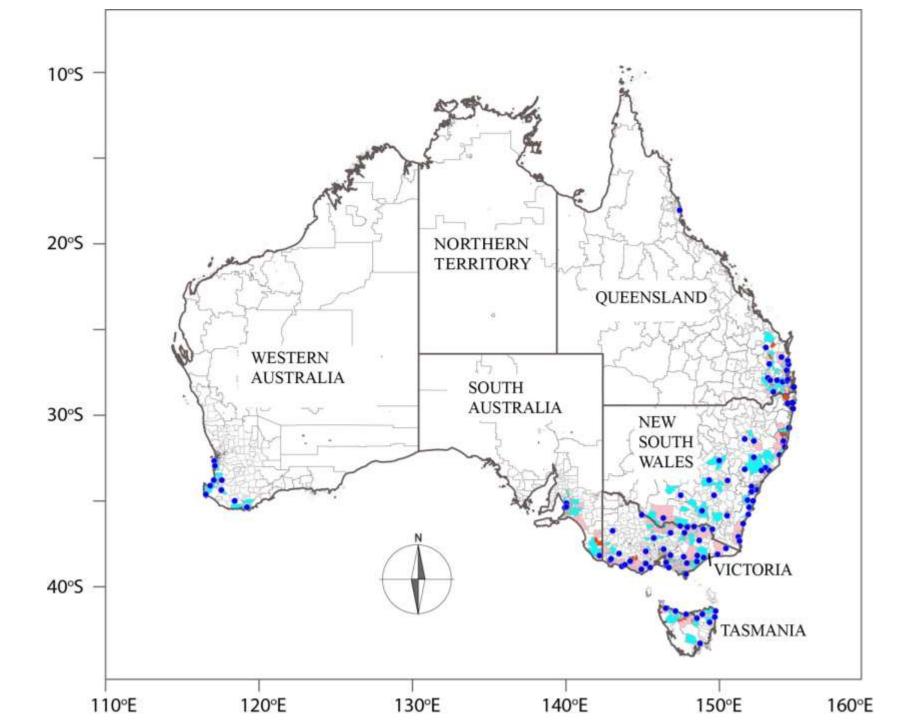


#### Herd recording

- Milk, fat, protein yields
- -1<sup>st</sup> -3<sup>rd</sup> parities
- Holsteins + Jerseys

#### B. of Meteorology

Temperature-Humidity
 Index (THI) (mean THI of test day + 4 days prior to test day)



## **Phenotypes and Genotypes**

Data	Holsteins	Jerseys
	1st parity	1st parity
Herds	1,762	519
Herd test dates	85,714	26,441
Number of cows	366,835	76,852

#SNP	Holsteins	Jerseys	
800K	1,620 sires	125 sires	Imputed
50K	1,115 sires 2,189 cows	585 sires 1,188 cows	using BEAGLE



## Models: using ASREML

Cow slope

•  $y_{ijklm} = \mu + HTD_i + YS_j + PAR_k + \sum_{n=1}^{3} A_n X_n + PAR_k \sum_{n=1}^{8} D_n Z_n + STG_l \sum_{n=0}^{1} T_n S_n + \sum_{n=0}^{1} C_{mn} W_n + e_{ijklm}$ 

Sire slope

Daughter trait deviation (averaged daughter slopes)

one slope

• 
$$\mathbf{y} = \mathbf{\mu} + \mathbf{Z}\mathbf{g} + \mathbf{e}$$
,  $\mathbf{g} \sim N(0, \mathbf{GRM}\sigma_g^2)$ 

**GBLUP** 

y = a vector of sire slope and cow slope

## **Genetic parameters**

Trait impacted	h <sup>2</sup> of cow slope		
	Holstein	Jersey	
Milk yield	0.22 ± 0.007	0.33 ± 0.018	
Fat yield	0.20 ± 0.007	0.26 ± 0.015	
Protein yield	0.23 ± 0.007	0.27 ± 0.016	



## Accuracy of genomic prediction

Breed	Reference	Validation	Trait affected by heat stress	A	ccuracy	1
Holsteins	2,300 sires	435 sires	Milk		0.43	
	2,189 cows		Fat		0.46	
			Protein		0.51	ı
						l
Jerseys	575 sires	135 sires	Milk		0.49	
	1,188 cows		Fat		0.55	
			Protein		0.52	



## Correlations of heat tolerance GEBV with EBV of fertility

Breed	Heat Tolerance	Fertility
Holsteins	HT Milk	0.39
	HT Fat	0.38
	HT Protein	0.29
Jerseys	HT Milk	0.27
	HT Fat	0.21
	HT Protein	0.15

## Validation study

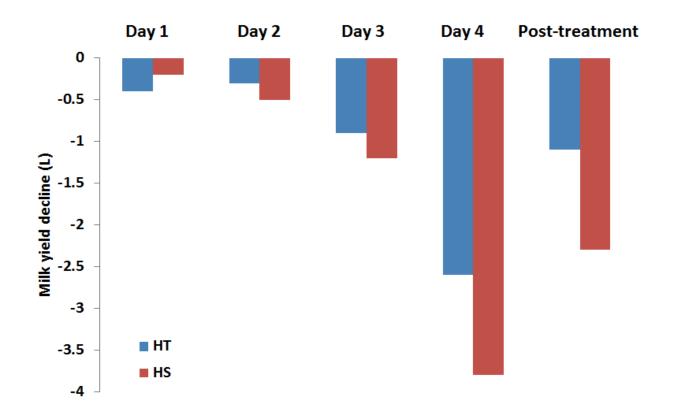
- 400 random heifers
- 24 predicted most heat tolerant, 24 predicted most heat susceptible based on GEBV
- Run through a simulated 4 day heat wave event in respiration chambers
- measure milk production, core temperature





## Validation study

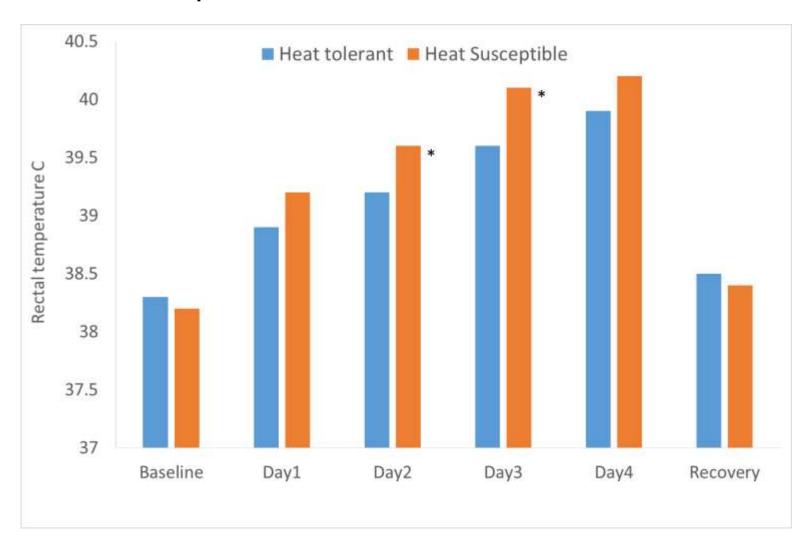
#### Decline in milk production





## Validation study

#### Rectal temperature



#### **IMPLEMENTATION**



#### **Economic Value**

Heat tolerance 
$$(\$) = \left(\begin{smallmatrix} EW_m * GEBV_{HTm} \\ EW_f * GEBV_{HTf} \\ EW_p * GEBV_{HTp} \end{smallmatrix}\right) * Heat Load$$

Heat Load varies by herd

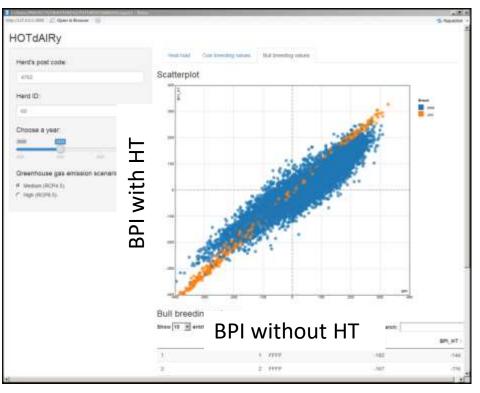


## **Bull breeding values**



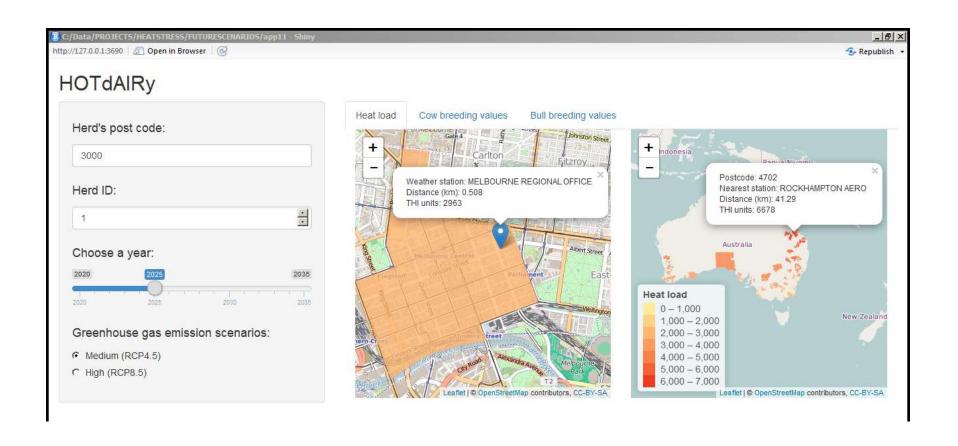
#### **HOTdAIRy** Herd's post code Scatterplot Hartl ID: -Choose a year. 노 **BPI** with F Venus (ROTAT) C NUMBER OF COMMUNICATION Bull breeding without BPI without HT

#### Queensland





#### **Heat load**





### **Implementation**

Location of herds (postcode) Projected future climate data (CSIRO) Marker-only EBV Location-specific economic value (index)



#### **Further work**

Effect of Heat on Fertility

Include heat data in routine analyses

Proper GEBV combining EBV + DGV

Include Heat tolerance in BPI



#### Conclusions

 Genomic selection can be used as a strategy to improve heat tolerance in dairy cattle

- Dairy industry will have validated genomic breeding values to improve heat tolerance
  - Favourable correlation with fertility
- Further work
  - Impact on fertility



## Acknowledgements

- The Department of Agriculture of Australia funded the project
- The Australia Dairy Herd Improvement
   Scheme provided production data
  - Test day data
  - BPI: Paul Koh & Gert Nieuwhof
- Dairy Futures CRC
- CSIRO and Bureau of Meteorology –
   Climate Change in Australia
  - Dr. John Clark











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