

FORUM

For the latest in red meat R&D

Improving within-breed genetic evaluation and developing multi-breed genetic evaluation with the Southern Multi-Breed Project

Brad Walmsley

Animal Breeding and Genetics Unit





Brad Walmsley



- B. Rural Science
 - UNE

• Phd

- Responsibilities:
 - BreedObject \$Indexes
 - Southern Multi-breed
 - Other things



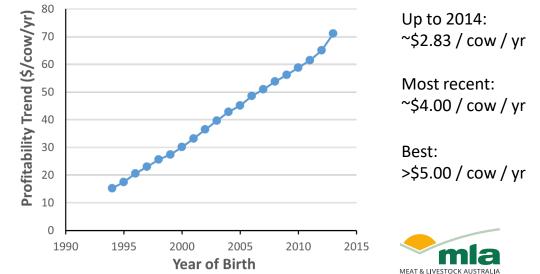


Commercial Profit

Profit = Income – Costs

As driven by genetics

Value of Genetic Improvement - South





Improving within-breed genetic evaluation and developing multi-breed genetic evaluation with the Southern Multi-Breed Project

Improving genetic evaluation





What drives genetic progress?

 $Response = \frac{selection\ intensity \times selection\ accuracy}{generation\ interval}$ Variation

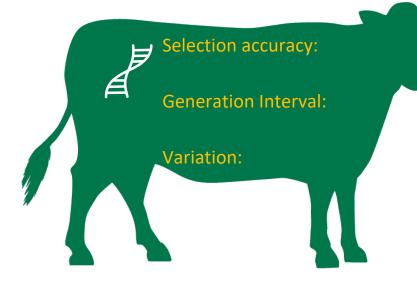
- pick only the best intensity
- make the right choice more often accuracy
- breed from them ASAP generation interval
- identify differences between animals variation



How fast you make genetic progress is dependent on how you balance these factors.



How does genomics help?



me

More information coming from "relatives"

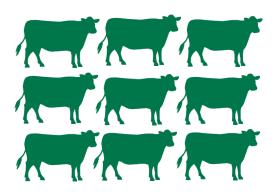
Identifying earlier who carries good genes

Traits that we can't measure any other way

Genomics can be used to drive faster rates of genetic gain.

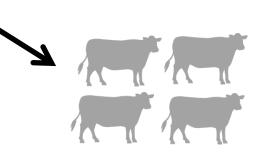


Genomics – basic principle



Reference population:

- measuring phenotypes and genotypes
- hard to measure traits
- late in life traits.



Industry animals:

- DNA tests on young animals
- predict breeding values based on genomic
- relationship and traits measured in reference.



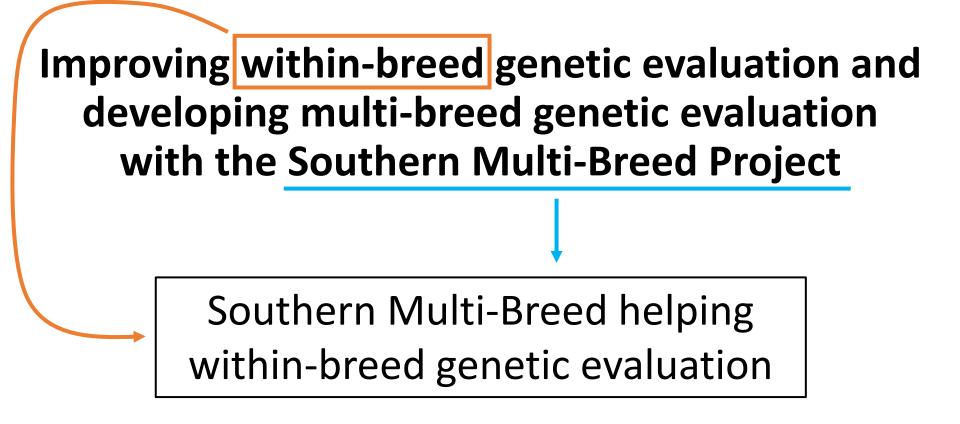
BREEDPLAN Developments

- BREEDPLAN includes genomics (single-step)
 - Brahman (2017)
 - Hereford (2017)
 - Angus (2017)
 - Wagyu (2018)
 - Santa Gertrudis (2021)
 - Droughtmaster (soon)













Research Stations

Site Diversity













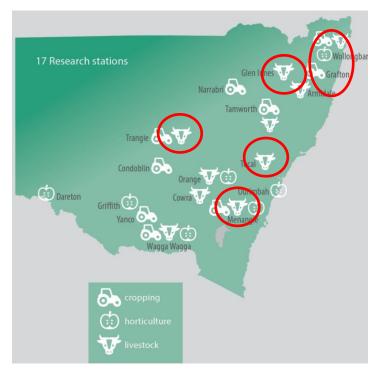


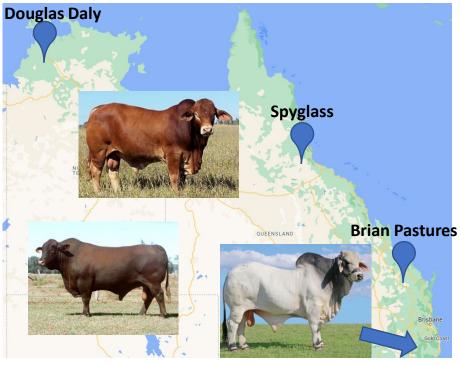




Southern Multibreed

Repronomics







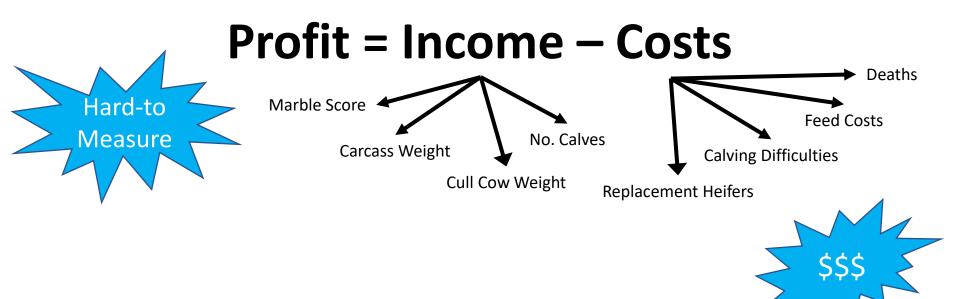








Commercial Profit







Current BREEDPLAN Traits

| Growth | Repro | Carcase | Others |
|-------------|------------------|-----------------|-----------------|
| Birth Wt | Gestation Length | Scan (live) | Feed Efficiency |
| Weaning Wt | Calving Ease | Carcase Wt | Temperament |
| Yearling Wt | Days to Calving | Marbling | Structure |
| Sale Wt | | Rump/Rib Fat | |
| Mature Wt | | Eye Muscle | |
| | | Tenderness (SF) | |





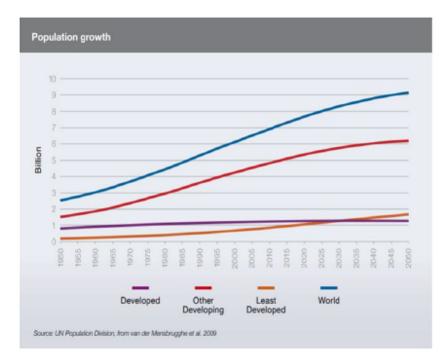


University of New England

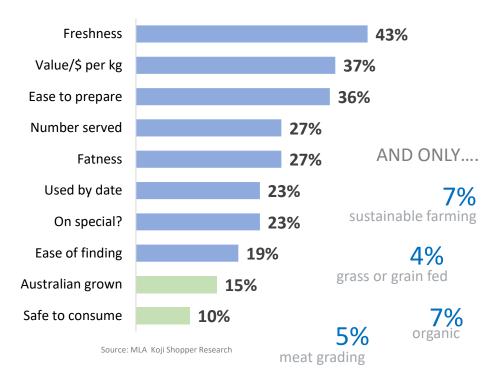




Growing Population



Purchasing decisions



An extra 1 BILLION people to feed every 15 years

MEAT & LIVESTOCK AUSTRALIA

https://news.un.org/en/story/2017/06/560022-world-population-hit-98-billion-2050-despite-nearly-universal-lower-fertility; UN Dept of Economic & Social Affairs, 2017

REDUCERS: Consumers who are reducing RM consumption not as big as 'noise' suggests. Price and health driving reduction.

- Less than **1** in **3** consumers are **reducing** their red meat consumption
- Proportion of reducers has **remained stable** for over a decade
- Price, health perceptions, environmental and animal welfare concerns are driving reduction

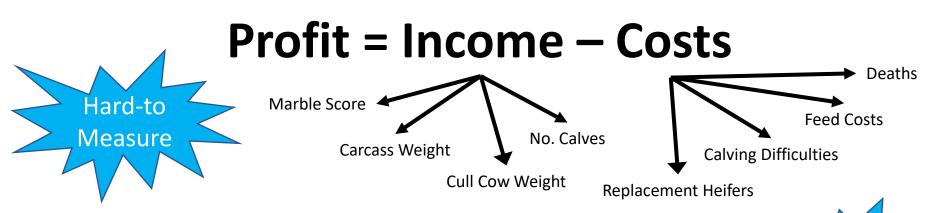


So what? Address concerns of reducers to help them feel good about eating red meat.





Commercial Profit



- What about future Profit?
 - Eating experience Welfare
 - Health benefits Health
- Horns
- Methane

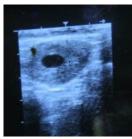




New Traits



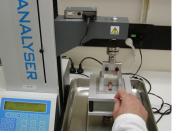
Age at Puberty 1st Calf Re-breed





Cow Composition

Meat Quality



Source: E. Toohey

Consumer Testing



Source: P. McGilchrist



Horn/Poll

Immune Competence





Methane













Improving within-breed genetic evaluation and developing multi-breed genetic evaluation with the Southern Multi-Breed Project Southern Multi-Breed helping multi-breed genetic development





Designed Research Program

• Breeds in Southern Australia with highest BREEDPLAN registrations + Brahman



Charolais



Hereford





Brahman

Shorthorn





Wagyu













Managed Head-to-Head















Designed Research Program

- Purebred matings = Purebred calves[#]
- Designed mating
 - Avoid inbreeding
- Produce comparable progeny
 - All in All out (No cull, no draft)











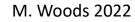




Grafton Matings[#]

| | | Cow breed | | | | | | | |
|---------------|----|--------------|--------------|--------------|--|--|--|--|--|
| | | AA | BB | НН | | | | | |
| Bull Breed | AA | \checkmark | \checkmark | | | | | | |
| | BB | \checkmark | \checkmark | \checkmark | | | | | |
| | HH | | \checkmark | \checkmark | | | | | |





S. Mortimer 2022











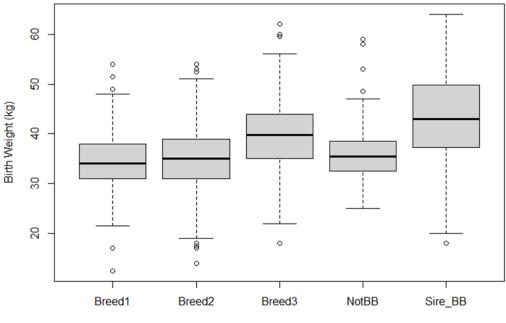
Improving within-breed genetic evaluation and developing multi-breed genetic evaluation with the Southern Multi-Breed Project

Key Learnings





Research Learnings - SMB



Breed



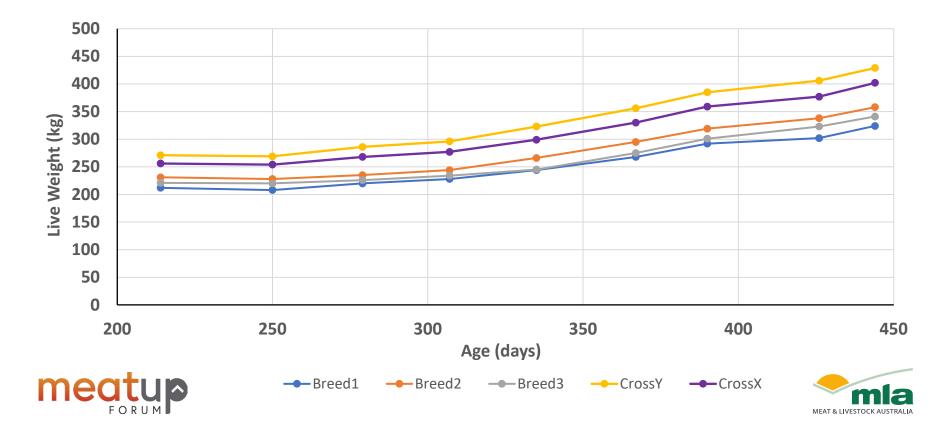
- Crossing breeding
 - Similar to Grafton 70s & 80s

• Important ramifications





Research Learnings - SMB



Research Learnings - Repronomics



Daughter Fertility?



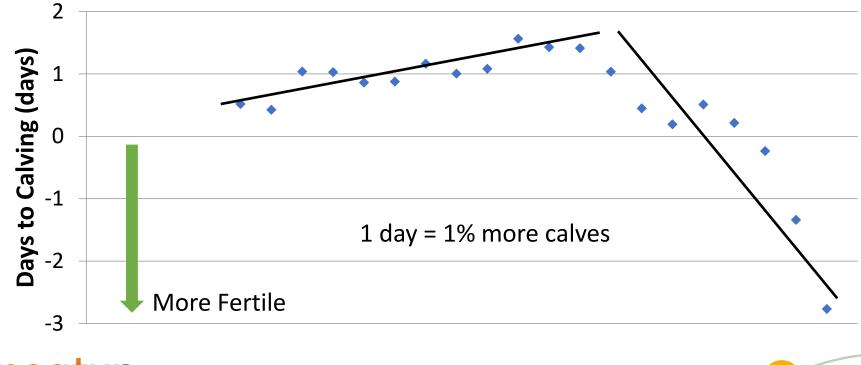
Puberty \rightarrow 8.9 months Recycle \rightarrow 4.4 months

20 day difference

Repronomics^{MT} -Johnston 2021



Research Impact - Brahman



Beef CRC/Repronomics



Research Impact - Brahman



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| | November 2022 Brahman BREEDPLAN | | | | | | | | | | | | | | | | |
|-----------|---------------------------------|------|------|------|------|------|---------|---------|---------|---------|------|------|--------|-----|---------|--------|-------|
| | | 200 | 400 | 600 | Mat | | | Days | | Eye | | | Retail | | Percent | | |
| Gestation | Birth | Day | Day | Day | Cow | | Scrotal | to | Carcase | Muscle | Rib | Rump | Beef | | Normal | Flight | Shear |
| Length | Wt. | Wt | Wt | Wt | Wt | Milk | Size | Calving | Wt | Area | Fat | Fat | Yield | IMF | Sperm | Time | Force |
| (days) | (kg) | (kg) | (kg) | (kg) | (kg) | (kg) | (cm) | (days) | (kg) | (sq.cm) | (mm) | (mm) | (%) | (%) | (%) | (secs) | (kgs) |
| +0.1 | +3.4 | +22 | +29 | +39 | +46 | +1 | +1.7 | -4.9 | +22 | +2.6 | -0.7 | -1.4 | - | 0.0 | - | -0.11 | +0.11 |
| 25% | 49% | 53% | 54% | 56% | 53% | 36% | 41% | 34% | 45% | 36% | 38% | 49% | - | 29% | - | 42% | 38% |

Traits Analysed: Genomics



| | November 2022 Brahman BREEDPLAN | | | | | | | | | | | | | | | | |
|-----------|---------------------------------|------|------|------|------|------|---------|---------|---------|---------|------|------|--------|------|---------|--------|-------|
| | | 200 | 400 | 600 | Mat | | | Days | | Eye | | | Retail | | Percent | | |
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| Length | Wt. | Wt | Wt | Wt | Wt | Milk | Size | Calving | Wt | Area | Fat | Fat | Yield | IMF | Sperm | Time | Force |
| (days) | (kg) | (kg) | (kg) | (kg) | (kg) | (kg) | (cm) | (days) | (kg) | (sq.cm) | (mm) | (mm) | (%) | (%) | (%) | (secs) | (kgs) |
| 0.0 | +4.3 | +21 | +27 | +38 | +53 | 0 | -0.3 | +8.3 | +26 | +3.1 | -1.2 | -1.4 | +0.9 | -0.4 | - | +0.03 | -0.08 |
| 44% | 55% | 58% | 59% | 60% | 56% | 36% | 45% | 34% | 49% | 35% | 37% | 47% | 25% | 30% | - | 43% | 38% |

Traits Analysed: Genomics



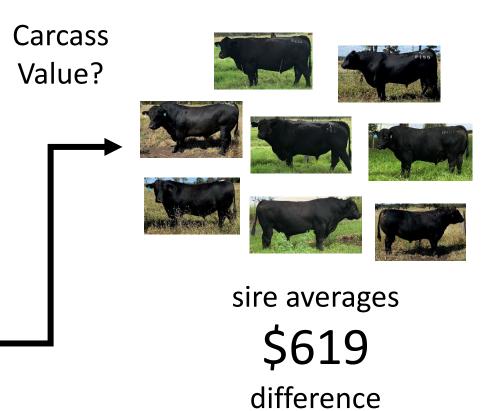
ABBA BREEDPLAN Database – November 2022

Others Findings



worst

\$2076 difference best



MEAT & LIVESTOCK AUSTRALIA

Beef Central & Angus Australia 2018

Acknowledgements

- David Johnston (AGBU Repronomics)
- Leadership Team: Kath Donoghue, Jason Siddell & Sam Clark
- Other Scientists (DPI, UNE, AGBU & CSIRO)
- Management and staff at Trangie, Grafton, Tocal, Glen Innes, EMAI, North Coast and Tullimba
- All technical staff (DPI, UNE & CSIRO)
- Project partners AI, DNA, Merchandise, Breeders, Breed Societies, Producers







Take home messages

- Work needed to capture benefits of genomics
- Investment in Southern Multibreed and Repronomics[™]
- Southern Multibreed benefits to emerge in the future
- Repronomics[™] benefits can be seen in:
 - Brahman
 - Santa
 - Droughtmaster







Tools and resources

• BREEDPLAN

- Tropical breeds already benefiting

BreedObject \$Indexes

<u>https://www.dpi.nsw.gov.au/animals-and-livestock/beef-</u> <u>cattle/breeding/southern-multi-breed-smb-project/project-</u> <u>overview</u>

• Google – Southern Multibreed



