

# Historical and future developments in the New Zealand genetic evaluation system for dairy cattle

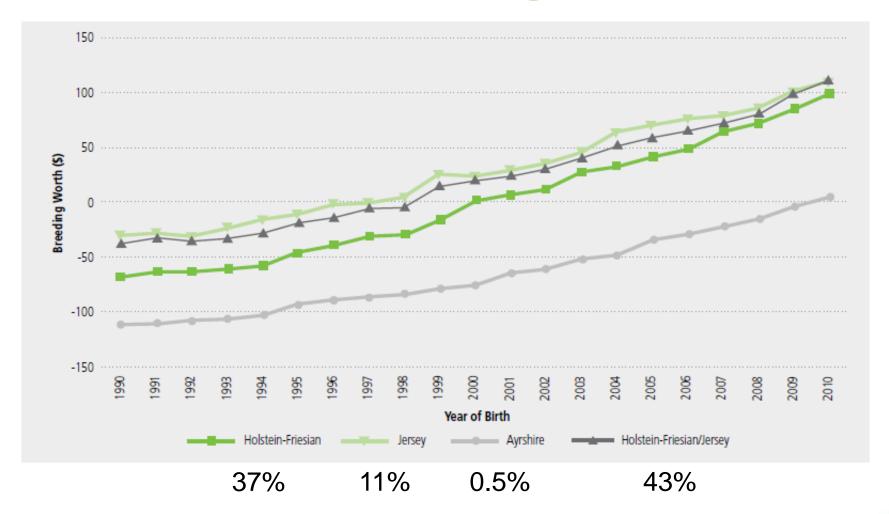
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## **Key points**

- Strong recording and AE history
- New database and shift of animal evaluation to NZAEL
- Traits and research of importance

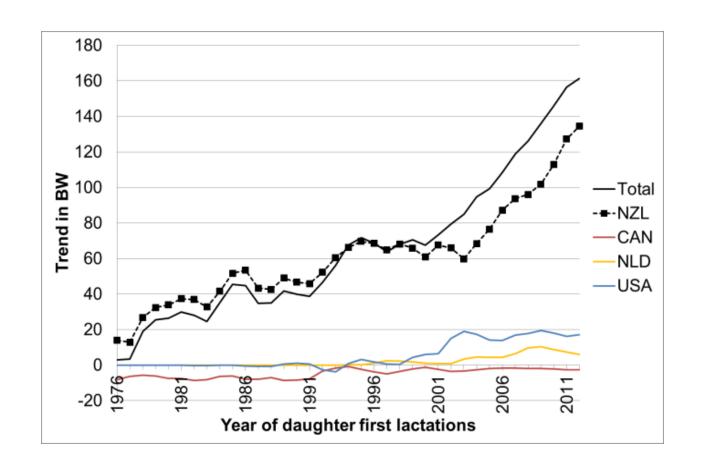


## **Trend - Cow Breeding Worth**





# Source of genetic trend - HF





## Why the success

- Strong history recording, herd testing, breeding infrastructure
- History of excellent quantitative geneticists
- Across breed, TDM, fertility BV, genomics
- Index that is economically relevant
- Investment industry good and company

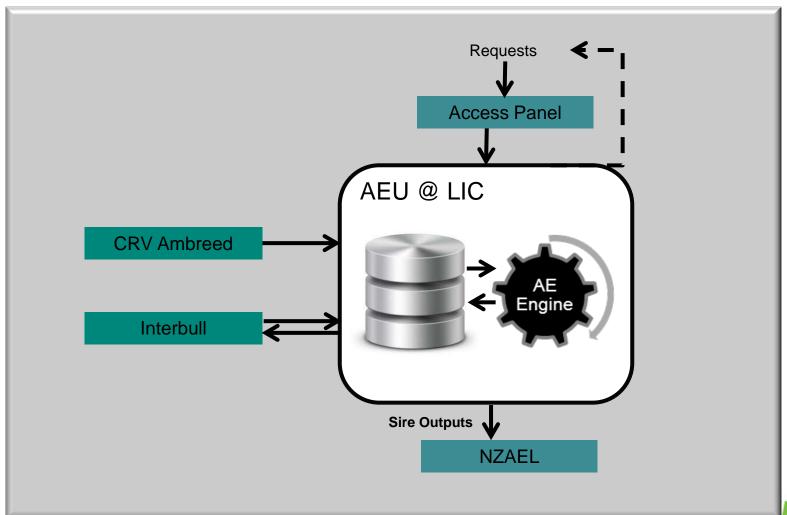






#### **Current State**

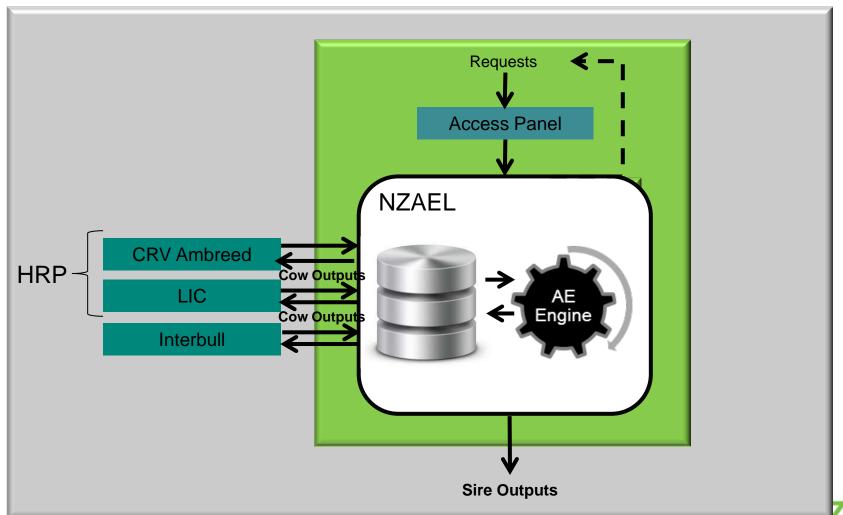






#### **Future State**





## **Principles - Transition**

- 1. Utilise national and international experts, organisations and systems
- 2. Highly modular
- 3. Utilise business intelligence tools
- Build deeper teams genetics and IT



# Parentage recording



- Parentage misidentification rates are high
- Likely deflates genetic merit of elite bulls
- Theory: Within herd heritability estimates provide an indication of parentage identification quality and phenotype quality
- Herd heritability vs. sire misidentification
  - r = -0.38 to -0.50 (Dechow et al. 2008)

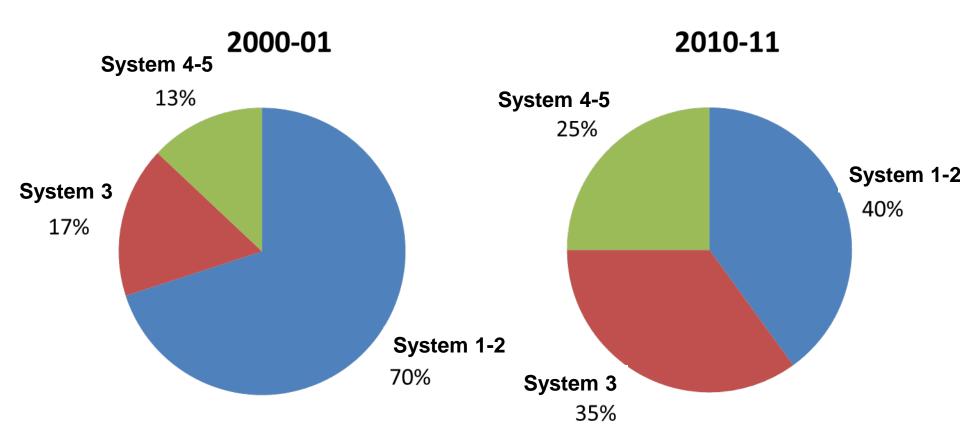


### **Body condition score**

- BCS of 5 at calving essential for reproduction
- Animal welfare and public perception
- Economic value (\$102/BCS)
  - Utilise cheaper feed in spring rather than winter for replenishment
  - Milk on longer
- Differences in BCS between breeds



# Farm system shifts



System 1-2: < 0.5 t DM/cow imported feed

System 3: 0.5 to 1.0 t DM/cow imported feed

System 4-5: >1.0 to 3.0 t DM/cow imported feed



#### Other initiatives

- Stronger links with universities, training institutes
- High quality and deeper phenotypic data
  - fertility BV enhancement, robotic and fixed in line herd testing, environmental traits
- Residual feed intake (RFI)
- Addressing genomic pre-selection



#### Conclusions

- Build on our strengths
- DIGAD and animal evaluation at NZAEL
- Addressing data quality and exploring new data sources
- System divergence
- Potential new traits for Breeding Worth
  - BCS, RFI

