

Methane Index Explorer: Optimising a Breeding Value Format for Simultaneous Inclusion of Enteric Methane Emissions in Breeding Schemes and National Inventories



Presented to AAABG June 2025 and
paper available soon from website

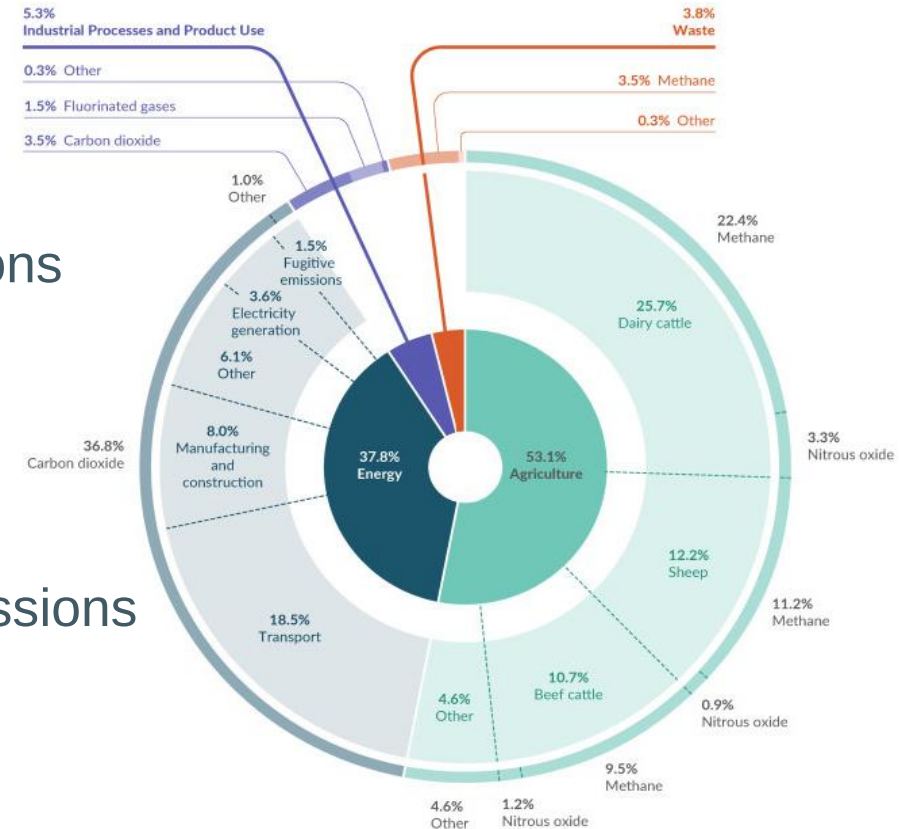
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Introduction

- Ruminants are a major source of NZ methane (CH₄) emissions
 - Agriculture (enteric fermentation from livestock) – 53%
 - Fossil fuel extraction and use – 38%
 - Landfills and industrial processes – 9%
- Methane is heritable (0.19 - 0.32) – selection can reduce emissions
- Antagonistic relationships (paper in preparation)
 - CH₄ – liveweight (0.61), feed intake (0.33)
 - Liveweight – weaning weight (0.91), carcass weight (0.85)
- Research problem: Would selection for CH₄ affect progress in the production index?



Research Objectives

- Develop a selection response evaluation format for reducing CH₄ using portable accumulation chamber breeding values (BVs) – (PACCH4BV)
- Aiming for:
 - Ram selection in breeding programs
 - Governmental inventory reporting
 - Methane Index Explorer (MIE) software integration
 - Allows breeder to explore consequences
 - With his animals
 - With his breeding scheme/recording/genotyping



Methods: Data Collection

- Animals: AgResearch Woodland Genetic Research Flock 2638 (birth year 2021 & 2022)
- BVs and genomic BVs (gBV) from the SIL database
- Animal evaluation (AE) 6 used more genotypes than AE5

	2021		2022	
Rams	744		605	
PAC recording	384 (51.6%)		324 (53.6%)	
Animal evaluation	AE5	AE6	AE5	AE6
Number genotypes used	1 (0.1%)	739 (99.3%)	304 (50.2%)	603 (99.7%)



Methods: Index Construction (MIE software integration)

- NZMW-plus Index = NZMW + FEC + DAGS + MEAT
- Dual-Purpose CH₄ index (DPCH₄)
 - PACCH4gBVs converted to PACCH4BV% : $(\text{g/day} \div \text{scaled mean PAC emissions (7.5)}) \times 100$
 - DPCH4 : $\text{PACCH4BV\%} \times -0.51 = \$100 \text{ tonne CO}_2\text{e GWP100}$
- PACCH4 assessed across 9 economic weights: \$0–\$200/t (GWP100)
 - 9 DPCH4 indices were calculated
- Economic weights of other traits are as per SIL Technical notes (2022)

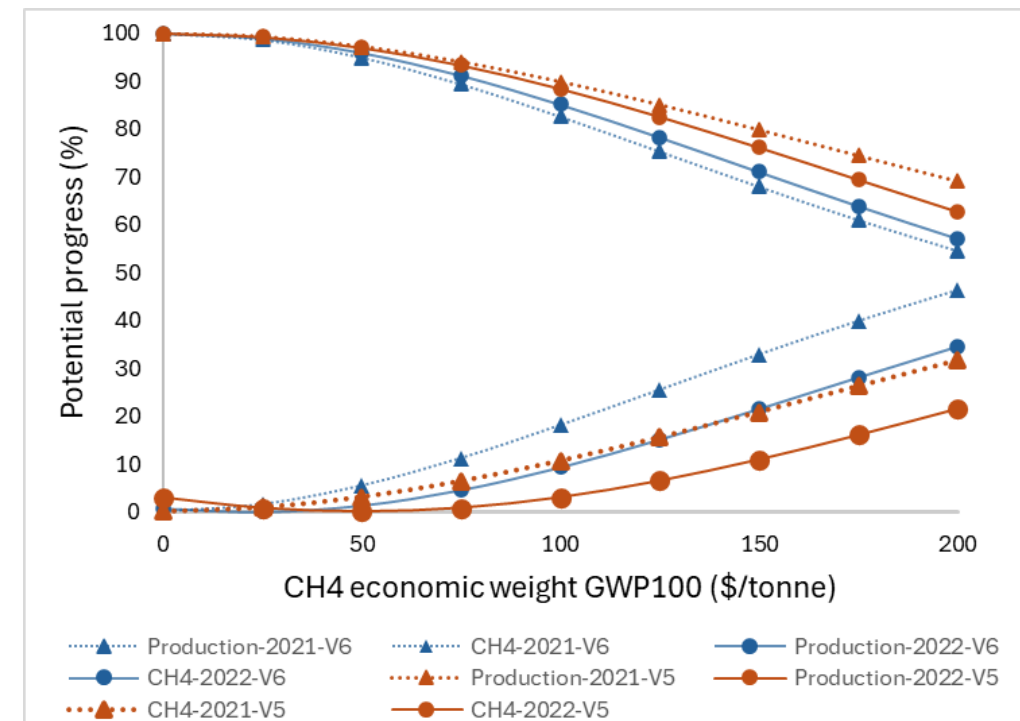
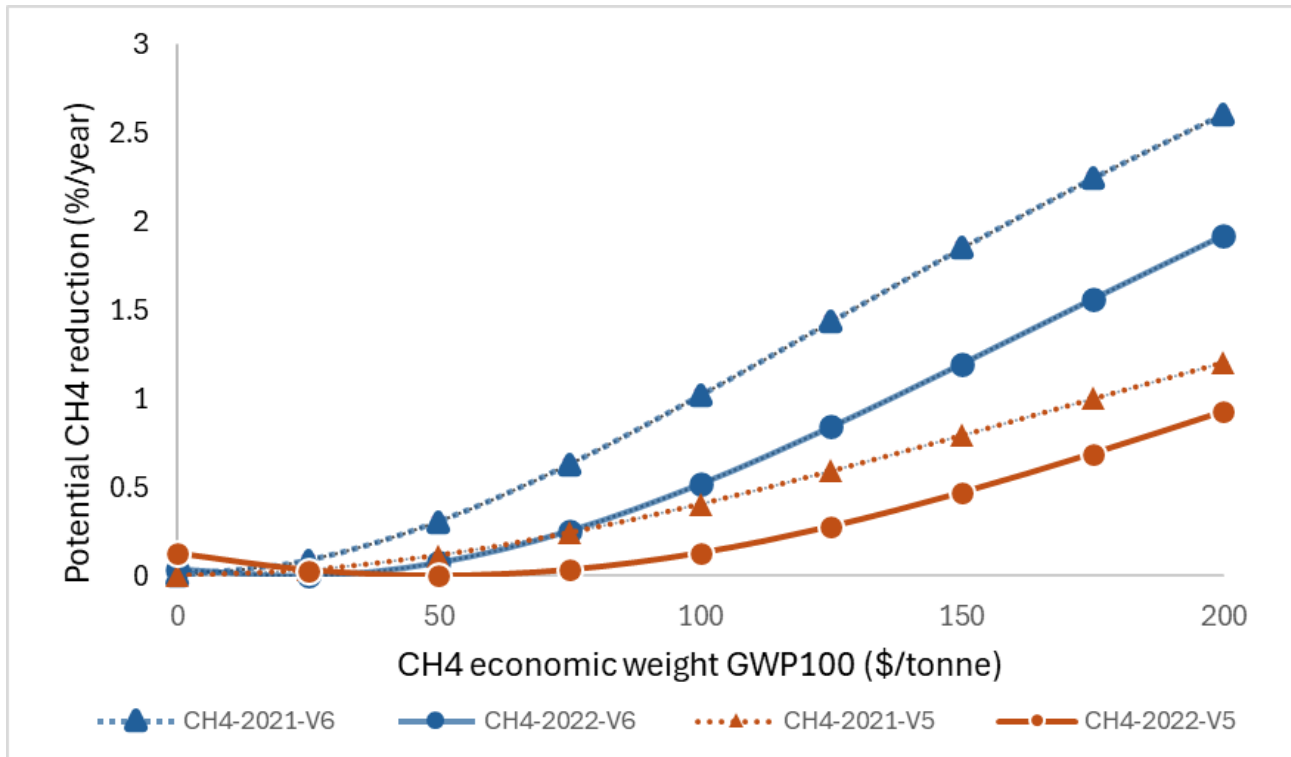
Methods: Evaluating Genetic Progress (MIE software integration)

- Progress: r^2 between indices for each economic weight
 - Production index progress: $r^2(\text{NZMW-plus Index} \ \& \ \text{NZMW-plus+DPCH}_4 \text{ index}) \times 100$
 - CH₄ index progress: $r^2(\text{NZMW-plus+DPCH}_4 \text{ index} \ \& \ \text{DPCH}_4 \text{ index}) \times 100$
 - but presented here as a % of potential maximum
- PACCH₄BV% reduction/year = CH₄ index progress $\times$ SD(PACCH₄BV%) $\times i/L$
 - i = selection intensity (2% rams, 30% ewes), L = generation interval (3.5 years)

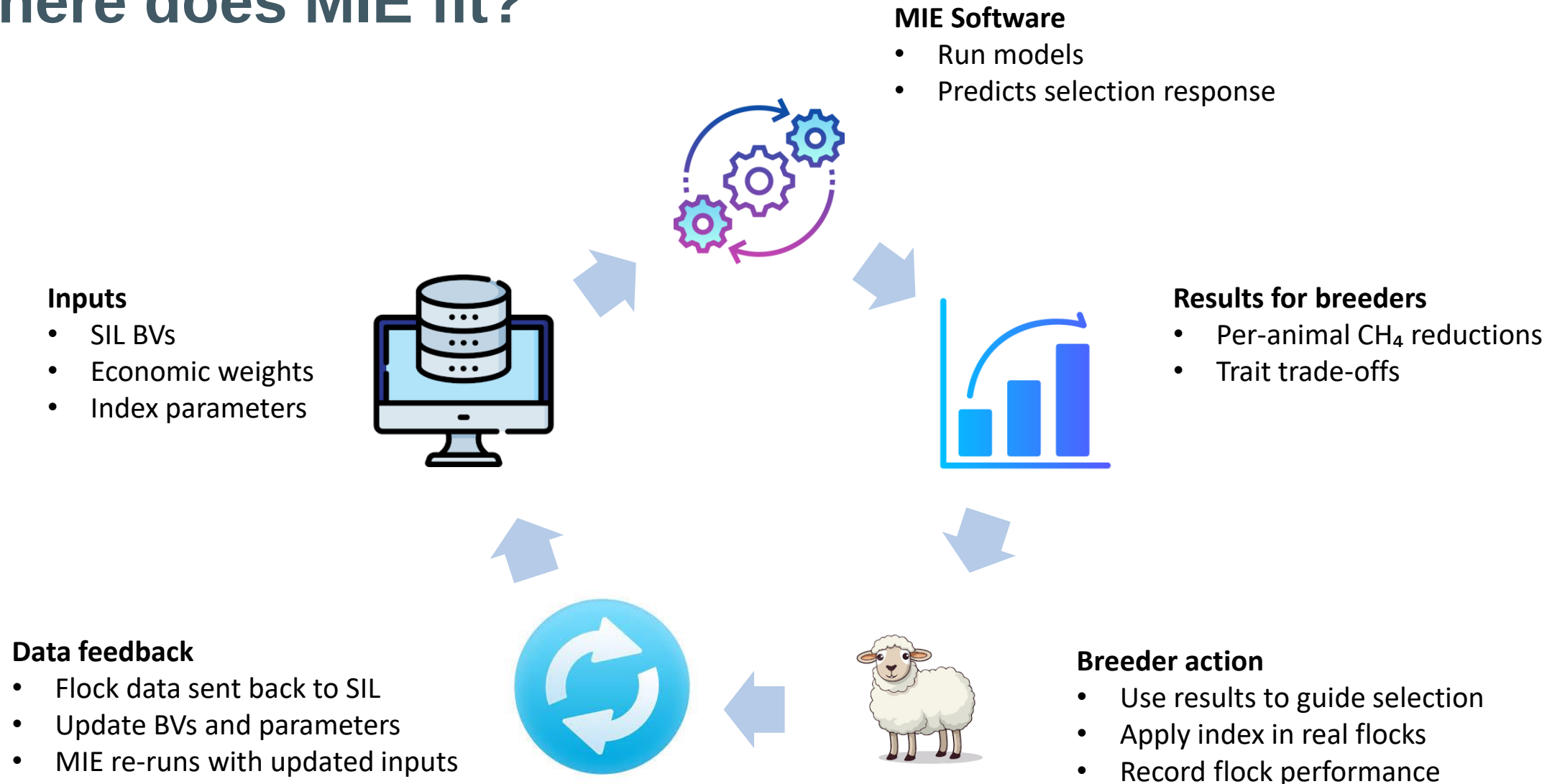
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Key Results

- AE6 (high genotyping) outperformed AE5
- Greater CH₄ reductions with \$50–\$100/t weighting
- Up to 2.6%/yr CH₄ BV reduction in AE6 @ \$200/t
- Low weight on PACCH₄ did not increase emissions
- Increased relative potential progress as well



Where does MIE fit?



Conclusions & Implications

- Breeding reduces CH₄ emissions per animal, while increasing per head productivity
- “sweet spot” ~\$50-\$100/tonne
 - CO₂e GWP100 >80% production gains
 - while reducing CH₄ by ~0.5%/year
- Genomic data is essential for accurate BVs
- PACCH4BV% is interpretable and effective
- Recommended for breeders and inventories



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