SUSTAINABILITY STARTS WITH SEMEX

INTRODUCING A GENETIC SOLUTION TO REDUCE METHANE EMISSIONS 20-30% BY 2050



ELEVATE® IS THE WAY TO BREED LOW METHANE COWS

At Semex, health, sustainability and the global 'Net Zero' discussion are high priority as we are all asked to reduce emissions by our global consumers and local governments.

Ten years ago, we started this conversation by changing the traditional trait discussions when we introduced Immunity+®. This technology genetically targeted breeding cows needing less antibiotics, that are healthier and last longer in herds. Now, we're excited to introduce a Methane Efficiency (ME) trait.





Methane Efficiency Available on Semex Holstein Bulls

- New with April 2023 proofs
- 70-80% Reliable
- Permanent & cumulative
- Independent from other traits
- Methane index available on all Holstein females tested through Elevate
- Depending on selection intensity, herds can reduce emissions 20-30% by 2050

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YOUR QUESTIONS ANSWERED

1. WHAT IS METHANE?

Methane (CH4) is a colorless, odorless gas composed of carbon and hydrogen that enters the atmosphere from both human related activities and natural sources. It is also a potent greenhouse gas contributing to climate change.

2. WHY IS METHANE IMPORTANT TO AGRICULTURE AND DAIRYING?

Methane's global warming potential is an estimated 25-35 times greater than carbon dioxide (CO_2) over a 100-year period, making methane emissions a climate change concern. Forty percent of global methane emissions are from agriculture. Dairy cows, as ruminants, produce methane from gut fermentation.

3. WHY IS METHANE IMPORTANT TO SEMEX?

Semex, our customers and all of production agriculture are under the same climate scrutiny as all companies or organizations. As the global dairy cow is a significant source of methane emissions, Semex and every dairy producer in the world should be very interested in reducing methane.

4. HOW DO COWS PRODUCE METHANE?

Approximately 90% of methane from cows is excreted by eructation (burping) rumen gases from their mouths. This is a rumen fermentation by-product and is often referred to as enteric methane emissions.

5. WHAT ROLE DOES THE ANIMAL ITSELF PLAY IN METHANE PRODUCTION?

Multiple studies show that the cow plays a driving role when it comes to methane production. In 2020, Zhang et al. discovered that host genetics and rumen microbiome jointly explained 31% of methane emission variation. Twenty-four percent of the variance was explained by host (cow) genetics, while the balance (7%) was explained by the rumen microbiome (rumen microbes). They also suggested that the host's genetics impact ruminal methane emissions via mechanisms other than composition of the rumen microbiome.

6. HOW DID WE MEASURE METHANE PRODUCTION?

Measuring methane is expensive and difficult. Collecting methane emission data in research herds was made possible through two largescale international projects led by Canadian researchers receiving funding from Genome Canada, and provincial genome centres and ministries. Methane emission was measured on +500 individual cows using the GreenFeed System.

7. WHY IS CANADA IN A UNIQUE POSITION TO DELIVER METHANE EVALUATION?

Canada has been recording milk Mid-Infrared (MIR) data from individual samples since 2013. The Canadian dataset includes more than 13M records from 1.6M cows. MIR spectroscopy relies on light absorption. The analysis produces a spectrum that passes infrared radiation through a sample, and measures the absorptions made by each type of milk compound bond. This can be used to predict different traits from milk samples including components, SCS and more recently methane production.

8. CAN WE USE MIR PREDICTED METHANE TO SELECT FOR METHANE REDUCTION?

Yes! The genetic correlation between MIR predicted methane and methane measured with the GreenFeed System is 0.85, demonstrating that the genomic value methane predictions using MIR data is a very efficient and costeffective way to select for methane reduction.

9. HOW WILL THE EVALUATION BE EXPRESSED?

The official evaluation will be expressed as "Methane Efficiency", i.e., production of methane at the same level of milk, fat and protein yield. In this way, the evaluation is independent of production and will not penalize productive, efficient animals. This importantly ensures the world will not need more cows to produce enough food, negating any individual cow methane reductions.

10. HOW MUCH OF AN IMPACT WILL THIS HAVE?

Dairy producers can expect to achieve 20-30% reduction in methane emissions from their herd by 2050 depending on their selection pressure.

11. WHAT ARE THE CORRELATIONS WITH OTHER TRAITS?

Methane efficiency is designed to be genetically independent from milk, fat and protein yield. The genetic correlations between methane efficiency and other economically important traits is low. Functional traits (i.e., daughter fertility, metabolic disease resistance) have low but favorable correlations with methane efficiency.

12. HOW IS THIS DIFFERENT FROM FEED EFFICIENCY?

The genetic correlation between Feed Efficiency and Methane Efficiency breeding values is low, confirming that these are genetically different traits. Ideally, milk producers would benefit by selecting for both traits.

13. WILL THE EVALUATION BE AVAILABLE FOR OTHER BREEDS?

Initially the evaluation will only be available for Holsteins. As Canada expands the reference population to other breeds, prediction equations will become available.

14. WILL SELECTING FOR METHANE EFFICIENCY REDUCE METHANE COMING FROM A METHANE DIGESTOR?

No. This evaluation only targets enteric methane. The methane produced from the manure is the result of anaerobic processes happening after excretion.

14. HOW DOES A PRODUCER MEASURE METHANE EMISSION IN THEIR HERD?

Producers that have already genomically tested with Elevate® or test before the April genetic evaluation, will automatically receive methane emission genomics. This allows them to benchmark and monitor their methane emissions for every animal tested with Elevate.





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