

Table 1. Estimated economic and greenhouse gas (GHG) emission benefits from adopting new genomic tools in commercial beef cattle assuming 100% adoption and five years post project.

Genomic tools	Benefit, (%/year)	Economic benefits, (\$million/year)	%GHG reduction/yr
1. Fertility haplotype test	0.5-1% increase in calf crop	\$25.8 - \$51.7	1%
2. Fertility haplotype test	0.5-1% decrease in culling	\$38.4 - \$76.9	1%
3. Female Fertility Indices	4-6% increased calf crop	\$24.8 - \$37.2	6-10%
4. Female Fertility Indices	10-15% decreased mortality	\$4.3 - \$6.5	1-2%
5. Female Fertility Indices	10-15% decreased morbidity	\$2.5 - \$3.8	<1%
6. Female Fertility Indices	1-3% feed saving-feeders	\$24.3 - \$73.0	1-3%
7. Female Fertility Indices	1-3% feed saving-cows	\$23.3 - \$70.1	1-3%
8. gEPD RFI, >0.45 acc.	0.5-1% feed saving	\$23.9 - \$47.8	1%
9. Female fertility indices	10-16% decrease in GHG	\$60.8 - \$97.3	10-16%
10. Feeding NOP	30-40% reduction in GHGs	-----	30-40%
Total gross benefit/year		\$228 - \$464	40-56%

Benefits at a national level

1. Base=4.27 million cows [includes bred heifers] x 0.84 calf crop x \$1210/calf weaned=\$4,340,028,000 gross income. Change would be a 0.5 to 1 percentile increase in calf crop or ((4.27 million x 0.845 x \$1210) – Base) = **\$25,833,500/yr.**
2. Base=4.27 million cows x 0.16 replacement rate x \$1800 to develop replacement=\$1,229,760,000 cost. Change would be a 0.5 to 1 percentile decrease in replacement rate or ((4.27 million x 0.155 x \$1800) – Base) = **\$38,430,000/yr.**
3. Base=4.27 million cows x 0.40 x 0.30 x 0.84 x \$1210/calf = \$520,803,360. Assumptions include 40% of cows could benefit from an increase in retained heterozygosity, a 30% replacement rate, and a base calf crop of 84%. Difference between a high and low hybrid vigor herd (30% different in retained heterozygosity) is 2-3% improvement in calf crop and 639 kg CO₂e/animal slaughtered (\$30/t carbon) or ((4.27 million cows x 0.40 x 0.30 x 0.88 x \$1210/calf) – Base) = **\$24,800,160/yr.**
4. Base=2.47 million feeder cattle x 0.01 x \$1750/feeder=\$43,225,000 lost due to death loss in feedlots across Canada. Change would be a 10-15% decrease in mortality or ((2.47 million x 0.009 x \$1750) – Base) = **\$4,322,500/yr.**
5. Base=2.47 million feeder cattle x 0.15 x \$70/treatment=\$25,935,000 in treatment costs in feedlots across Canada. Change would be a 10-15% decrease in morbidity or ((2.47 million x 0.135 x \$70/treatment) – Base) = **\$2,593,500/yr.**
6. Base=2.47 million feeders x 9 kg DM/day x 365 days x \$0.30/kg DM = \$2,434,185,000 in feed cost in feedlots across Canada. Change would be a 1-3% improvement in feed efficiency from increased retained heterozygosity in feeders or ((2.47 million x 8.91 kg DM/day x 365 days x \$0.30/kg) – Base) = **\$24,341,850/yr.**
7. Base=4.27 million cows x 15 kg DM/day x 365 days x \$0.10/kg DM = \$2,337,825,000 in feed costs in cows across Canada. Change would be a 1-3% improvement in feed efficiency from increase retained heterozygosity in cows or ((4.27 million x 14.85 kg DM/day x 365 days x \$0.10/kg) – Base) = **\$23,378,250/yr.**
8. Base = (4.27 million cows x 15 kg DM/day x 365 days x \$0.10/kg DM) + (2.47 million feeders x 9 kg DM/day x 365 days x \$0.30/kg DM) = \$4,772,010,000. Change would be a 0.5-1% improvement in feed efficiency from selection for ((4.27 million cows x 14.925 kg DM/day x 365 days x \$0.10/kg DM) + (2.47 million feeders x 8.955 kg DM/day x 365 days x \$0.30/kg DM)) – Base) = **\$23,860,050/yr.**
9. Canada emits 732,000 kt CO₂e; Beef cattle (cattle, non-dairy) are responsible for 2.77% of the total, or 20,276 kt CO₂e. Reduced GHG of 16%/yr = 20,276,000 t CO₂e x 0.16 x \$30/t CO₂e = **\$97,324,800/yr**
Environment Canada, 2016, at:
http://unfccc.int/national_reports/annex_i_ghg_inventories/national_inventories_submissions/items/9492.php
Other assumptions: 550 lb wean calf at \$2.20/lb = \$1210/weaned calf; \$70/calf treated; each 10% increase in genomic retained heterozygosity improves RFI by 0.08 kg DM/d; 375,732t CO₂e = \$11 million/yr.

Costs at a national level:

Number of heifer calves available=(4.27 million cows and bred heifer x 0.84 calf crop x 0.5 female:male ratio) = 1,793,400

1. Year 1 genotyping

Replacement heifers=(1,793,400 female calves x 0.30 selected for replacements x \$50/genotype) = \$26,901,000

Bulls = (4.27 million cows and bred heifers/30:1 cow bull ratio) = 142,333 bulls x \$90/genotype=\$12,809,970

2. Year 2 genotyping

Replacement heifers=(1,793,400 female calves x 0.30 selected for replacements x \$50/genotype) = \$26,901,000

Bulls = 142,333 bulls x 0.25 replacement rate x \$90/genotype=\$3,202,493

3. Year 3 genotyping

Replacement heifers=(1,793,400 female calves x 0.30 selected for replacements x \$50/genotype) = \$26,901,000

Bulls = 142,333 bulls x 0.25 replacement rate x \$90/genotype=\$3,202,493

4. Year 4 genotyping

Replacement heifers=(1,793,400 female calves x 0.30 selected for replacements x \$50/genotype) = \$26,901,000

Bulls = 142,333 bulls x 0.25 replacement rate x \$90/genotype=\$3,202,493

5. Year 5 genotyping

Replacement heifers=(1,793,400 female calves x 0.30 selected for replacements x \$50/genotype) = \$26,901,000

Bulls = 142,333 bulls x 0.25 replacement rate x \$90/genotype=\$3,202,493

Total over 5 years =\$160,124,942 or \$32.0 million/year; highest costs in first year.

At a 15% adoption rate the cost would be \$24.0 million over 5 years; cost ranges from 6.9 to 14.0% of gross benefit (ROI ranges from 7:1 to 14:1). Reducing the cost of genotyping is a key factor to adoption of genomic tools.