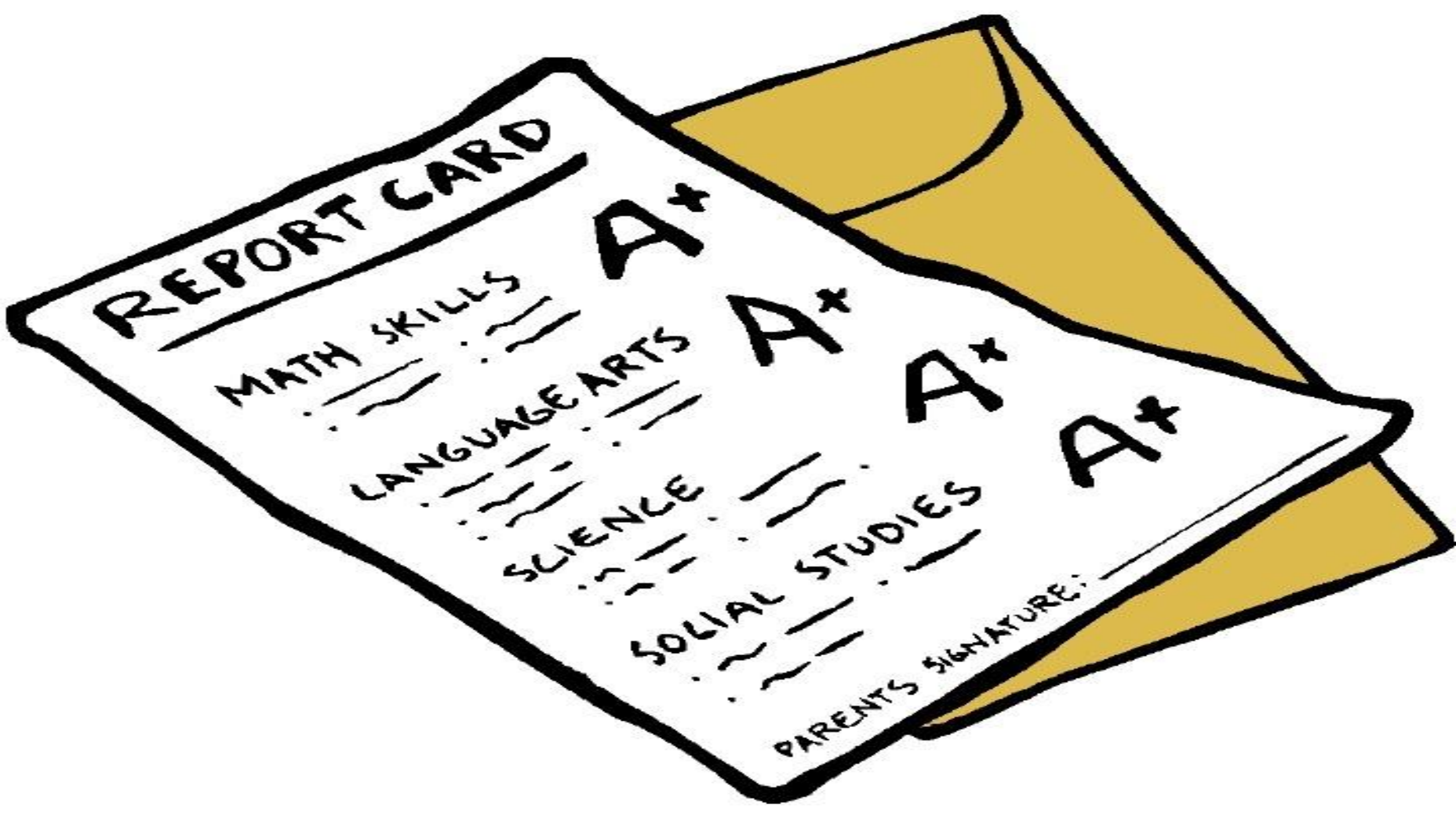




The Ten-Year Genomics Report Card

Dr. Michael Lohuis
VP, Research & Innovation, Semex

National Genetics Conference. Appleton, Wisconsin (June 26-27, 2019)



REPORT CARD

MATH SKILLS

A+

LANGUAGE ARTS

A+

SCIENCE

A+

SOCIAL STUDIES

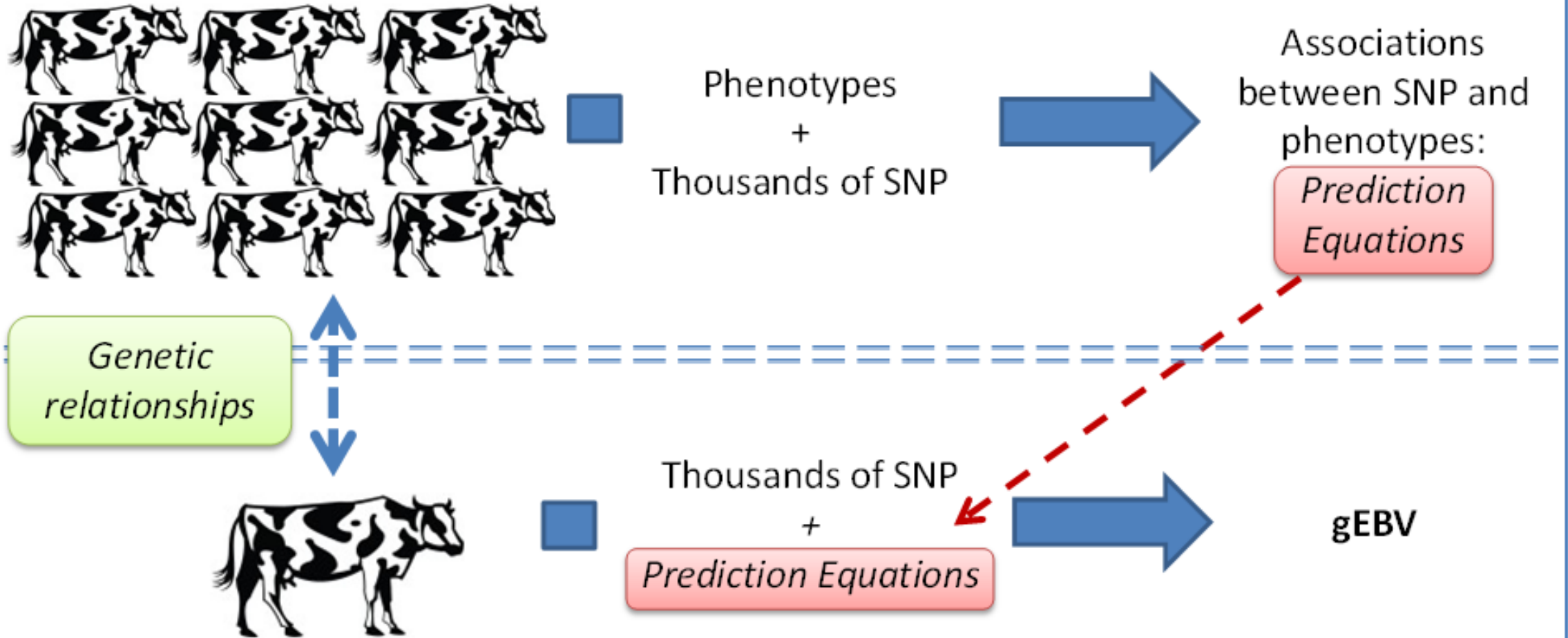
A+

PARENTS SIGNATURE: _____

Subject	Credits	Objective
Genomic Evaluation	6	<ul style="list-style-type: none">• Provide <i>relatively</i> accurate genomic rankings
Genetic Improvement	6	<ul style="list-style-type: none">• Make faster genetic improvement
On-Farm Testing	6	<ul style="list-style-type: none">• Develop genomic tests for on-farm use
Genetic Diversity	3	<ul style="list-style-type: none">• Preserve genetic variation for future use
Understanding Genotype to Phenotype	3	<ul style="list-style-type: none">• Identify genotypes that change phenotype
Meeting Consumer Expectations	6	<ul style="list-style-type: none">• Produce products that consumers value

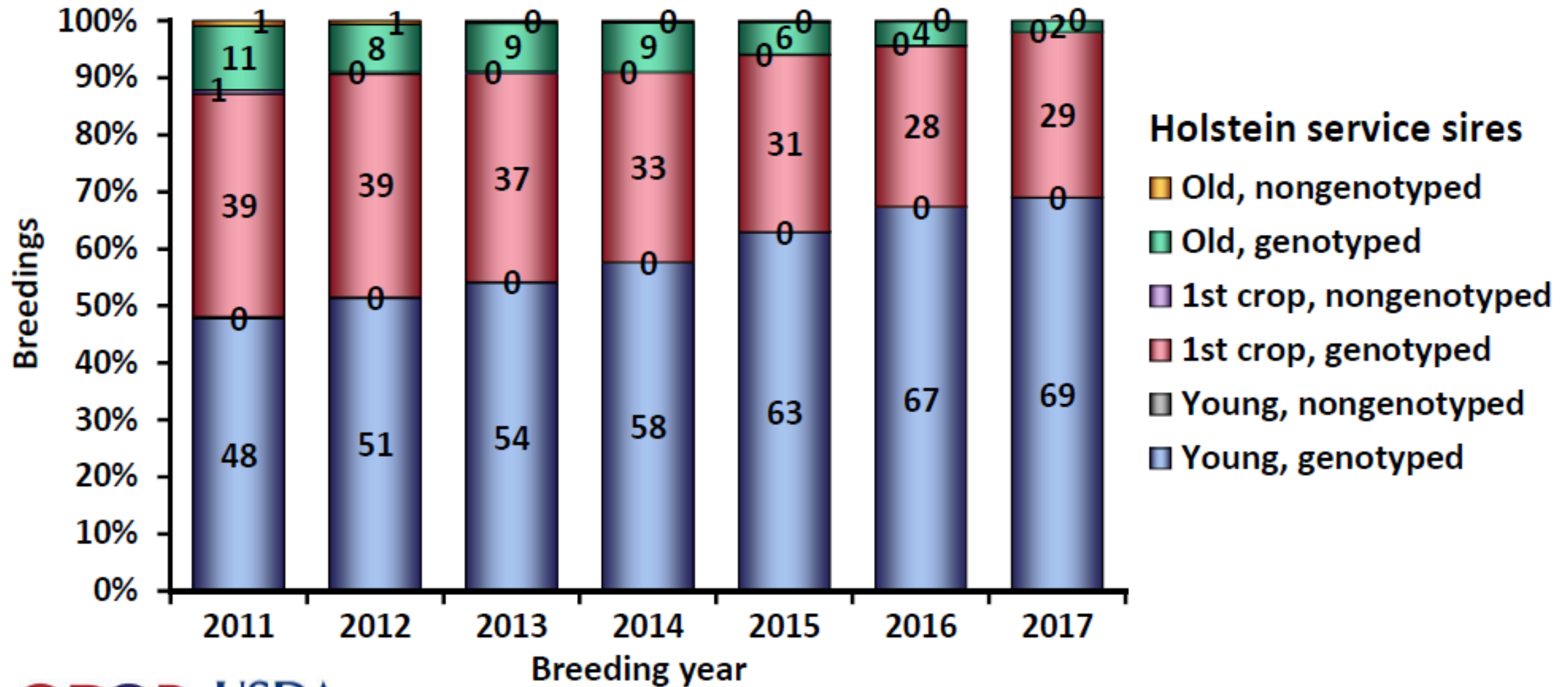
Subject	Grade	Comments
Genomic Evaluation		

Reference population: Development of prediction equations

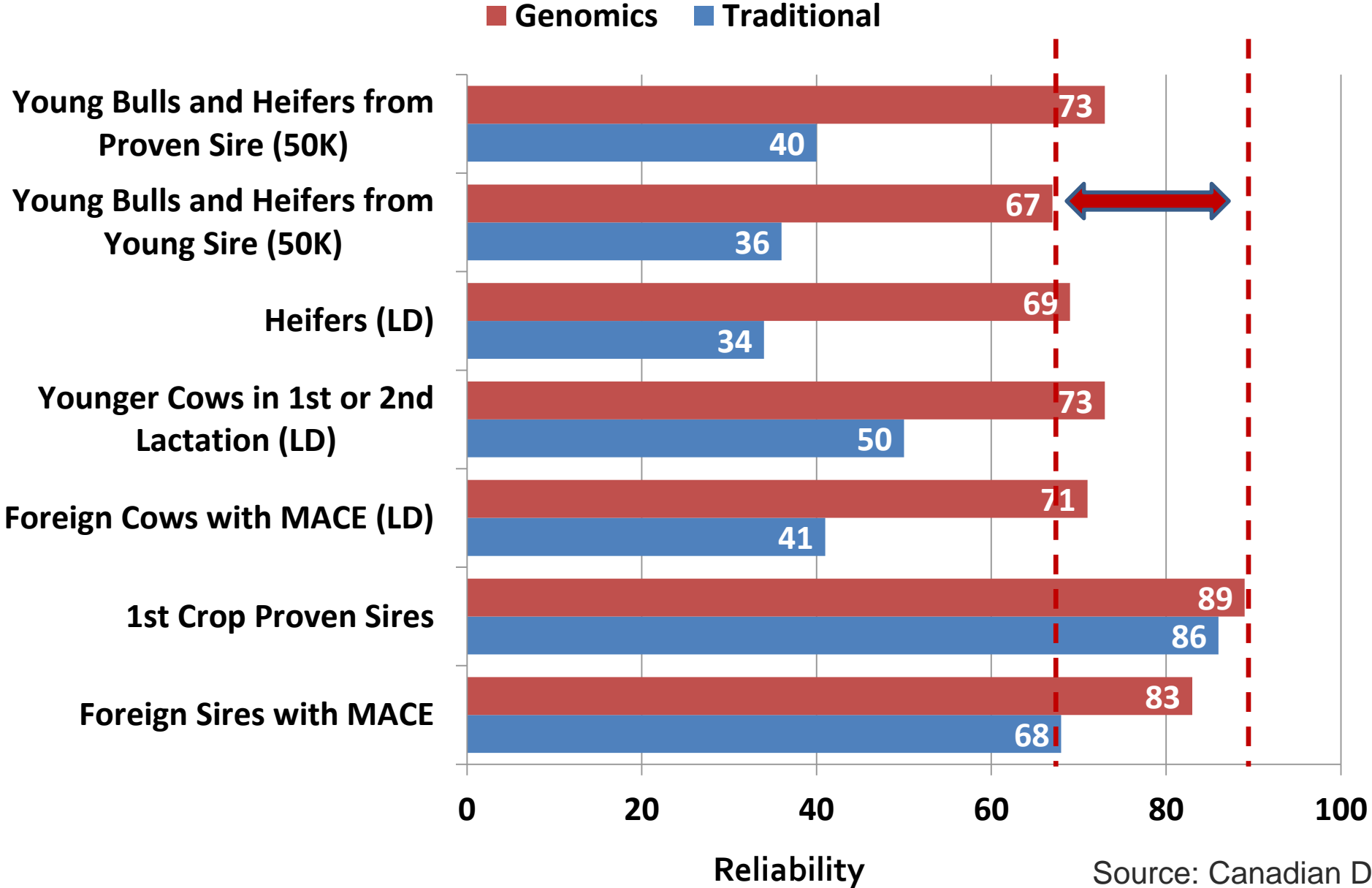


Main population: Application of prediction equations

AI breedings to genomic bulls

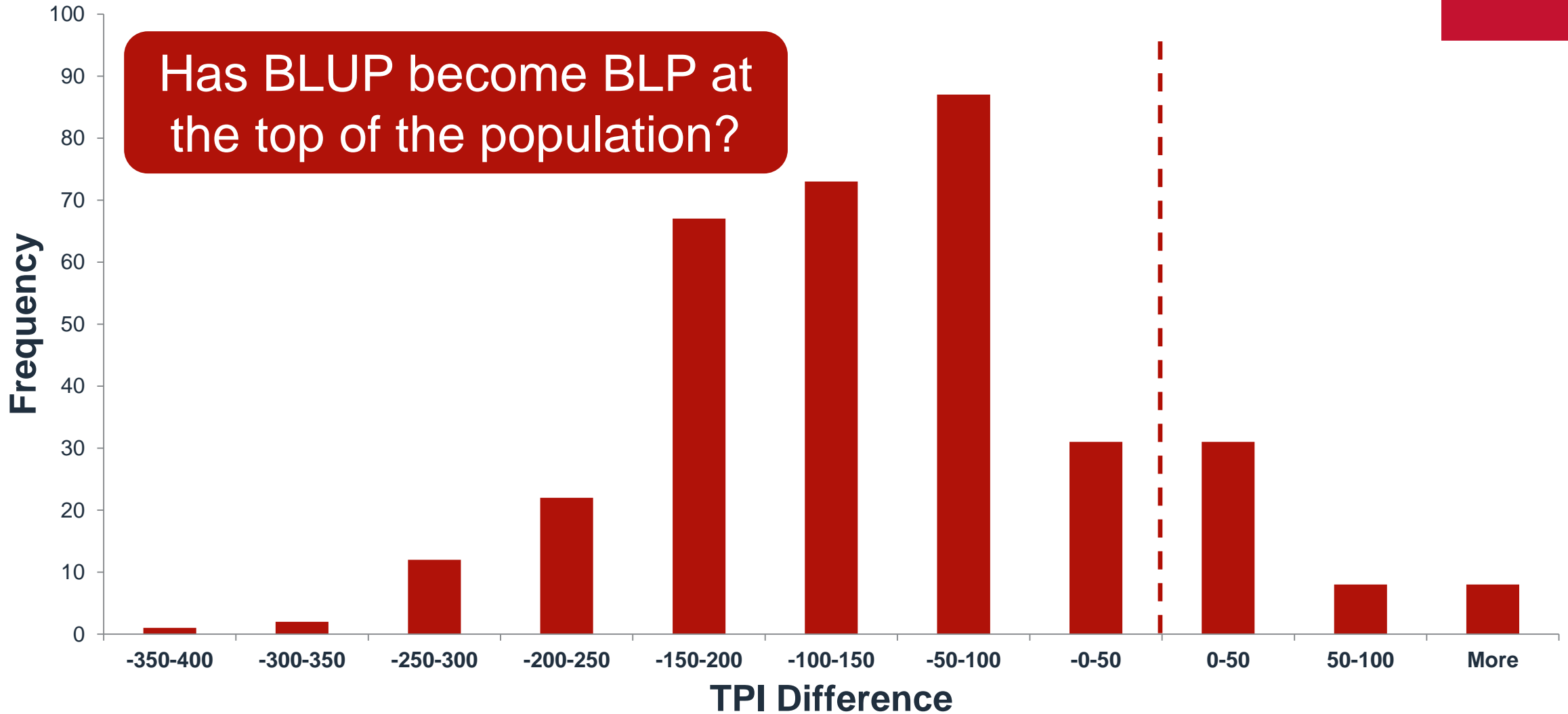


Gain in Reliability with Genomics

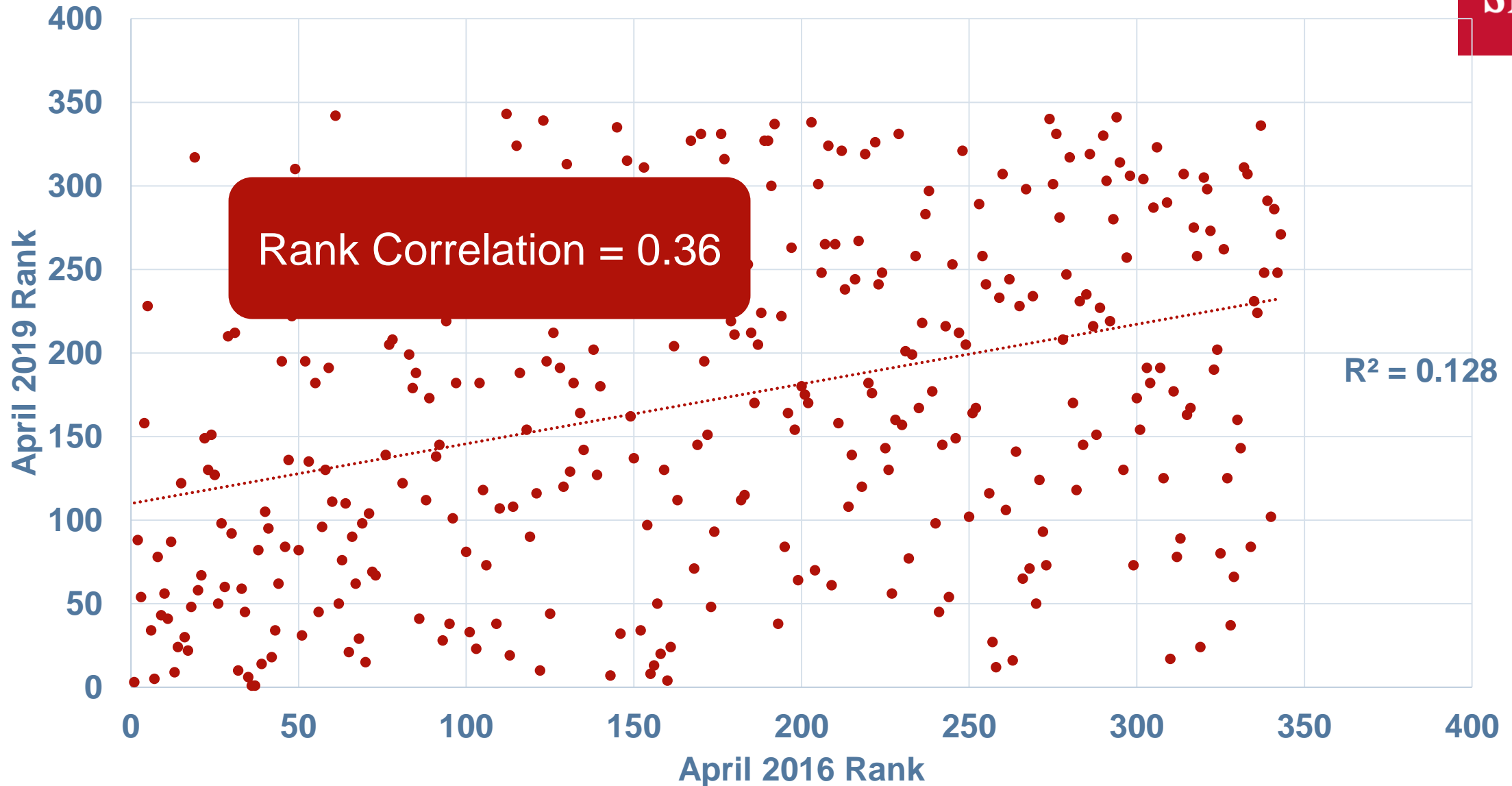


Change in TPI (April 2016 to April 2019) of top 400 Bulls

Has BLUP become BLP at the top of the population?



Top 400 TPI bulls in Apr. 2016 (vs. Apr. 2019 rank)



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Genomic Evaluation	A-	<ul style="list-style-type: none">• Application of theory relatively quick• Very quick uptake of new technology!• Perhaps too much instability in top animals

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Genetic Improvement		

Genetic Improvement

$$\Delta G = i * r * \sigma g / L$$

ΔG = genetic progress per year

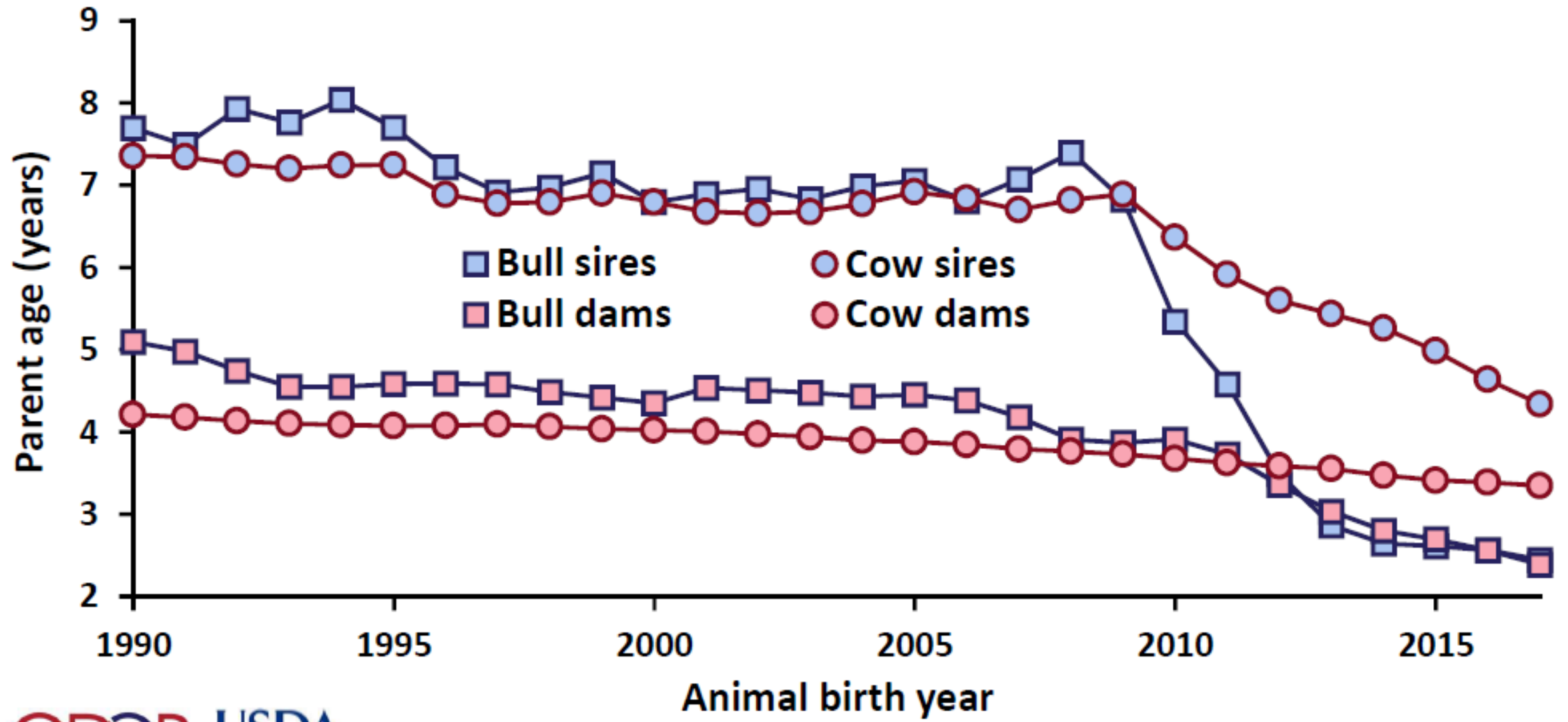
i = intensity of selection (% of selection candidates retained as parents)

r = accuracy of selection (the square root of reliability)

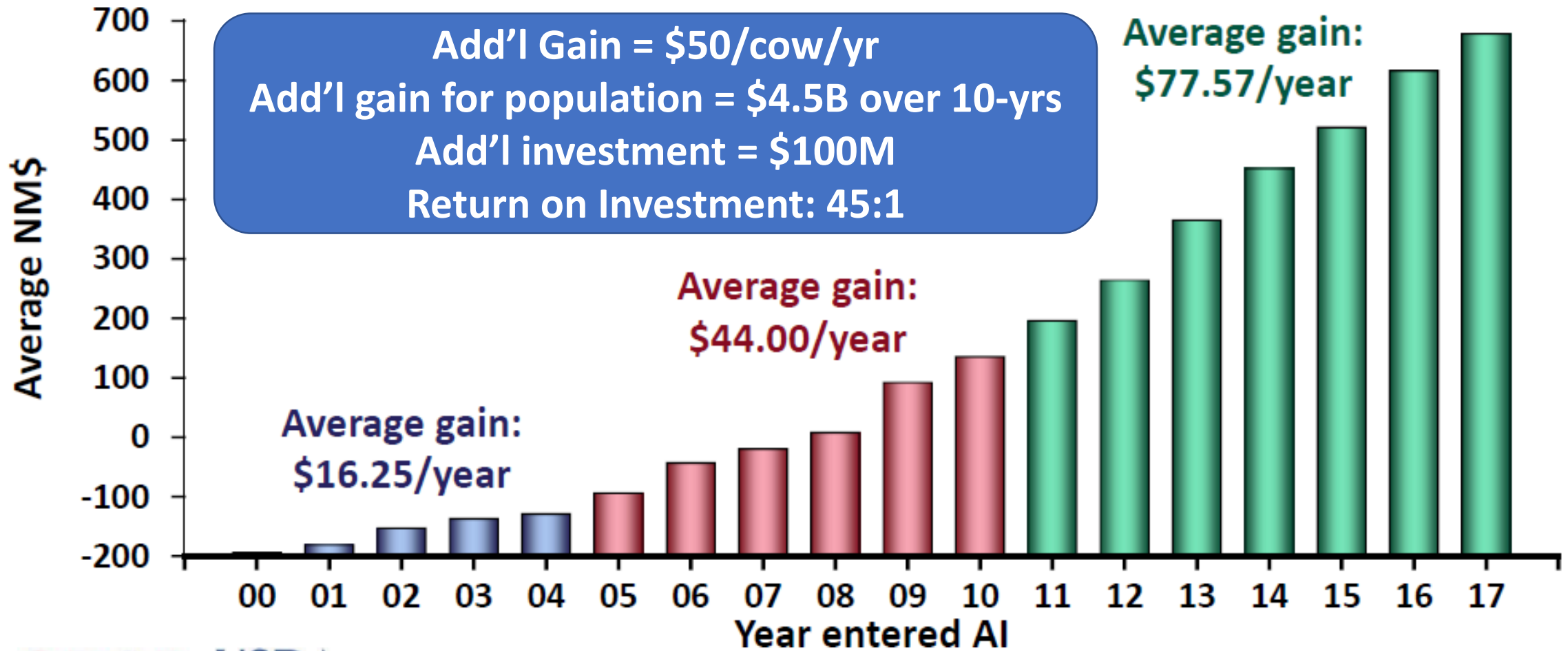
σg = amount of genetic variation in the population

L = generation interval (age of parents when replacement progeny are born)

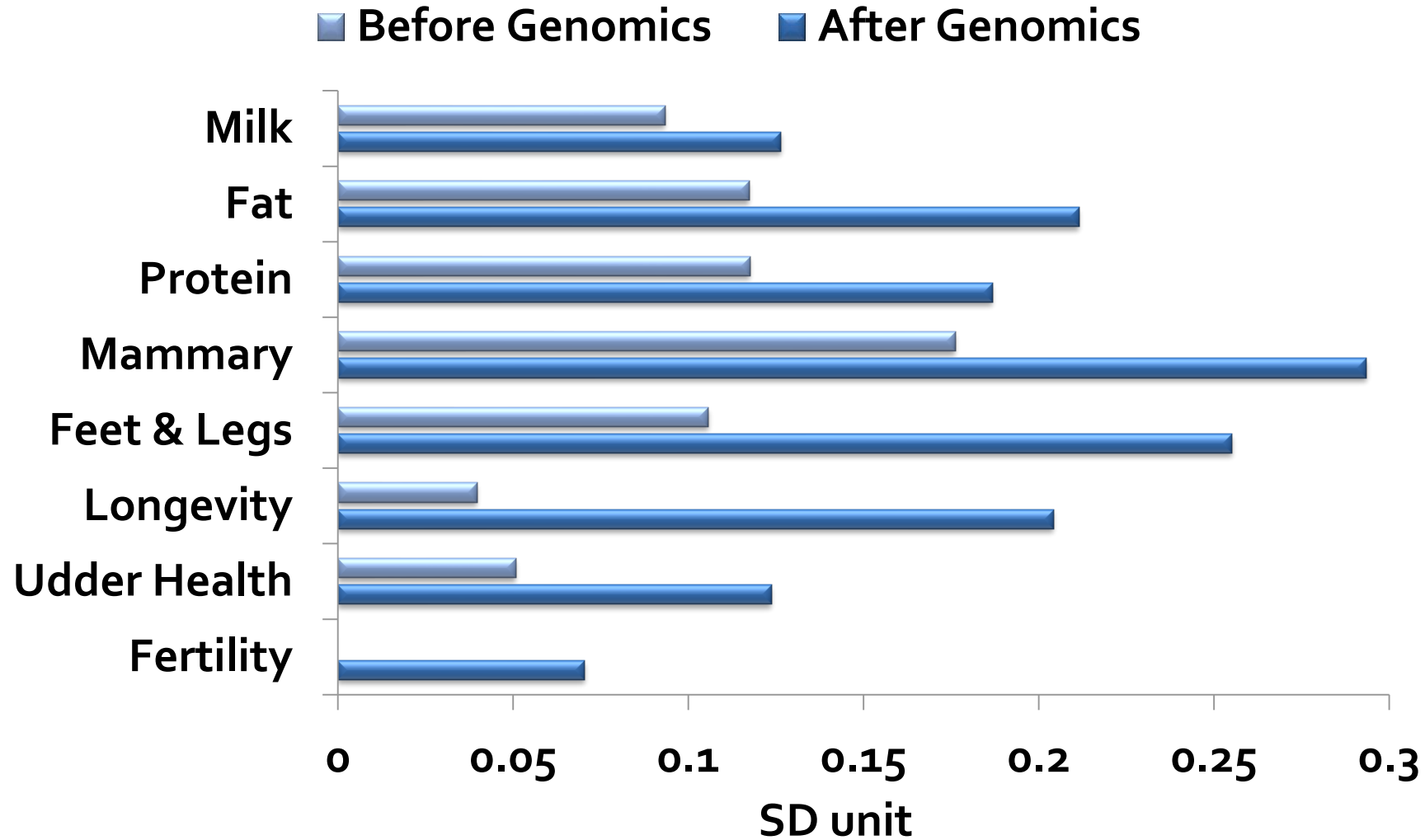
Generation interval – Holstein



Genetic merit of marketed Holstein bulls



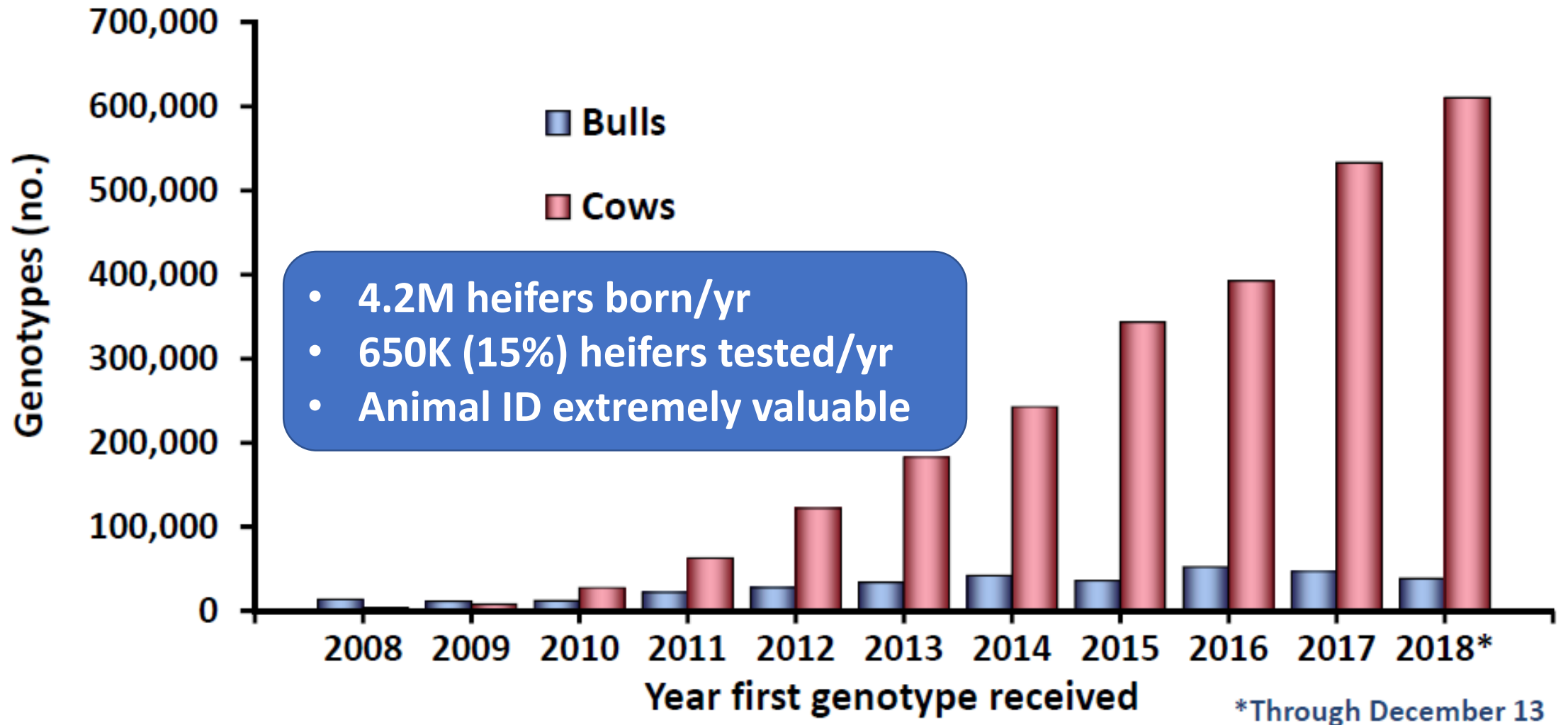
Annual Genetic Progress – Major Traits



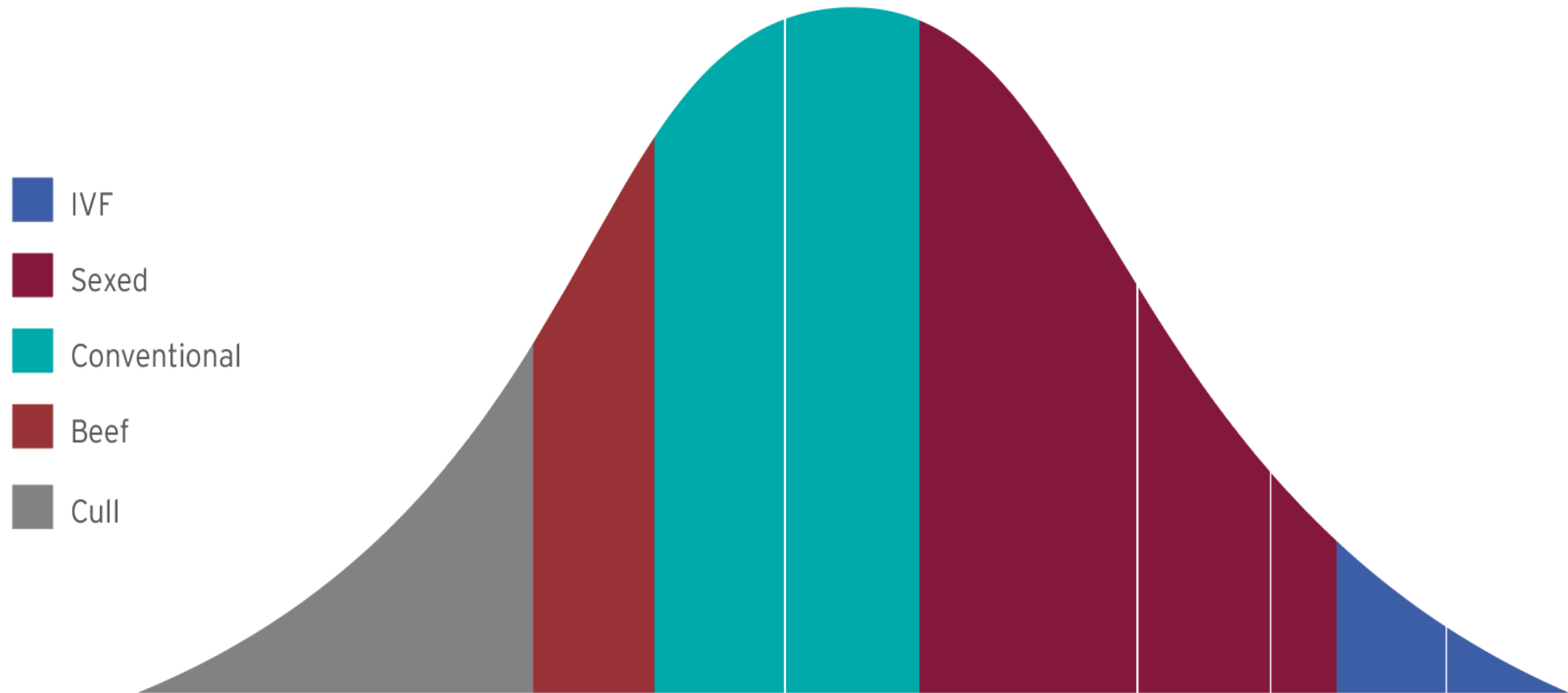
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Genomic Evaluation	A-	<ul style="list-style-type: none">• Application of theory relatively quickly• Very quick uptake of new technology!• Perhaps too much instability in top animals
Genetic Improvement	A+	<ul style="list-style-type: none">• Doubled or tripled genetic progress!• Most significant progress on low h^2 traits• Great return on investment!

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On-Farm Testing		

CDCB usable genotype counts/year by animal sex



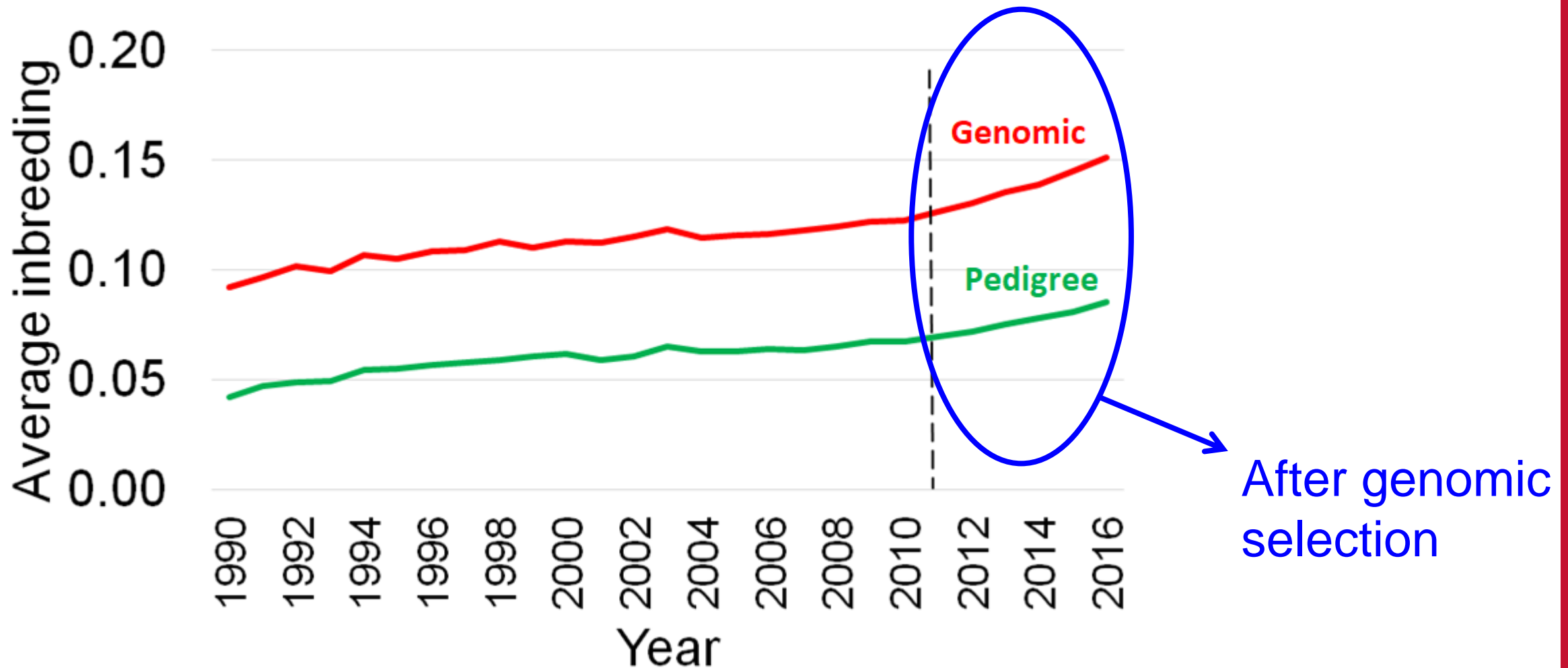
Genotype-Aided Decisions



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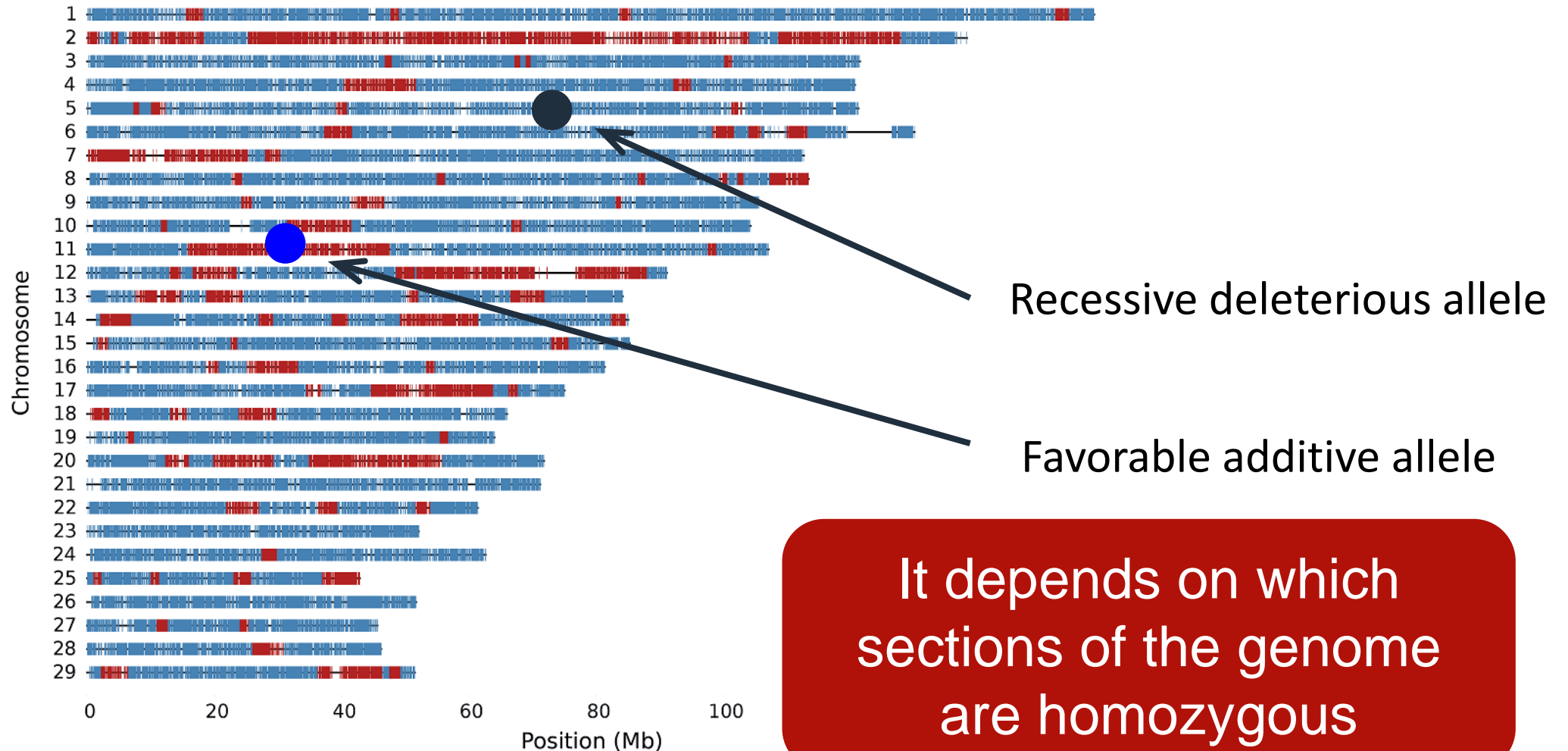


Inbreeding Depression

Trait	Loss per 1% inbreeding
Fat yield (kg)	1.1
Protein yield (kg)	0.5
Conformation (points)	0
Days open (days)	1.4
Calf survival 1 st calving (%)	0.5
Productive life (days)	13

(Van Doormaal, 2008. CDN report. March 2008)

Is Inbreeding Always Bad?



Is Inbreeding always bad?

Bad

- Inbreeding depression
- Reduced fertility & production
- Higher probability of genetic defects and disease
- Loss of between-family genetic variation

Good

- More uniformity in best regions
- Most desirable alleles are “fixed”
- Most undesirable alleles are “purged”
- More potential for hybrid vigor in crosses

Effective Pop. Size $< 50 \rightarrow$ (20% less long-term gain)

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Understanding Genotype to Phenotype		

Understanding Genotype to Phenotype

- Genomic improvement still essentially a “black box”
- The genomic SNP profile is only part of the story
- DNA → RNA → Protein pathway variation not well understood (epigenetic, GxE, etc.)
- Non-additive genetic variation is difficult to predict (heterosis/inbreeding depression, GxG interactions)
- Few additional causative mutations have been found

Reducing Freq. of Undesirable Haplotypes

Haplotypes affecting fertility

Name ¹	Chromosome ²	ARS-UCD location (Mbp) ³	Current carrier frequency (%)	Earliest known genotyped ancestor
HH1	5	62.8*	2.6	Pawnee Farm Arlinda Chief
HH2	1	93.5 – 95.6	2.4	Willowholme Mark Anthony
HH3	8	93.8*	5.3	Glendell Arlinda Chief, Gray View Skyliner
HH4	1	2.0*	0.5	Besne Buck
HH5	9	91.8 – 91.9	4.8	Thornlea Texal Supreme
HH6	16	29.0 – 29.1	0.9	Gray View Skyliner
JH1	15	15.4*	18.4	Observer Chocolate Soldier
BH2	19	10.8*	13.3	Rancho Rustic My Design
AH1	17	63.7*	22.3	Selwood Betty's Commander
AH2	3	51.1	13.3	Oak-Ridge Flashy Kellogg

¹BH1 and JH2 discontinued ²*Bos taurus* (BTA) ³Mbp = megabase pairs; * = causative mutation known

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Understanding Genotype to Phenotype	D	<ul style="list-style-type: none"> • Overpromised and underdelivered • Genotype to phenotype path is VERY complex • Some genes identified for disease traits

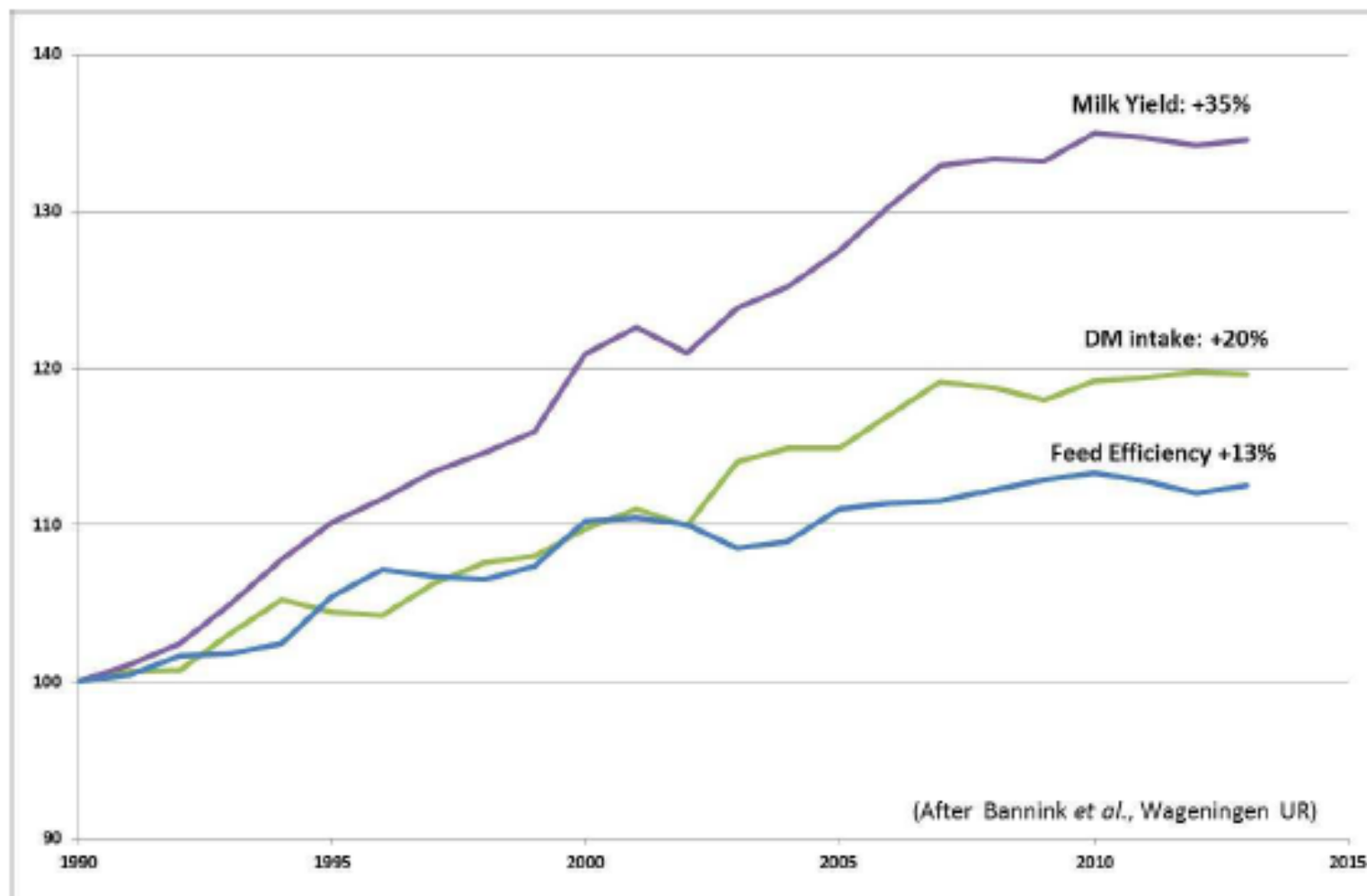
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Meeting Consumer Expectations		

Meeting Consumer Expectations



- Safe & affordable ✓
- Good for environment ✓
- Hormone/antibiotic-free ?
- Not cruel to animals ?
- More choice
 - Taste, Variety, Local
 - Digestibility & health claims ?
 - Production methods

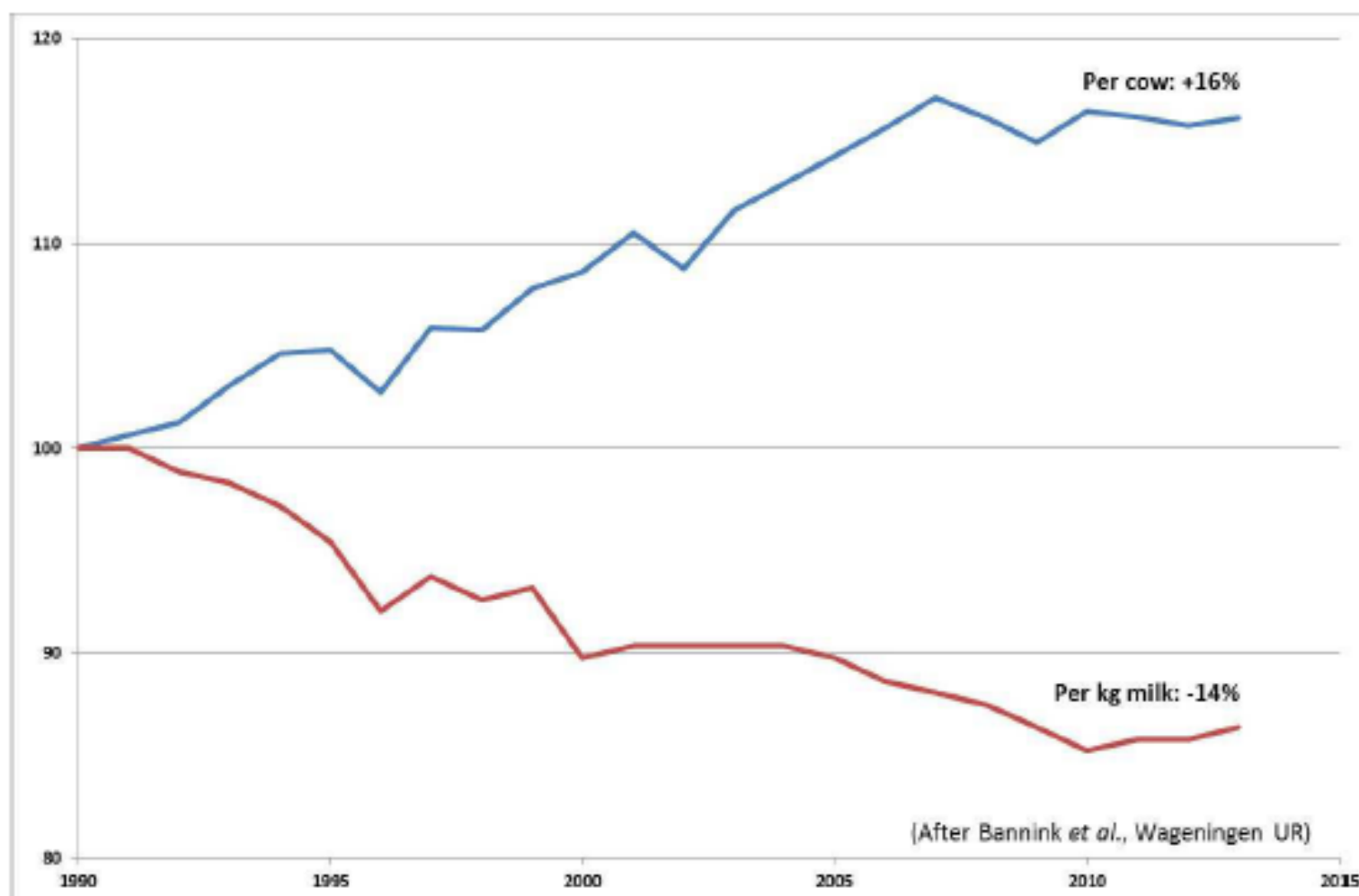
Dairy – production efficiency



Milk yield increased faster than DM intake

-> better feed efficiency

Dairy – enteric CH₄ emission

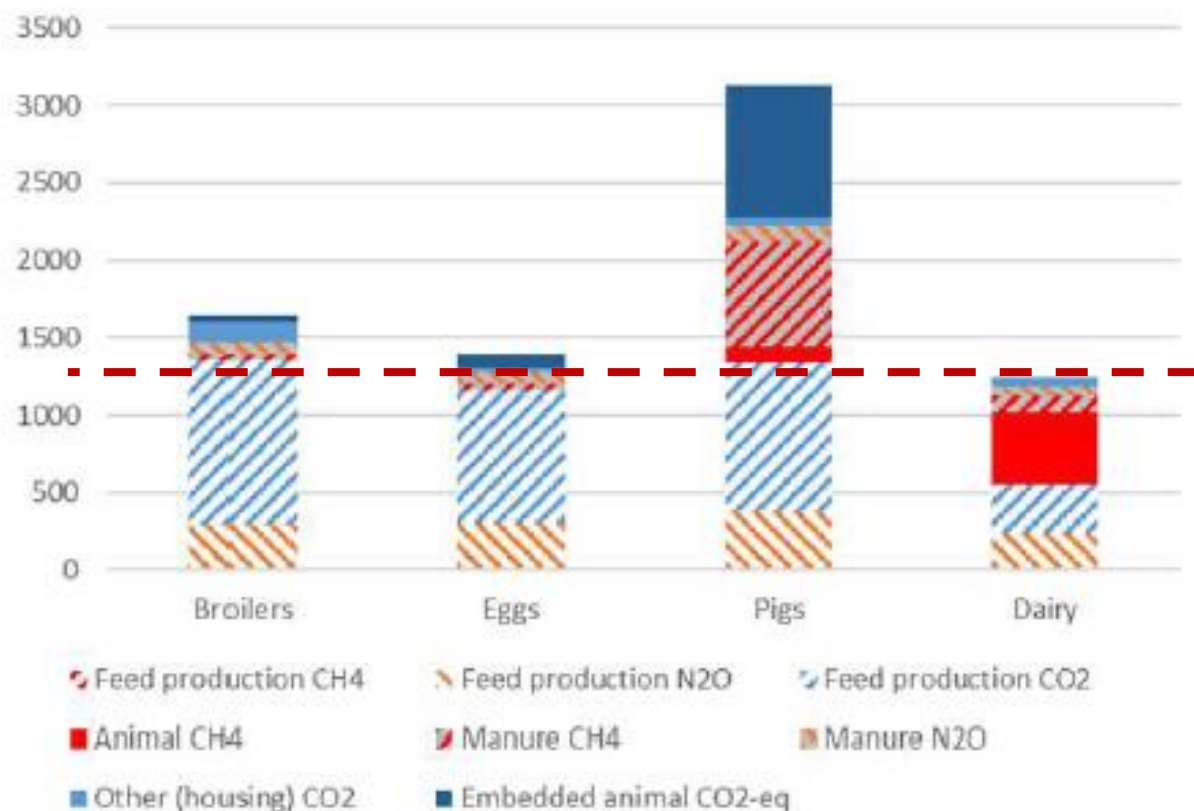


Increased per cow

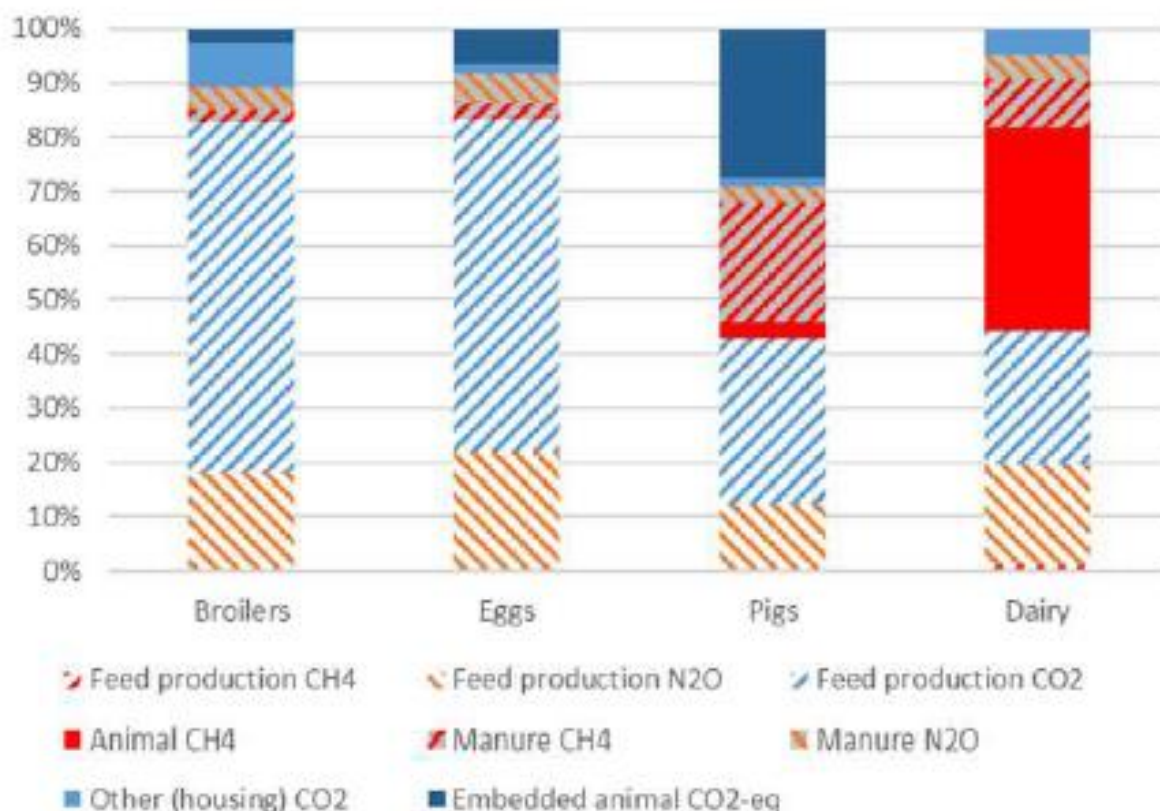
Decreased per kg milk

GHG emissions of different species

Absolute (kg CO₂-eq / ton product) contribution to GHG emissions (without LULUC)



Percentage contribution to GHG emissions (without LULUC)



Dairy – reduction of GHG¹

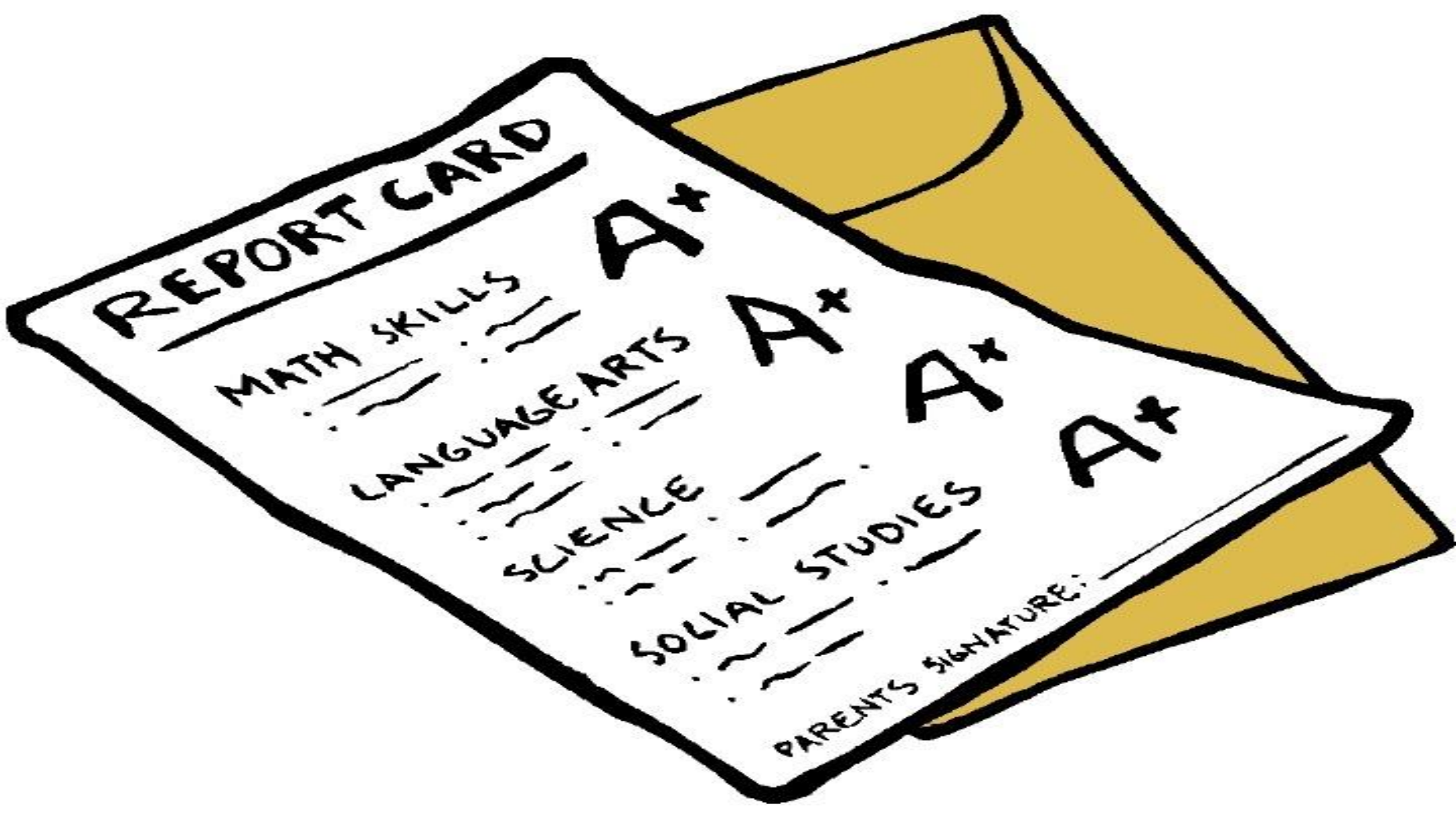
Carbon footprint dairy

1990: 2.06 kg CO₂-eq. / kg milk
2012: 1.42 kg CO₂-eq. / kg milk } -31%

Similar reductions (30-50%) are possible via nutrition
(e.g. 3-nitrooxypropanol (3NOP))

K. Beauchemin, AAFC

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REPORT CARD

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LANGUAGE ARTS

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SOCIAL STUDIES

A+

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Thank You



Han Hopman