

# USDA AGIL Research Updates:

*Improving dairy animals by increasing accuracy of genomic prediction, evaluating new traits, and redefining selection goals*

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**Animal Genomics & Improvement Laboratory**

**USDA Agricultural Research Service**

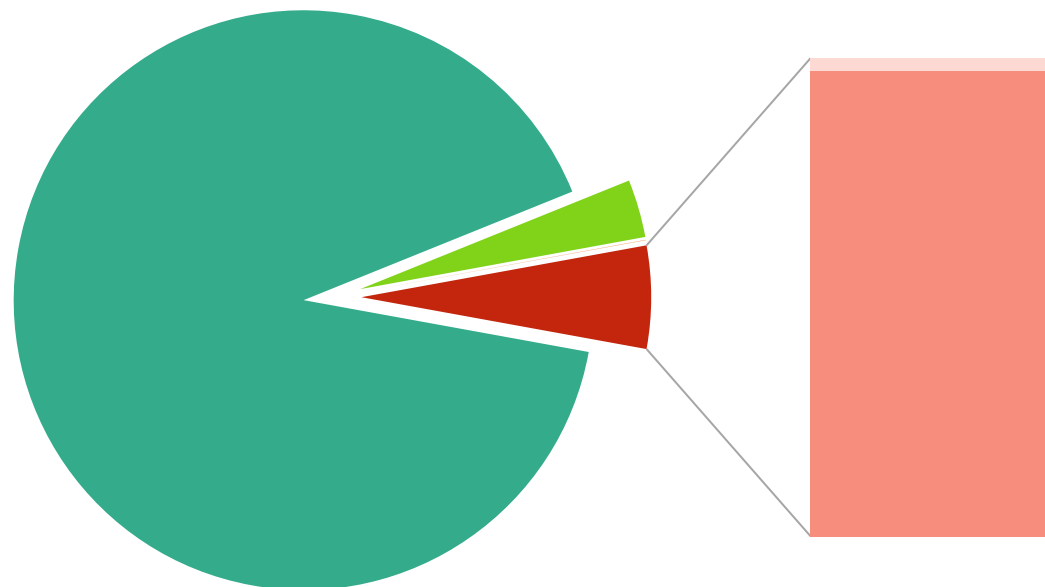
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# Embryo Transfer & Fertility Evaluations

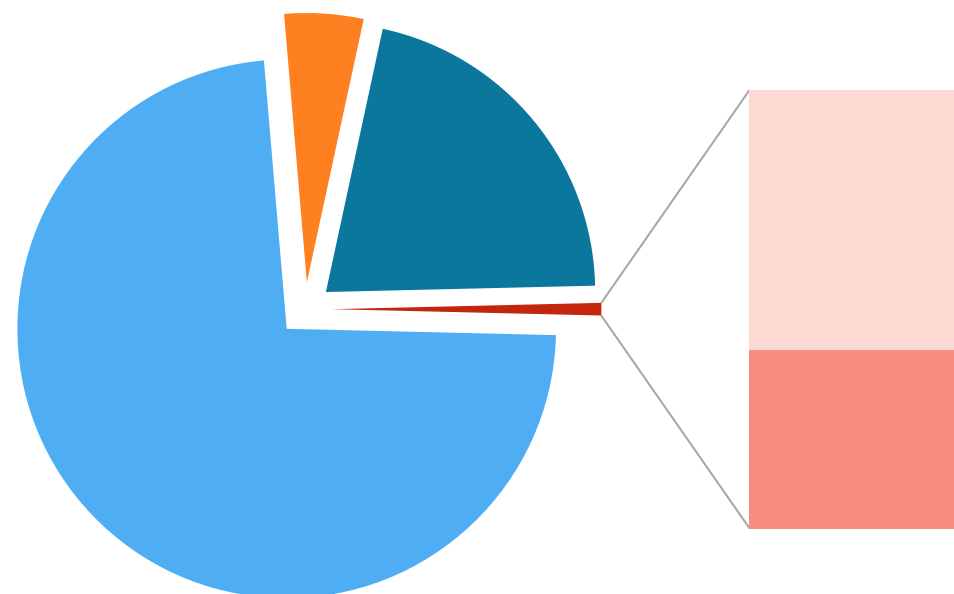
Dr. Asha Miles  
Jana Hutchison  
Dr. Paul VanRaden

## CALVING EVENTS



- Single Birth
- Multiple Birth (not ET)
- Split Embryo (artificially)
- Clone from Nuclear Transfer
- Embryo Pedigree (implantation stored as birth date)
- Birth from ET

## BREEDING EVENTS

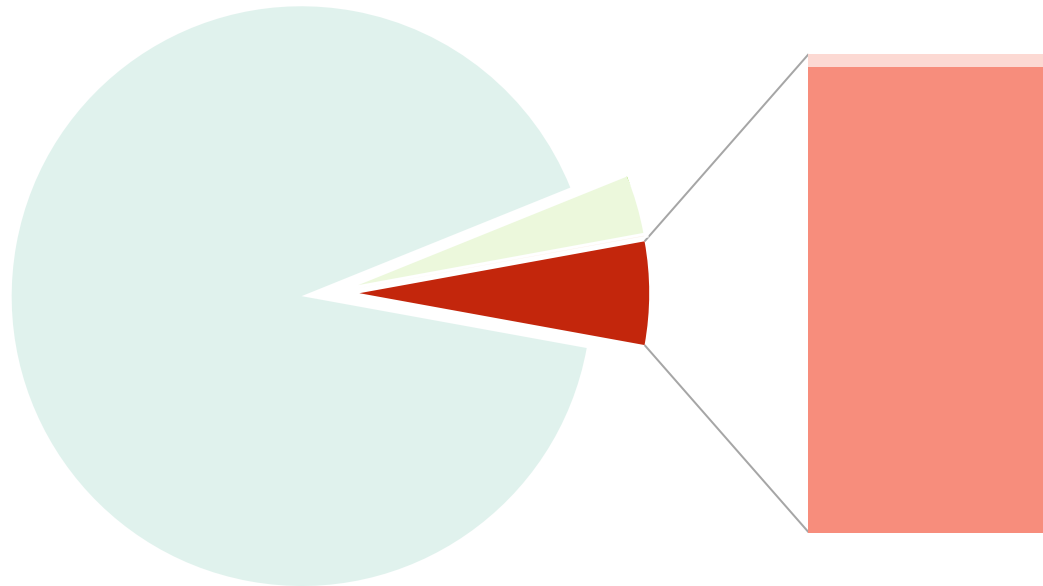


- AI Breeding
- Natural Service Breeding
- AI Sexed Semen
- Embryo Donation
- Embryo Implantation (reporting embryo sire)
- Embryo Implantation (reporting embryo dam)

# Embryo Transfer & Fertility Evaluations

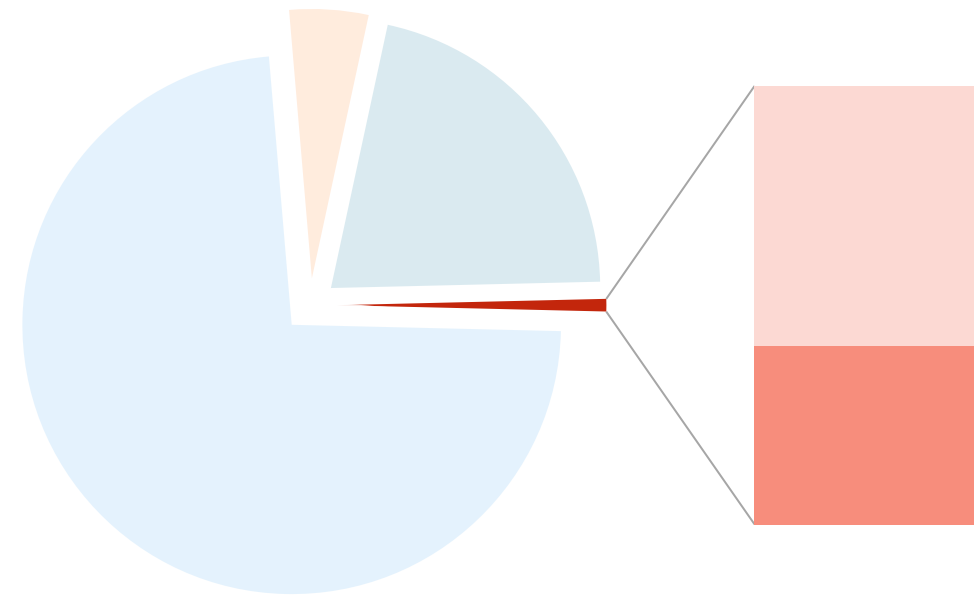
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Format 5:  
Code E  
Code I  
Code J



# Embryo Transfer & Fertility Evaluations

Dr. Asha Miles  
Jana Hutchison  
Dr. Paul VanRaden

*Is this discrepancy because ET is incorrectly being coded as AI?*

Code	Mating Type	ET Births	All Other Calvings
A	AI	35,100	11,060,000
G	AI (sexed semen)		
N	Natural Service		
E	Embryo Donation		
I/J	Embryo Implantation		

**0.32%**

The rate at which ET is **incorrectly** recorded

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*Is this discrepancy because ET is incorrectly being coded as AI?*

Code	Mating Type	ET Births	All Other Calvings	
A	AI	35,100	11,060,000	<b>0.32%</b>
G	AI (sexed semen)	14,833	1,175,795	<b>1.25%</b>
N	Natural Service	2,035	584,971	<b>0.35%</b>
E	Embryo Donation	0	13	<b>0.00%</b>
I/J	Embryo Implantation	372	29,416	<b>1.25%</b>

The rate at which ET is **incorrectly** recorded

The rate at which ET is **correctly** recorded

**This suggests ET is not being reported at all, rather than being reported incorrectly**

# Embryo Transfer & Fertility Evaluations

Dr. Asha Miles  
Jana Hutchison  
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- **Some solutions:**
  - **Cleaning-up historical data**
  - **Breed associations report ET**
  - **Better on-farm ET recording (VAS, DC305)**
  
- **This has implications for:**
  - **On-farm reports about fertility**
  - **Fertility evaluations (e.g., SCR, HCR, and CCR)**
  - **Bulls whose 1<sup>st</sup> calves may all be ET but are not reported as such**

- Official factors for adjusting lactation records to mature equivalent were last estimated in 1994 by Mike Schutz
- At that time, additive adjustment factors were included in the animal model by George Wiggans to automatically adjust future data for changes in maturity rates

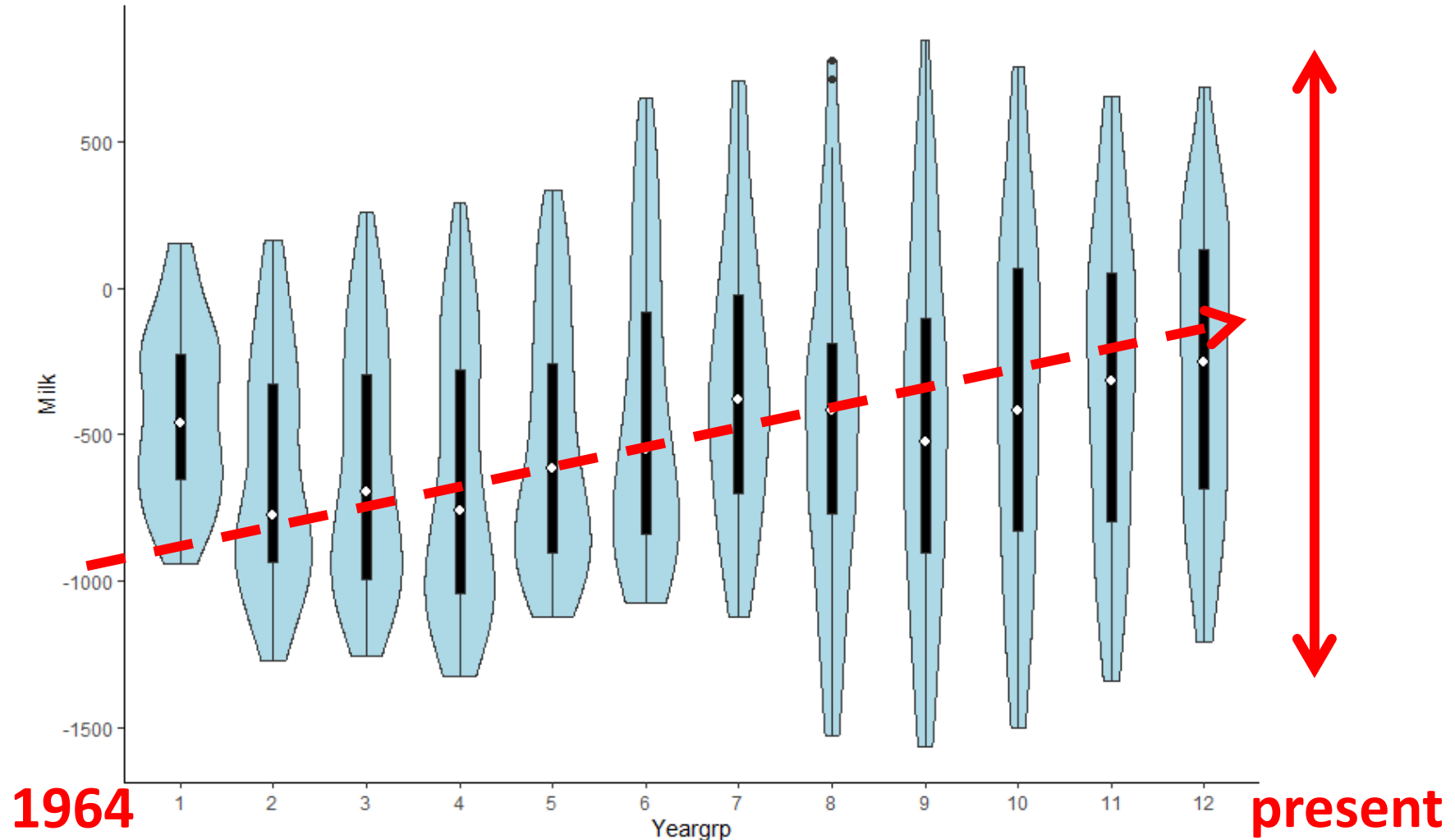
**1. Look at size of additive corrections to assess how well preadjustment factors are working**

# Mature Equivalent & Age Adjustment

Dr. Asha Miles  
Dr. Paul VanRaden

Milk

Factors are indeed changing over time



1964

present



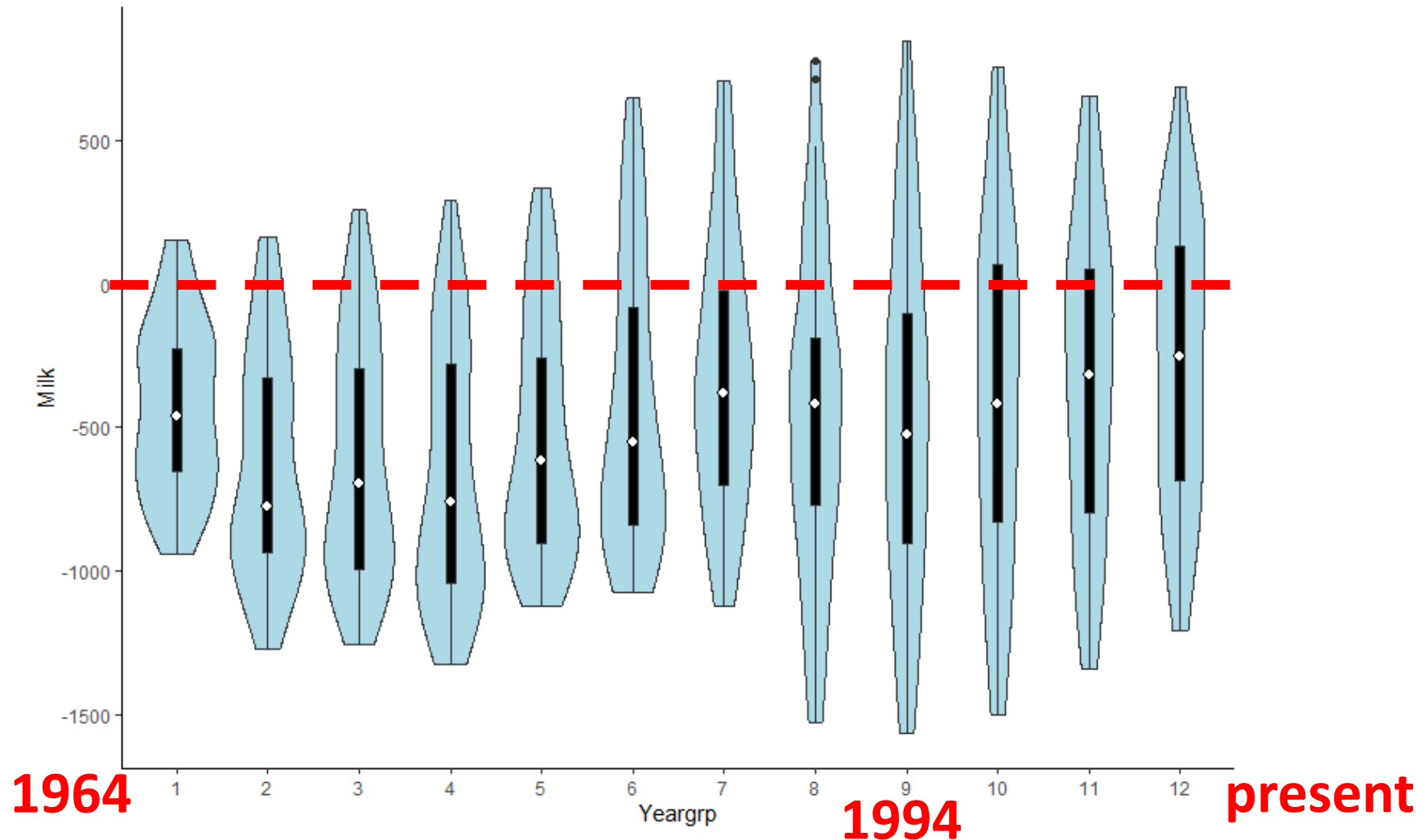


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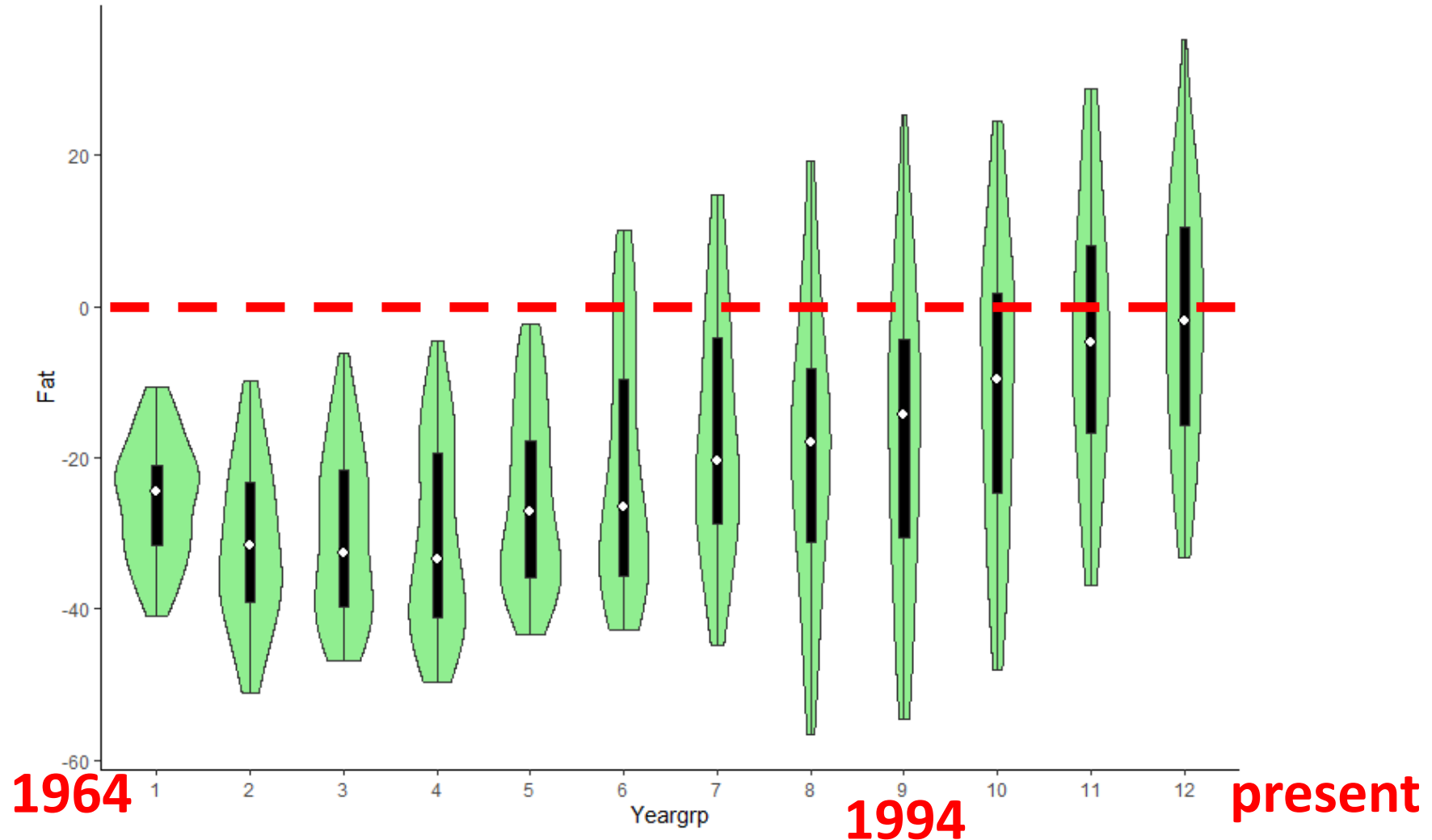


# Mature Equivalent & Age Adjustment

Dr. Asha Miles  
Dr. Paul VanRaden

## Fat

Factors are indeed changing over time

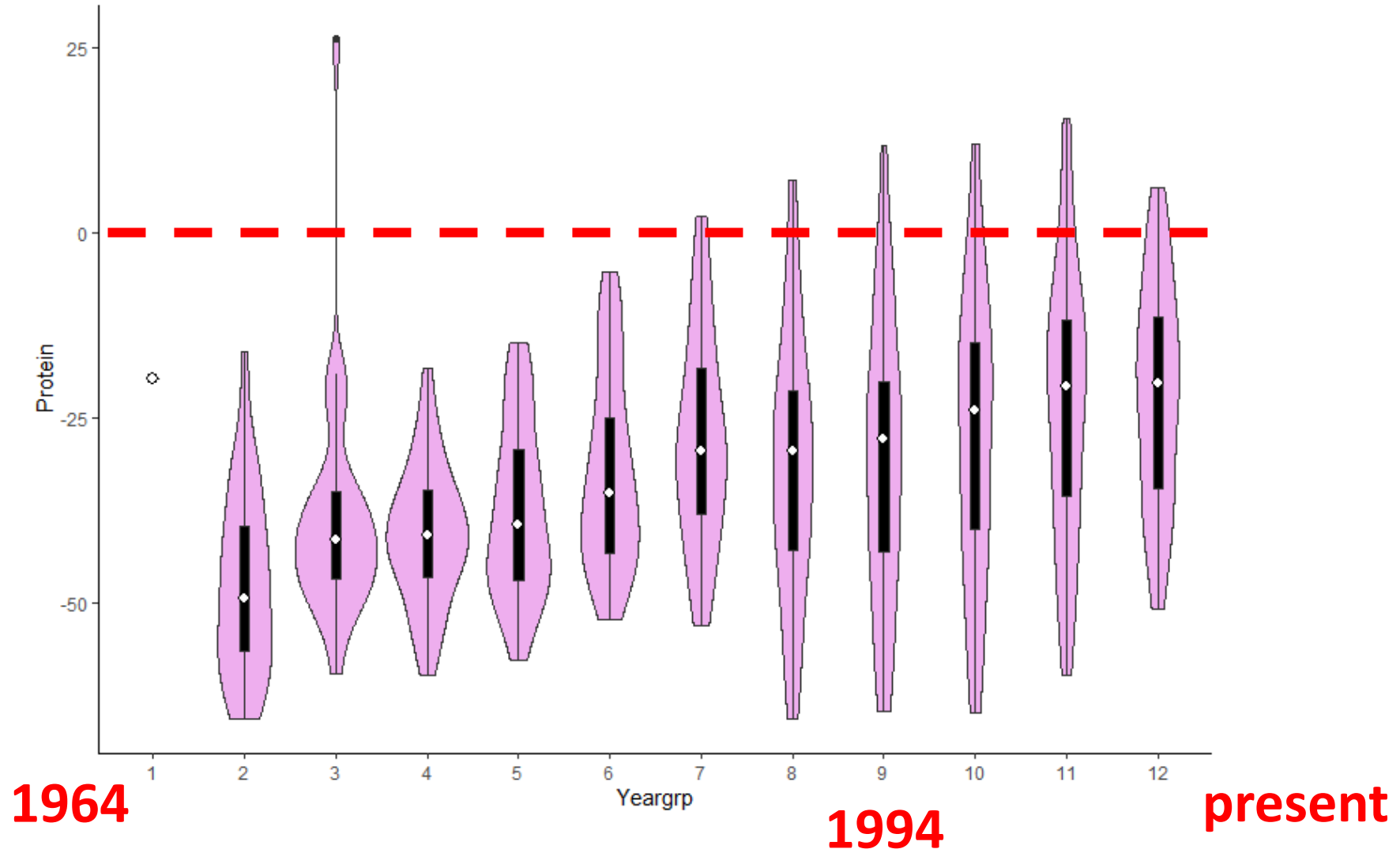


# Mature Equivalent & Age Adjustment

Dr. Asha Miles  
Dr. Paul VanRaden

## Protein

Factors are indeed changing over time



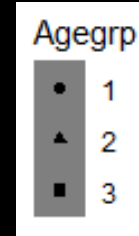
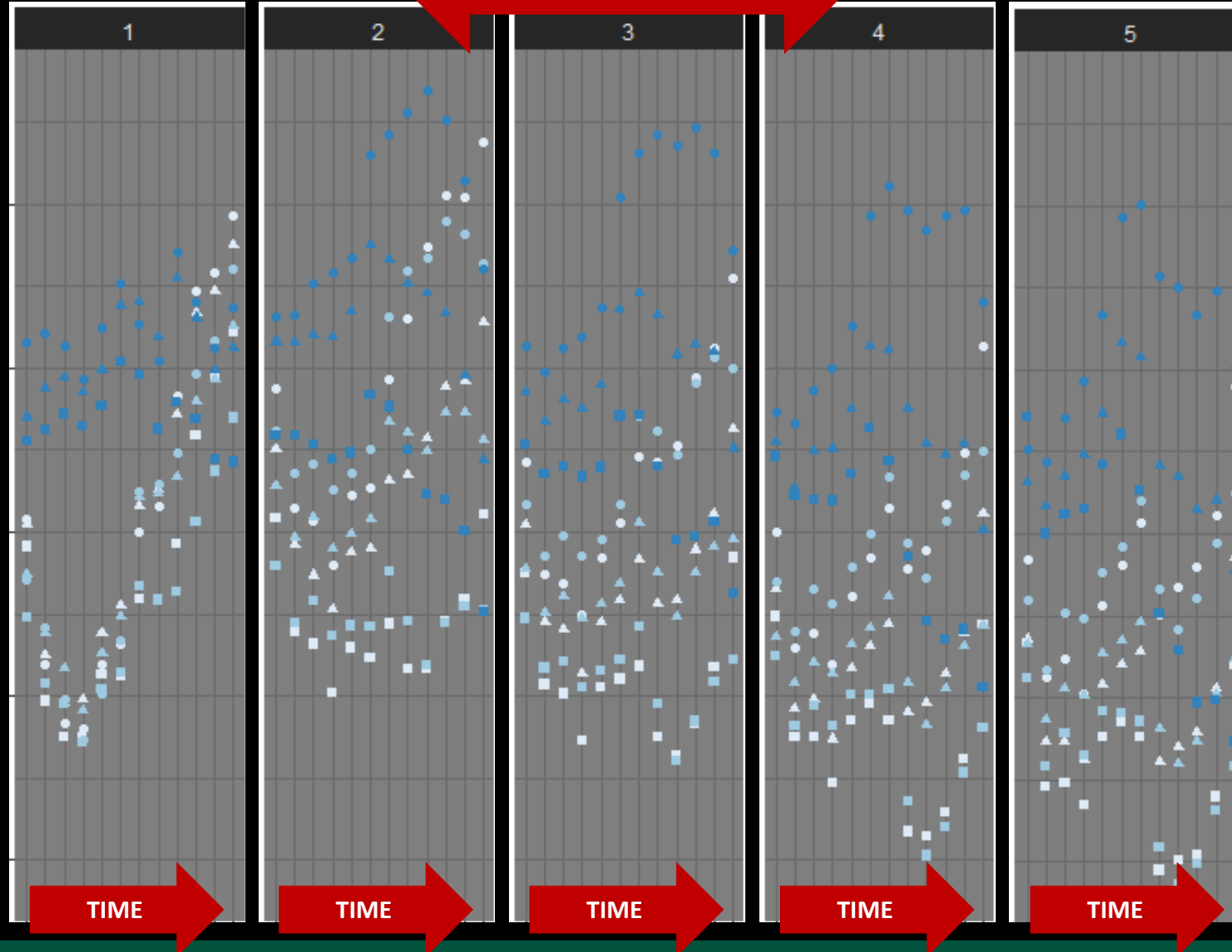
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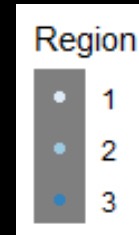
Milk

Additive adjustment factors

PARITY



AGE W/IN  
PARITY



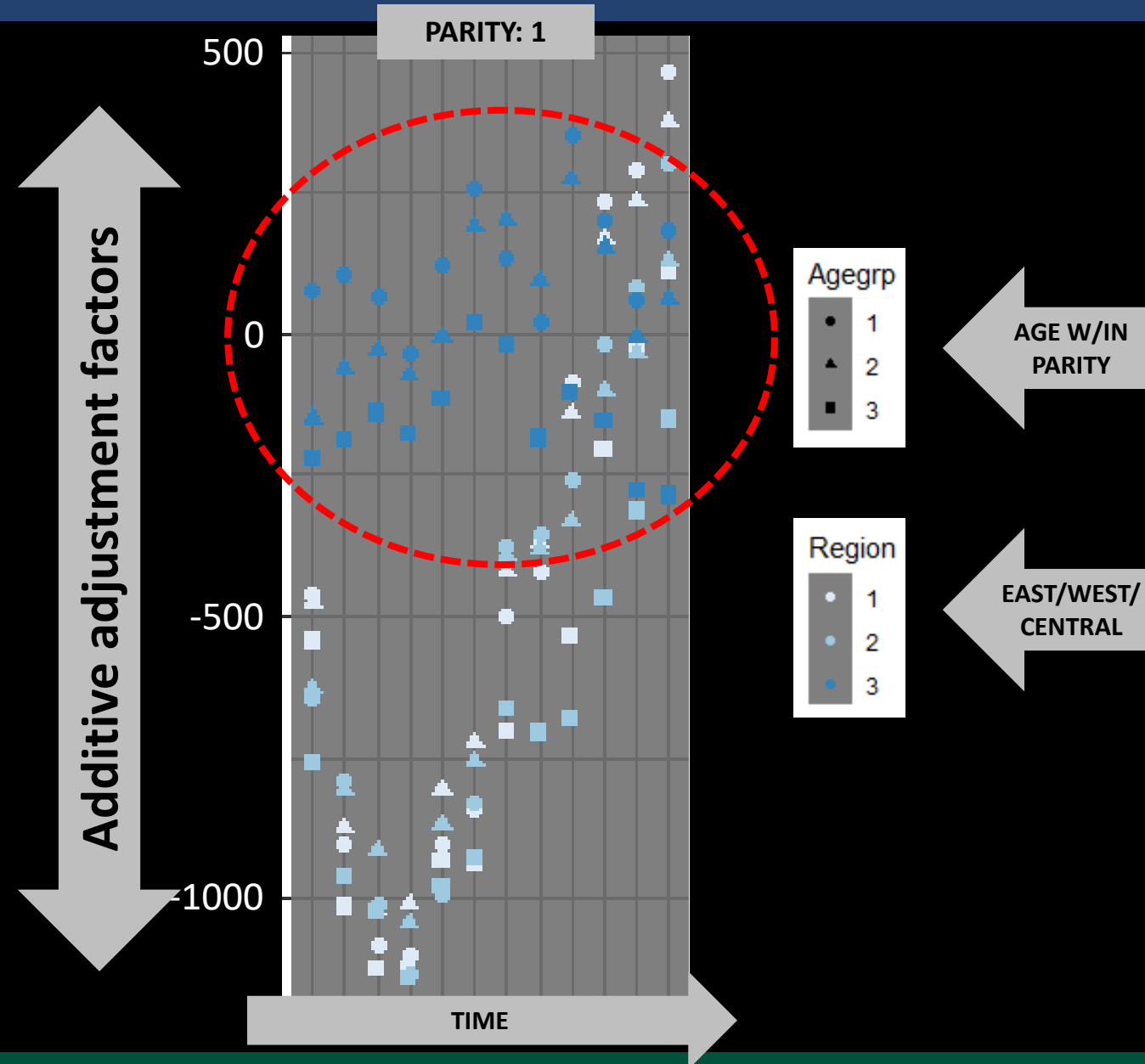
EAST/WEST/  
CENTRAL



# Mature Equivalent & Age Adjustment

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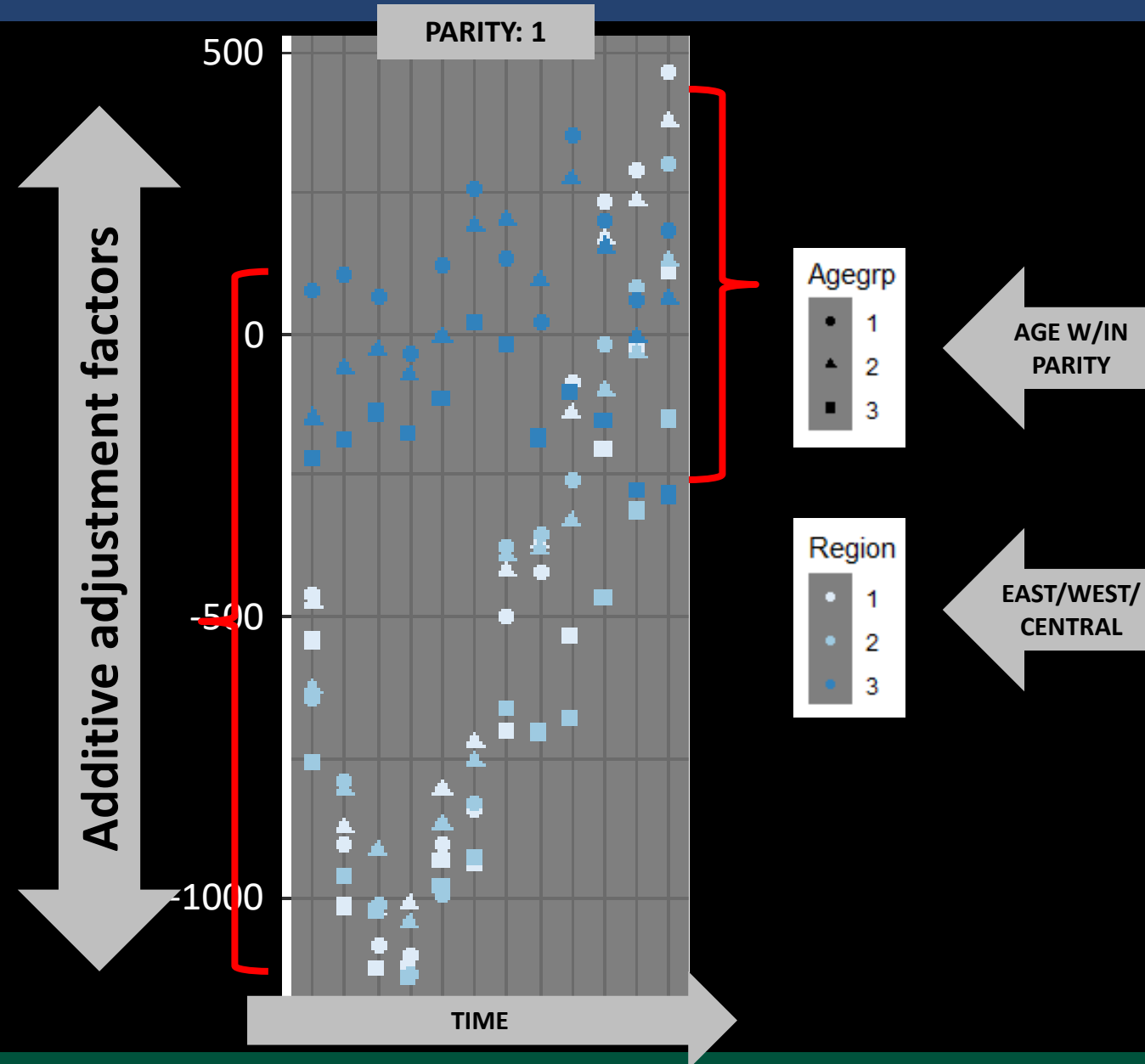
Milk



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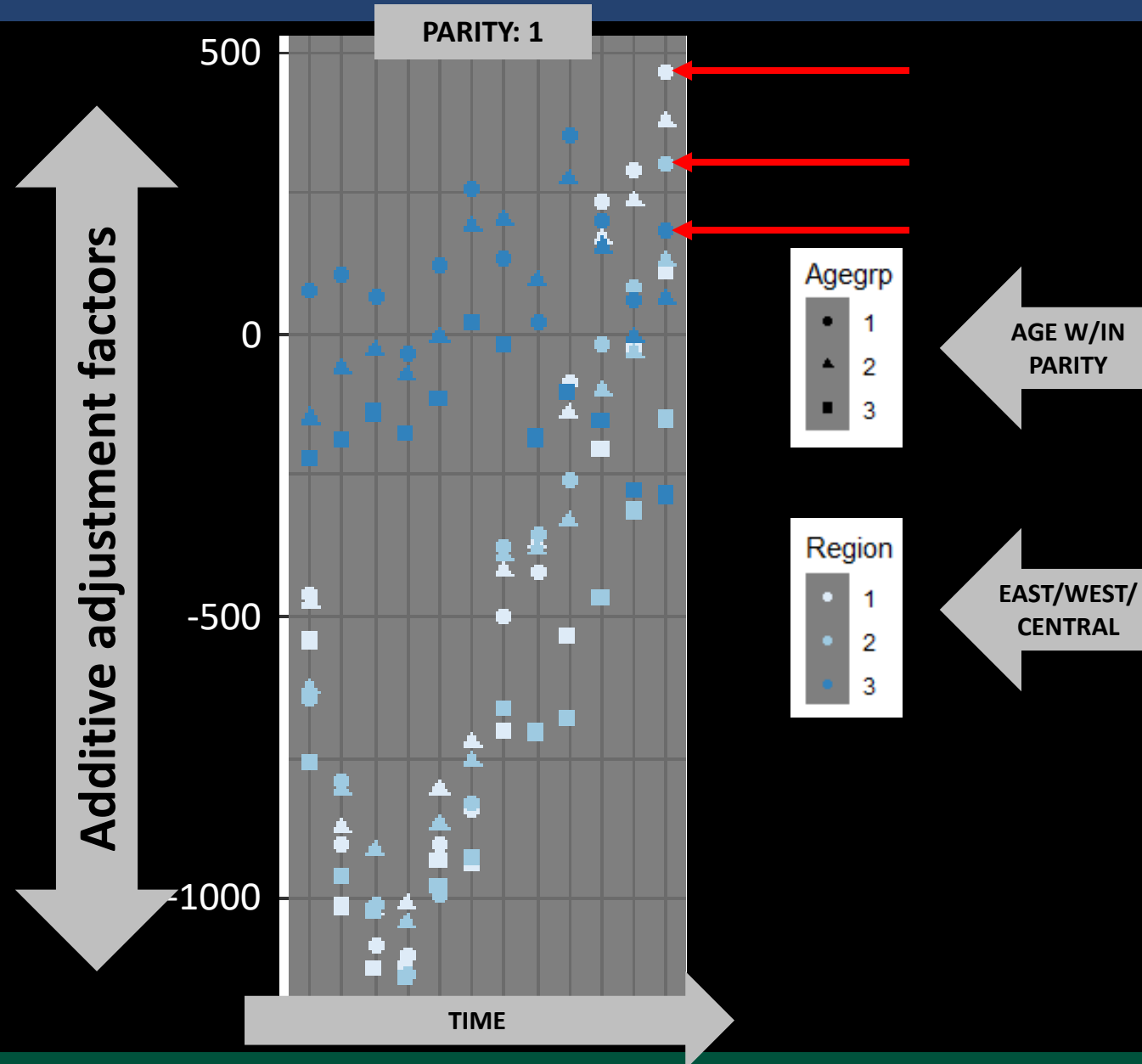
Milk



# Mature Equivalent & Age Adjustment

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Dr. Paul VanRaden

Milk



## 2. Improve preadjustment corrections

- In 2005, PTAs were adjusted to 36 months instead of mature age to make predicted yield differences more similar to actual average yields and to make breed comparisons fairer in the all-breed animal model (est. 2007)

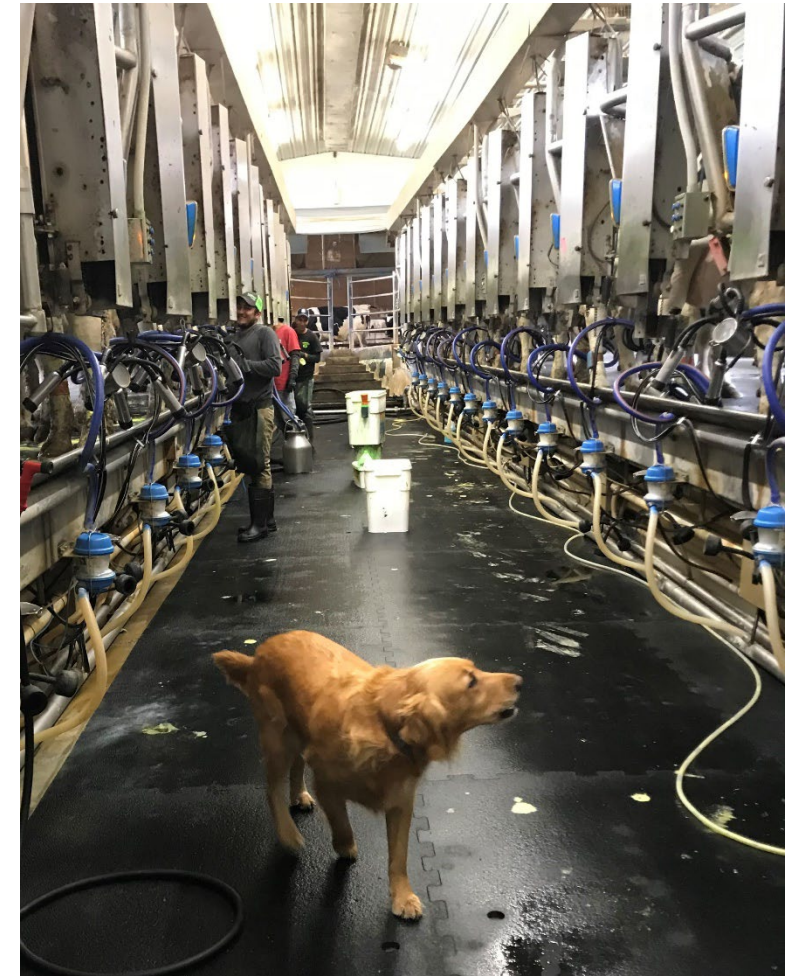


# Flexible Testing & Milk-Only Records

Dr. Paul VanRaden  
Gary Fok

- Some farms with in-line milk meters do not participate in DHI testing; they record milk weights but not fat and protein
- National genetic evaluations exclude milk-only records; these records are stored but never extracted
- There is not a need for more milk records, but removing the censorship of herds without approved component sampling will allow us to use all of their other traits

**This will improve the accuracy of prediction for lower heritability traits**



*Jack, a Pennsylvania Herd Manager, surveys morning milking*

- Top bull lists obtained using Multitrait Across-Country Evaluations (MACE) on each country's scale

**Table 1.** Actual percentages of foreign sire use and expected use based on the top 100 or top 1000 proven Holstein sires in each country's ranking

Foreign%	AUS	CAN	CHE	DEU	DFS	ESP	FRA	GBR	IRL	ISR	ITA	JPN	NLD	NZL	USA
Expected <sup>2</sup>	96	83	100	94	90	100	98	100	81	91	100	98	87	24	8
Expected <sup>3</sup>	98	88	100	91	90	99	95	99	86	95	98	96	92	54	23

<sup>1</sup> Percentages of milk-recorded cows with foreign sires born since 2008.

<sup>2</sup> Percentages of top 100 proven sires born 2005-2013 that are foreign.

<sup>3</sup> Percentages of top 1000 proven sires born 2005-2013 that are foreign.

**Most countries should increase their use of foreign sires**

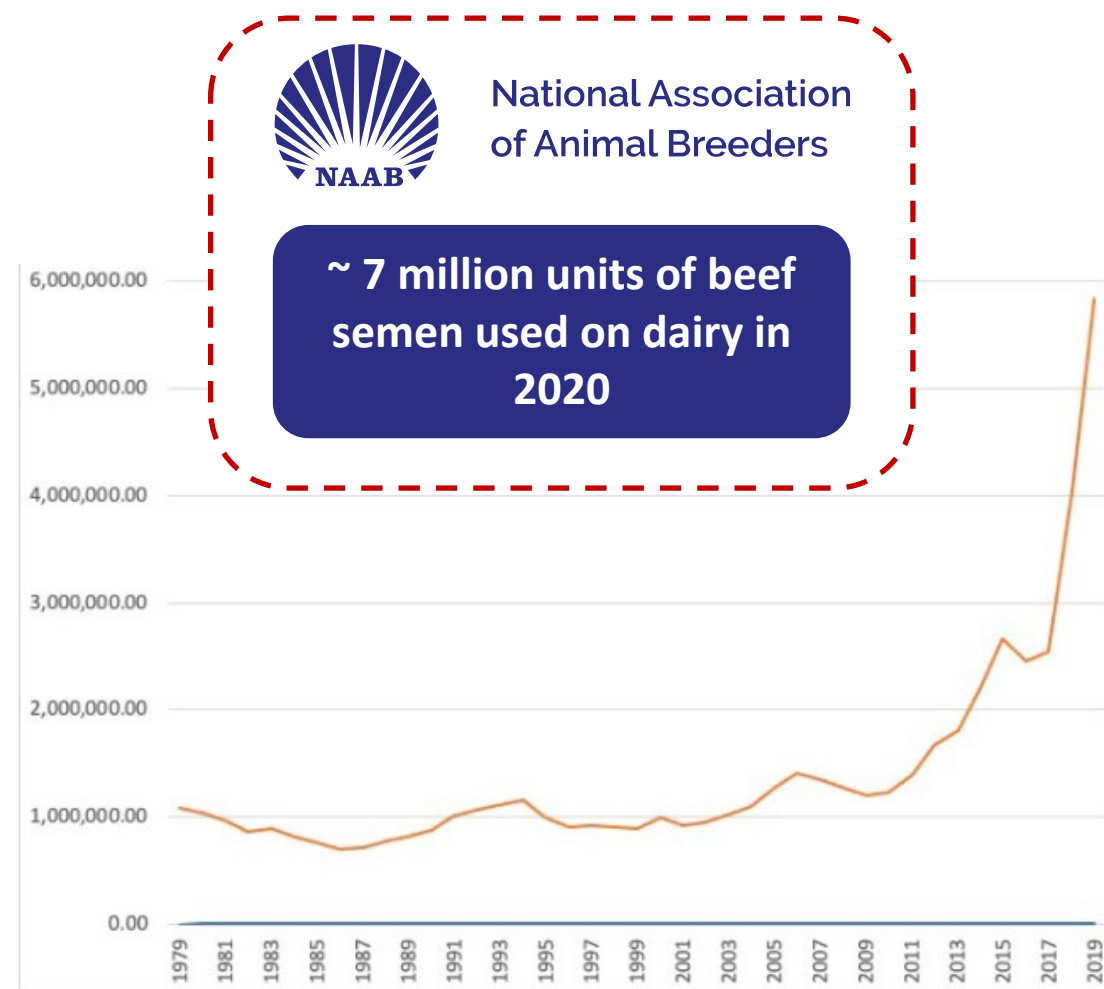
# Long term considerations

**USDA is writing our 5-year plan for submission this November**



# Beef on Dairy: Effects on Dam Productivity

- With recent low milk and heifer prices, producers have turned to beef supply chain as an alternative revenue source
- In 2017, up to 20% of fed cattle were Holstein<sup>1</sup>
- Cow conception rate does not appear to be affected by breed of service sires (McWhorter et al., 2020)



From: Geiger, C. Beef on dairy more than doubled in two years. <https://hoards.com/article-27667-beef-on-dairy-more-than-doubled-in-two-years.html>

<sup>1</sup>NBQA. National Beef Quality Audit: Steer and Heifer. (2016).

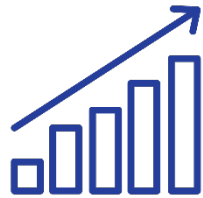
**Question: how does carrying a crossbred calf affect milk production?**



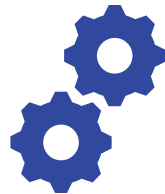
**A recent study found dairy cows do have decreased milk production if bred to beef service sire, but the higher value of her calf makes this loss negligible**

*Berry and Ring 2020. J Dairy Sci 103(9)*

**This will not always be the case depending on variable meat and milk markets**



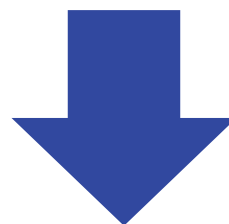
**Quantify effects of beef service sire merit on milk yield & quality**



**Develop decision-making tool for producers based on this data and current market trends**



$$\text{Phenotype (P)} = \text{Genotype (G)} + \text{Environment (E)}$$

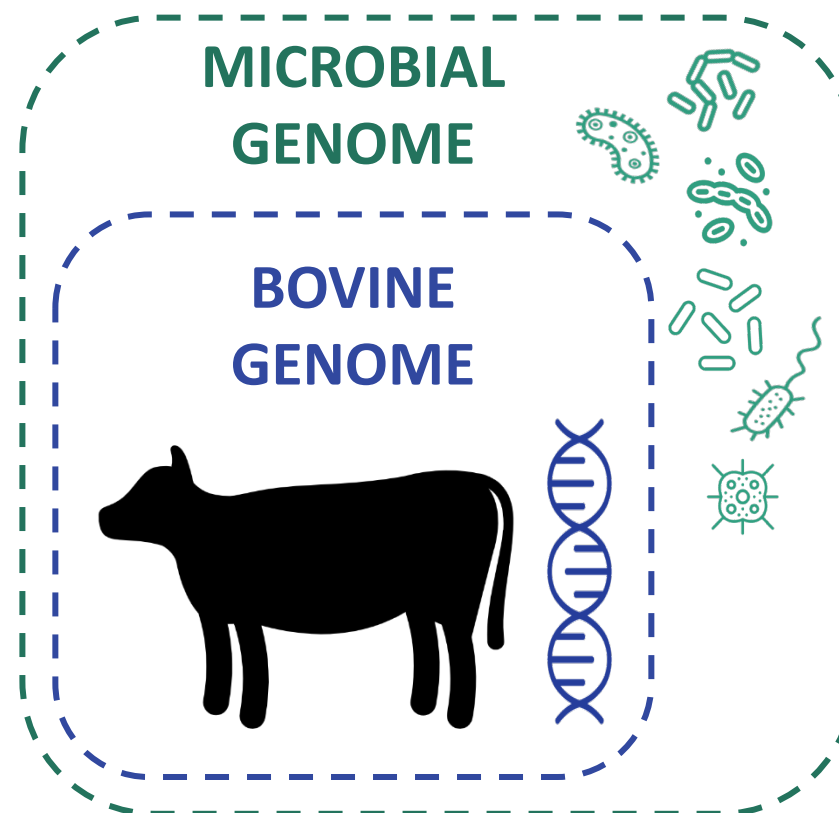


Metagenomics can provide insight into metabolic efficiency



We need solutions to mass-produce this information

- buccal swabs & pooling
- representative environmental sampling
- high covariance with other traits





- **How is the rumen microbiome related to:**
  - Heifer growth and feed efficiency?
  - Lactating cow feed efficiency?
  - Milk yield & quality?
  - Enteric methane emissions?  
-6% ingested energy
- **Do low efficiency heifers necessarily turn into low efficiency cows?**
- **Can cows be well-adapted to both high efficiency and high forage diets?**

# Stakeholder Input Summary

## Dairy Research Priorities

- **Milk Spectral Data**
  - Add new measures (i.e., P, minerals, DMI, pregnancy)
  - currently done in research settings; has not been widely applied yet
- **Herd Management Tools**
  - Review/optimize methods and prediction factors used for estimating lactation yields
- **Sire Fertility**
  - Improved genomic prediction/evaluations for bulls
- **High-throughput Data Systems**
  - Evaluate the use of metabolic activity indicators for selection purposes
  - Incorporate high throughput phenotypes available @ commercial dairies into national evaluation systems
  - Assess the use of high throughput phenotypes as proxy for traits difficult or costly to collect
  - Explore application of deep learning techniques for decision-making tools
- **Animal Welfare & Sustainability**
  - Identify & develop new traits for animal welfare & sustainability
  - Quantify their long-term economic impact on dairy breeding goals
  - Develop long-term selection and mating strategies to maintain diversity
- **Major Theme: Data Availability & Reporting**
  - a lot of data is available; what will it take to get it flowing
  - we need a path for data reporting and standardizing
  - we need better communication with the data providers on what the return on their investment is

National Program 101 ABBL/AGIL  
Stakeholder Meeting 2021

Breakout session report



- **Milking Speed**  
**21 reports of MSPD**  
**from 2006-2010**

## Dairy Research Priorities

- **DNA Sequencing**
  - Assess the impact of low pass sequencing on genomic predictions
  - Facilitate the identification of functional genes and pathways for traits of economic importance
- **Quantitative Genetic Tools**
  - Develop methods & tools to monitor economic weights of traits to select for
  - Assess the impact of predicting retained heterosis in crossbred animals
  - Assess the impact of genotype by environment interactions on selection programs (e.g., heat stress)
  - Investigate techniques to manage increasing volumes of data and improving accuracy of prediction

# Thank you. Questions?



Contact: [asha.miles@usda.gov](mailto:asha.miles@usda.gov)

<https://www.ars.usda.gov/northeast-area/beltsville-md-barc/beltsville-agricultural-research-center/agil/>