### Development, implementation, and future perspectives of health evaluations in the U.S.

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### DEVELOPMENT



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## Changes in emphasis over time

### **1971 PD\$**



COUNCIL ON DAIRY CATTLE BREEDING

### 2018 NM\$



<sup>™</sup> Milk Fat Protein PL SCS 🕷 HTH\$ DPR CA\$ **HCR** CCR **BWC** UC **FLC** 

### Since the 1980s

- Evidence that selection for health events could be successful
  - E.g., Scandinavian countries direct recording of health events
- Within U.S. calls for a unified system of reporting health events
  - Possibility for improvement through selection
  - Since 1994 Indirect selection through traits SCS and PL, and later LIV

 Introduction of genomics in 2009 – feasible to select for lowly heritable traits that are expensive and/or difficult to measure





### U.S. hurdles

- No mandated reporting system
- Need a single repository to collect and store data
- No unified way to record health events
  - Standardization critical





(https://www.thesun.co.uk/news/3420620/showjumpingcow-jumps-hurdles-pictures/)

### Data flow

- Cooperation from the Dairy Records **Processing Centers** 
  - Flow through DHI system
- Necessary standardization performed by DRPCs •











### Format 6

### Includes 20 health event codes + 4 management codes

				Health Event Segments (up to 20 segments)			
Health Event Segment Block (# 1)							
4	AAAA	СН	170	Health event code			
8	XXXX	СН		Health event date (YYYYMMDD)			
1	Α	СН		Health event date type (A = actual; $E = estimated$ )			
6	AAAA	СН		Health event detail			
19	AAAA	CH		Health event segment block # 2			
19	AAAA	CH		Health event segment block # 3			
19	AAAA	CH		Health event segment block # 4			
	8 1 6 19 19	8    XXXX      1    A      6    AAAA      19    AAAA      19    AAAA	4    AAAA    CH      8    XXXX    CH      1    A    CH      6    AAAA    CH      19    AAAA    CH      19    AAAA    CH	4    AAAA    CH    170      8    XXXX    CH			





### INPLEMENTATION







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## Health trait implementation

- April 2018: Official genomic evaluations for 6 direct health traits from CDCB for Holstein
  - Milk fever (MFEV)
  - Displaced abomasum (DA)
  - Ketosis (KETO)
  - Mastitis (MAST)
  - Metritis (METR)



- August 2018: Inclusion of health trait sub-index (HTH\$) in net merit indices (NM\$, FM\$, CM\$, GM\$)
  - 2.3% total emphasis within NM\$



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## Data processing

- Two levels of editing at CDCB
  - <u>General edits</u> date checks, parent checks, herd checks, etc.
  - <u>Constraints to be included for genetic evaluation</u> parities
    1 to 5, Holstein (currently), minimum/maximum incidence

restrictions, etc.



## Phenotypes used for evaluation

	Number of Records	Number of Cows		
Milk fever	1.2 M	0.7 M		
Displaced abomasum	1.9 M	1.1 M		
Ketosis	1.4 M	0.8 M		
Mastitis	2.4 M	1.4 M		
Metritis	2.0 M	1.1 M		
Retained placenta	2.2 M	1.3 M		

\*As of April 2019 evaluation





### **Evaluation models**

- Single-trait linear animal repeatability models
- Additional details available

	Heritability (observed)
Milk fever	0.6%
Displaced abomasum	1.1%
Ketosis	1.2%
Mastitis	3.1%
Metritis	1.4%
Retained placenta	1.0%



### https://www.uscdcb.com/



### **CDCB Health Traits**

As of August 2018, Net Merit \$ includes the six health traits launched in April.





### Cost considerations

- Direct costs of each event used in development of HTH\$
  - Considers veterinary and treatr
  - Excludes costs that are accounted for by other traits in NM\$ (e.g., declines in fertility, decreased production)
- Yield traits designated as "sick" test days are adjusted
  - These test days are accounted for with an additional adjustment (in

parentheses above)



	Event	Direct cost
	MFEV	\$34 (38 – 4)
	DA	\$197 (178 + 1
	KETO	\$28 (28 + 0)
	MAST	\$75 (72 + 3)
	METR	\$112 (105 + 7
ment costs	RETP	\$68 (64 + 4)



## Variance adjustments

- Linear model used with binary trait
- Phenotypic pre-adjustments applied to all health events
  - Phenotypes are adjusted based on calving year, parity, and heritability of the trait prior to genetic evaluation
- Similar to methodology described by Wiggans and VanRaden, 1992 and the adjustment applied to livability
- Implemented April 2019





## Variance adjustments

- Most health traits had PTA correlations ranging from 0.92 to 0.98 for bulls with > 70% reliability born since 2000
  - Exception milk fever
- For all traits first lactation trends agreed with the new trends more closely than with the old trends.





### Interbull validation

- trait group
- Validation of genetic trends •
- Only see on average a 1 point increase in reliability
- also have genotypes available in the US



### MAST now sent along with SCS PTA to Interbull for Udder Health

Minimal foreign bulls from countries supplying MAST directly that

## FUTURE PERSPECTIVES





### Future developments

- Health evaluations for Jersey
  - Genomic evaluations for the 6 health traits

on average







### Reliability approximately 10-15 points lower than Holstein

### See L. Jensen's talk – ADSA Tuesday 10:30 AM Room

### Future developments

- Multiple trait evaluations
  - Approximate genetic correlations
  - Mastitis & SCS
  - Groups of traits metabolic, reproductive?

	Protein	PL	LIV	SCS	DPR	CCR	HCR
MFEV	-0.21*	-0.10	0.08	-0.02	-0.07	-0.08	-0.01
DA	0.15	0.40*	0.41*	-0.14	0.30*	0.30*	0.12
KETO	0.20*	0.39*	0.31*	-0.25*	0.41*	0.39*	0.19*
MAST	0.06	0.52*	0.39*	-0.68*	0.32*	0.31*	0.10*
METR	0.27*	0.47*	0.33*	-0.21*	0.44*	0.45*	0.29*
RETP	0.02	0.21*	0.16*	-0.13	0.19*	0.19*	0.19*







### Potential health traits

- - Lameness or locomotion
    - conformation, metabolic, infection
    - How to differentiate between these?
  - Johne's



### Continued investigation on economically important health traits

# Events represent a variety of reasons for lameness – injury,



(https://vetextension.wsu.edu/researchprojects/lameness/)

### Potential health traits

- Calf health & calf termination
  - Dairy calf death losses estimated at \$327.3 million in 2015 (Lombard et al., 2019)
  - Possible to include calf/heifer health records with Format 6 •
  - Lombard et al., 2019 proposed death loss categorization scheme •
    - Pursuing Data Quality group of CDCB working with this scheme and termination reasons already collected by CDCB
    - Goal: expand termination codes to include calves/heifers





(https://hoards.com)

### Maintenance of data pipelines

- Expand current pipelines to capture additional informationMonitor data being submitted, accepted, and rejected
- Two-way communication with data providers
- Updates to standardization "dictionaries" as needed



/ What We Can Do For You / Service Documentation / Error Documentation

### Number of Health Event Segments Errors



### Code Description

- Number of health event segments does not agree with length of 9Ab record. Length of record corrected
- Cow already has 50 health events. New event is ignored. 9Ac

Action Returned Data Change

Reject Event date

Updated 08/22/2007

01/17/2008

Error Codes Complete Error Lists CSV/Excel Tab Separated

- 0 General Record
- 1 Animal Identification
- 2 Sire Identification
- 3 Dam Identification



## New functional traits

- Feed efficiency
  - Project funded by Foundation for Food and Agriculture Research (FFAR) and CDCB
  - Institutions include Michigan State University, University of Wisconsin, Iowa State University, University of Florida, and USDA Animal Genomics and Improvement Laboratory
  - Continuing the work of USDA NIFA grant
  - Projected that breeding for more efficient dairy cows could save the U.S. dairy industry \$540 million per year
  - Inclusion of feed efficiency in Net Merit \$





## Creation of data pipelines

- New data types
  - E.g., feed intake data, sensor data
    - Different systems at various institutions
    - Protocol needs to be developed to streamline data processing
    - Need for standardization







### **Evaluation sources**

- - Published methodologies •
    - Health \$ (CDCB)
    - Clarifide Plus (Zoetis)
  - Proprietary evaluations / indices
    - TransitionRight index (ABS)
    - Better Life Health index (CRV)
    - Ideal Commercial Cow index (Genex)



### Increasing number of similar evaluations from different sources



## Differing results

- Traits with limited data + low heritabilities
  - Different populations
  - Different editing
  - Different statistical model
  - Different presentation
  - Different economic assumptions



## Handling multiple sources

- Producers have to consider the source of information Critical to not focus selection on only a few traits
- What does the future hold?







## Continued progress

- More data available than ever before
  - Make better selection decisions
    - Phenotypes are critical
    - Quality control standards
    - Unbiased science and research
    - Establishment and maintenance of data pipelines

### Communication and Cooperation









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CDCB

AGIL







### Thank You!

