

# GENETIC & GENOMIC EVALUATIONS OF QUANTITATIVE MILKING SPEED PHENOTYPES

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# PROPOSED RESEARCH

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- OBJ. 1:** Assemble a high-resolution dataset pertinent to MS representing different dairy breeds, equipment manufacturers, parlor types, and milking management strategies
- OBJ. 2:** Characterize MS for herds grouped by equipment manufacturer and parlor type and assess the impact of additional system effects on the phenotype
- OBJ. 3:** Characterize any biological effects that impact MS, especially concerning udder health
- OBJ. 4:** Standardize MS trait definition and estimate heritability to determine its suitability for selection

# AVAILABLE DATA

## Demographics

~300 herds  
>230,000 cows  
>300,000 lactations  
>40 million observations

31 States  
6+ Breeds  
11 OEMs

DeLaval	80
GEA	75
Lely	47
Boumatic	46
AfiMilk	45
SCR	13
DairyMaster	10
AIC Waikato	5
AMS Galaxy	3
Jantec	2
Universal	2



# PRELIMINARY PTAS

## Different Trait Definitions

1. **Average MS (lbs/min) over all available data**
  - a) *Fixed effects: breed, parity, lactation length, OEM*
  - b) *n = 20,000 cows with complete lactations (1 year)*



## PRELIMINARY RESULTS

$$h^2 = 0.37$$

### Genetic Correlations

SCS	0.39
Milk Yield	0.14
NM\$	0.08

Mean REL	0.67
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# PRELIMINARY PTAS

## Different Trait Definitions

1. Average MS (lbs/min) over all available data

Trait	<i>PTA</i>				<i>REL</i>			
	Min	Mean	SD	Max	Min	Mean	SD	Max
<b>MSPD</b>	<b>-0.80</b>	<b>0.12</b>	<b>0.30</b>	<b>1.00</b>	<b>50.10</b>	<b>67.05</b>	<b>11.84</b>	<b>97.80</b>
<b>SCS</b>	<b>-0.72</b>	<b>-0.17</b>	<b>0.18</b>	<b>0.67</b>	<b>50.00</b>	<b>92.95</b>	<b>10.50</b>	<b>99.90</b>

**\*Detailed results for 772 HO bulls**

# PRELIMINARY PTAS

## Different Trait Definitions

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  - b) *n = 20,000 cows with complete lactations (1 year)*
2. Average MS (lbs/min) from test-days only
3. Primiparous cows only



***A hypothetical 3X cow would have  $3 * 305 = 915$  phenotype records***

# PRELIMINARY PTAS

## Different Trait Definitions

1. Average MS (lbs/min) over all available data
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***A hypothetical 3X cow would have  $3 * 10 = 30$  phenotype records***

***(97% reduction in data!)***

# PRELIMINARY PTAS

## Different Trait Definitions

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2. **Average MS (lbs/min) from test-days only**
3. Primiparous cows only



## PRELIMINARY RESULTS

$$h^2 = 0.28$$

### Genetic Correlations

SCS	0.43
Milk Yield	0.16
NM\$	0.06

Mean REL	0.64
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# PRELIMINARY PTAS

**Genetic Correlations (upper diagonal)**  
**Phenotypic Correlations (lower diagonal)**

	Avg_all	Avg_TD	Avg_all_P1	Avg_TD_P1
Avg_all		0.968	0.916	0.976
Avg_TD	0.821		0.944	0.991
Avg_all_P1	1.000	0.819		0.924
Avg_TD_P1	0.820	1.000	0.819	

# ENSURING DATA FLOW

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## Minimum Required Data Novel to MSPD

Observation date (YYYYMMDD)

Milking Session Number (1, 2, ... 6)

Milking Frequency (01X, 02X, 03X, 04X, AMS)

Robotic or Manual Attachment (R or M)

**Original Equipment Manufacturer (OEM) Code**

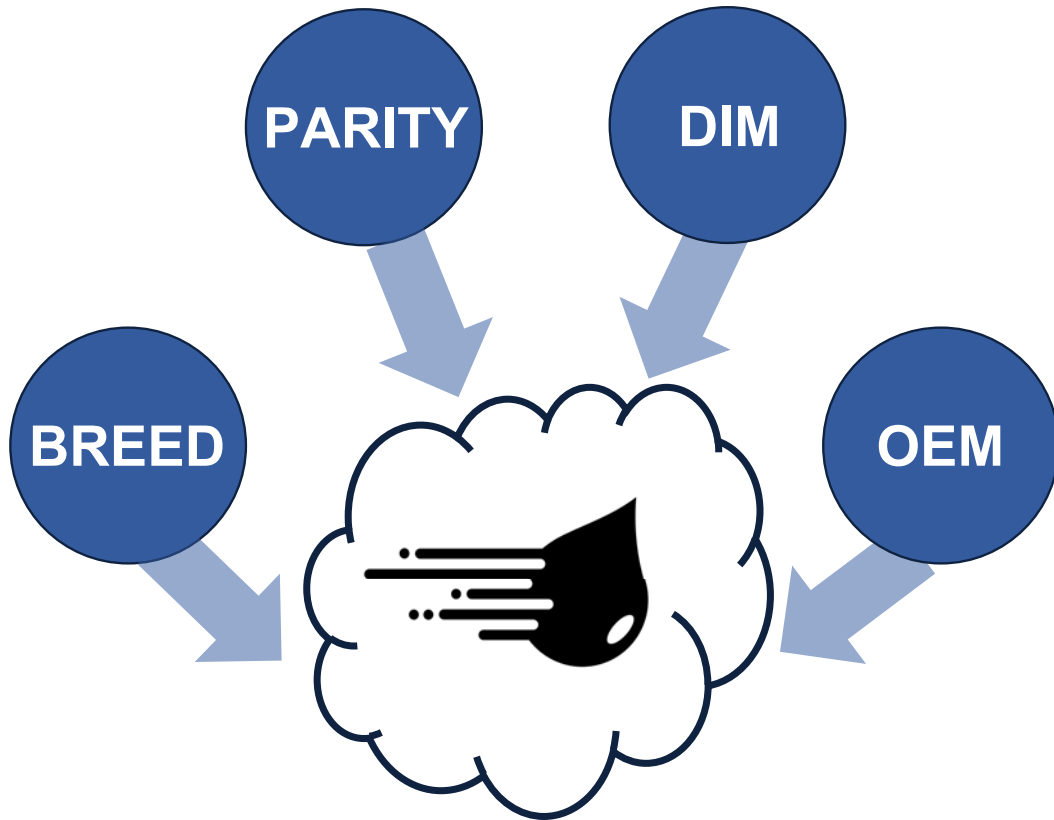
Milk Yield from Individual Milking (lbs \* 10)

Milking Duration of Individual Milking (minutes \* 10)

Abnormal Flags (Y or N)



**ICAR Device  
Reference IDs**



**Many factors influence  
quantitative MSPD measurements**

## THE BOTTOM LINE

- Genetic and genomic prediction methodology for milking speed has been developed
- We are targeting delivery of a new trait in December 2024
- Routine data flow is a key hurdle
- Next steps include incorporation of partial lactation records and exploring the use of AMS data

# THANK YOU

Data were available to the authors from CDCB under USDA Agricultural Research Service Material Transfer Research Agreement #58-8042-8-007. While CDCB offers data stewardship, sole ownership and rights pertaining thereto remain with the producer and we thank U.S. dairy producers for sharing their data for research use.

This work was supported by USDA-ARS project 8042-31000-113-000D, “Improving Dairy Animals by Increasing Accuracy of Genomic Prediction, Evaluating New Traits, and Redefining Selection Goals”.

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