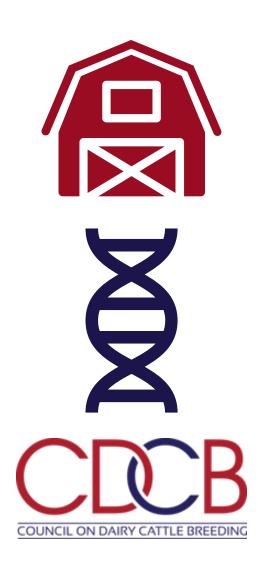
## CDCB Evaluations

Courtney K Hoff, Data Support Specialist
2024 CDCB Genomic Nominators and Laboratories Workshop
Tuesday, July 30<sup>th</sup> 2024



#### Overview

- What is an evaluation?
- Qualifying for an evaluation
- Three CDCB Evaluation Types
- Checking to see if an animal qualifies





#### What is an evaluation?

- An evaluation is a report that contains traits and indices used for breeding selection for an animal
- Traditional evaluations utilize pedigree and phenotypic data (production, composite/type)
- Genomic evaluations utilize pedigree, phenotypic, and genotypic data



#### What is an evaluation?

- CDCB genomic evaluations are expressed as GPTAs
  - Genomic Predicted Transmitting Ability
- We evaluate for ~50 individual traits and 4 selection indices Net Merit (NM\$), Cheese Merit (LCM\$),
   Fluid Merit (LFM\$), and Grazing Merit (LGM\$)



#### What is an evaluation?

CDCB produces genomic evaluations for 5 dairy breeds:

- Ayrshire (AY)
   Guernsey (GU)
   Jersey (JE)

- Brown Swiss (BS)
   Holstein (HO)
- CDCB also produces genomic crossbred (XX) evaluations for mixed breed animals thanks to Breed Base Representation (BBR) calculations



## Qualifying for an evaluation:

#### Genomic Evaluations



The animal's genotype must Nomination must be Animal receives preliminary, pass a robust process to successful with a CDCB fee unofficial weekly genomic determine which genotypes evaluation code assigned for the animal. qualify for evaluation

- 1. Successful nomination 2. Genotype loaded and and fee code assignment
- usable
- 3. Animal gains a Use Weekly Timestamp



#### **Females**

Code	Fee amount	Description
1	\$0.00	US birth & yield & health & type records
2	\$1.00	US birth & yield & (health or type records)
3	\$2.00	US birth & yield records
4	\$3.00	US birth & (type or health records) (health records are probably not possible w/o yield records)
5	\$4.00	US birth
6	\$6.00	CAN birth
7	\$8.00	No exemptions
8	\$1.00	Parentage only

#### Males

Code	Fee amount	Description
Р	\$0.00	International exchange partner (code P)
S	\$25.00	Controller is a collaborator stud
1	\$25.00	US herd providing yield & health & type
2	\$50.00	US herd providing yield and (health or type)
3	\$75.00	US herd providing yield
4	\$75.00	US herd providing type or health
5	\$100.00	US herd
7	\$200.00	No exemptions
8	\$1.00	Parentage only



- Fee Code Assignments: 1–7
- •1–5 = US Fee codes, assigned based on performance data contribution attached to DHIA herd codes
- 6 = Animals with Canadian (CAN) IDs only
- 7 = All other foreign animals



- Fee Code Assignments: Other
- 8 = Parentage Verification Only (PVR)
- S = Collaborator Stud Fee, can only be used for bulls owned by collaborator studs





- Fee Code Assignments: N
- Animal will not receive an evaluation if they are assigned fee code N
- Previously used for genotypes exchanged with Lactanet –
   a topic to be covered in Day 2 of the Workshop



- CDCB will reassign a fee code N on a nomination if:
  - · A fee code wasn't assigned in the nomination
  - If an animal has a sex conflict between the pedigree and genotype after genotype loading



## Qualifying for an evaluation: Usable Genotype

- A Usable genotype (Use=Y) is a genotype that:
- 1. Has passed quality control checks
- 2. Does not have a sex conflict with the animal's pedigree
- 3. Does not have genomic conflicts with close relatives that make the animal's pedigree appear unreliable



## Qualifying for an evaluation: Use Weekly

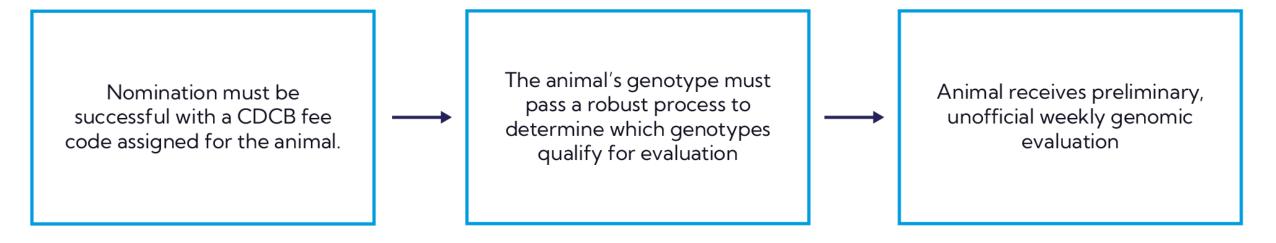
- Usable Genotype ≠ Use Weekly Timestamp!
- Genotype looks correctly assigned, but there are some discrepancies in its pedigree data
- Two common cases:
  - Maternal grandsire (MGS) unlikely
  - · Animal's BBR results do not agree with the animal's ID breed



### Qualifying for an evaluation:

#### Genomic Evaluations





After qualifying for a weekly evaluation a *female* animal will continue receiving genomic evaluations as long as it has a usable genotype



### Qualifying for an evaluation: Female Animals

- A Use Weekly Timestamp has been applied
- At least one usable genotype
- Animal must meet *all* of the criteria before evaluation deadlines!
- Requester on nomination should match requester of genotype (unless a share file for results has been submitted)



### Qualifying for an evaluation: Male Animals

- Must meet the female qualifications, + additional requirements revolving around NAAB status and domestic vs foreign
- Easier to explain what would prevent an evaluation:
  - Foreign bull\* over 16 months of age without an AI service fee paid
  - An active AI bull (only receive triannual evaluations)
  - Some additional rules based on bull breed and ownership
  - \*Foreign bulls do not have *public* evaluations unless they are in Al



#### CDCB Evaluations

- Weekly: Preliminary genomic evaluations released every Tuesday
- Monthly: Genomic evaluations released on the first Tuesday of a month
- Triannual: Genomic and traditional evaluations released three times a year in April, August, and December



#### CDCB Evaluations: Triannual

- Incorporate updates to trait and index calculations
- Inclusion of new phenotypic data
- Have an earlier cutoff date than monthly evaluations to allow inclusion of traditional evaluations and evaluations from Interbull
- Triannual evaluations are public and published via the SFTP\* and WebConnect
  - \*SFTP files are password protected, unlike monthly and weekly

#### CDCB Evaluations: Triannual

- Reminders for bulls:
  - Evaluations for bulls in AI service released in the triannual
  - Bulls enrolled in NAAB (status other than C) released only in triannual evaluations
  - Foreign bulls that are not in AI do not have public evaluations



## CDCB Evaluations: Monthly

- · SNP effects are updated each month
- New phenotypic data is not applied to monthlies
- · All animals (except for some bulls) receive updated evaluations each month
- Qualify as official evaluations



# CDCB Evaluations: Weekly

- Simplified evaluations
- Preliminary because new SNP effects are not applied and BBR is not included
- Intended to be used for quick herd management decisions: culling and breeding
- Not meant to be used for marketing animals or semen

## CDCB Evaluations: Weekly

- Includes all usable genotypes that have a Use Weekly
   Timestamp from 6PM EST of the preceding Sunday
- · Limited to one weekly evaluation per animal
- After inclusion in the first weekly extraction the fee code cannot be reverted to N and genotype can no longer be withdrawn



# Checking to see if an animal qualifies

Queries Data Exchange 

Special Section 

Top Animal Listing Summary Stats 

National Performance Metrics **QUERIES** Animal ID (17 bytes)/ Animal ID + Sex Code (18 bytes) HO840003282837717 **Run Query** Clear



https://webconnect.uscdcb.com/#/queries

Q Search

Requested ID Searches

HO840003282837717

#### HO840003282837717

Blend Code:

ID/Pedigree Evaluation Genotype Progeny Errors Lactations Evaluation Breed: HO - Holstein Sire -Animal ID: HO840003218556317 Preferred ID: SIEMERS RZ PAZZLE 34954-HO840003282837717 Name: COLDSPRINGS JEWEL Name: Maternal Grandsire 2021-02-09 14624-ET DOB: Source Code: B DOB: 2024-03-07 Genotyped: Yes ID: HOUSA000074024998 Sex: F Name: PINE-TREE SPLENDID-P-ET Multi-Birth 3 DOB: 2015-08-05 Code: Dam Source Code: B **ID Source** В Genotyped: Yes Code: ID: HO840003149596913 <u>Pedigree</u> **COLDSPRINGS SPLENDID** Name: В Source Code: 10898

2018-09-07

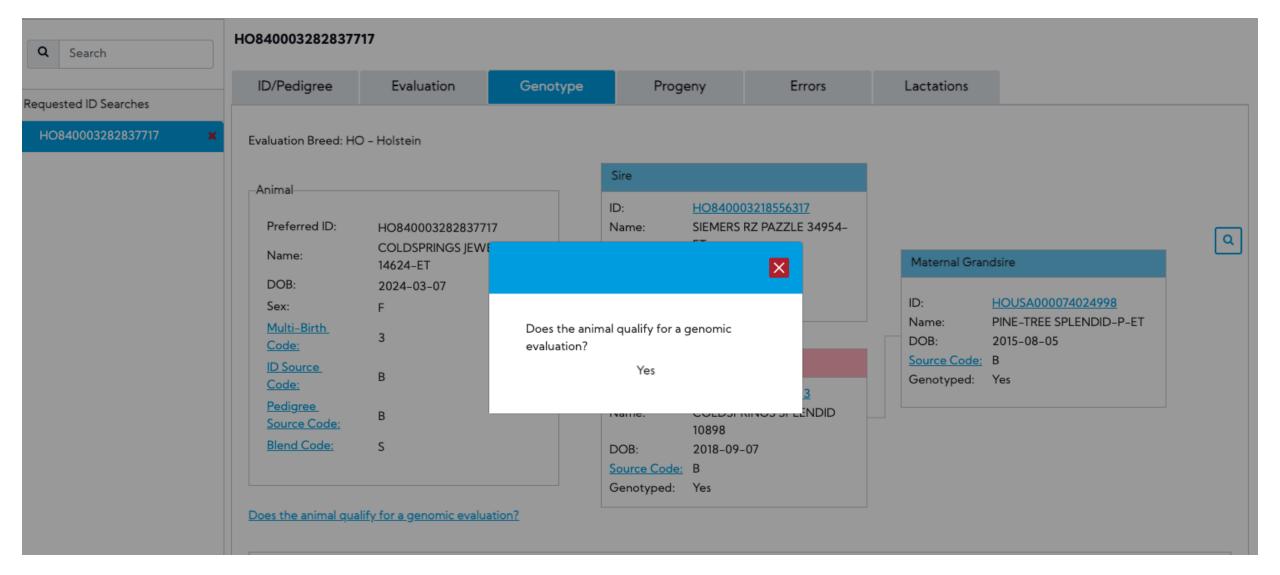
DOB:

Source Code: B
Genotyped: Yes

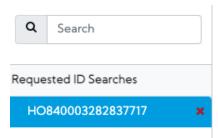


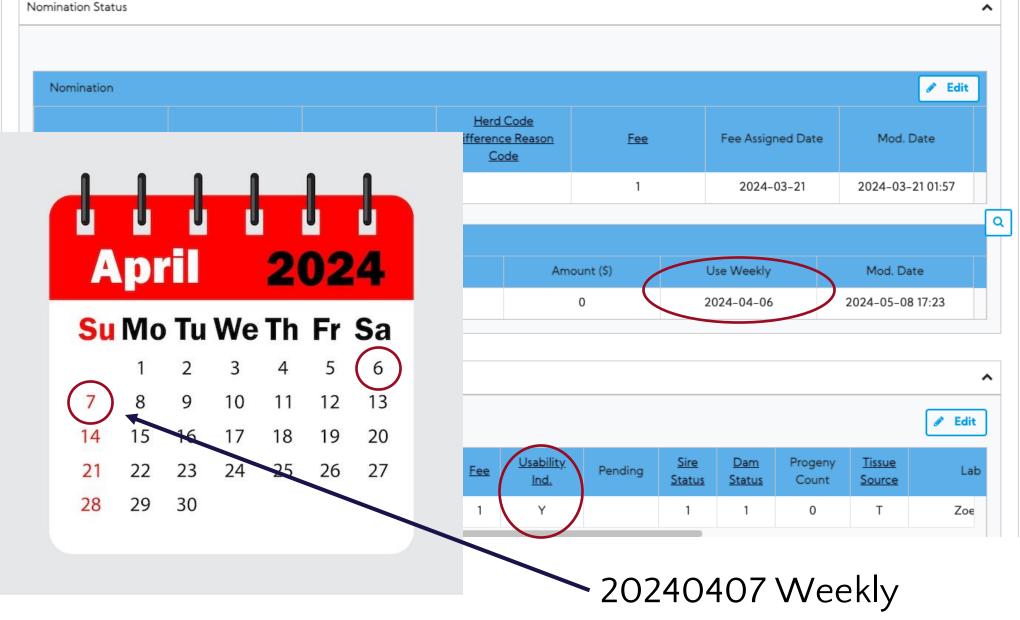
S















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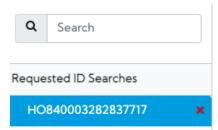
#### Q

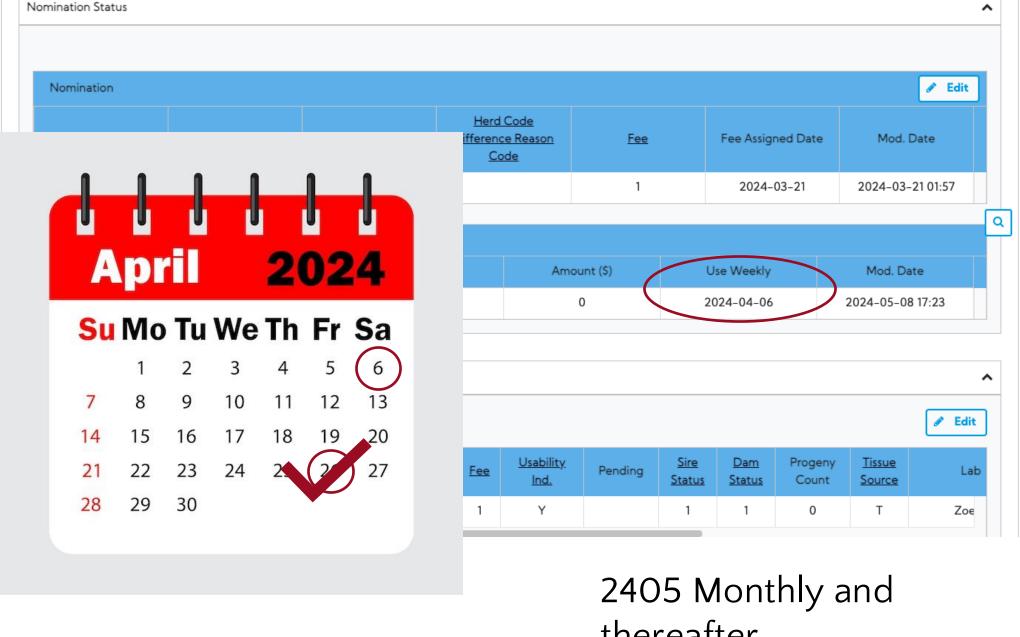
#### May 2024 genomic evaluations

April 22, 2024	International genotype exchange data due (for CDDR exchange partners)
April 26, 2024	Genomic data due
May 7, 2024	Evaluation release (8:00am ET)

https://uscdcb.com/genetic-evaluation-schedule/2024/

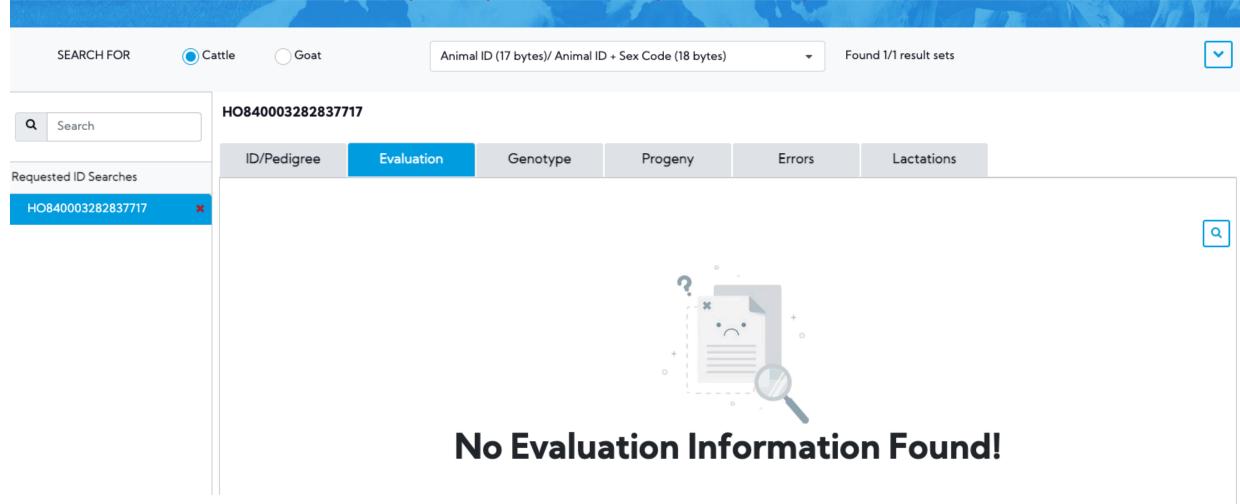








thereafter



If animal has a usable genotype on the 2408 deadline, a 2408 triannual evaluation should be published on WebConnect on 2408 release date



#### HO840003218556317

Evaluation ID/Pedigree Genotype Progeny **Errors** Lactations Run: April 2024 or History Evaluation Breed: HO - Holstein Animal Sire Preferred ID: HO840003218556317 ID: HO840003143986308 Long Name: SIEMERS RZ PAZZLE 34954-ET SIEMERS RENEGADE ROZLINE-ET Name: Short Name: **PAZZLE** DOB: 2019-04-15 NAAB DOB: 2021-02-09 Source Code: B Primary NAAB Code: 250HO16115 Genotyped: Yes Sex: Μ **Current Status:** G Multi-Birth Code: 3 2022-04 Entered Al: Recessive Codes: TP TL TD TR TC TY TV TE Control Stud: 250 Cross References: No Dam Original Stud: 0 Pedigree Comp %: 100 Sampling Status: Genomic Indicator: ID: HO840003141495692 Name: SIEMERS LMDA PARIS 27856-ET Herd with most Daughters DOB: 2017-07-14 Herd: 0 Source Code: B Number of Daughters: 0 Genotyped: Yes Merit Table and Inbreeding ^ Genotyped: Yes

History

**≛** Export CSV

^

Eval Date	NM\$		MACE	Gen Ind	DAUS	HERDS	%US									
Eval Date	PTA	REL	MACE	Gen ind	DAUS	HERD3	/ <sub>0</sub> U3	Mlk	Fat	Pro	PL	SCS	DPR	HCR	CCR	
2024-04	851	77%		1	0	0	100%	547	95	48	4.4	2.75	0.0	1.8	1.2	
2023-12	867	76%		1	0	0	100%	605	92	47	4.8	2.76	0.4	1.9	1.7	
2023-08	849	76%		1	0	0	100%	609	89	46	4.7	2.75	0.7	1.9	1.9	
2023-04	861	74%		1	0	0	100%	574	90	47	4.8	2.76	0.6	1.6	1.5	
2022-12	838	74%		1	0	0	100%	546	89	43	4.6	2.75	0.5	1.4	1.3	
2022-08	888	73%		1	0	0	100%	614	93	46	4.9	2.73	0.4	1.5	1.3	
2022-04	884	73%		1	0	0	100%	647	95	48	4.5	2.73	0.4	1.7	1.4	
2021–12	822	73%		1	0	0	100%	539	90	43	4.5	2.73	0.5	1.3	1.2	
2021-08	840	72%		1	0	0	100%	505	92	41	4.7	2.72	0.6	1.2	1.4	
2021-04	922	74%		1	0	0	100%	631	96	44	4.6	2.72	0.7	1.2	1.3	

<< 1 > >>

1 - 10 of 10 items

#### Links for evaluation related content

- General evaluation information: <a href="https://uscdcb.com/services/">https://uscdcb.com/services/</a>
- Genomic fee codes: <a href="https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/REFERENCES#Ref164">https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/REFERENCES#Ref164</a>
- 2024 evaluation deadlines: <a href="https://uscdcb.com/genetic-evaluation-schedule/2024/">https://uscdcb.com/genetic-evaluation-schedule/2024/</a>
- Reasons that fee code would be reassigned to N by CDCB: <a href="https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/Reasons\_that\_the\_fee\_code\_would\_be\_reassigned\_to\_N\_by\_CDCB">https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/Reasons\_that\_the\_fee\_code\_would\_be\_reassigned\_to\_N\_by\_CDCB</a>
- Reasons that a genomic evaluation was not released: <a href="https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/Reasons\_that\_a\_genomic\_evaluation\_was\_not\_released">https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/Reasons\_that\_a\_genomic\_evaluation\_was\_not\_released</a>
- DOB limitations for foreign bulls: <a href="https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/DOB\_limitation\_for\_foreign\_bulls">https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/DOB\_limitation\_for\_foreign\_bulls</a>
- Genomic evaluation distribution formats: <a href="https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/Genomic\_evaluations\_format\_(CSV)">https://redmine.uscdcb.com/projects/cdcb-customer-service/wiki/Genomic\_evaluations\_format\_(CSV)</a>



# Thank you for your attention!

