

Council on Dairy Cattle Breeding CDCB future developments

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Topics

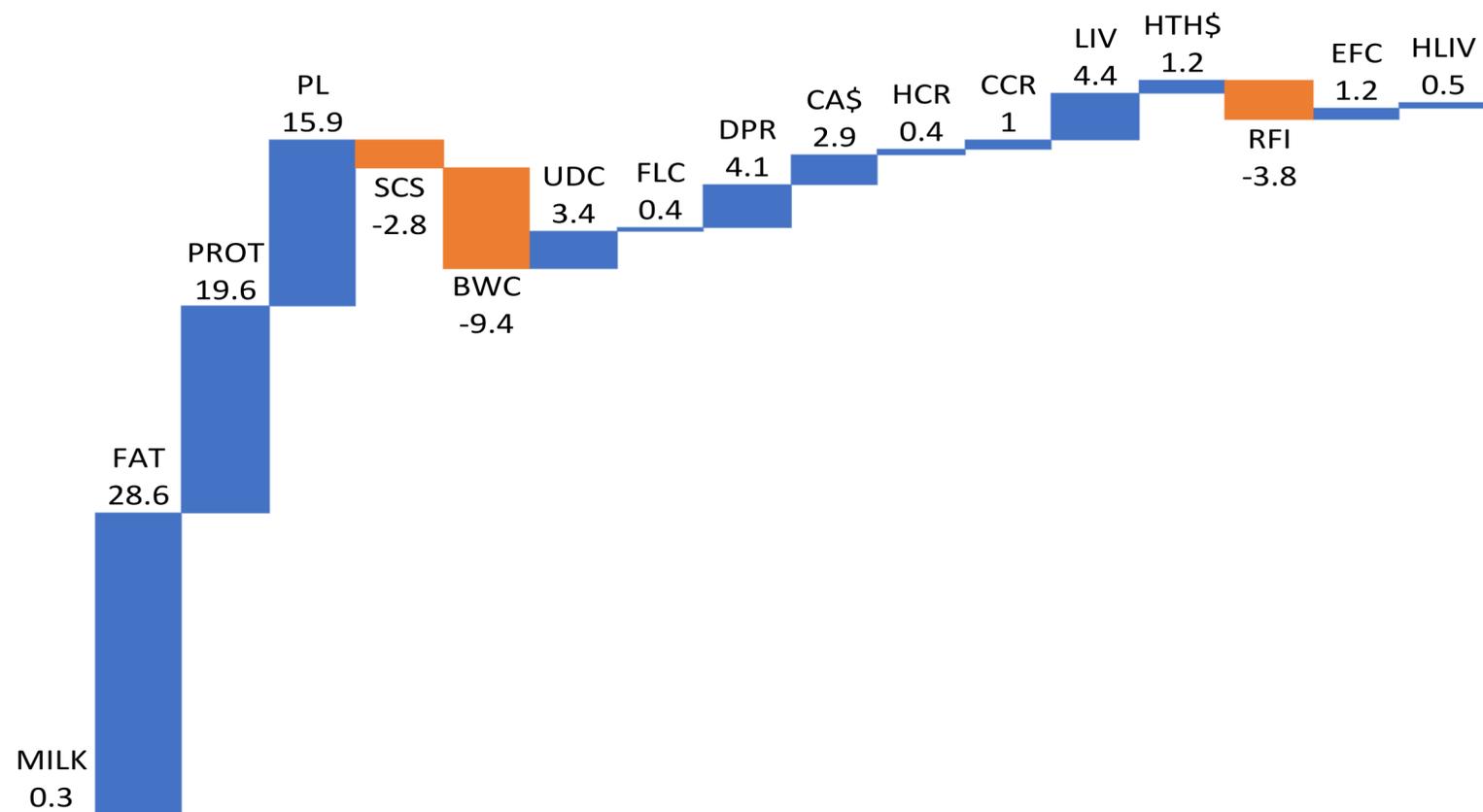
- Net merit update (&co.)
 - New trait being published (RFI)
- Slightly modified evaluation SNP set (+ yearly usability update)
- Genomic weighting
- Heterosis enhancement
- Jersey recessives
- Ancestor discovery and virtual dam project (evaluation perspective)
- Weekly breed ID updates
- Weekly/Monthly processing update
- Publication and distribution rules review

Net merit update

- New NM\$, same goal
 - maximizing cows' lifetime profitability
- Most significant revision in a long time
 - Inclusion of Residual Feed Intake (RFI), Early First Calving (EFC) and Heifer Livability (HLV)
 - Revision of parameters used in existing traits

What changes?

Relative emphasis (%) on traits included in Net Merit - 2021 revision



Relative emphasis: *contribution of each trait when ranking the animal in the NM\$ scale (rel + genetic variation)*

Reduced RE	FAT	Revised estimated costs of milk components
	UDC	Gains in udder conformation reduced milking labor
	FLC	Not well correlated with hoof health or lameness
	CA\$	Progress made reduced phenotypic average and SD
	LIV	Death rates and cull cow prices declined
Increased RE	PL	Better accounts for maturity effects
	BWC	Maintenance costs higher than previously assumed

Overview and take-home messages (I)

- Body weight (BWC) receives much more RE than before
 - BWC estimates are now based on effective measurements feed costs for maintenance (parameters were estimated indirectly before)
- BWC includes multiple traits.
 - Strength is mostly associated with body weight and feed intake.
 - Dairy form (thin cows) will still be penalized because of upward selection of PL, fertility and hth\$, combined with a downward selection for size.

Bottom line: selection for more efficient and robust animals

Overview and take-home messages (II)

- Residual Feed Intake currently has lower reliability so its RE is low.
 - Evaluation validated using single/multi step approaches and 5-way cross-validation ([VanRaden et al., 2018](#); [VanRaden and Hutchison, 2018](#); [Li et al, 2020](#))
 - As (much) more data is collected, its RE will likely increase.
- Overall NM\$ reliability will drop slightly

Overview and take-home messages (III)

- NM\$ remains an index that maximizes lifetime profitability on all breeds
 - Market conditions, production costs and consumer trends
 - RFI is available for HO only
 - Other breeds receive 0 emphasis from RFI
 - Its economic weight distributed among the other traits
 - Not just RFI...

Webinar with more in-depth details

- Train the trainer session (technical session)

June 7th, 2021 at 2 pm ET

Target:
August 2021

Slightly modified evaluation SNP set

- Will “affect” usability on 18,000 SNPs in various chips.
 - Usability checks (yearly now)
 - Will likely affect a few animals in the database when reprocessed (clear/add errors)
- 100 SNPs difference in evaluation set (negligible impact expected)
 - JNS area (better results and prep for lab tests when available)
- New imputation options (skipping long-range haplotypes)
 - Accuracy only marginally lower (>97% matching – avg. 200/79,000 SNP difference in JE)
 - Impact on 3k SNP chip animals, negligible in all others.
- **Yearly** update of usability from now on

Target:
August 2021
(?)

Genomic weighting

- Research ongoing
- Expectation (first indications):
 - Recent research indicates that increasing genomic vs. traditional weighting improves accuracy of prediction (even if it increases slightly bias)
 - Changes will be depending on the difference of DGV vs TRAD and different by breed/trait group
 - Only minimal (+) effect on reliability.

Target:
August 2021

Heterosis enhancement

- Research ongoing
- Expectation (first indications):
 - Mostly a performance enhancement package (few hours saved)
 - Minor changes to logic for complex pedigrees
 - Heterosis in peculiar XX ancestors
 - Single-animal changes, but 0 impact on population

Jersey recessives

- Currently a standard process for Holstein
- Aim: test results from registered JE animals
 - Polled, etc..
 - JNS tests when available
- Better haplotype calling stability
- Talks with AJCA in progress

Ancestor discovery and virtual dam project (evaluation perspective)

- Including more information on pedigree connections will impact evaluations
 - Better parent averages
 - Better inbreeding/heterosis calculations
 - Less missing information (less use of “phantom groups”)

Virtual dam project (evaluation perspective)



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Discovering ancestors and connecting relatives in large genomic databases

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Table 2. Traditional and genomic EBV value means, SD, and reliabilities for yield traits of 295,136 animals with newly found ancestors, before and after pedigree completion

EBV	Trait	Incomplete pedigree			Complete pedigree		
		Mean (kg)	SD (kg)	Reliability (%)	Mean (kg)	SD (kg)	Reliability (%)
Traditional	Milk	1,948	720	26.6	2,064	811	32.6
	Fat	72.2	15.1	25.0	76.7	18.2	32.0
	Protein	59.4	14.6	26.9	63.0	17.0	32.9
Genomic	Milk	2,186	492	76.2	2,258	513	77.1
	Fat	74.9	19.7	76.0	77.6	20.5	76.9
	Protein	63.6	13.4	76.3	65.7	14.1	77.3

Target: after
August 2021

Weekly breed ID updates

- Similar to what is being done in monthlies
- Weeklies still affected
- Strategy identified
 - Only testing (e.g. time/resources) needed
 - Unlikely before end of August run

Target: after
August 2021

Weekly/Monthly processing update

- Mostly a performance enhancement package
 - Better management (less processing) of crossbred evaluations.
 - Implemented in April for weeklies (1/3 time cut)
 - No impact on evaluations
 - Likely implemented after August for monthlies (~ logic)

Target: after
August 2021

Publication and distribution rules review

- Stage one of a larger effort to increase frequency of full evaluations (6 times / year)
- **Steps :**
 - 1) Document all publication and distribution rules currently in place, for all files published and released in weeklies, monthlies and triannuals (30% done)
 - 2) Write a proposal of modification
 - 3) Discuss with "Evaluation frequency task force"
 - *repeat*

THANK YOU FOR YOUR ATTENTION

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