Standardised Labeling for Genetic Trait Coding

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on behalf of the WHFF Registration Working Group



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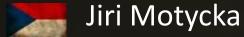
World Holstein Friesian Federation Objectives

- *Harmonise* technical and administrative matters with regard to the Holstein Breed
- *Represent* the common interest of breeders worldwide in developing and promoting the Holstein Breed
- Exchange Information on important issues concerning the breed
- Assist emerging herdbook associations
- Co-operate with Research Centres and other recognised International Organisations involved in animal improvement

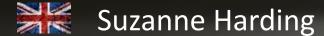


Members of WHFF Registration Working Group

















Members of WHFF Registration Working Group <u>Assigned Tasks</u>

- Harmonize a process for the categorization / recognition of major recessives.
- Standardize labeling nomenclature and codes tested carrier, tested free and not conclusive.
- Publishing haplotypes and standardize the way they are reported.



Why Standardise?

- Easy for all organisations to reference the same Genetic Traits
- Useful on farm when making breeding decisions
 - Farmers can choose to minimise the impact or increase the likelihood of the effect
- Recommended to report on official documents
- Harmonization of codes and nomenclature is imperative for overall accuracy and international data exchange.



Genetic Trait Names

Gene Name	Description	Gene and Expression Code
BLAD	Bovine Leukocyte Adhesion Deficiency (deficiency of a normally occurring protein needed for white blood cells or leukocytes, which are body's infection fighters)	BLC = tested carrier of BLAD BLF = tested non-carrier of BLAD
Mule foot	Mule-Foot (toes of foot are joined, giving animal a single hoof, instead of cloven ones)	MFC = tested carrier of Mule foot MFF = tested non-carrier of Mule foot
DUMPS	Deficiency of Uridine Monophosphate Synthase (one of many enzymes contributing to normal metabolic processes)	DPC = tested carrier of DUMPS DPF = tested non-carrier of DUMPS
CVM	Complex Vertebral Malformation (causes still-born calves, abortions, and early embryonic losses)	CVC = tested carrier of CVM CVF = tested non-carrier of CVM
Factor X1	Factor X1 (blood clotting disorder)	XIC = tested carrier of Factor X1 XIF = tested non-carrier of Factor X1
CIT	Citrullinemia (accumulation of ammonia and other toxics in blood in baby calves)	CNC = tested carrier of Citrullinemia CNF = tested non-carrier of Citrullinemia
	Brachyspina (causes abortion and	DVC - tosted corrier of Drachyonina



Genetic Trait Names

Coat Colour Carrier Gene	Description	Gene and Expression Code
Red	Red gene	RDC = carrier of red gene RDF = tested non-carrier of red gene
Red	Variant Red gene	VRR = not tested/determined by lineage VRS = tested true (homozygous) VRC = VRC =tested carrier (heterozygous) VRF = tested free
Black/Red	Black/red gene	BRC = carrier of black / red gene
Black	Black gene	BKC = carrier of black gene



Gene Test Differences

- Direct gene test:
 - o reliability: very close to 100%, excluding technical errors / issues
 - o are marker-based tests
 - o result from presence of mutated allele
- Indirect gene test:
 - o reliability: very high, can be as high as 98%
 - o risk of false positive / false negative results
 - does not detect causal allele; are looking for alleles in close proximity to causal nucleotide / genome variation



An Example of a direct test code

- Cholesterol Deficiency
 - Originally indirect test
 - Causal mutation found so direct test available
 - Naming can be completed



An Example of an indirect test code

• Haplotypes?

- Only indirect test available for HH2 Still looking for causal mutation
- HH1, HH3, HH4 now have direct tests
- HH5 recently detected



Expression Codes

- For many years WHFF has adopted two Alpha characters assigned for monogenetic inherited traits
- New proposal will facilitate the differentiation between direct and indirect traits
- Codes to be used following the WHFF two (alpha) characters
- No previously named traits will be renamed
- Naming of traits will continue to evolve as research continues



Expression Codes

• Codes to be used following the WHFF two (alpha) characters assigned for the monogenetic trait.

	Direct Tests		Indirect Tests
F	Tested Free	0	Tested Free/non-carrier.
С	Tested Carrier / Heterozygous	1	Tested Carrier/Heterozygous/Confirmed with pedigree info.
S	Tested / Homozygous	2	Tested True/Homozygous/Confirmed on both sides of pedigree.
		3	Additional Characteristics e.g. suspect carrier origin could not be confirmed from pedigree.
		4	Additional Characteristics e.g. suspect homozygous origin could not be confirmed from pedigree.
		5	As required should an additional characteristic be identified.



An Example of naming a direct test code

- Cholesterol Deficiency
- CDF tested non-carrier / free of cholesterol deficiency
- CDC tested carrier of cholesterol deficiency (heterozygous)
- CDS tested true carrier of cholesterol deficiency (homozygous)



An Example of naming an indirect test code

- Currently there are no common codings of indirect tests using the WHFF recommended nomenclature
- Each country has named using their own coding



Reporting Procedure

- Industry partner advises WHFF that there is a newly discovered Genetic Trait
- Fill in 'Request for information' form
 - Describe new genetic trait
 - What is the evidence
 - Who is reporting
 - O When was it first observed?
 - Which animal / family was it observed in
 - Additional information
 - Contact details
- Send back to WHFF (worldholstein@gmail.com)
- Four weeks later the WG will deliver the standardised label for coding.



BUT.....

- Practical problem of informing all herd books of genetic codes
- Proposal from WHFF President Jos Buiting to ICAR for better dissemination of Bulls genetic codes
- New procedure:
 - Every herd book sends genetic codes for all bulls to Interbull when send evaluation data
 - Interbull can then send this data back to members
 - Could link in with plans to add data to the Interbull data exchange
- Procedure standardized Internationally
- ICAR / Interbull considering proposal



Conclusions

- Important to promote nomenclature to scientific community
- Talk to ICAR with regard to proposing new Guidelines
- Encourage reporting of new genetic traits
- Communicate new genetic traits
- Harmonisation reduces farmer and industry confusion
- Farmer can choose to use when breeding
- WHFF proposal to ICAR for automatic data dissemination
- www.whff.info for full list of Holstein Genetic Traits



Thank you for your Attention.

Any Questions?

