

Standardised Labeling for Genetic Trait Coding

Suzanne Harding

on behalf of the WHFF Registration Working Group



Contents

- WHFF Registration Working group
- Why Standardise?
- Genetic Traits
- Gene Test Differences
- Expression Codes
- Reporting Procedure
- Conclusions



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



Linda Markle (*Chair*)



Jiri Motycka



Christa Kühn



Suzanne Harding



Peter Cole



Tom Lawlor



Liliana Chazo



Members of WHFF Registration Working Group

Assigned Tasks

- 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 | BVC = tested carrier of Brachyspina |



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 VRR = not tested/determined by lineage |
| Red | Variant Red gene | 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:
 - reliability: very close to 100%, excluding technical errors / issues
 - are marker-based tests
 - result from presence of mutated allele
- Indirect gene test:
 - reliability: very high, can be as high as 98%
 - 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. |
| C | 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
 - 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 ?

