

Martin-Luther-University Halle-Wittenberg Institute of Agricultural and Nutritional Sciences Animal Breeding



Health traits and lactation survival in relation to conformation traits in German Holstein Cows

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- conformation is a well recorded trait
 - it is still important to many breeders and breeding organizations
 - it corresponds to longevity
- health traits become more and more important
 - but health recording is a challenge within farms

The objectives:

- 1. analyze relationships between health traits and conformation traits
- 2. analyze differences in conformation traits comparing cows culled within the first lactation to cows reaching four or more lactations



Material and methods

- data originated from large contract herds in North-Eastern Germany
 - all cows are scored for conformation during first lactation (19 traits recorded on a 1 to 9 scale, except for stature, measured in cm)
 - health traits for all sorts of diseases were recorded under a supervised scheme
- data set I: 56,273 first lactation records with information on health and conformation,
 years of first calving from 2008 to 2015
- data set II: 8,272 cows culled within the first lactation and 24,933 cows that finished at least three lactations

> years of first calving from 2008 to 2013



two different health trait definitions:

- 1) number of new cases over all main groups (from 0 to n)
- 2) number of different groups of disease cases during lactation (from 0 to 4)
- main groups of diseases are: mastitis, claw disorders, reproductive disorders, metabolic disorders

- definition for longevity:
 - culled within first lactation (LS1) or survived third lactation (LS4+)



Inear mixed models were used for data analysis in SAS 9.4 :

 $y_{ijklmno} = \mu + HYSC_i + RANK_j + AFC_k + HT_l + \beta LL_m + s_n + e_{ijklmno}$

- *y_{ijklmno}* conformation trait
 - μ overall mean
- *HYSC*_i herd*year*season*classifier (i=1,2,...,2025)
- *RANK_j* rank for milk yield (j=1,2,3)
- AFC_k age at first calving (k=1,2,...,5)
- HT_l health trait (l=0,1,...,5)
- βLL_m regression coefficient for lactation length
 - *s_n* random sire effect
- *e*_{*ijklmno*} random residual effect

Health traits:

- 1. number of new cases
 - from 0 (=healthy) to 5+ across diseases
- 2. number of different groups of disease cases
 - from 0 to 3 or 4 different groups of diseases



Material and methods – Data set II

Inear mixed models were used for data analysis in SAS 9.4 :

 $y_{ijklmn} = \mu + HYC_i + RANK_j + AFC_k + LS_l + s_m + e_{ijklmn}$

- *y*_{*ijklmn*} conformation trait
 - μ overall mean
- *HYC*_i herd*year*classifier (i=1,2,...611)
- $RANK_j$ rank for milk yield (j=1,2,3)
- AFC_k age at first calving (k=1,2,...,5)
- LS_l lactation survival (l=1,2)
- *s_m* random sire effect
- *e*_{*ijklmn*} random residual effect

Longevity:

- defined as *lactation survival* two categories/classes:
 - culled during first lactation (LS1)
 - survived at least three lactations (LS4+)



Results – Data set I – Frequency within health traits





MARTIN-LUTHER-UNIVERSITÄT HALLE-WITTENBERG **Table 1**LSMEANS for conformation traits with a highly significant (p<0.0001 and F-value>10) effect of
number of new disease cases

Class	Dairy form	Rump width	Rear leg set	Hock quality	Foot angle	Loco- motion	Fore udder attachment	Udder depth	Body condition
0	5.38	5.23	5.17	5.17	5.00	5.16	5.48	5.64	5.07
1	5.46	5.25	5.22	5.12	4.97	5.05	5.40	5.57	5.00
2	5.52	5.27	5.26	5.10	4.94	4.98	5.30	5.50	4.92
3	5.59	5.32	5.30	5.05	4.91	4.85	5.24	5.45	4.84
4	5.64	5.32	5.36	5.03	4.87	4.75	5.14	5.40	4.78
5+	5.75	5.41	5.45	4.94	4.83	4.55	5.07	5.37	4.63
Diff	-0.37	-0.17	-0.28	+0.23	+0.17	+0.61	+0.41	+0.27	+0.44

SE ranged from 0.02 to 0.03

• further important traits: stature, chest width, rump angle, rear leg rear view, rear udder height, central ligament



Results – Data set I – Number of different groups of diseases

Table 2LSMEANS for conformation traits with a highly significant (p<0.0001 and F-value>10) effect of
number of different groups of diseases

Class	Dairy form	Rump width	Rear leg set	Hock quality	Rear leg rear view	Foot angle	Loco- motion	Rear udder height	Central liga- ment	Fore udder attach- ment	Udder depth	Body con- dition
0	5.38	5.23	5.18	5.17	5.03	4.99	5.15	5.31	5.51	5.48	5.64	5.06
1	5.49	5.26	5.24	5.11	4.98	4.95	5.01	5.26	5.45	5.38	5.55	4.96
2	5.57	5.31	5.31	5.04	4.93	4.91	4.85	5.23	5.43	5.21	5.44	4.85
3+	5.72	5.40	5.39	5.04	4.82	4.87	4.70	5.17	5.36	5.08	5.32	4.70
Diff	-0.34	-0.17	-0.21	+0.13	+0.21	+0.12	+0.45	+0.14	+0.15	+0.40	+0.32	+0.36

SE ranged from 0.02 to 0.03

• **further important traits**: stature, rump angle



Results – Data set II – Differences in body capacity traits contrasting (LS1 - LS4+)



* p<0.0001



Results – Data set II – Differences in feet and legs scores contrasting (LS1 - LS4+)



* p<0.0001



Results – Data set II – Differences in udder scores contrasting (LS1 - LS4+)





Conclusion





- there are conformation traits which are valuable indicators of health and longevity in dairy cattle
- important traits differ hardly between the two data sets and should be the focus of breeding for improved conformation
- it could be discussed if other conformation traits are really needed



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Zucht

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Thank you for your attention!





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Material and methods – Definition of health traits



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Figure 1 LSMEANS for conformation traits with a highly significant (p<0.0001) effect of *number of new disease cases*

Class	Stature	Dairy form	Chest width	Rump angle	Rump width	Rear leg set	Hock quality	Rear leg rear view	Rear udder height	Central liga- ment	Fore udder attach- ment	Udder depth	Foot angle	Body con- dition	Loco- motion
0	147.42	5.38	5.34	4.95	5.23	5.17	5.17	5.04	5.31	5.51	5.48	5.64	5.00	5.07	5.15
1	147.49	5.46	5.35	4.96	5.25	5.22	5.12	4.99	5.27	5.44	5.40	5.57	4.97	5.00	5.05
2	147.52	5.52	5.31	4.96	5.27	5.26	5.10	4.98	5.24	5.44	5.30	5.50	4.94	4.92	4.98
3	147.53	5.59	5.30	4.98	5.32	5.30	5.05	4.91	5.25	5.43	5.24	5.45	4.91	4.84	4.85
4	147.60	5.64	5.25	5.02	5.32	5.36	5.03	4.88	5.23	5.41	5.14	5.40	4.87	4.78	4.75
5+	147.77	5.75	5.25	5.08	5.41	5.45	4.94	4.80	5.16	5.39	5.07	5.37	4.83	4.63	4.55
Diff	-0.35	-0.37	+0.09	-0.13	-0.17	-0.28	+0.23	+0.24	+0.15	+0.21	+0.41	+0.27	+0.17	+0.44	+0.60

SE ranged from 0.02 to 0.03 (0.05-0.07 for stature)



Figure 2 LSMEANS for conformation traits with a highly significant (p<0.0001) effect of number of different groups of diseases

Class	Stature	Dairy form	Rump angle	Rump width	Rear leg set	Hock quality	Rear leg rear view	Rear udder height	Central liga- ment	Fore udder attach- ment	Udder depth	Foot angle	Body con- dition	Loco- motion
0	147.42	5.38	4.95	5.23	5.18	5.17	5.03	5.31	5.51	5.48	5.64	4.99	5.06	5.15
1	147.49	5.49	4.96	5.26	5.24	5.11	4.98	5.26	5.45	5.38	5.55	4.95	4.96	5.01
2	147.56	5.57	4.99	5.31	5.31	5.04	4.93	5.23	5.43	5.21	5.44	4.91	4.85	4.85
3+	147.74	5.72	5.06	5.40	5.39	5.04	4.82	5.17	5.36	5.08	5.32	4.87	4.70	4.70
Diff	+0.32	-0.34	-0.11	-0.17	-0.21	+0.13	+0.21	+0.27	+0.34	+0.40	+0.32	+0.12	+0.36	+0.45

SE ranged from 0.02 to 0.03 (0.05-0.07 for stature)



















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Results – Data set II – Frequency of scores for body condition and stature





Results – Data set II – Frequency of scores for hock quality and locomotion









Results – Data set II – Frequency of scores for fore udder attachment and udder depth





Results – Data set II – Frequency of scores for rear udder height and central ligament









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Rear teat position



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Frequencies of number of new disease cases



Number of new disease cases

