INVESTIGATION OF EXTERIOR TRAITS DEPENDENCE ON THE GENOTYPE OF LITHUANIAN BLACK AND WHITE COWS ACCORDING TO THE DEGREE OF HOLSTEIN GENES

Vida Juozaitienė1, Lina Anskienė2, Antanas BaNAS3, Audronė Rekštiūtė1, Arūnas Šileika3, Aleksandras Muzikevičius1, Jone Kantautaitė1, Virginija Žoštaitienė1, Arūnas Juozaitis3
1Animal Breeding Department, Veterinary Academy, Lithuanian University of Health Sciences
Tilžės 18, LT-47181, Kaunas, Lithuania
Phone (8-37) 36 35 75; E-mail: biometrija@lva.lt, anskiene@lva.lt
2Department of Non-Infectious Diseases, Veterinary Academy, Lithuanian University of Health Sciences
Tilžės 18, LT-47181, Kaunas, Lithuania; Phone +370-36 34 02
3Department of Animal Nutrition, Veterinary Academy, Lithuanian University of Health Sciences
Tilžės 18, LT-47181 Kaunas, Lithuania; Phone (8-37) 36 34 08

Abstract. The conformation traits of cows are heritable and linked with functionality and longevity. The dominant dairy breed in Europe is the Holstein. The objectives of this study were to evaluate exterior traits dependence on genotype of Lithuanian Black and White cows according to the degree of Holstein genes. For analysis, the conformation traits data of 53160 cows (in average of 2.7±0.01 lactation) were evaluated, for the degree of Holstein genes estimation records of cows with complete pedigree information of 3 ancestor generations from the national database were used. The data were grouped by exterior evaluation scores. Although Lithuanian Black and White breed nowadays is intensively improved by Holstein breed, the average degree of Holstein genes has been determined to be 63.1±0.09%. The results of this study showed dependence of cows’ exterior traits on genotype according to the degree of Holstein genes. The biggest positive influence of Holstein breed is produced on the height and udder depth of Lithuanian Black and White cows (P<0.001). The average content of Holstein genes of cows evaluated by optimal score based on body traits varied from 60.8±0.36% (rump width) to 71.6±0.19% (height), on extremities traits from 61.1±0.12% (rear leg set angle) to 67.4±0.30% (rear leg form), and on udder traits from 61.6±0.73% (teat length) to 72.8±1.22% (udder depth) (P<0.001). The data of Lithuanian Black and white cows included in the present study suggest that cows with high proportion of Holstein genes are taller, with deeper and higher attached udders.

Keywords: exterior traits, degree of Holstein genes, Lithuanian Black and White cattle

Introduction. Farm animals have been undergoing human-managed selection ever since their original domestication. Initially, selection was probably limited to docility and manageability, but in the last 60 years breeding programs have focused on the genetic improvement of production traits (Oltenacu and Broom, 2010). Although selection for yield traits has received primary emphasis in the selection goals of dairy cattle, substantial emphasis has been placed on other traits, particularly in North America. Many of these non-yield traits are related to the outward appearance of cows, such as overall conformation or ‘type’, udder-type traits, body size (including height, chest width and body depth), and angularity (Oltenacu and Broom, 2010).

A primary reason for collecting information on type is to aid breeders in selecting profitable, functional cows in order that early culling for causes unrelated to yield can be avoided (Strapak et al., 2005). Selection emphasis on type traits associated with increased herd life may be beneficial to decrease involuntary culling and increase profitability (Caïster et al., 2010; Geourgescu, et al., 1998; Larroque et al., 2001; Lavrinović et al., 2009; Rogers, 1989). Conformation traits are recorded in many dairy cattle breeds (Boelling et al., 1998; de Haas et al., 2007; Foster et al. 1988; Harris et al., 1992; Koenen et al., 1998).

In recent years, a considerable number of scientific studies have been carried out and it has been determined that the dairy cow productivity and culling reasons are very closely related to the genotype of animals (Chirinos et al., 2007; Martinez et al., 2004; Meszaros et al., 2008; Pachova et al., 2005; Peterson et al., 2005; Tsuruta et al., 2004) and conformation traits of cows (Auldist et al., 1996; Dadpasand et al., 2008; Forsbäck et al., 2010; Sanders et al., 2009; Shearer et al. 1996; Urech et al., 1999).

Today, the Holstein breed of cows is known in Lithuania as the world’s highest-production dairy animals (Jukna, Pauliukas, 2001). This breed has been used in Black-and-White cattle breeding improvement programs.

The objectives of this study were to evaluate the dependence of the exterior traits on the genotype of Lithuanian Black and White cows according to the degree of Holstein genes.

Material and methods

The average % of Holstein genes in cows genotype has been estimated according to records of cows with complete (3 ancestors generations) pedigree information from the database of State Agricultural Information and Rural Business Centre. For the analysis of conformation traits the data of 53160 Lithuanian Black and White cows in average of 2.7±0.01 lactation were used.

The conformation assessment system, which has been applied in Lithuania, follows the recommendations of international organizations ICAR and INTERBULL and The State Animal Breeding Supervision Service founded under the Ministry of Agriculture of the Republic of
Lithuania, Order No. 1A-15, 2011 05 26.

Traits, all scored on a 9-point scale, were for height, stoutness, body depth, chest width, dairy type, rump width, rump angle, rear leg set angle, rear leg form, heel joint, hoof height, hoof tarsus angle, udder attachment, udder height, udder cleft, udder depth, teat thickness, and teat length (from 1 to 9 point scale). Optimal scores of cows conformation traits (Saikevičius and Juozaitiene; 2004; European council directives 77/504, 86/130, 87/328, 94/515 and recommendations of international organization ICAR) are presented in the Table 1.

**Table 1. Optimal scores of cows conformation traits**

<table>
<thead>
<tr>
<th>Trait</th>
<th>Optimal scores of body assessment traits</th>
<th>Optimal scores of extremities assessment traits</th>
<th>Optimal scores of udder assessment traits</th>
</tr>
</thead>
<tbody>
<tr>
<td>Height</td>
<td>9</td>
<td>5</td>
<td>9</td>
</tr>
<tr>
<td>Stoutness</td>
<td>8</td>
<td>6</td>
<td>9</td>
</tr>
<tr>
<td>Body depth</td>
<td>8</td>
<td>8</td>
<td>9</td>
</tr>
<tr>
<td>Chest width</td>
<td>7</td>
<td>8</td>
<td>8</td>
</tr>
<tr>
<td>Dairy type</td>
<td>8</td>
<td>7</td>
<td>9</td>
</tr>
<tr>
<td>Rump width</td>
<td>7</td>
<td>7</td>
<td>6</td>
</tr>
<tr>
<td>Rump angle</td>
<td>5</td>
<td>5</td>
<td>6</td>
</tr>
</tbody>
</table>

Statistical characteristics in the sample (n) – arithmetic mean (M), standard error (SE), and indicator of statistical significance (P) – were calculated using R 2.1.0” package (http://www.r-project.org/).

**Results and discussion**

Analyzing our data set, we estimated that the average proportion of Holstein genes in Lithuanian Black and White cows population is 63.1 ± 0.09% (degree of Holstein genes amounts to 50% in 21% of Lithuanian Black and White cows, from 50 to 75% in 44%, and 75% and more in 35% of Lithuanian Black and White cows).

Conformation was one of the first nonproduction traits scored and included in the selection indices of dairy cattle populations around the world (Powell, 2006; Shook, 2006; White, 1974).

Type traits are recorded relatively early in life, most often in the first lactation, and are more heritable than longevity (Boettcher et al., 1997; Kadarmideen et al., 2003; Strapak et al., 2010; VanRaden and Wiggans, 1995) which makes selection relatively more efficient.

In cattle, body conformation traits such as height and body depth affect feed intake and thus milk production (Hiendleder et al., 2003).

**Table 2. The average % of Holstein genes of cows evaluated by body assessment traits**

<table>
<thead>
<tr>
<th>Score</th>
<th>Height</th>
<th>Stoutness</th>
<th>Body depth</th>
<th>Chest width</th>
<th>Dairy type</th>
<th>Rump width</th>
<th>Rump angle</th>
</tr>
</thead>
<tbody>
<tr>
<td>n</td>
<td>M±SE</td>
<td>n</td>
<td>M±SE</td>
<td>n</td>
<td>M±SE</td>
<td>n</td>
<td>M±SE</td>
</tr>
<tr>
<td>1</td>
<td>1507 51.2±0.50</td>
<td>1556 49.9±0.56</td>
<td>2 43.8±0.00</td>
<td>17 45.4±7.64</td>
<td>93 53.9±2.34</td>
<td></td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>1499 51.3±0.54</td>
<td>935 54.7±0.66</td>
<td>14 65.9±4.12</td>
<td>29 79.9±3.43</td>
<td>85 63.0±2.31</td>
<td>21 70.8±4.93</td>
<td>457 64.8±0.88</td>
</tr>
<tr>
<td>3</td>
<td>3868 53.1±0.33</td>
<td>1600 55.1±0.53</td>
<td>149 61.1±1.44</td>
<td>326 68.7±1.41</td>
<td>581 60.8±0.90</td>
<td>263 63.0±1.16</td>
<td>1834 64.7±0.45</td>
</tr>
<tr>
<td>4</td>
<td>4826 57.0±0.30</td>
<td>2803 56.9±0.41</td>
<td>1413 61.7±0.63</td>
<td>1991 64.7±0.55</td>
<td>2256 61.5±0.49</td>
<td>4549 62.9±0.32</td>
<td>8853 64.9±0.22</td>
</tr>
<tr>
<td>5</td>
<td>6537 59.3±0.24</td>
<td>3858 60.5±0.34</td>
<td>8177 61.2±0.25</td>
<td>10255 61.7±0.22</td>
<td>6183 61.4±0.30</td>
<td>25394 64.3±0.13</td>
<td>31301 61.5±0.12</td>
</tr>
<tr>
<td>6</td>
<td>7434 61.4±0.26</td>
<td>6171 63.3±0.27</td>
<td>19975 62.7±0.15</td>
<td>17016 61.0±0.16</td>
<td>10724 61.3±0.21</td>
<td>19022 62.1±0.15</td>
<td>8043 65.2±0.23</td>
</tr>
<tr>
<td>7</td>
<td>8794 64.6±0.21</td>
<td>6029 63.3±0.27</td>
<td>19179 64.8±0.15</td>
<td>16431 64.0±0.16</td>
<td>15217 61.6±0.17</td>
<td>3480 60.8±0.36</td>
<td>2216 69.6±0.41</td>
</tr>
<tr>
<td>8</td>
<td>8307 68.2±0.22</td>
<td>6225 62.1±0.28</td>
<td>3714 62.8±0.35</td>
<td>6769 67.8±0.24</td>
<td>13699 64.7±0.17</td>
<td>379 57.6±0.13</td>
<td>337 67.0±1.11</td>
</tr>
<tr>
<td>9</td>
<td>10388 71.6±0.19</td>
<td>25983 66.2±0.13</td>
<td>539 56.8±0.94</td>
<td>341 66.8±0.86</td>
<td>4398 71.8±0.27</td>
<td>52 60.0±1.80</td>
<td>26 80.9±3.58</td>
</tr>
</tbody>
</table>

Note: the optimal exterior trait assessment score is bold.
As presented in Table 2, the average proportion of Holstein genes of cows evaluated by height ranged from 51.2±0.50 % (score 1, n = 1507) to 71.6±0.19 % (score 9, n = 1038), by stoutness from 49.9±0.56 % (score 1, n = 1556) to 66.2±0.13 % (score 9, n = 23983), and by body depth from 61.1±1.44 % (score 3, n = 149) to 65.9±0.42 % (score 2, n = 539) (P<0.001).

Malgorzata Morek-Kopeć et al. (2012) investigated the highest impacts on functional longevity were for body depth, chest width, and dairy type.

In our study, the average degree of Holstein genes of cows evaluated by chest width varied from 43.8±0.00 % (score 1, n=2) to 79.0±3.43 % (score 2, n=29) (P<0.4) and by dairy type assessment from 45.4±7.64 % (score 1, n=2) to 79.0±3.43 % (score 2, n=29) (P<0.4).

The average proportion of Holstein genes of cows evaluated by rump width varied from 57.6±0.13 % (score 1, n=17) to 71.8±0.27 % (score 9, n=4398) (P<0.001).

The average proportion of Holstein genes of cows assessed by rump angle ranged from 56.1±0.36 % (score 1, n=37) to 70.8±1.08 % (score 1, n=173) (P<0.001) (Table 3).

Caravello et al., (2004) found traits such as udder depth, rear legs side view and foot angle and showed an intermediate optimum with respect to longevity and extreme scores of these traits which increased the risk of involuntary culling. Every year in Lithuania an average of 4 % of the cows (of all culled cows) are culled for feet and 13–14 % (of all culled cows) and udder diseases (Annual report of milk recording No. 74 and No. 75).

In our investigation, the average degree of Holstein genes of cows evaluated by the shape of rear legs ranged from 55.6±1.90 % (score 1, n=154) to 67.4±0.30 % (score 6, n=4610) (P<0.001).

Table 3. The average % of Holstein genes of cows evaluated by extremities assessment traits

<table>
<thead>
<tr>
<th>Score</th>
<th>Rear leg set angle M±SE</th>
<th>Rear leg form M±SE</th>
<th>Heel joint M±SE</th>
<th>Hoof height M±SE</th>
<th>Hoof tarsus angle M±SE</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>37</td>
<td>77.7±3.50</td>
<td>154</td>
<td>55.6±1.90</td>
<td>173</td>
</tr>
<tr>
<td>2</td>
<td>228</td>
<td>65.1±1.43</td>
<td>493</td>
<td>59.8±0.98</td>
<td>375</td>
</tr>
<tr>
<td>3</td>
<td>1344</td>
<td>67.7±0.59</td>
<td>3565</td>
<td>63.8±0.33</td>
<td>1518</td>
</tr>
<tr>
<td>4</td>
<td>6906</td>
<td>61.3±0.27</td>
<td>23134</td>
<td>62.7±0.13</td>
<td>5460</td>
</tr>
<tr>
<td>5</td>
<td>27775</td>
<td>61.1±0.12</td>
<td>18850</td>
<td>62.1±0.16</td>
<td>9429</td>
</tr>
<tr>
<td>6</td>
<td>13108</td>
<td>67.2±0.18</td>
<td>4610</td>
<td>67.4±0.30</td>
<td>8828</td>
</tr>
<tr>
<td>7</td>
<td>3233</td>
<td>66.1±0.36</td>
<td>1898</td>
<td>66.9±0.47</td>
<td>14705</td>
</tr>
<tr>
<td>8</td>
<td>463</td>
<td>66.1±0.89</td>
<td>331</td>
<td>65.6±1.00</td>
<td>10195</td>
</tr>
<tr>
<td>9</td>
<td>66</td>
<td>54.3±2.85</td>
<td>125</td>
<td>66.7±1.39</td>
<td>2477</td>
</tr>
</tbody>
</table>

Rear leg set angle, dry, not swollen heel joint are desirable for dairy type cows. Heel joint scored 1 to 4 is undesirable, the average score being 5. (Saikievicius and Juozaiene, 2004).

According to the results of our research, the average proportion of Holstein genes of cows by hell joint assessment ranged from 60.8±0.17 % (score 7, n=14705) to 70.8±1.08 % (score 1, n=173) and by hoof height assessment from 58.6±0.42 % (score 7, n=2530) to 65.8±1.38 % (score 9, n=30) (P<0.001).

Zavadilova et al. (2011) found that foot and leg traits showed substantially lower effect on longevity, and the effect of foot angle was minimal.

In our investigation, the average % of Holstein genes of cows evaluated by hoof tarsus angle varied from 53.6±6.04 (score 9, n=26) to 68.1±0.31% (score 6, n=4256) (P<0.001).

Hoof tarsus angle disorders affect the welfare of the animal and have economic implications due to costs of treatment, earlier culling, and production losses (Koenig et al., 2005; Schöpke et al., 2013; Swalve et al., 2008; Van der Linde et al., 2010; Van der Waaij et al., 2005).

Dadpassand et al. (2008) found the strongest relationship between the length of productive life and type traits of the udder attachment and udder depth.

In our research (Table 4), the average degree of Holstein genes by udder attachment assessment varied from 61.8±1.09 (score 1, n=314) to 68.0±0.94 % (score 9, n=264) (P<0.001).

In accordance to Rupp et al. (1999), Tilki et al. (2005), the score udder attachment and udder depth are among the most important conformation traits with regard to udder health.
The average proportion of Holstein genes by cows udder height assessment varied from 53.5±1.10 % (score 1, n=421) to 71.6±0.22 % (score 9, n=6261) (P<0.001). Strapak et al. (2005) and Vacek et al. (2006) suggest a well attached fore udder, high attached rear udder, strong cleft, close front teat placement and moderately long teats as important traits for a long productive cow’s life.

In our investigation, the average degree of Holstein genes evaluated by cows udder cleft ranged from 60.3±0.48 % (score 3, n=1856) to 67.0±0.20 % (score 9, n=225) (P<0.001). As presented in (Fig.1), the average % of Holstein genes of cows evaluated by teat length ranged from 68.0±0.94 % (score 2, n=6261) to 71.6±0.22 % (score 9, n=225) (P<0.001).

It is convenient when teats of milk cows are cylindrical, uniformly located, not deviating too much from side to side or forward. It is undesirable if the space between the teats is very large, teats are located in the bottom quarter edge or if the space is very small and teats are close to each other. Teats which are turned outward or extremely inward also make milking very difficult.

Our data records showed that the average degree of Holstein genes by cows teat placement assessment ranged from 58.7±0.85 % (score 2, n=652) to 69.3±0.44 % (score 7, n=2151) (P<0.001). Too short or too long teats often do not enable correct placing of teat cups and teat cup liner causing the interruption of vacuum pressure and penetration of air into milking system, contaminating the milk by microorganisms from the environment (Pantelic et al., 2010).

Teat length and thickness is important in milking cows. It is convenient when teat length of heifers is 5.0–5.5 cm and of cows 5.5–6.0 cm and thickness is 2.4–2.5 cm in heifers and 2.7–2.8 cm in (Bardakcioglu H.E. et al., 2011).

The average proportion of Holstein genes of cows by teat thickness assessment varied from 49.2±0.00 % (score 1, n=12) to 76.2±6.10 % (score 9, n=359) (P<0.001).

Visual evaluation and recognition of milk characteristics of cows are preliminary indicators of milk production, and partially also of longevity and reproductive ability of cattle, which is also very important in terms of economic efficiency of milk production. Detailed knowledge of the cow’s exterior features may provide valuable information about the genetic life potential of cows (Pantelic et al., 2007).

In a considerable number of scientific researches it has been determined that cows with optimal conformation traits are more productive and liable to increased longevity (Chirinos et al., 2007; Forsbäck et al., 2010; Meszats, 2008; Sanders et al., 2009).

The data of Lithuanian Black and White cows included in the present study suggest that cows with high proportion of Holstein genes are taller and have deeper and higher attached udders.

Similar changes have occurred in other Holstein-Friesian populations. In the Dutch Holstein-Friesian dairy population, from 1981 to 2007, the average height of heifers increased by 14 cm (Oltenacu and Broom, 2010).
Teat length scores

The amount of Holstein genes

Fig. 1 The average % of Holstein genes of cows evaluated by teat length

The results of our study showed dependence of cows exterior traits on genotype according to the degree of Holstein genes. The biggest positive influence of Holstein breed has been produced on height and udder depth of Lithuanian Black and White cows (P < 0.001). The average content of Holstein genes of cows (Fig. 2) evaluated by optimal score based on body traits varied from 60.8 ± 0.36% (rump width) to 71.6 ± 0.19% (height), on extremities traits from 61.1 ± 0.12% (rear leg set angle) to 67.4 ± 0.30% (rear leg form), and udder traits from 61.6 ± 0.73% (teat length) to 72.8 ± 1.22% (udder depth) (P < 0.001).

Fig. 2 The average % of Holstein genes of cows evaluated by optimal exterior traits score

The Holstein is the dominant dairy breed in Europe (Oltenacu and Broom, 2010). Atkins G. et al. (2008) reported that cattle of Holstein breed perform high production and desirable conformation. The functional conformation necessary for a cow to express her productive and reproductive potential is the ability to maintain adequate body condition and the ability to move with sound locomotion (Atkins et al., 2008). Since conformation traits are heritable and are linked with functionality, selection for conformational traits is an effective tool to predict genetic improvement in functionality.

Conclusions
1. According to our investigation data set, the estimated average proportion of Holstein genes in Lithuanian Black and White cattle population was 63.1 ± 0.09%.
The obtained results showed dependence of cows exterior traits on genotype according to the degree of Holstein genes. The biggest positive influence of Holstein breed has been produced on height and udder depth of Lithuanian Black and White cows (P< 0.001). The average content of Holstein genes of cows evaluated by optimal score based on body traits varied from 60.8 ± 0.36% (rump width) to 71.6±0.19% (height), on extremities traits from 61.1 ± 0.12% (rear leg set angle) to 67.4 ±0.50% (rear leg form), and on udder traits from 61.6± 0.73% (teat length) to 72.8±1.22% (udder depth) (P< 0.001).

References
24. Małgorzata Morek-Kopeć, Andrzej Zaremba. Relationship between conformation traits and longevity in


Received 22 November 2013
Accepted 27 May 2014