Multi-trait animal model estimation of genetic parameters for linear type and gait traits in the Belgian warmblood horse

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Summary

Genetic parameters for the height at withers, 27 linear type and six linear gait traits were estimated for the Belgian warmblood horse. Observations on 987 mares, mostly 3 years old, were analysed using a multi-trait animal model. The statistical model included appraiser, age and location (date \times place of appraisal) as fixed effects. Genetic parameters were estimated using a canonical transformation and an expectation-maximization restricted maximum likelihood algorithm with an additional deceleration step. Estimates of heritability for the 33 linear traits were between 0.15 and 0.55. Heritability of the height at withers was 0.34 ± 0.06 . Estimated genetic correlations ranged from -0.60 to 0.98 with an average SE of 0.10. The highest positive correlations were found among traits of walk and among traits of trot. Volume and the quality of legs were the most negatively correlated. Estimated genetic parameters indicated that the linear scoring system is a valuable tool to assess conformation. The full (co)variance matrix is now available for breeding value estimation to support selection for conformation and gaits.

Introduction

The Belgian warmblood horse (BWP) is a saddle horse, used primarily in show jumping and dressage at recreational and international level. The breeding organization of the BWP has introduced conformation and gaits recording in its breeding program because of their influences on horse abilities. In fact, conformation and gaits are important for aesthetics, wellness, durability and functionality of horses (Forest 1996). The success in many types of horse shows depends partly on these characteristics. Conformation and gaits have also been proposed as indirect indicators of performance capacity, since heritability coefficients of conformation traits were often higher than those found for performance traits (Saastamoinen & Barrey 2000).

Therefore, functional conformation and gaits are important selection criterions in horse breeding. This is reflected in the breeding objective of the BWP which mentions the achievement of a correctly-built horse with a rectangular frame, big outlines and good basic paces.

Conformation and gaits of the BWP mares are appraised in two ways: the conformation and gaits contests (i.e. halter presentation) and a linear scoring system.

In the conformation and gait contests, mares are ranked by a jury based on aesthetics ('expressiveness', 'brightness', etc.) and on functional (correctness, etc.) aspects of type, legs, walk and trot. This system has two important drawbacks, namely the fact that conformation and gaits are appraised in a global manner and that preferences of judges play a role.

Linear scoring could improve on this because body and gaits are subdivided into several distinct traits, assessed separately. Each trait of the horse is scored on a linear scale between the two biological extremes (Mawdsley *et al.* 1996). By assessing traits individually rather than in combination and by describing rather than evaluating, this method easily reveals differences between animals (Koenen *et al.* 1995). A linear scale can also be considered as a continuous scale. Therefore, this kind of traits can be analysed by a linear mixed-model which is nowadays the standard tool to estimate unbiased breeding values.

Even if linear scoring is used less in horses than in other species like cattle; genetic parameters for linear traits have already been reported in other horse breeds. Using a derivative-free restricted maximum likelihood algorithm and an animal model. Koenen et al. (1995) estimated genetic parameters for the linear scoring of the Dutch warmblood riding horse. Heritabilities, obtained in univariate analyses, ranged between 0.09 and 0.28. Genetic correlations within four trait groups were estimated using bivariate models and varied between -0.69 and 0.98. Samoré et al. (1997), working with the Haflinger Horse, found higher heritabilities, using a multi-trait animal model and an expectation-maximization restricted maximum likelihood algorithm. Values were between 0.02 and 0.53. Genetic correlations (within 10 groups of traits) ranged between -0.32 and 0.99. Sampling errors for the estimates of heritability or genetic correlations were not available for these studies. Nevertheless, Van Bergen & Van Arendonk (1993) have reported standard errors from 0.04 to 0.07 for heritabilities varying between 0.10 and 0.39 (data on Shetland ponies).

The objective of this research is to estimate genetic parameters (heritabilities and genetic correlations) for linear type traits in the BWP horse population as the necessary first step to develop a genetic evaluation system.

Materials and methods

Scoring system and data

Linear scoring was introduced by the BWP studbook in 2003. During the assessment, the height at withers is measured and 33 linear traits are scored. Description of the traits is given in Table 1. Of these traits, 17 refer to type, 10 to legs and feet and six to gaits. Linear traits are scored between -20 and 20, these endpoints corresponding theoretically to biological extremes in the BWP population. Scores between the two extremes are multiples of 5. The 0 of this scale corresponds to the theoretical mean of each linear trait in the BWP population. Descriptive statistics and test of normality were conducted using the Univariate procedure of SAS software (SAS 1985).

From 2003 to 2007, 1215 mares have been classified by 15 classifiers at 28 locations. Mares are scored only once, when they are 3 or 4 years old. Only very few mares aged over 4 years have been recorded. Most horses were between 3 and 3.5 years old (only 8 % of the 1215 mares were over 3.5 years). Mares were classified in seven age classes (Table 2). Six mares aged over 44 months were not included in the set for estimation of variance components. The number of mares varied between 5 and 372 per appraiser and between 1 and 39 per location of assessment. For the estimation of variance components, a minimum of three records per date *location of assessment was set.

Statistical model

In the present study, the following multi-trait animal model was used to describe the data:

$$y = (S \otimes I_t) \text{datloc} + (H \otimes I_t) \text{age} + (F \otimes I_t) \text{app} + (Z \otimes I_t) a + e$$

y, the vector of records; datloc, the vector for the combined fixed effect of the date and the location of assessment (1–94); age, the vector for fixed effect of the age (1–6); app, the vector for fixed effect of appraiser (1–15); *a*, the vector for animal random effect and *e*, for random residual effect. *S*, *H*, *F*, *Z* were the common incidence matrices for all traits that associate datloc, age, app and *a*, with *y* and *I*_t was an identity matrix for the traits 't'.

These fixed effects included in the model were found to be significant for most traits (p < 0.05) in a preliminary least squares analysis with the general linear models procedure of SAS (SAS 1985).

(Co) variance components estimation

Estimates of variance components were obtained by a multi-trait REML procedure following a method developed by Misztal *et al.* (1995) for the analysis of linear scores in dairy cattle (Gengler *et al.* 1997). The method consists of a canonical transformation and an expectation-maximization REML algorithm (EM-REML) with an additional deceleration step. The canonical transformation reduces the computing effort by transforming the 34 correlated traits to 34 uncorrelated traits. Therefore, complexity of the

Table 1 Description of the linear scoring system for Belgian warmblood horses

		Scale		
Traits	Definition	-20	20	
Туре				
1. Frame	The shape of the body (ratio of body length to body height)	Square	Rectangle	
2. Condition	General muscularity and overweight	Poor	Heavy	
3. Head	Size of the head, distance between top of the head and the mouth	Small	Tall	
4. Head neck connection	Length of the junction between the poll and the throat latch	Light	Heavy	
5. Length of neck	Distance between the beginning of the withers and the top of the head	Short	Long	
6. Muscularity of neck	Development of the neck	Poor	Heavy	
7. Orientation of neck	Angle formed by the neck and the head	Vertical	Horizonta	
8. Height of the withers	Height between the highest and the lowest point of the withers	Flat	Prominent	
9. Length of withers	Length of the line formed between the highest and the lowest point of the withers	Short	Long	
10. Orientation shoulders	Angle between the corner of the crest of the shoulder and an horizontal axis	Straight	Sloping	
11. Length of shoulders	Distance between the top of the shoulder-blade and the breast	Short	Long	
12. Length of back	Distance between the end of the withers and the beginning of the loins	Short	Long	
13. Back force	Form of the upper line	Deep	Straight	
14. Loins	Form of the line formed by the loins	Weak	Straight	
15. Slope of croup	Angle formed by the line going from upper side of the haunch to the point of the buttock and an horizontal axe	Straight	Sloping	
16. Length of croup	Distance from front side of the haunch to rear side of the point of the buttock	Short	Long	
17. Muscularity of hind legs	Muscularity of hind legs above the hock	Poor	Heavy	
Limbs				
18. Position of forelegs 1	Position of leg relative to a straight line passing trough the canon bone and the forearm (side view)	In roe-deer	Sunken	
19. Position of forelegs 2	Seen from the front, O or X position formed by the legs	Flemish	French	
20. Position of hind legs	Angle formed by the hock	Sickle-hocked	Straight	
21. Position of hocks	Seen from the back, O or X position formed by hind legs	Outside	Inside	
22. Volume of hocks	Volume and quality of hocks	Light	Heavy	
23. Form of hoof	Form of the hoof	Narrow	Broad	
24. Position of pasterns	Angle between pasterns and an horizontal axis	Weak	Straight	
25. Heels	Distance from the coronet to the front of the foot	Low	Higher	
26. Volume of legs	Volume of legs in proportion of the rest of the body	Light	Heavy	
27. Quality of legs	Contour of articulation	Blurred	Lean	
Gaits (walk + trot)				
28. +31. Stride length	Length of step	Short	Long	
29. +32. Impulsion	Place of hind legs under the body during movements	Lack of power	Powerful	
30. +33. Suppleness	Transmission of movement to the body	Stiff	Supple	

 $\label{eq:table_$

Year	Number of records
2003	201
2004	259
2005	264
2006	254
2007	237
Age class	
age ≤36 months	76
36 <age months<="" td="" ≤38=""><td>406</td></age>	406
38 <age months<="" td="" ≤40=""><td>481</td></age>	481
40 <age td="" ≤42months<=""><td>150</td></age>	150
42 <age months<="" td="" ≤44=""><td>17</td></age>	17
44 <age months<="" td="" ≤48=""><td>6</td></age>	6
48 months <age< td=""><td>79</td></age<>	79

multi-trait analysis decreases and simultaneous estimation of (co)variance components of the 33 linear traits and the height at withers is feasible. The program guarantees that the covariance matrix is positive semi-definite. The deceleration step improves the stability of estimates and the convergence in small data sets (Wiggans *et al.* 2006). However, all traits must be recorded on every animal to apply the method of Misztal *et al.* (1995). Therefore, only mares with complete data were kept. This resulted in a subset of 977 mares with scores. The pedigree file contained 13 592 ancestors (up to 20 generations).

A method to approximate SE of the heritabilities was integrated in the canonical transformation process.

After convergence of the EM-REML algorithm, the SE of transformed genetic variances was estimated for all 34 traits using the expectation maximization average information REML matrix (Jensen et al. 1997; Johnson & Thompson 1995). In the next step, we computed a maximal and minimal genetic variance matrix by adding and subtracting the corresponding SEs to the (diagonal) matrix of genetic variances. After back transformation, maximal and minimal genetic variances on the original scale were obtained and from these, we computed maximal and minimal heritabilities using the intermediate phenotypic variance. Finally, the deviation between maximal and minimal heritabilities and the estimated intermediate heritabilities were used to approximate the SE of the heritabilities.

Standard errors for the genetic correlation between the trait *x* and *y*, $r_g(x,y)$, were computed

Table 3 Mean, standard deviation (SD),skewness and kurtosis of linear type traitsand gaits; and heritabilities [standarderrors(SE)] in Belgian warmblood horses

using the estimated $r_g(x,y)$, the estimated heritabilities of each trait (h^2) and their respective SE (σ_{h^2}), in the following formula (Falconer & Mackay 1996):

$$\sigma_{r_g}(x, y) = \frac{[1 - {r_g}^2(x, y)]}{\sqrt{2}} \sqrt{\frac{\sigma_{h_x^2} \sigma_{h_y^2}}{h_x^2 h_y^2}}$$

Results and discussion

Scoring system

Means and SD of the linear scores are given in Table 3. Mean scores for the linear traits were in the range from -1.5 to 4.8 and the SD of records within linear traits were in the range from 6.7 and 2.7. For most traits, the mean of the assessed population was

	Mean	SD	Skewness	Kurtosis	h ²	SE
1. Height at withers (cm)	165	4	0.03	2.83	0.34	0.06
2. Frame	4.8	6.2	-0.63	-0.42	0.34	0.04
3. Condition	3.7	5	0.02	0.01	0.15	0.03
4. Head	0.1	4.6	0.07	0.53	0.45	0.06
5. Head neck connection	-1.5	4.9	0.10	-0.29	0.26	0.05
6. Length of neck	0.6	5.4	0.06	-0.62	0.27	0.04
7. Muscularity of neck	0.2	5.3	0.03	-0.32	0.31	0.05
8. Orientation of neck	0.4	4.7	0.02	-0.26	0.24	0.05
9. Height of withers	2.6	5.1	-0.05	-0.39	0.34	0.05
10. Length of withers	1.5	5.8	-0.14	-0.65	0.33	0.06
11. Orientation shoulders	0	5.3	0.07	-0.69	0.20	0.04
12. Length of shoulders	2.7	5.1	-0.15	-0.48	0.31	0.06
13. Length of back	2.4	4.4	0.09	-0.16	0.34	0.05
14. Back force	-1.2	4.6	0.17	0.49	0.55	0.07
15. Loins	1.1	5.4	0.06	-0.59	0.47	0.06
16. Slope of croup	2	5.1	0.04	-0.32	0.30	0.05
17. Length of croup	0	5.3	0.00	-0.51	0.30	0.06
18. Muscularity of hind legs	2.6	5.3	-0.10	-0.24	0.20	0.04
19. Position of forelegs 1	-0.2	2.9	-0.47	3.99	0.35	0.05
20. Position of forelegs 2	-0.7	3.6	-0.03	3.01	0.36	0.07
21. Position of hind legs	0.2	4.2	0.03	0.51	0.22	0.04
22. Position of hocks	0	2.7	-0.18	5.88	0.24	0.04
23. Development of hocks	0.9	5.2	0.15	-0.55	0.22	0.05
24. Position of pasterns	0.3	3.7	0.29	2.05	0.25	0.05
25. Form of hoof	0.1	4.4	-0.07	0.86	0.25	0.05
26. Heels	-2.2	4.5	-0.02	0.38	0.25	0.04
27. Volume of legs	-0.1	4.5	0.02	-0.16	0.46	0.07
28. Quality of legs	2.8	5.7	-0.09	-0.36	0.22	0.05
29. Walk stride length	1.9	5.6	-0.20	-0.31	0.38	0.06
30. Walk impulsion	1.3	5.5	-0.11	-0.28	0.52	0.07
31. Walk suppleness	1.2	5.6	-0.08	-0.28	0.41	0.06
32. Trot stride length	1.9	6.2	-0.14	-0.41	0.47	0.05
33. Trot impulsion	0.5	6.7	-0.12	-0.59	0.36	0.05
34. Trot suppleness	0.9	6.5	-0.07	-0.44	0.33	0.04
Mean of all traits	-	-	-	-	0.32	0.05

close to 0. However, most SD were lower than the expected value of 7. Skewness of some traits (Table 3) indicated slightly asymmetric distributions (e.g. frame, back force, stride length of walk). Negative kurtosis (Table 3) was observed for many traits indicating a flatter distribution than the normal distribution. On the other hands, height at the withers, position of forelegs and position of hocks have a peaked distribution. However, formal tests (SAS 1985) did not reject the assumption of sampling from a normal distribution for all traits.

Differences in the distribution of scores between appraisers (>100 records) can be observed. Appraiser 1 assigned more positive scores, appraiser 4 more negative scores and appraiser 3 scored the majority of mares with 0. Scores given by appraiser 2 were well distributed along the scale. The recent start of the system and a lack of formal training of the appraisers might explain the different use of the linear scale. A study of the repeatability between and within appraisers would provide more insight about the lack of similarity. Some of the problems could partially be solved by practical and/or statistical interventions. In dairy cattle, it is common practice to train appraisers to be as similar as possible and often heterogeneity of means and variances due to appraisers are included in the statistical model. In our model, mean differences between appraisers were already modelled and future enhancements of the model should include adjustments for heterogeneity variances Figure 1.

Heritabilities

Estimates of heritabilities and associated SE are presented in Table 3. Heritabilities ranged from 0.15 to 0.55 with a mean of 0.32. Approximate SEs varied from 0.03 to 0.07 with a mean of 0.05. These results show that many linear scored traits were moderately heritable with reasonably low SEs. Traits with a heritability over 0.45 were force of the back, loins, the volume of legs, the impulsion of the walk and the length of stride in trot. Seventeen traits had moderate heritability (0.30–0.45), namely the height at withers, frame, head, muscularity of the neck, length of withers, length of the shoulders, length of the back, croup, forelegs, impulsion and suppleness of the trot, stride length and suppleness of the walk.

The lowest estimates of heritability (<0.25) were found for condition, orientation of neck and shoulders, muscularity and position of hind legs, position and volume of hocks and quality of the legs. In general, traits of hind legs and; 'feet' had lower

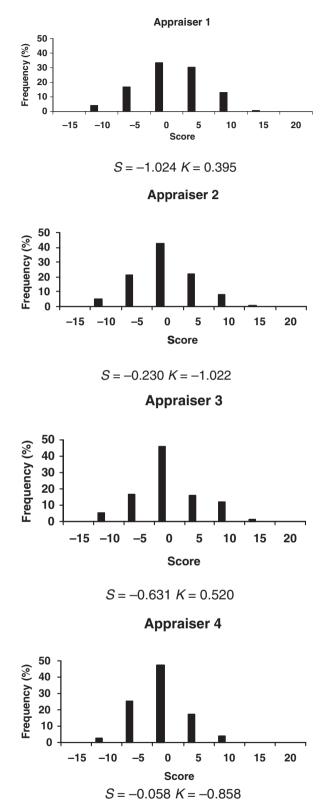


Figure 1 Distribution of scores (proportion of observed mares) for the four major appraisers in the linear scoring system for Belgian warmblood horses.

Multi-trait animal model estimation

heritabilities than others (back, loins, croup, head and withers), with a mean of 0.23 for hind legs and 'feet' traits compared to 0.35 for all the other traits. Our results on the heritability of hind leg traits do not support the general belief of horsemen that leg conformation in horses is highly heritable (Saastamoinen & Barrey 2000). Similar findings are also often reported in cattle (e.g. Gengler et al. 1997) where feet and leg traits have lower heritability than other conformation traits. Several reasons have been suggested for the low heritabilities of hind leg and feet traits, such as the difficulty of assessment, the training status of the horse or other environmental factors not included in the model (Arnason 1984; Preisinger et al. 1991: Saastamoinen 1993: Saastamoinen et al. 1998).

Heritabilities (and the SEs) for linear scores in the BWP were slightly different from values reported in the literature, nevertheless, the same general tendency was observed as in the study of Van Bergen & Van Arendonk (1993); Koenen *et al.* (1995) and Samoré *et al.* (1997). Differences with other studies could be related to differences between linear systems in term of definitions of traits and scales. Also, the method of estimating variance components was different (e.g. multi-trait versus single trait).

Heritabilities for linear scores in the BWP were lower than for zoometrical measurements (i.e. by metre or reference point on photography). For example, Molina *et al.* (1999) found heritabilities between 0.35 and 0.95 for seven zoometrical measurements in the Andalusian horse. On the other hand, the mean heritability of linear traits of the BWP was higher than the value of 0.24, obtained for ranking in 'conformation and gaits contests' (Janssens 1995) and this improvement was achieved without the need for specific or expensive equipment often needed to perform zoometrical measurements.

Genetic correlations

Genetic correlations ranged between -0.60 and 0.98 (Table 4) with an average SE of 0.10. Genetic correlations for many traits were low and rather close to zero, between -0.30 and 0.30, indicating virtually independence of these traits. Due to this independence, low correlated selection effects would be expected for many pairs of traits. Some high (≥ 0.60) and moderate correlations (0.40–0.59) were also obtained.

The highest genetic correlations, varying from 0.88 to 0.98, were found between stride length,

suppleness and impulsion within each gait (walk and trot). High positive correlations were also observed between walk and trot. Gaits were moderately to highly correlated with the height at withers and frame. Traits of the trot were positively correlated with muscularity and length of the neck and more moderately with the length of shoulders and the volume of the hocks. The stride length in trot was also moderately correlated with the length of the back. Interpretation of genetic correlations depends in the trait definitions For example, the impulsion, suppleness and stride length increase with taller mares having a more rectangular frame. A short neck with a poor muscularity, light developed hocks and a short back and shoulder is related with low impulsion, short stride length and low suppleness of the trot. Due to the correlations, selection for one of the traits will result in indirect responses in other traits. Consequently, selection for general riding ability could be quite effective. In fact, a long back permits the horse to move smoothly, the neck functions as a balancing tool and long shoulders permit a good development of muscles.

The high correlation (0.87) between the frame and the length of the back is logical because these traits partly overlap. A genetic correlation of 0.61 was found between frame and length of the neck. Lengths of the back and of the neck were also highly correlated (0.72). Lengths of two other traits of the upper line (the withers and the croup) were moderately correlated with the frame. The volume of the hocks was moderately linked with frame and the length of back.

Condition was moderately to highly correlated with muscularity and volume traits. The condition of the horse, reflecting muscularity and overweight, is important. In fact, a good muscularity is necessary to provide sufficient power to get over an obstacle or to perform in dressage. Contrary, a fatty or a bony horse will have some difficulties to perform in competitions. Condition was also highly correlated with the head-neck connection between the head and the neck. Furthermore, a 'large' connection was linked with heavy muscularity of the neck. The muscularity of the neck and the connection between the head and the neck are important for flexing the neck and correct positioning of the neck and head during riding (Van Der Veen 2002).

High and moderate correlations within traits of the shoulders and between shoulders and withers were observed; sloping shoulders and prominent and long withers were related with long shoulders. The importance of shoulders for horse abilities in

Genetic correlation	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33
0. Height at withers (cm)		+	- 0	+	+	+	+	+	+	+	+	+	+	0	0	+	+	0	+	+	+	+	+	0	+	+	+	-	+	+	+	+	+	+
1.Frame			+	+	-	+	+	0	-	+	-	+	+	0	0	0	+	-	-	+	-	+	+	-	+	-	+	-	+	+	+	+	+	+
2.Condition				+	+	<u>ا د ا</u>	+	-	-	+	0	0	0	+	+	-	+	+	0	0	+	+	+	+	-	+	+	-	+	+	+	+	+	+
3.Head					+	+	0	+	+	0	-	-	+	0	-	-	+	+	+	+	+	-	+	0	+	-	+	-	-	-	-	-	-	-
4.Head neck connection						-	+	-	-	+	+	+	0	+	+	0	+	+	+	+	+	-	+	+	-	+	+	-	+	+	0	+	+	+
5.Length of neck							+	+	+	+	+	+	+	+	-	0	+	-	+	-	+	+	+	+	+	-	+	-	+	+	0	+	+	+
6.Muscularity of neck								-	-	0	+	+	0	+	+	0	+	+	+	+	+	+	+	+	-	+	+	-	+	+	+	+	+	+
7.Orientation of neck									+	+	-	+	0	+	0	-	-	-	-	+	0	-	+	-	+	-	0	0	+	-	-	-	-	-
8.Height of withers										+	+	+	-	-	-	0	0	-	+	+	+	-	+	-	+	-	0	+	+	0	0	-	-	-
9.Lenght of withers											-	+	+	+	+	0	+	0	-	+	0	0	+	-	+	-	+	0	+	+	+	+	+	+
10.Orient. shoulders												+	0	+	+	+	+	-	+	+	0	+	0	+	+	+	+	+	0	+	0	+	+	+
11.Lenght of shoulders													0	-	+	+	+	0	+	+	+	0	+	-	+	-	+	0	+	+	+	+	+	+
12.Lenght of back														0	-	-	+	-	-	-	-	+	+	-	+	-	+	-	+	+	+	+	+	+
13.Back force															+	-	+	0	-	-	+	+	+	+	-	+	0	-	-	-	-	+	+	+
14.Loins																-	+	+	-	+	0	+	+	+	-	+	-	0	0	0	0	+	+	+
15.Slope of croup																	0	+	0	+	+	+	0	-	-	-	0	+	-	-	-	0	+	0
16.Lenght of croup																		+	-	+	0	+	+	0	+	+	+	-	+	+	+	+	+	+
17.Muscul. of hind legs																			-	+	+	0	+	-	-	+	+	-	-	-	-	-	-	-
18.Posit. of forelegs 1																				+	+	-	+	-	0	0	+	-	+	+	+	0	+	0
19.Posit. of forelegs 2																					+	0	+	-	0	0	+	0	+	0	0	0	+	0
20.Posit. of hind legs																						-	+	+	-	+	+	0	-	-	-	-	+	-
21.Posit. of hocks																							0	0	-	+	0	+	-	-	-	+	+	+
22.Volume. of hocks																								0	+	-	+	-	+	+	0	+	+	+
23.Position of pasterns																									-	+	+	-	-	-	-	+	+	+
24.Form of hoof																										-	+	-	+	+	+	-	0	0
25.Heels																											0	-	0	+	+	+	+	+
26.Volme of legs																												-	+	+	0	+	+	+
27.Quality of legs																													-	-	-	-	-	-
28.Walk stride length																														+	+	+	+	+
29.Walk impulsion																																+	+	+
30.Walk suppleness		1																														+	+	+
31.Trot stride length																																	+	+
32.Trot impulsion		1																																+
33.Trot suppleness																																		
											•																							

Table 4 Magnitude and direction of genetic correlations between linear traits and height at withers in Belgian warmblood horses

0 - 0.19 0.2 - 0.39 0.4 - 0.59 0.6 - 0.79 0.80 - 1 + positive correlation - negative correlation

dressage has been recognized (Holmström *et al.* 1994). A good orientation and length of the shoulders are necessary to have sufficient stride length in the movements and sufficient muscular force. The moderate correlation between these two traits would be favourable for a selection towards a long and sloping shoulder. A prominent withers was also correlated with a long withers, a poor condition, a horizontal and poor muscled neck and a poor muscled hind leg. Even if the withers seem to be less important for riding in a direct way, it is related to other conformation traits.

The neck was associated with other traits. Its muscularity was moderately correlated with the muscularity of hind legs. Its orientation was negatively correlated with its muscularity and positively correlated with the withers. A selection for a long neck would also increase volume of the hocks.

Loins were moderately correlated with the muscularity of hind legs, the length of the croup, the position of forelegs (1) and strongly correlated with the force of the back. The loins are an important part of the body as they link the hindquarters with the forehand and permit the transmission of energy developed by the hind legs. Moreover, the loins were positively correlated with the length of the croup which should permit an optimal stretching of muscles and increases stride length (Van Der Veen 2002). Loins were also positively correlated with the position of forelegs. A bad stance of the forelegs predisposes to a low soundness and a short durability (Stashak 1987). In conclusion, the loins are very important for their direct role in riding abilities and improvement by selection would be facilitated due to their correlations with other conformation traits.

The effectiveness of selection for suppleness of the horse can be increased due to: (i) the high correlation observed between heels and the position of pasterns; (ii) the moderately high correlations between heels and the back force; and (iii) between the length of back and the position of pasterns. In fact, heels and pasterns function as shock absorbers of movements and a correct line of back is also necessary to have supple horses (Van Der Veen 2002).

The correlation between volume and the quality of legs was negative with a value of -0.60. In fact, a horse with heavy legs can not have blurred legs and inversely. Volume of the legs had a particularly high correlation with volume of the hocks which are

important for the solidity of legs and for the length of stride in gaits. Volume of the legs was also moderately correlated with length of the back.

In other studies (Koenen et al. 1995; Samoré et al. 1997), linear traits were grouped. For example, linear traits were grouped in traits of the body, traits of legs and traits of the front and different groups were evaluated separately. By doing so, no correlations were obtained between traits belonging to different groups. This results in a loss of information about the possible relationships between traits and consequently, indirect selection responses might be incorrectly evaluated. In this study, we found relative high genetic correlations between traits which would have ended up in different groups, if we had followed the classification of Koenen et al. (1995) and Samoré et al. (1997). For example, frame and gaits, muscularity and orientation of the neck and trot, volume of the hocks and the length and muscularity of the neck, the position of hind legs and the length of the back are pairs that would cross group borders. Therefore, the analysis of all traits jointly was preferred to grouping traits.

A joint analysis of all traits will eventually result in estimated breeding values (EBV) for each linear trait. EBV for separate traits will provide more information to the breeders than aggregated once. On the other hand, too much information could make interpretation difficult and might cause confusion. If required by the studbook or breeders, a principal component analyses could be considered to reduce the information into a limited number of factors (Arnason 1984). Also, indices could be constructed to avoid confusion of producing too much information. However, more important for the success of the system is the collection of more data, so that a larger part of the population is covered.

Conclusion

The genetic parameters obtained in this study indicate that the linear scoring system is able to generate quantitative and detailed information on distinguishable conformation and gaits traits of the BWP. Magnitudes of heritabilities and genetic correlations indicate that selection is feasible and horse riding abilities could be improved. Even if the current data quality seems adequate, training of appraisers could improve uniformity.

The next step will be to implement a procedure for breeding values estimation. These breeding values will be more effective to support the selection of horses with respect to conformation and gaits.

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