

The effect of the gene for polledness on conception rate and litter size in the *Damascus* goat

A. CONSTANTINOIU (*), A. LOUCA (*) and A.P. MAVROGENIS (**)

(*) *Ministry of Agriculture and Natural Resources
Department of Agriculture, Nicosia, Cyprus*

(**) *Agricultural Research Institute, Nicosia, Cyprus*

Summary

Data on 1709 matings and 1247 parturitions were used to study the effect of the gene for polledness on conception rate and litter size in the *Damascus* breed of goat in Cyprus.

With one exception all polled sires used in the study proved to be heterozygous *Pp* for the polled factor *P*.

There was consistent excess of males in the offspring of all mating types with an overall proportion of males (53.1 p. 100) significantly different from the 50:50 ratio. Intersexes (6.0 p. 100) were found only when both parents were polled (*Pp* × *Pp* matings).

The percentage of intersexes corresponds to an expressivity $\gamma_2 = .48$ of the *P* genes for that trait in the *PP* females. Among alpine breeds γ_2 was between .44 and .57.

The percentage of males in excess corresponds to an expressivity of $\gamma_1 = .16$ of the *P* gene for that trait in the *PP* females ($\gamma_1 = .43$ to .56 in alpine breeds). Theoretically this excess must not be found in horned × polled and horned × horned crossings where no *PP* genotypes are produced. This discrepancy, as well as the low value of γ_1 in the *Pp* polled × *Pp* polled matings has not been explained.

Conception rate for polled (*Pp*) females was 74.7 p. 100, non significantly different from that of *pp* ones (76.4 p. 100). Litter size at birth for resp. all born and born live kids was 1.88 and 1.82 in polled goats (*Pp*) while it was resp. 1.75 and 1.71 in horned ones. This gives coefficients of surprolificacy of 1.074 and 1.066 in the *Pp* females compared to the *pp* ones for a value of 1.06 reported in alpine breeds.

There is a light but significant fertility disadvantage of the *Pp* (polled) males compared to the *pp* (horned) ones which gives a r_2 coefficient of compared fertility of .91 instead of 1.007 observed in alpine stocks. Such a disadvantage is also found at the conception stage.

These slight discrepancies indicate one needs more precise analysis of the behaviour of *P* gene in the goat.

Introduction

The inheritance of polledness (absence of horns) and its relationship to intersexuality and other forms of genital malformation have been extensively studied in goats, especially in strains of alpine origin (see BOYAJEAN, 1969 for a review).

It has been well established that practically all polled sterile males from polled parents are homozygous *PP* for the polled gene, the spermiostasis due to bilateral epididymal stenosis being the main cause of total sterility. Sterile males with hypoplastic testicles on the other hand were identified as masculinized genetic females, homozygous *PP* for the polled gene (RICORDEAU *et al.*, 1972). Polled heterozygous *Pp* females were found to be more prolific than horned homozygous *pp* females (SOLLER & KEMPENICH, 1964) and fertile homozygous *PP* polled males as well as heterozygous *Pp* more prolific than homozygous horned *pp* males RICORDEAU & LAUVERGNE, 1967 ; RICORDEAU, 1969).

A list of thirteen parameters has been proposed by LAUVERGNE (1969) in order to give a full account of various effects of the polled *P* gene in male and female goats.

In a study on the inheritance of polledness and its relationship to intersexuality in *Damascus* goat of Cyprus HANCOCK & LOUCA (1975) found a similar genetical behaviour regarding the polled factor, and concluded that the *P* gene, as in alpine breeds, was present in this breed.

The purpose of the present work was to further investigate the effect of the gene for polledness on sex ratio, conception rate and litter size in the *Damascus* breed of goat in Cyprus.

Materials and methods

The study utilized data on 1709 matings and 1287 parturitions recorded at the *Goat Breeding Station* of the *Cyprus Department of Agriculture* over a six-year period (1972 to 1977), including the years (1972 and 1973) which were previously utilized in the study of HANCOCK & LOUCA (1975).

All available information (returns to service, abortions and kidding dates) was used to classify matings as successful (terminated by pregnancy) or unsuccessful (failure to conceive). Litter size and sex of kids were determined at birth. In view of the occasional difficulty in determining intersexes at an early age it may be possible that some intersexes, which died soon after birth, had been erroneously classified as normal males or females.

The data were pooled across years and chi-square was used to test differences among the various classes. The data on conception rate (conceived = 0, not conceived = 1) and litter size were, in addition, analyzed by least squares procedures using the following statistical model :

$$Y_{ijk} = \mu + m_i + f_j + (mf)_{ij} + e_{ijk}$$

where :

μ = an overall mean ;

m_i = effect of the i^{th} phenotype (horned or polled) in the male ;

f_j = effect of the j^{th} phenotype (horned or polled) in the female ;

TABLE 1

Percentages of male, female and intersex offspring by parental phenotype
Pourcentages de mâles, de femelles et d'intersexués selon les types d'accouplements

Parental phenotype		Progeny No.	Males		Females		Intersexes		χ^2 (1)
Male	Female		No.	%	No.	%	No.	%	
polled	polled	684	356	52.0	287	42.0	41	6.0	1.15
polled	horned	400	221	55.3	179	44.7	—	—	4.41
horned	polled	515	271	52.6	244	47.4	—	—	1.42
horned	horned	730	389	53.3	341	46.7	—	—	3.16
Overall		2 329	1 237	53.1	1 051	45.1	41	1.8	9.02 *

(1) Ho : male to female ratio 50:50 (intersexes considered as genetic females).

* Significant at the 5 p. cent level.

$(mf)_{ij}$ = interaction effect of the i^{th} phenotype (horned or polled) in the male with the j^{th} phenotype (horned or polled) in the female ;

e_{jik} = random element assumed normally and independently distributed with zero mean and variance σ_e^2 .

Results

All males used in the present study had a least some offspring, which shows that none of them was totally infertile. A simple microscopic test of sperm obtained by electroejaculation at the beginning of the mating season may have substantially contributed to this end, since males having ejaculates with no sperm, or with no live sperm, were discarded.

All polled males but one had horned offspring, which shows that they were heterozygous for the *P* gene. The only male with no horned progeny served only 5 goats of which two conceived, giving birth to 4 polled kids.

The percentage of horned kids in the offspring of polled \times polled and horned \times horned matings was 27.4 and 100.0, respectively. The corresponding value for matings having one parent horned was 49.5 p. 100. The percentage of male, female and intersex offspring in the progeny of the various mating groups is given in table 1. Intersexes (6.0 p. 100) were found only when both parents were polled. There was a consistent excess of males in the sex ratios, the overall proportion (53.1 p. 100) being significantly different from the 50 : 50 ratio of the null hypothesis.

Table 2 refers to the conception rate by parental phenotype. Conception rate was significantly higher ($p < 0.01$) in goats mated to horned males than in those mated to polled males.

TABLE 2
Conception rate by parental phenotype
Taux de conception selon le type d'accouplement

Parental phenotype		Matings No.	Conceptions		Failures		χ^2 (1)	
Male	Female		No.	%	No.	%		
polled	polled	578	374	72.2	144	27.8	11.25 **	
polled	horned	325	233	71.7	92	28.3		
horned	polled	346	271	78.3	75	21.7		
horned	horned	520	413	79.4	107	20.6		
polled		843	607	72.0	236	28.0		
horned		866	684	79.0	182	21.0		
	polled	864	645	74.7	219	25.3		
	horned	845	646	76.4	199	23.6		
Overall		1 709	1 291	75.5	418	24.5		0.75

(1) H_0 : Complete independence of the two classifications.

** Significant at the 1 p. cent level.

The variation of litter size according to parental phenotypes is given in table 3. Polled goats gave significantly larger litters than horned goats. Goats mated to horned males produced more live kids at birth, but differences in actual litter size, i.e. stillbirths included, were insignificant. Horned \times polled matings yielded 1.92 and polled \times polled matings 1.73 kids per kidding.

TABLE 3

Means and standard errors of litter size at birth according to parental phenotypes
Moyennes et écart-types de la taille de la portée à la naissance selon les types d'accouplement

Parental phenotype		Litter size at birth					
		all born kids			live born kids		
Male	Female	Observ. No.	Mean	S.E.	Observ. No.	Mean	S.E.
polled	polled	372	1.84	0.034	364	1.77	0.033
polled	horned	233	1.73	0.043	224	1.69	0.042
horned	polled	271	1.92	0.039	264	1.86	0.038
horned	horned	411	1.78	0.032	399	1.73	0.031
polled		605	1.78	0.026	588	1.73	0.026
horned		682	1.85	0.025	663	1.80	0.024
	polled	643	1.88	0.025	628	1.82	0.025
	horned	644	1.75	0.025	623	1.71	0.025
Overall		1 287	1.82	0.018	1 251	1.76	0.018

The significance of the effect of the horn phenotype on conception rate and litter size is tested in table 4. The conception rate of horned females was similar to that of polled females. In addition, no interaction effects among parental phenotypes were found.

TABLE 4

Mean squares and tests of significance for the effect of parental phenotype on conception rate and litter size

Test de signification de l'influence du type d'accouplement sur le taux de conception et la taille de la portée

Source of variation	Conception rate		Litter size at birth			
			All born		Live born	
	d.f.	M.S.	d.f.	M.S.	d.f.	M.S.
Polled or horned male (A)	1	1.9467**	1	1.2889	1	1.5627*
Polled or horned female (B)	1	0.0039	1	4.8427**	1	3.3302**
Interaction (A \times B)	1	0.0263	1	0.0248	1	0.1504
Error	1 705	0.1840	1 283	0.4220	1 247	0.3907

* $p < 0.05$ ** $p < 0.01$

Discussion

The results of the present study show that all polled males were probably heterozygous for the *P* gene. Since the fertile polled females can be assumed heterozygous (*Pp*), the frequency of the gene in the various matings is well established.

If we name γ_1 and γ_2 the expressivity of resp. the pseudomales and the intersexes among *PP* females (LAUVERGNE's terminology) a frequency of $.5 \times .25 \gamma_1$ of excess males to the .5 sex ratio and of $.5 \times .25 \gamma_2$ of intersexes is expected in polled \times polled (*Pp* \times *Pp*) matings.

The estimation of γ_1 (.16) is far lower from the range of values in alpine breeds : .43 to .56, according LAUVERGNE (1969) after SOLLER & ANGEL (1964) and RICORDEAU & LAUVERGNE (1967). Moreover a similar discrepancy in the sex ratio was also observed in the other mating types where no excess of males was expected. In line with the results of HANCOCK & LOUCA (1975) the excess of males was consistent in all mating types.

Contrary to γ_1 the value of γ_2 , already reported by HANCOCK & LOUCA (1975) fits well the previous observations.

The difference in litter size between horned and polled does is in agreement with the findings of SOLLER & KEMPENICH (1964) and RICORDEAU (1969), who reported larger litter size in polled (*Pp*) than in horned goats (*pp*), and concluded that the *P* gene in the heterozygous condition may have heterotic effects on female fertility. The superiority of the *Pp* female in terms of prolificacy can be measured in terms of the *t* parameter of LAUVERGNE (1969). The values of *t* for, total kids born and kids born live only, were 1.074 and 1.064 (tables 3, 4), respectively, and are in good agreement with the value of 1.066 reported by LAUVERGNE (1969) for alpine breeds. The value at the conception level not being significantly different (table 2).

Such an advantage, however, was not found for fertility of heterozygous polled males. The coefficient of fertility (r_2) for *Pp* genotypes compared to *pp* genotypes was 0.96 for all kids born and 0.96 for kids born live (table 3). These values are somewhat lower than the value of 1.007 given by RICORDEAU (1969). Since conception failures where the polled males are also handicaped (see table 2) were determined using information on returns to service, abortions and kidding dates only, early embryonic death could be considered as contributing to the differences found between the two groups.

Conclusions

The behaviour of the *P* gene in the female *Damascus* goat fits grossly the theory of the behaviour of the *P* gene established earlier. Nevertheless, the consistent excess of males in all matings cannot be explained on the basis of the current theory. Moreover, an advantage in fertility of horned over polled males was found in this study.

These discrepancies indicate the need for a more accurate description of the complex behaviour of the P gene in the goat.

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Résumé

L'influence du gène motte sur le taux de conception et la taille de la portée de la chèvre Damasquine

L'influence du gène motte sur le taux de conception et la taille de la portée a été étudié à Chypre en race caprine *Damasquine* à partir de données de 1 709 accouplements et de 1 247 parturitions.

A une exception près, on a pu montrer que tous les mâles mottes considérés dans l'étude étaient hétérozygotes *Pp* pour le facteur d'absence de cornes *P*.

Il y avait un excès significatif de mâles dans la descendance de tous les types de croisement (53,1 p. 100) mais on a trouvé des intersexués seulement dans les descendance de parents mottes tous les deux (*Pp* × *Pp*).

Le pourcentage d'intersexués correspond à une expressivité $\gamma_2 = 0,48$ du gène *P* pour cette manifestation parmi les femelles *Pp*. Dans les races alpines, la valeur de γ_2 était comprise entre 0,44 et 0,57.

Le pourcentage de mâles en excès correspond à une expressivité $\gamma_1 = 0,16$ du gène *P* pour cette manifestation chez les femelles *PP*. Dans les races alpines, γ_1 est compris entre 0,43 et 0,56. Théoriquement, ces mâles en excès ne devraient pas se retrouver dans les croisements cornu × motte ou cornu × cornu, qui ne produisent aucun génotype *PP*. Cette anomalie, de même que la faible valeur de γ_1 n'a pas reçu d'explication.

Le taux de conception des femelles *Pp* était de 74,7 p. 100, pas significativement différent de celui des *PP* (76,6 p. 100). Les tailles des portées à la naissance pour respectivement tous les chevreaux et les chevreaux nés vivants, étaient de 1,88 et 1,82 pour les chèvres mottes *Pp*, alors qu'il était de 1,75 et 1,71 pour les chèvres cornues. Cela donne un coefficient de surprolificativité de resp. 1,074 et 1,006 pour les femelles *Pp* comparées aux *pp*. Ce chiffre est à rapprocher du chiffre de 1,06 observé en races alpines.

Il existe un léger mais néanmoins significatif désavantage des mâles mottes *Pp* comparés aux mâles cornus *pp* quant à la fertilité, le coefficient de fertilité comparé étant $r_2 = 0,91$ alors qu'en race alpine l'avantage est dans le sens contraire $r_2 = 1,007$. Cet écart se manifeste déjà lors de la conception.

Ces légers écarts entre les comportements du gène en races alpines et en race *Damasquine* indiquent que des études plus précises sont nécessaires.

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