

Genetic determinism of maternal behaviour in sheep and relationship with lamb mortality

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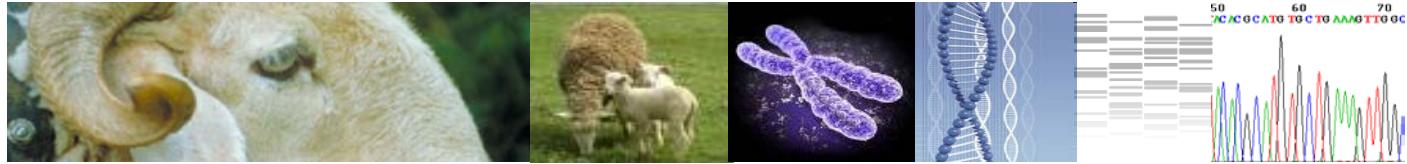
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CONTEXT



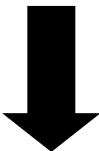


Context

Global overview

- 1) Lamb mortality remains a major concern for sheep production.
- 2) Evolution of sheep farming systems includes :

- increased size of flock
 - reduced support provided by human
 - use of harsh environments
- A teal curly brace is positioned to the right of the last two items, spanning vertically from the bottom of the word "environments" up to the end of the list, with the text "unfavourable for lamb survival" written next to it.



Survival depends on the
successful interaction
between ewe and lamb.



- Improving of maternal behaviour can contribute to increase behavioural autonomy and lamb survival



Context

Genetic for maternal behavior (previous work in sheep)

- maternal behavior varies markedly between breeds in sheep
(Dwyer, 2008, Boissy et al., 2005)
- genetic variability within breed poorly characterized
 - maternal behaviour score : $h^2 < 0.15$
(Lambe et al. 2001,
Everett-Hinks et al, 2005)

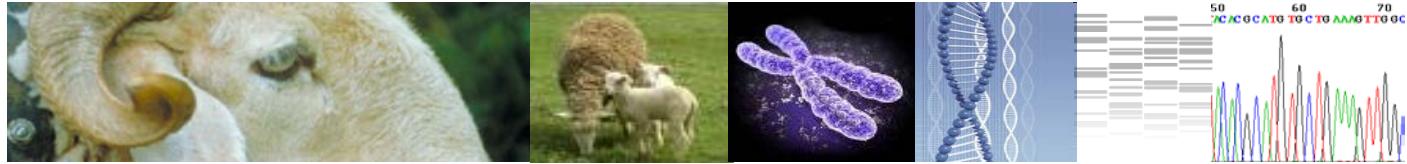
Genetic selection for maternal behaviour should offer opportunities for improving the behavioural autonomy of ewe and lamb survival.

Purpose of this study

To investigate genetic variability and to map QTL for maternal behavioural traits



Materials & Methods





Materials & Methods

Animal material:

- Romane breed (Berrichon x Romanov)
- flock raised outdoors all along the year
fed on rangelands (1 ewe/ha)
at the **experimental farm La Fage** (Roquefort)



Causse du Larzac

- Genetic parameters :
 - 1285 phenotyped ewes (along 10 years)
 - recorded for 1 to 3 lambing

- QTL mapping:
 - 9 half-sib families (103 individuals per family)
 - n=933 genotypes and 470 phenotyped ewes



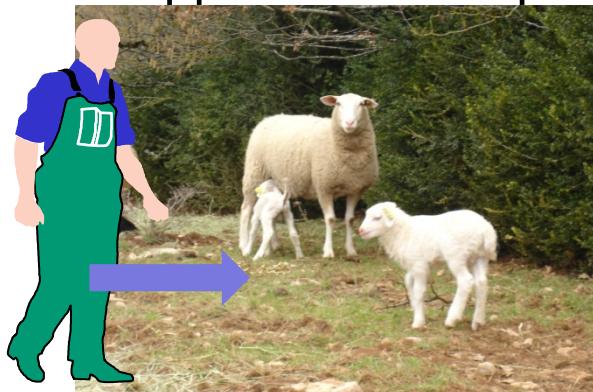
Materials & Methods

Two experimental tests for behavioural measurement :

Handling test (+2h)

Both tests aim to evaluate maternal attachment to the lamb

1. Approach of shepherd

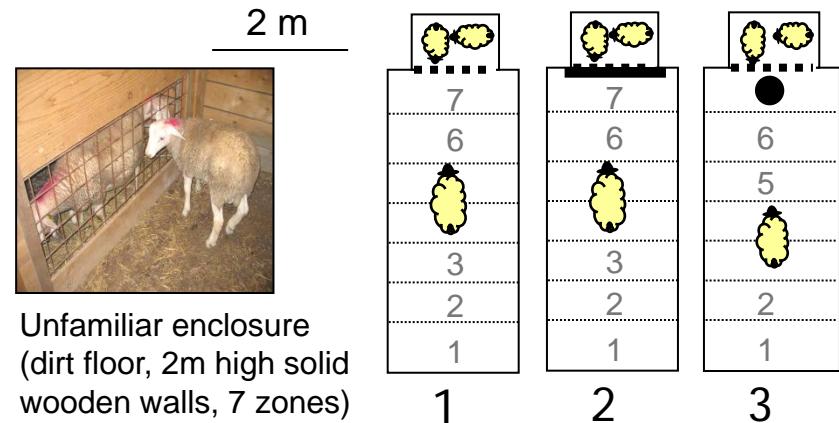


2. Handling of lambs



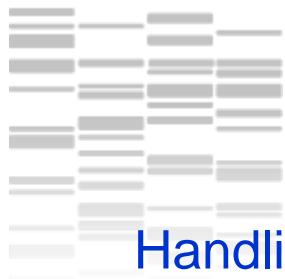
Shepherd leaves with lambs.

Arena test (+24h)



- 1: litter behind a grid-barrier, ewe introduced
- 2: opaque panel pulled down to separate litter and ewe.
- 3: panel pulled up and a human enters and stays in front of the grid

Behaviours recorded during 1min (each phase).



Materials & Methods

Behavioural traits

Handling test

Approach (of the shepherd)

- Maternal Behaviour Score (MBS1)

- 1- ewe flees and does not return
- 2- ewe retreats and comes back
- 3- ewe retreats with lamb and comes back
- 4- ewe retreats and returns repeatedly
- 5- ewe stays close to the lambing area

Handling (of lambs by the shepherd)

- Maternal Behaviour Score (MBS2)

- 1- ewe flees
- 2- ewe stays close to the lambing area
- 3- ewe follows but keeping a distance
- 4- ewe follows staying close to the shepherd

Arena test 1/2/3

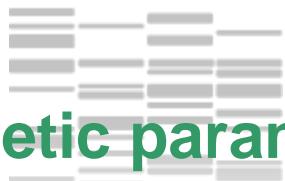
- High Bleats (number)

- Low Bleats (number)

- Locomotion (number of zones entered)

- Proximity score with the litter and the human

(score computed depending upon time spent
in different zones)



Materials & Methods

Genetic parameters

➤ **Methodology:** Variance and covariance components were estimated by restricted maximum likelihood (ASREML)

➤ **Animal model:** $Y_{ijklmn} = \mu + \underbrace{\text{year}_j + \text{lsr}_j + \text{age}_k + \text{parity}_l}_{\text{Fixed effects}} + \underbrace{\text{ewe}_m}_{\text{Environmental effect}} + \underbrace{\text{animal}_n}_{\text{Additive genetic effect}} + e_{ijklmno}$

QTL mapping

➤ **Genotyping :** - Illumina OvineSNP50K beadchip



- SNP quality control (call rate, HW disequilibrium, MAF...)
- 43820 SNPs for analysis

➤ **Phenotypes:** corrected for fixed effects, took into account repeated measurements

➤ **Methodology:** likelihood computation every 0.1 cM (QTLMAP)

3 methods: - linkage analysis (LA)

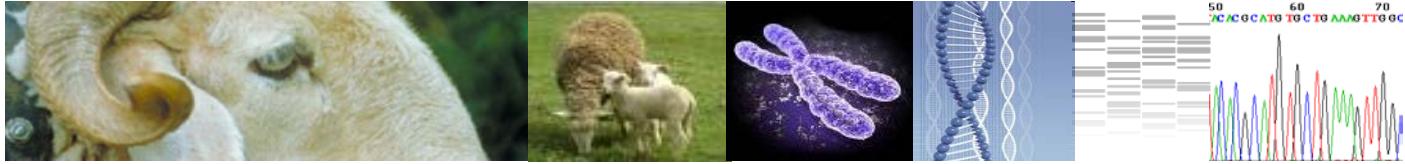
- association analysis (GWAS)

- linkage and association joint analysis (LDLA)

➤ **Thresholds** computed with permutations

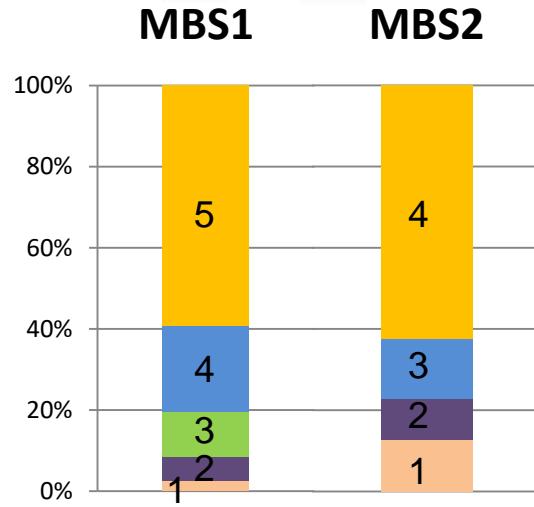


Results



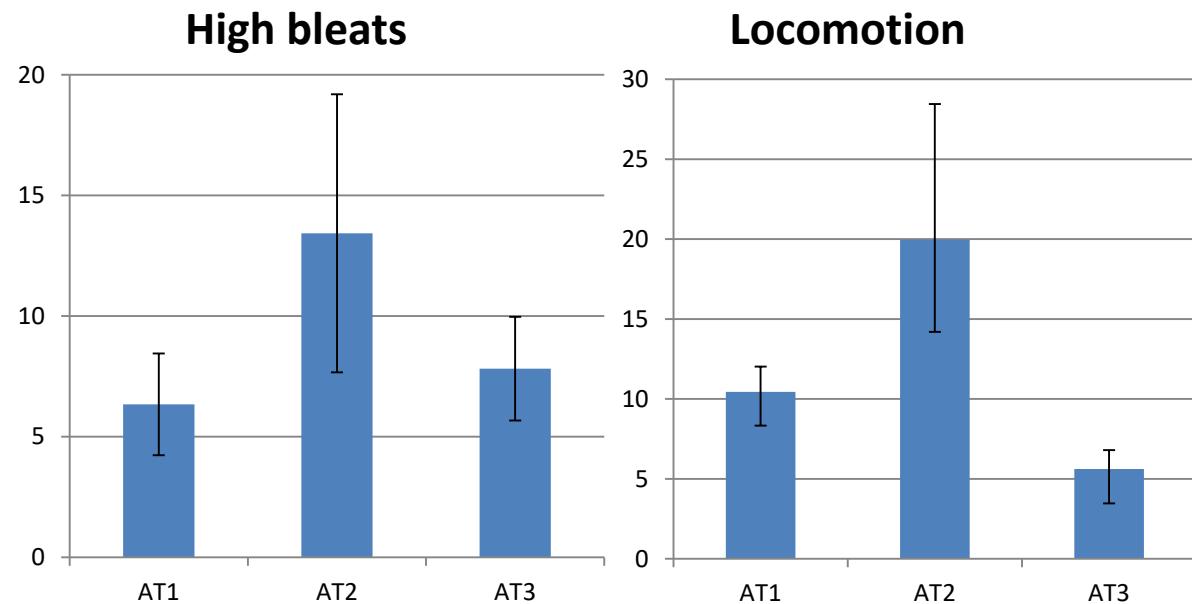
Results

Handling test



➤ Higher MBS represent the “better mothers”.

Arena test



➤ Higher reactive ewes to social separation are expected to be the “better mothers”.

➤ Phenotypic variability for maternal reactivity to social separation from litter.



Results Genetic parameters

Handling test

	MBS1	MBS2
MBS1	0,23 +/- 0,06	0,82 +/- 0,13
MBS2		0,19 +/- 0,05
r	0,38 +/- 0,06	0,30 +/- 0,06

Diagonal: heritability; r, repeatability

➤ Moderate to high heritability estimates for traits of maternal attachment.

Arena test

	LB (1)	HB (1)	HB (2/3)	Locom (2/3)	Prox (3)
LB(1)	0,44 +/- 0,05	-0,74 +/- 0,05	-0,24 +/- 0,10	NS	-0,29 +/- 0,12
HB(1)		0,44 +/- 0,05	0,70 +/- 0,07	NS	NS
HB(2/3)			0,46 +/- 0,05	NS	0,30 +/- 0,12
Locom (2/3)				0,15 +/- 0,04	NS
Prox (3)					0,27 +/- 0,04
r	0,58 +/- 0,06		0,60 +/- 0,06	0,30 +/- 0,06	0,42 +/- 0,06

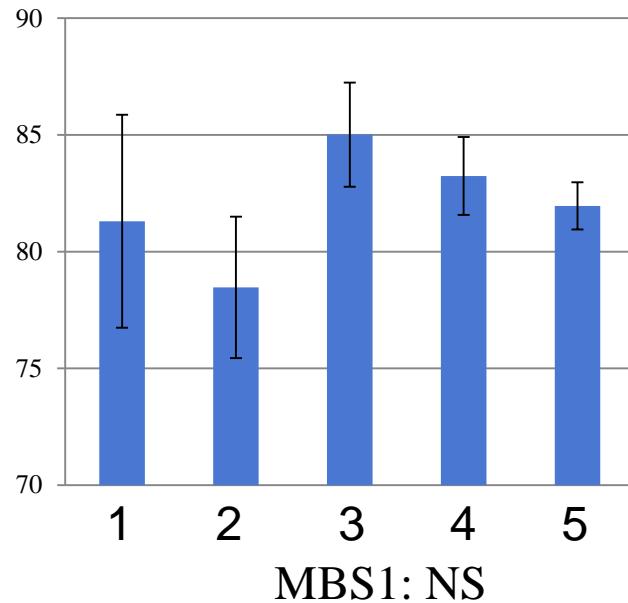
Diagonal: heritability; r, repeatability



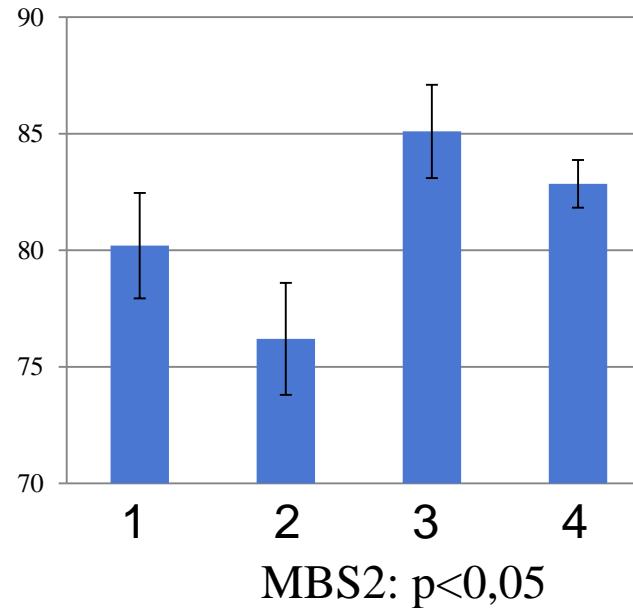
Results

Relationships with lamb mortality

Litter survival (lsmeans, %)



Litter survival (lsmeans, %)

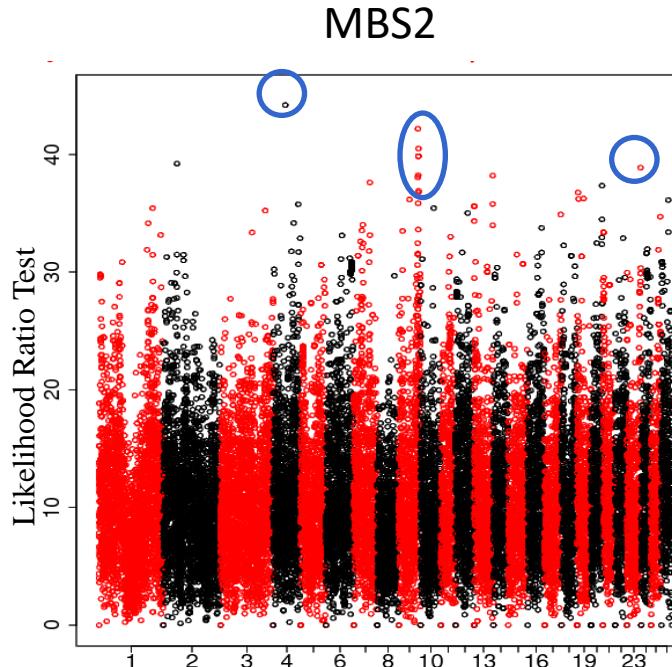


➤ Litter survival increased as MBS increased.

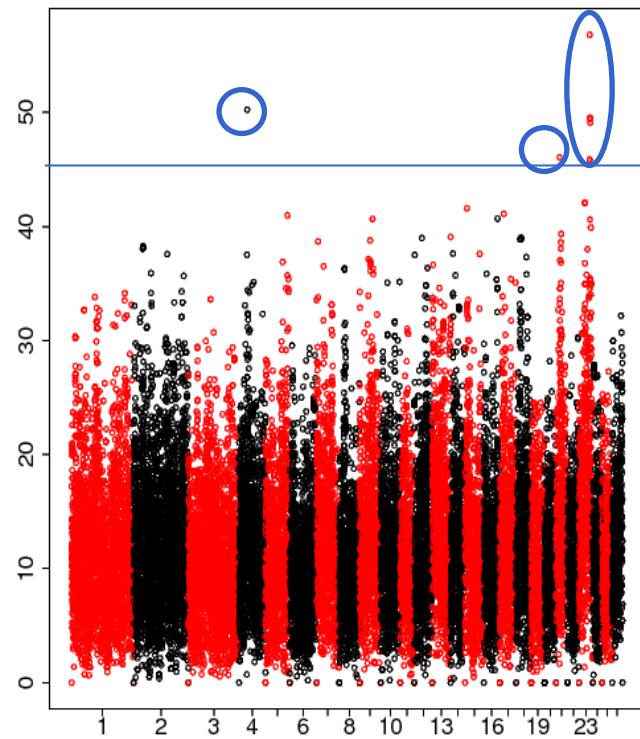


Results QTL mapping

Trait	OAR
MBS1	4, 9
MBS2	4, 9, 23
LB(1)	4, 21, 23
HB(1)	16, 21, 23
HB(2/3)	4, 16
Locom(2/3)	1, 11, 19
Prox (3)	1, 5, 18

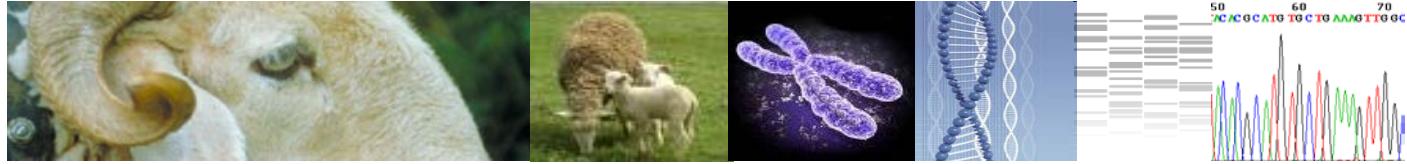


Low bleat (Arena test 1)





Conclusion





Conclusion

Analytical issue:

- Several moderate to highly heritable traits associated with maternal reactivity for litter
- Several significant QTL associated with maternal reactivity

These results will:

- lead to a better knowledge of genetic variability of maternal behaviour in sheep
- help to understand behavioural interaction between ewe and lamb

Applied issue: This work presents a relevant approach that could be used to improve maternal behaviour of the ewes by genetic selection and to increase lamb survival

Perspectives: Further work required

- to investigate relationship between handling test and arena test
- to dissect underlying genetic mechanisms (to perform multitraits QTL analyses, to search candidate genes ...)



Acknowledgements

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- **INRA** for funding phenotypes



Thank you for your attention !

