

Genetic structure of the French Blonde d'Aquitaine, Charolais and Limousin beef cattle populations

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Aug. 24th 2009

60th EAAP, Barcelona

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Three major beef breeds in France



- 77% of beef cows in France in 2007
- Huge changes in beef cattle populations
 - Demography
 - Artificial Insemination (AI) use
 - Breeding programs
- Assessment of genetic diversity by pedigree analyses
 - Inbreeding trends
 - Probability of gene origin approach

Pedigree analyses carried out with Pedig (Boichard, 2002)

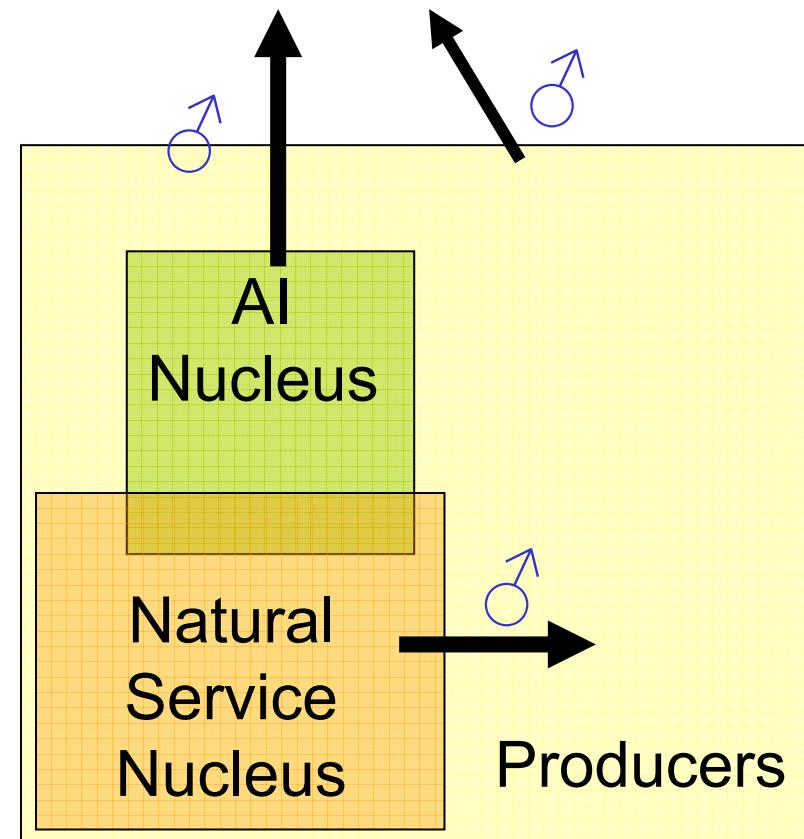
http://www-sgqa.jouy.inra.fr/article.php3?id_article=110

Presentation of populations

	Recorded calves	Pedigree
Blonde d'Aquitaine (BLA)	807 221	7 500 000
Charolais (CHA)	2 992 246	20 700 000
Limousin (LIM)	2 142 788	12 500 000

↓
Recorded
population =
~15% of the whole
French population

AI breeding program



Pedigree knowledge

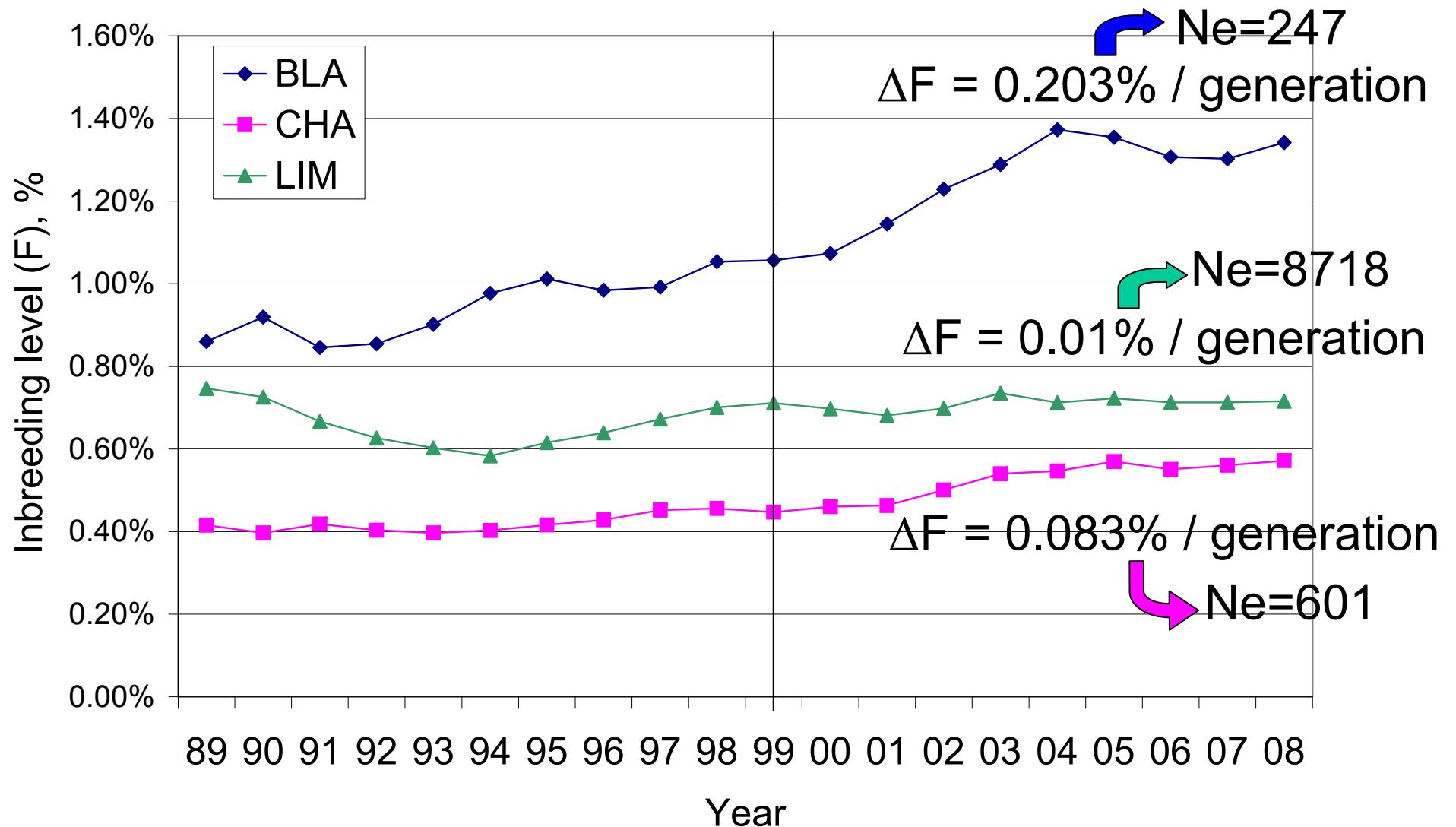
Complete Generation Equivalents : CGE = $\sum_{\text{Generations}}$ %known ancestors

	BLA	CHA	LIM
Recorded Pop	6.0	8.7	7.2
Producers	5.9	8.4	7.2
NS Nucleus	6.9	10.2	8.3
AI Nucleus	6.6	9.8	8.3

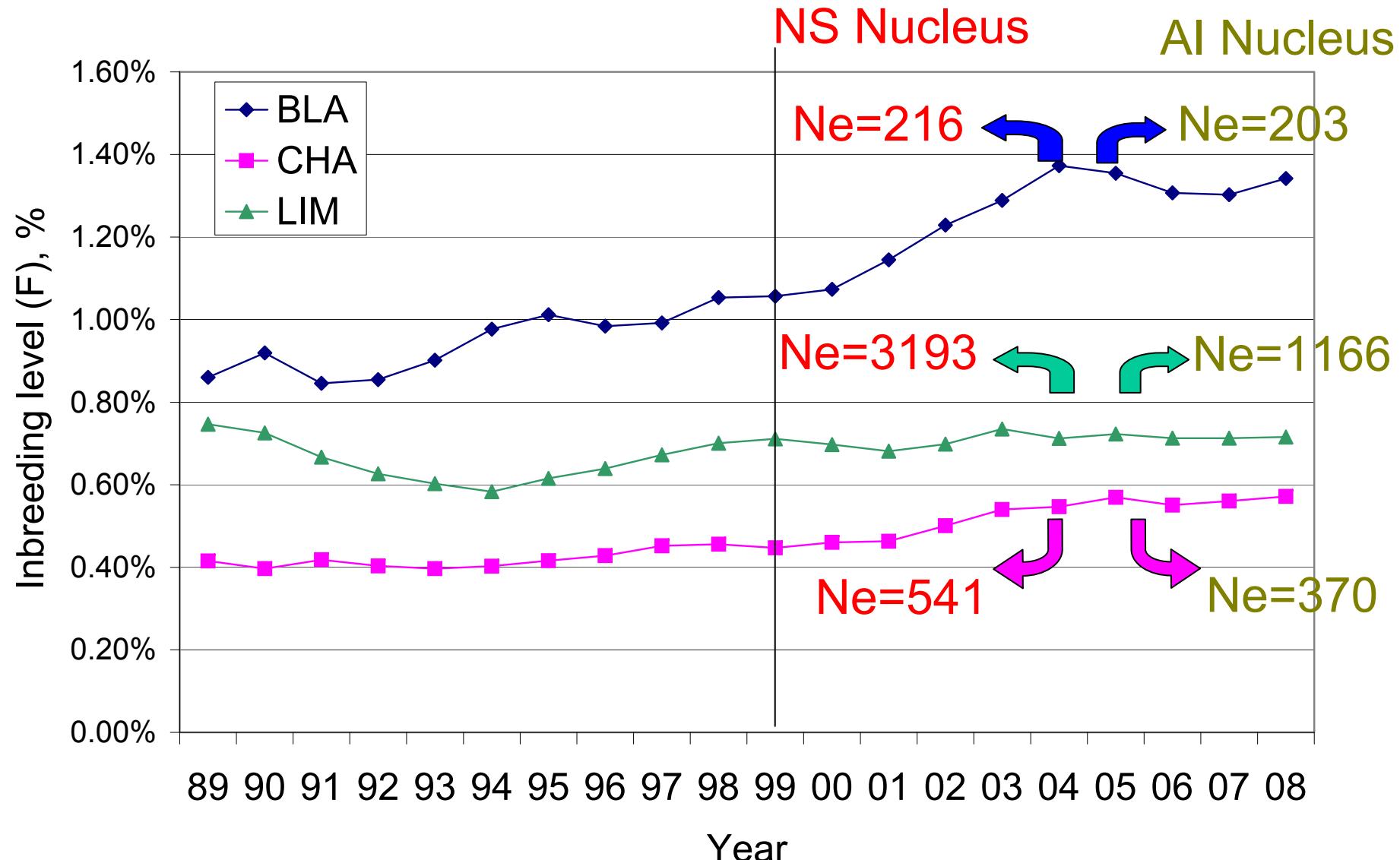
Inbreeding (F) & effective size (Ne)

- Inbreeding calculated with Meuwissen & Luo 's algorithm
- Annual rate of increase in inbreeding calculated by regression on the birth year of calves (t):
 $\ln(1-F)=t*\ln(1-\Delta F_a) + k$ (Perez-Enciso, 1995)
- Effective size: $Ne= 1/(2 L\Delta F_a)$, with L=generation interval

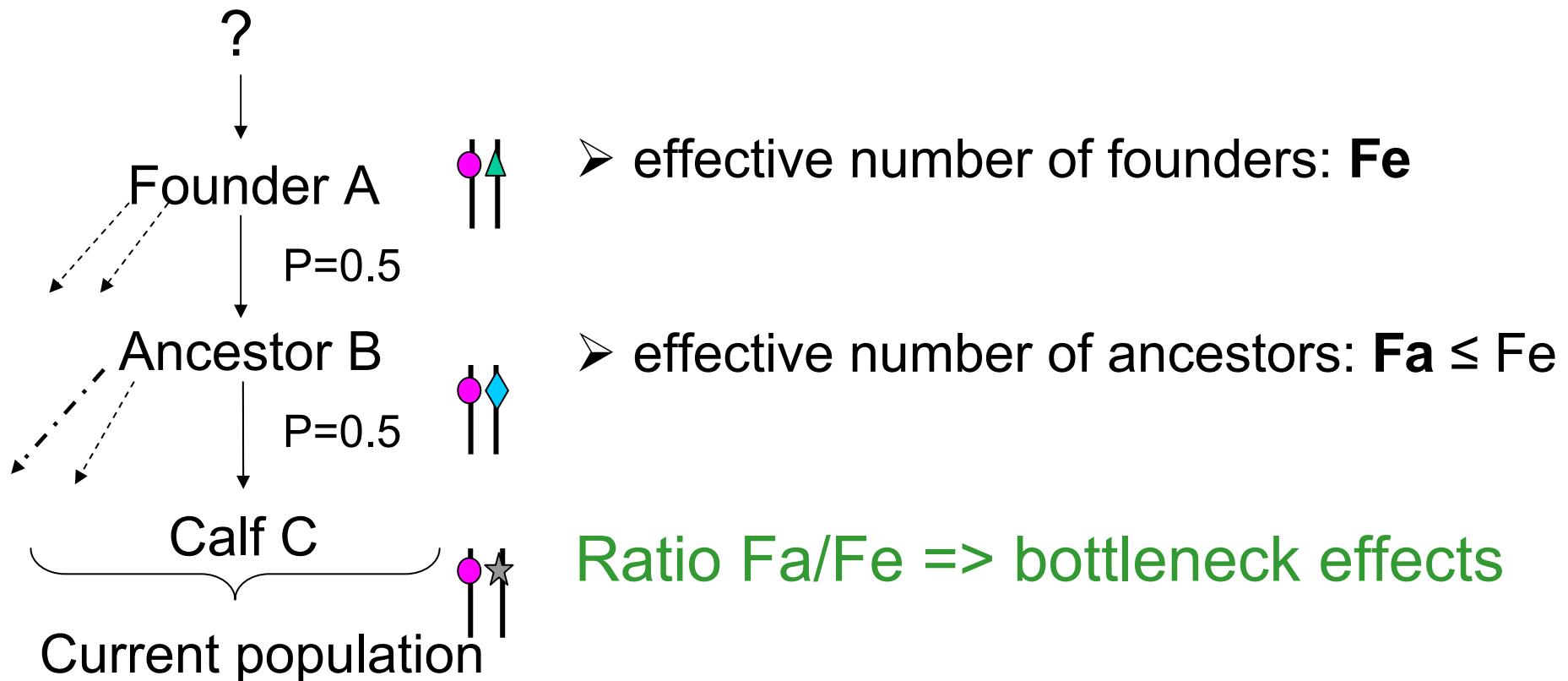
Inbreeding trends (ΔF) and effective size (Ne)



Inbreeding trends (ΔF) and effective size (Ne)



The probability of gene origin approach



Genetic diversity within populations

- Statistics calculated for calves born
 - between 1984-1988, 1994-1998 & 2004-2008
- Ratio Fa / Fe:
 - CHA: 79% > 63% > 48%
 - LIM: 47% > 39% > 28%
 - BLA: 41% > 39% > 37%

Conclusions

- Low inbreeding trends & high effective sizes
- Existence of strong bottlenecks
 - BLA: old bottlenecks but stable
 - LIM: old bottlenecks getting worse
 - CHA: recent bottlenecks
- Take care for the future !



Thank you for your attention!

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Evolution of the AI rate

