

Are there limits to selection in poultry ? theoretical, biological, ethical, environmental ?

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Selection is at the beginning of the poultry chain : Which types of animals are needed ? how are the breeding goals defined ? What diversity of genotypes should be offered ?

Does selection always reach its goals ? Litterature on selection 'plateaux' : →Selection may exhaust genetic variability But →Evolutive mechanisms maintain variability

Selection may reduce fitness \Rightarrow unrealized selection response





Main issues

- Selection theory and possible limits
 - Principles and Experimental evidence on poultry
- Biological limits and decreased fitness
 - Physiological constraints
 - Unfavorable correlated responses
- Ethical limits
 - Welfare issues and social acceptance
- Environmental limits
 - Climate change and limited natural resources



Quantitative genetics theory

An infinite number of genes with additive effects: simple and powerful !

Selection response:

R = h² **S** (h² heritability = σ_a^2 / σ_p^2) S selection differential (function of σ_p^2 and fitness)

Additive genetic variance at a locus $\sigma_a^2 = 2 \sum_i p_i q_i (a_i - b_i)^2$ depends on allelic effects and allelic frequencies (a - b) = effect of substituting one allele by the other True for a population at *equilibrium* (no selection, mutation, migration)

⇒ Selection is modifying allelic frequencies





Effects of selection on genetic variance

Change in p: $\Delta p = i pq (a - b) / \sigma_p$ i = selection intensity,

Selection effect will be maximal for intermediate values of p and for large substitution effects (major gene) When allelic frequencies are very unbalanced, selection will be less and less effective

- \rightarrow complete fixation seems very unlikely or very slow
- \rightarrow AND a large number of loci is controlling selected traits





Deviations from the simple model of additivity

Dominance effect :

the genetic value of an heterozygous is different from the mean of the values of both homozygous \rightarrow d may be negative or positive $\sigma_d^2 = 4 \sum_i (\mathbf{p}_i \mathbf{q}_i \mathbf{d}_i)^2$

Interaction between genes : epistasis

Additive by additive or additive by dominance, or dominance by dominance

Interactions between genetic effects and environmental effects :

statistical G x E epigenetic mechanisms

\Rightarrow These sources of variation influence response to selection





Genome dynamics

Mutation: the typical evolutive force maintaining diversity, takes place at a low frequency, can generate genetic abnormalities

Recombination : regular event, 'the life of the genome' !
Genes may act as clusters, some regulating the action of others, even at distance ; recombination may break such clusters and create new variation



This has been proven in plants, by genotyping molecular markers across generations

Mobile elements, endogenous retroviruses
 Selection limits may not be observed



Importance of population size for predicting selection response

Population size limits the gene pool

Small population size increases the impact of random events : genetic drift may become as important as selection pressure

Ne: effective population size represents the size of a population in equilibrium showing the same Δp as the observed population \Rightarrow Prediction of ultimate selection response R according to Ne

	N males N females		Ne ≈	50% R at
Approximate value of Ne	50	200	160	224 gen
	10	50	33	47 gen





Selection in practice

Several factors are preventing gene fixation during selection :

- > Evolving selection objectives, adding new traits
- Moderate selection pressure, slow process
- Large base population
- New lines arising from crossbreeding
- > at the opposite, inbreeding (F) is limiting the selection response
 - \blacktriangleright Within population σ^{2}_{a} is linearly decreasing with F
 - Increased frequency of genetic abnormalities

A higher ratio of deleterious mutations/neutral mutations in the most inbred of 3 layer lines (Derks et al., 2018) ⇒ may limit selection response by decreasing fitness





First take-home message

Selection is unlikely to exhaust all components of genetic variation

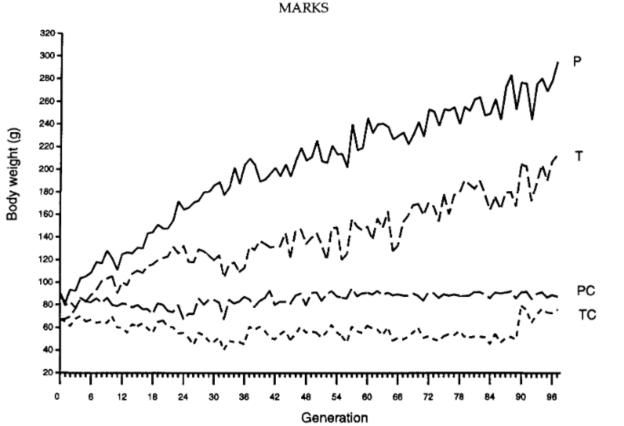
but inbreeding is the main risk for long-term selection response





Evidence from experimental lines

The longest selection experiment on 4 wk body weight in quails (Marks, 1996)



97 generations

P = control diet 28% CP T = 20% CP PC and TC : controls

Selection response Still observed in P Much lower in T

Realised heritability h² 0.29 \rightarrow 0.11 in P h² 0.22 to 0 in T 7 ×

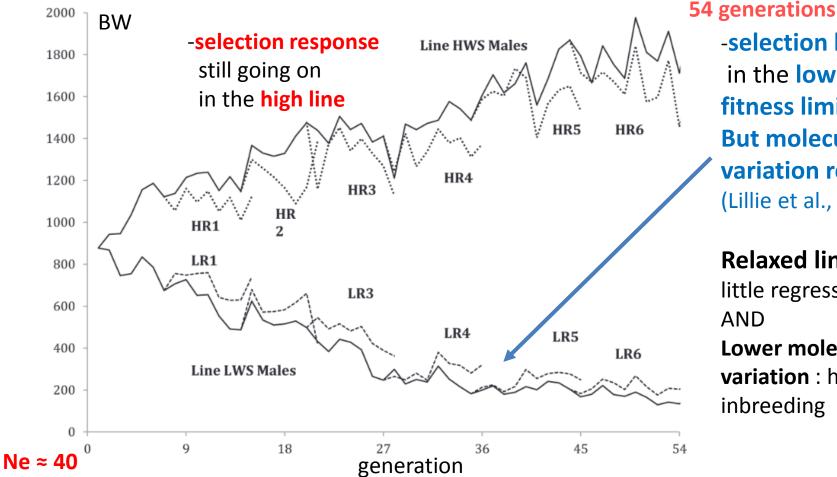
Ne ≈ 72





Evidence from experimental lines

Long term selection experiment on 8 wk BW in chickens (Dunnington et al., 2013)

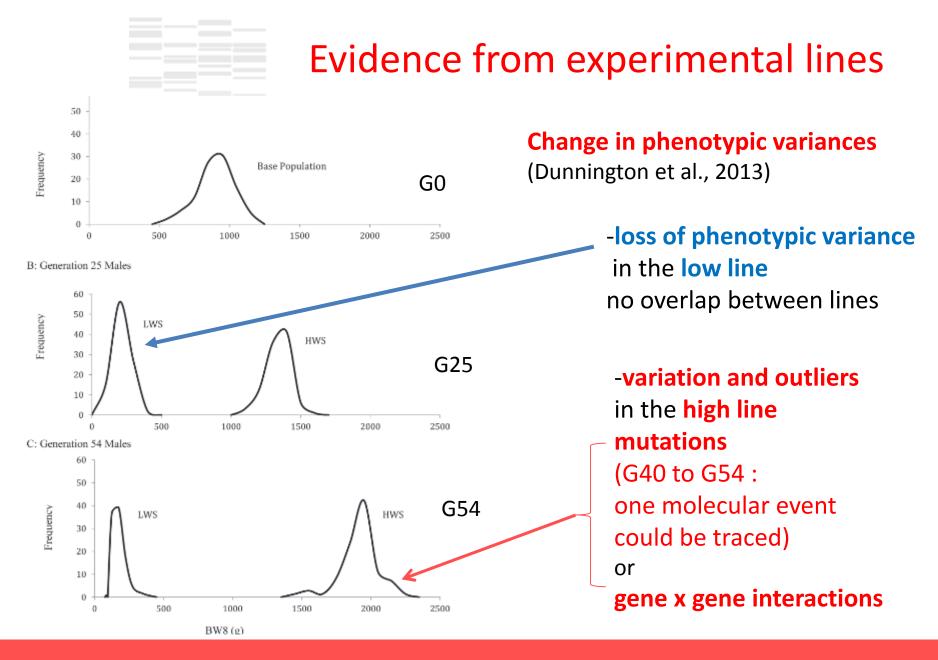


-selection limit in the **low line** fitness limits R **But molecular** variation remains (Lillie et al., 2018)

Relaxed lines little regression AND Lower molecular variation : higher inbreeding



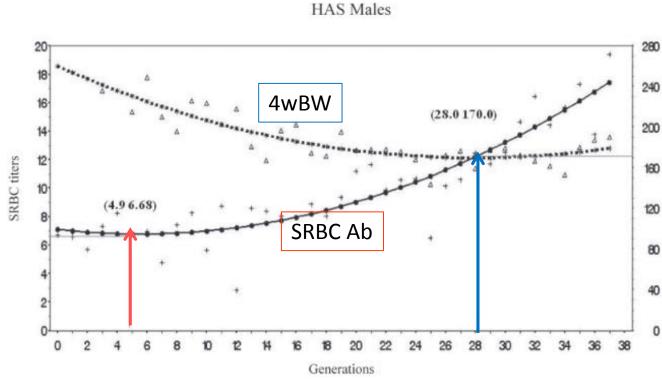
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Evidence from experimental lines

Long term selection experiment on immune response in chickens (Zhao et al., 2012)



37 generations

divergent selection on SRBC antibodies : effect on 4 wk BW in males

Trade-off with 4wk BW In males

different inflexion points according to trait

Ne ≈ 25





Lessons from selection experiments

Divergent selection experiments generally exhibit dissymetry : Limit is reached more easily in one direction (physiological limit) Whereas molecular variation remains

No evidence for selection plateau in lines selected for a high value, in spite of a decrease in additive genetic variance But number of generations generally still < 1.4 Ne

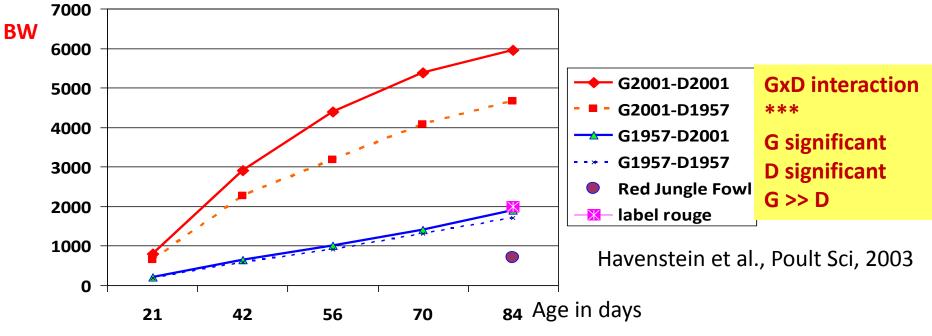
Long term selection show evidence of mutations

Changing environments can 'reveal' hidden variation





Selection response has been taking place for > 50 generations still < 1.4Ne



However, variation assessed by molecular markers indicates some loss of genetic variation particularly in layers



Within-population variability

(microsatellites Leroy et al., 2012)

Mean values	Unbiased Hetero- zygosity	Observed Hetero- zygosity	Population History	Mean Number of alleles	Effective Number of alleles
Broiler (3 lines)	0.487	0.490	Commercial GP	3.26	2.21
Layer (2 lines)	0.351	0.328	Conserved breeds	2.55	1.76
Taïwan (6 breeds)	0.489	0.488		3.33	2.20
France (14 breeds)	0.527	0.505	Traditional breeds	3.68	2.45
West Africa (23 provinces 450 birds)	0.588	0.580	Village Chickens	4.64	2.96



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Impacts of selection on genome diversity patterns: linkage disequilibrium

LD: Does SNP1 provide information on SNP2 ?

Calculate correlation coefficient r^2 between adjacent markers haploblock size for association studies = segment size where $r^2 > 0,3$

Differences reflect not only selectionWhite-egg layers

Brown-egg layers broilers

20-30 kb

SNP2

Village chickens

SNP1

Qanbari et al., 2010, Muir et al., 2008 Mwacharo et al., 2013 Qanbari et al., 2014



1-5kb



Biological limits to selection

✓ Physiological constraints :

 limiting value of a trait for survival or reproduction as illustrated by selection experiments:
 →selection limits on decreased BW or decreased antibody level
 →decreased fitness : limits the selection intensity

✓ Unfavorable correlated responses to selection:

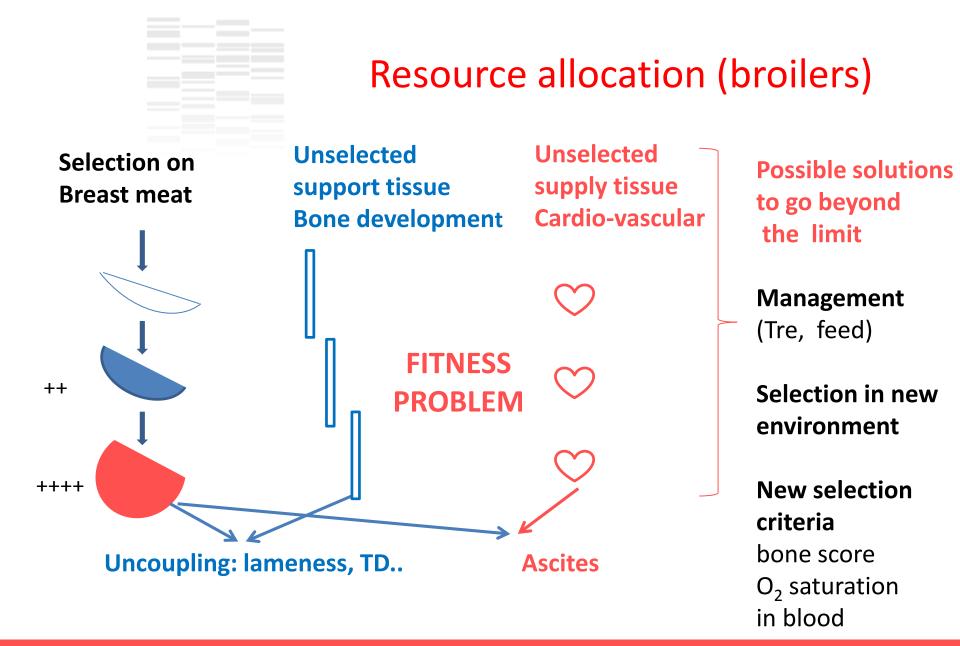
question of resource allocation

→ opposition high growth/immune response high growth/reproduction

→ decoupling between 'muscular growth' and 'supply organs'

Limit in the expression of a trait: ex = ovulation rate





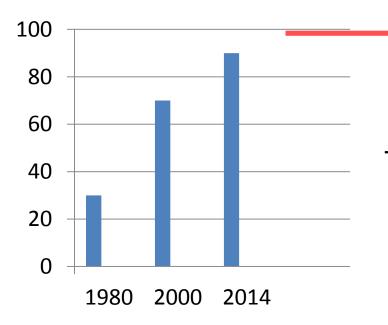




Antagonistic functions (broilers)

Fast early growth / excessive appetite /fatness / poor reproduction

→ Management solution : severe feed restriction of future breeders



Physical limit to feed restriction !

Welfare issue: hunger

The genetic correlation between growth and reproduction is not linear: too small or too large BW : poor reproduction



Antagonism growth/reproduction → genetic solution : modify growth curve (broilers)

2b. BW at 16 wk (Vg_{BW16}). 1b. BW at 4 wk (Vg_{BW4}). 1.81.8 X-+ Males A-+ Males 1.4 1.4 → X+- Males ▲— X+- Males 1.01.0 0.6 → X++ Males X++ Males VgBW16 0.6 0.2 X-- Males --- X--- Males Vgawa 0. -0.2 - X-+ Females -D- X-+ Females -0.2 -0.6-0.6 -1.0→ X+- Females -1.0-1.4 → X++ Females → X++ Females -1.8-1.4 -O-X-- Females -O-X-- Females -1.8 0 2 -3 4 5 8 9 10 11 12 13 14 6 0 12 13 14 Generation Generation 2c. BW at 20 wk (Vg_{BW20}). 1c. BW at 8 wk (Vg_{BW8}). 1.8 1.8X-+ Males A X-+ Males 1.4 1.4 → X+- Males ★ X+- Males 1.01.0 -+- X++ Males 0.6 --- X++ Males Vg_{BW20} 0.6 0.2- X-- Males --- X--- Males Vgaws 0.2-0.2 -0.2-O-X-+ Females -0.6 -0.6→ X+- Females → X+- Females -1.0-1.0 -1.4→ X++ Females → X++ Females -1.4 -1.8→ X-- Females -O-X-- Females -1.80 9 10 11 12 13 14 2 8 0 9 10 11 12 13 14 -8 Generation Generation

Mignon-Grasteau et al., 2001

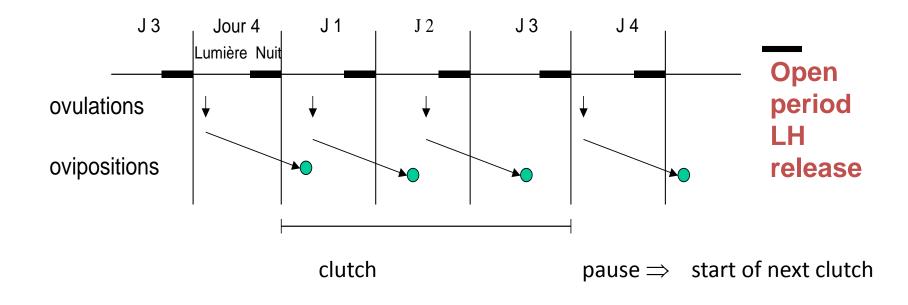
towards a biological compromise ?



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Biological limit (layers) 24h cycle and ovulatory cycle

Endogenous cycle is constrained by the day-light external cycle

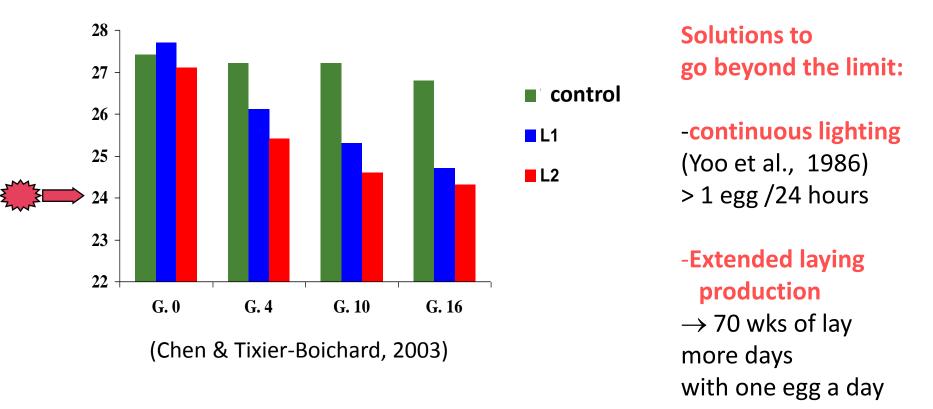






Biological limit (layers) 24h cycle and ovulatory cycle

Selection on clutch length in a dwarf brown egg-layer \rightarrow Effect on interval between ovipositions with a 16 L/ 8 D external cycle







Take-home messages about biological limits

Several examples of such biological limits illustrate that selection response can be limited by a decreased fitness

Increased frequency of metabolic disorders :

Prevents selection of individuals with high breeding value for growth

Management solutions: short term

Genetic solutions (new trait in the breeding goal):

cumulative, long term, more complex programme with a price to pay on the progress of the other traits





Ethical limits

Ethics raises questions, does not impose answers moral values and a more general view on the living world There are ethical issues raised by intensive selection : welfare : hunger in fed-restricted broiler breeders Cannibalism / debeaking in layers

⇒ Who does feel responsible for this in the production chain ? breeder? Producer? Slaughter-house, retailer ? Consumer ?

⇒ What is the main justification : to feed human population

Reactions from society : example in The Netherlands with the 'Wakker Dier' welfare association against the 'Plofkip'







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From the 'Plofkip' to the Chicken of to-morrow

Media campaign + Pressure on supermarkets

Agreement on

Maximal growth rate allowed for the chicken: **50 g/day** Has increased its market share (50% retailer) in a few years

Real limit to the selection goal

Leave room to select on other traits





Ethical limits

Relative perceptions according to countries and culture

- Technology often considered as the solution, indeed, but
 - « Higher technology calls for higher responsability »

to be considered for genome editing ! Just take some time to think of consequences !





Bird metabolism and ambient temperature :

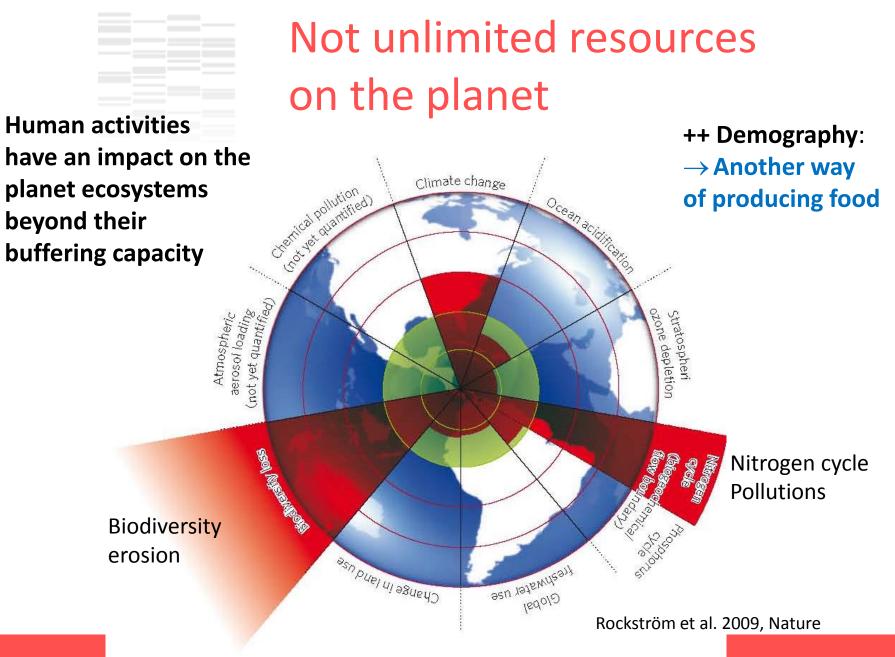
A very fast growing broiler ⇒ high metabolic heat **Confort zone** : temperature range where the animal does not need to use evaporation or adapt its metabolism to maintain its normal body temperature 20 years ago : confort around 22-23°C ambient **Nowadays: confort zone of fast growing broilers ≈ 16-18 °C ambient Higher ambient T**^{res} will limit growth

What to do $? \rightarrow$ decrease body weight objective

- \rightarrow use naked animals (Sc mutation)
- \rightarrow air conditionning (energy cost,

(humans don't get it ...)





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Green-house-gas emissions: poultry on the safe side

Feed efficiency selected with high quality feed
⇒ dependency on arable lands for high quality feed,
⇒ select on ability to digest other feeds

Competition food/feed : stronger for poultry than for ruminants

Optimization of animal proteins in human diets: 15 to 45% proteins (Van Zanten et al., 2016; Van Kernebeek et al., 2016) Data on cattle or pig production, but **few studies on chickens**





Conclusions

Selection response is still taking place

Acceleration with genomics

 \Rightarrow careful monitoring of effects on genetic variance and deleterious variants

Limit 1: extreme performance level is decreasing fitness

Genetics should be more predictive of such consequences Selection objectives should consider trade-offs between fitness and production

Limit 2 ethical responsability regarding the living world ? awareness of dependency on environment ?

> Which food systems do we want ?

- update breeding goals
- manage diversity, gene bank or live populations
- > propose a Portfolio of genotypes





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