



Molecular genetics of the adrenocortical axis and breeding for robustness

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Diet, agriculture, environment

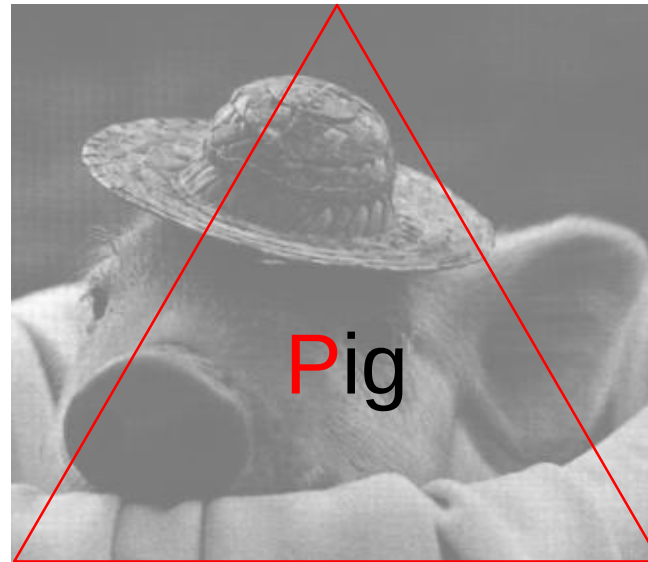
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New challenges for farm animal breeding

Sustainability

Reduced production costs
-feed efficiency
-less labour (independent animals)
-adaptability to environmental constraints

Profit



Pig

-environmental footprint
(GHG, N, P)
-antibiotics, heavy metals
& hormones

-food safety
-product quality
-animal welfare
-rural lifestyle

Planet

People

Main breeding objectives: feed efficiency, robustness, product quality

ROBUSTNESS

- ❖ The concept of robustness refers to the ability of the animals to express their production potential in a wide range of environmental conditions without compromising their physical health and welfare (Knap 2005)
- ❖ Includes several components
 - Sensitivity of the production potential
 - ✓ to diverse commercial environments
 - ✓ and to external stressors (temperature)
 - Functional traits
 - ✓ quality of supportive tissues (e.g. leg soundness)
 - ✓ newborn survival
 - ✓ resistance to diseases (efficiency of the immune system)
 - ✓ productive longevity

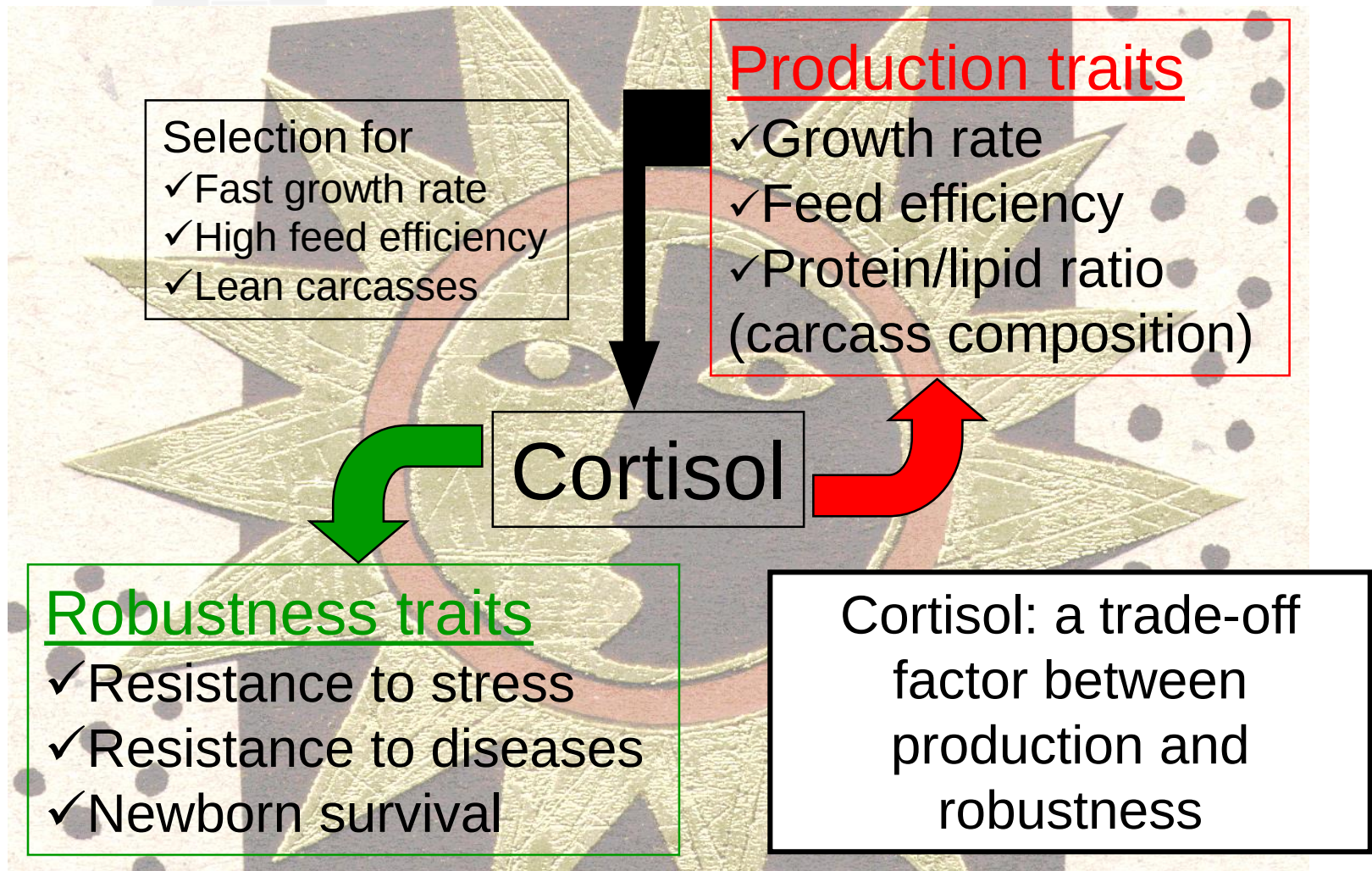
Robustness as a breeding objective

‘Sustainable breeding goals combine robustness traits with production traits to such an extent that selection balances genetic change in production potential with genetic change in environmental sensitivity’ (Knap 2009)

Selection strategies

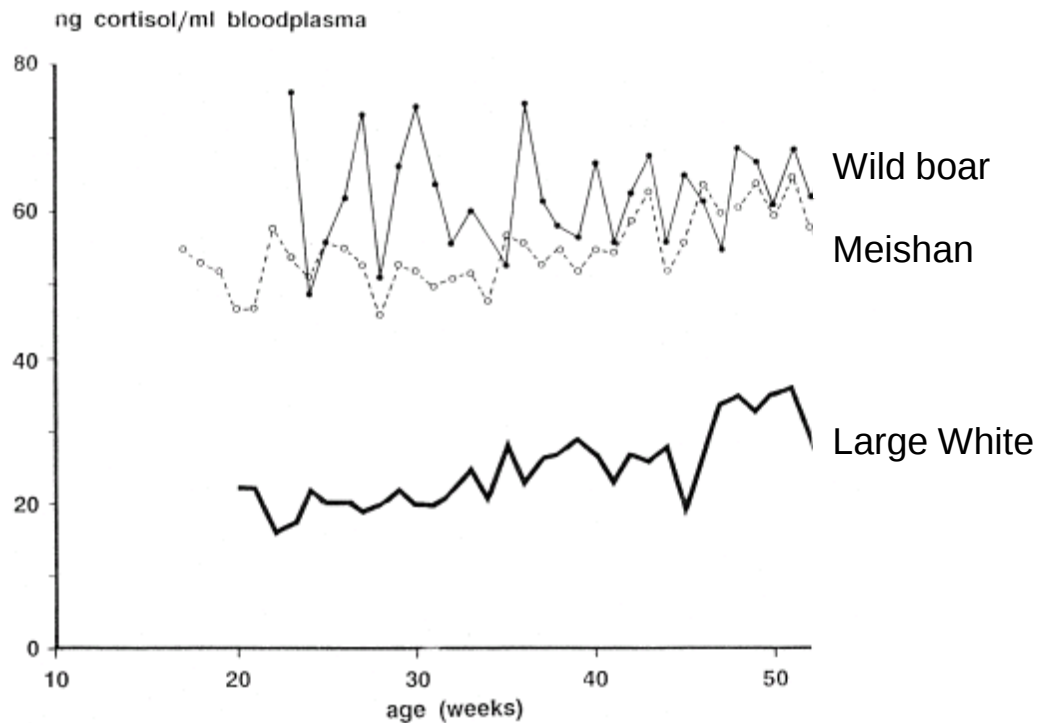
1. Inclusion of directly measurable **functional traits** in the breeding objective and in the selection index (BLUP) – leg soundness, newborn survival rate, fertility, cellularity of milk ...
2. **Reaction norm** analysis – global sensitivity to the environment (Lynch and Walsh 1998) – low heritability
3. **Canalizing selection** for a reduced variance
4. Genetics of **stress responses**
behavioural and **neuroendocrine**

Effects of cortisol on production and robustness traits



Domestication and the HPA axis

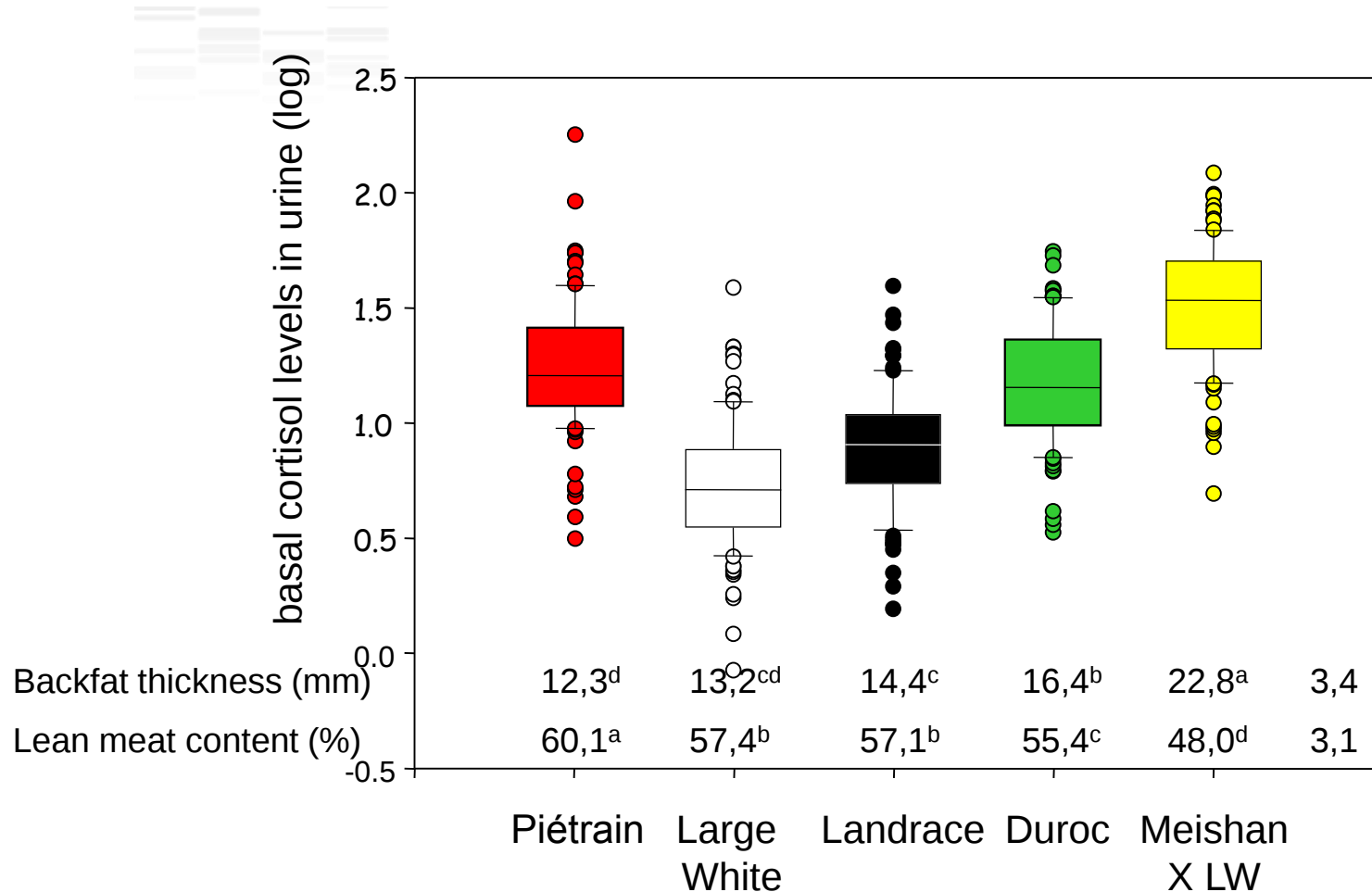
U. Weiler et al. / Livestock Production Science 54 (1998) 21–31



Breeding objectives to increase robustness by selection of a stronger HPA axis

- ❖ Selection for fast growth rate, high feed efficiency and lean carcasses may have reduced HPA axis activity with negative consequences on robustness
- ❖ Working hypothesis : selection for a more active / reactive HPA axis could improve robustness with (hopefully) limited effects on production traits.
- ❖ Strong genetic influences on HPA axis activity
 - Differences between genetic stocks (e.g. Large White / Meishan)
 - Divergent genetic selection in chicken (ACTH, social stress), turkey (cold), duck (suspension), trout (confinement), mouse (restraint).
- ❖ Several sources of genetic variation
 - Cortisol production (sensitivity to ACTH)
 - Bioavailability
 - Receptor function

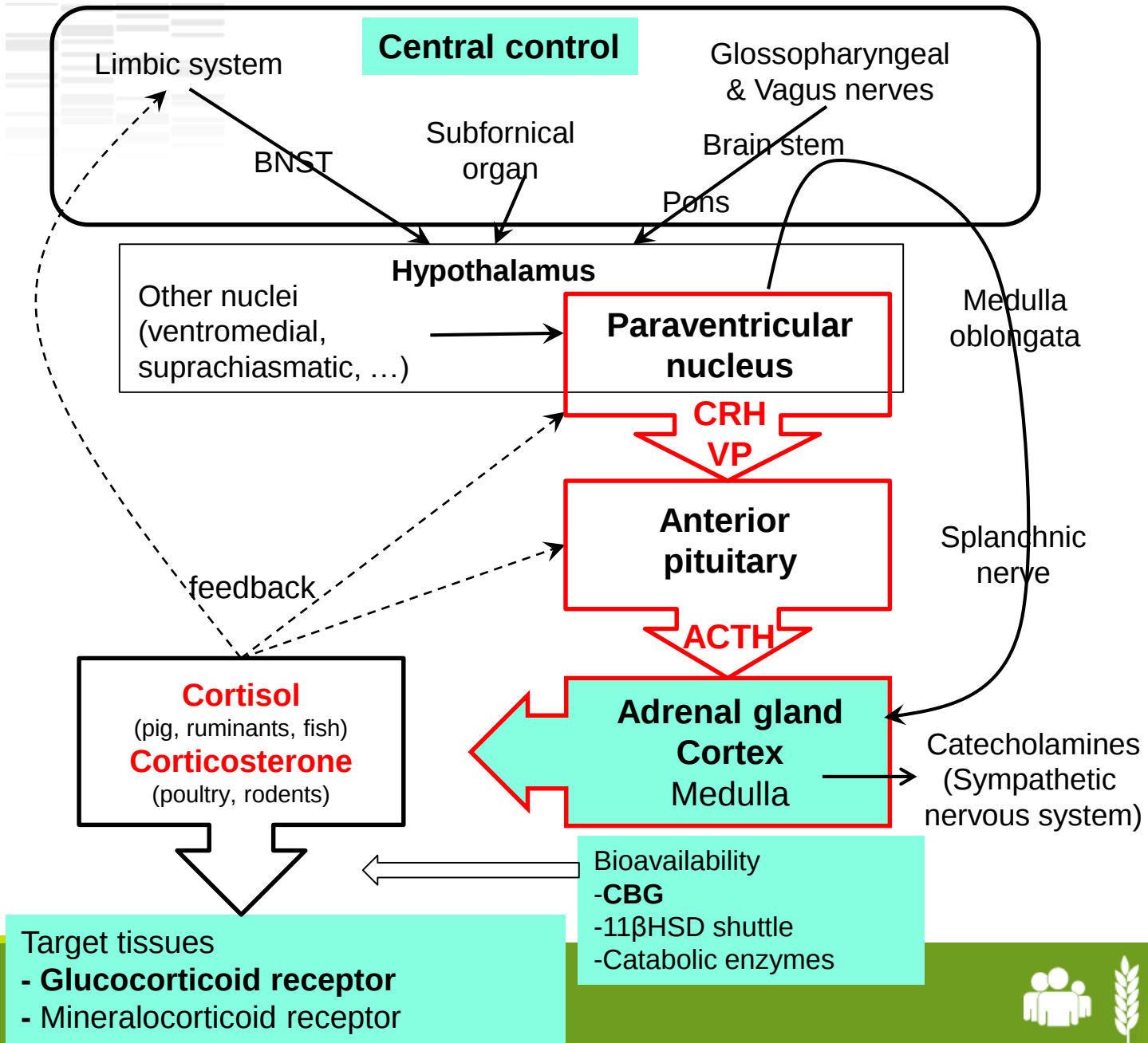
Between- and within-breed variation of cortisol production



QualityPorkGenes – EU FP5
Foury et al. Animal 1:973, 2007

N = 100 female pigs/ breed

HPA axis, sources of genetic variability



Sources of genetic variability in the HPA axis

CORTICOSTEROID HORMONE PRODUCTION

- Response of the adrenal cortex to ACTH

- ❖ Is the main source of genetic variation in cortisol production
- ❖ Is an individual trait (stable across time)
 - Pigs [Hennessy et al. 1988](#)
 - Humans - 'Adrenal phenotype' [Bertagna et al. 1994](#); [Coste et al. 1994](#)
- ❖ Is heritable
 - $h^2 = 0.68$ - analysis in 30 litters of Large White pigs
 - [Larzul et al. Animal 9:1929, 2015](#)
 - Divergent selection
 - Direct selection in chicken, response to ACTH ([Edens et Siegel, 1975](#))
 - Main trait selected in response to stress (trout, duck)
- ❖ Molecular genomics
 - [Hazard et al. BMC Genomics 9:101, 2008](#)
 - [Jouffe et al. BMC Proceedings 3:S14, 2009](#)



Adrenals: differentially expressed genes



❖ No difference

- In the **ACTH signaling pathway** (i.e. ACTH receptor, G protein, Adenylate Cyclase, Protein Kinase A and MAPK, ERK1, ERK2) except *Crem* (cAMP response element modulator) that activates genes involved in steroidogenesis
- In **steroidogenic enzymes** (i.e. *Cyp11a1*, *Cyp17*, *3βhsd*, *Cyp21*, *Cyp11b*)

❖ Differentially expressed

- **Cholesterol transport** (Star, Ldlr) and several kinases involved in phosphorylation of StAR (Snf1lk)
- **Tricarboxylic acid cycle** (Mdh2, Sdha and Suclg2) involved in heme biosynthesis (Alas1)
- Still to be characterized transcripts / genes

Hazard et al. *BMC Genomics* **9**:101, 2008

Sources of genetic variability in the HPA axis

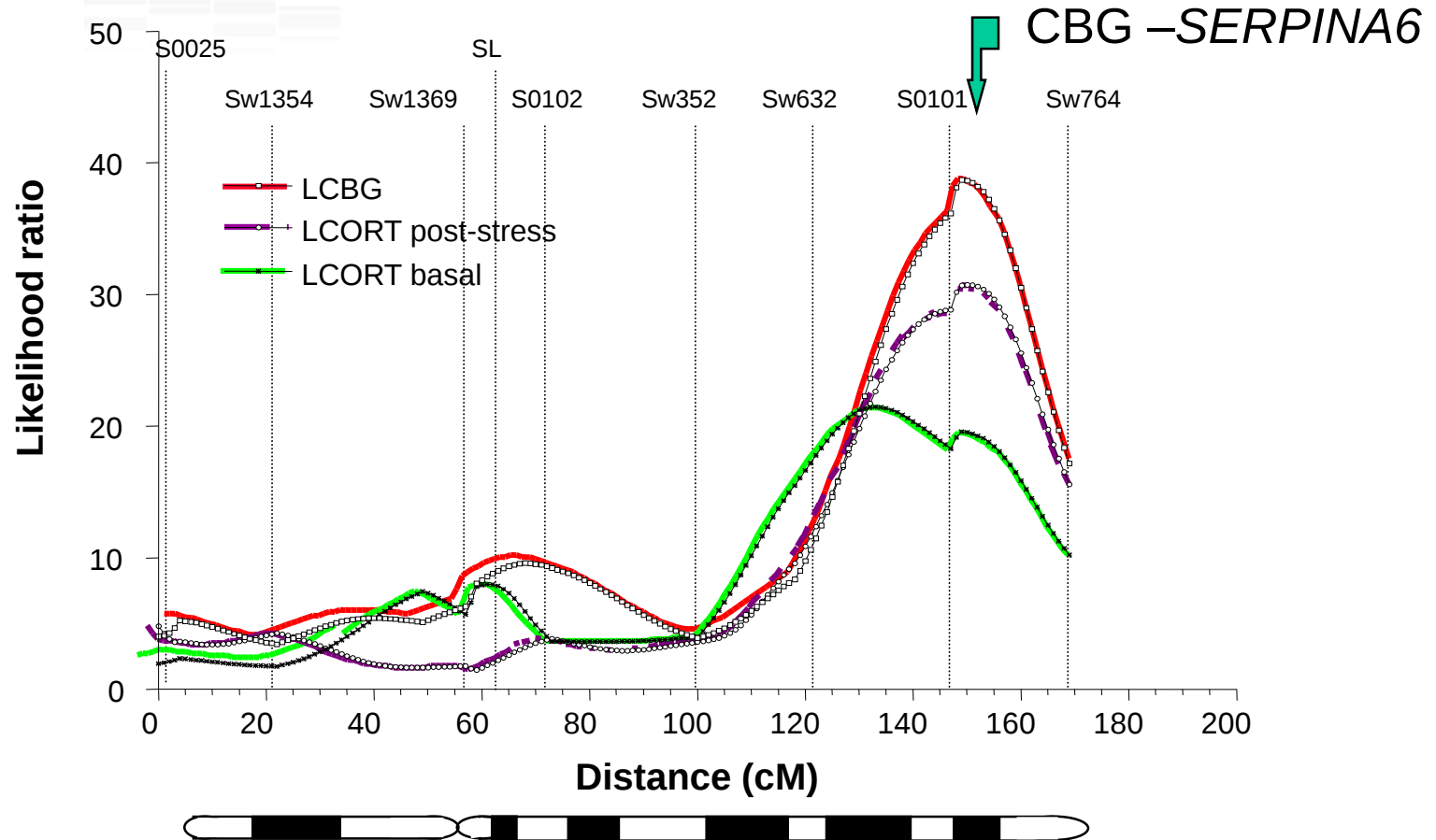
BIOAVAILABILITY

- Corticosteroid-binding globulin (CBG)
- 11 β -hydroxysteroid dehydrogenases

❖ Corticosteroid-binding globulin (CBG or transcortin)

- Glycoprotein synthesized in the liver and carrying cortisol in plasma
Plays a dual role of sequestration of cortisol in plasma and delivery to the target tissues
 - Moisan Mol Cell Endocr 316:35, 2013
- Genetic determinant of cortisol levels by linkage study in a Large White x Meishan F2 population
 - Désautés et al. J Anim Sci 80:2276, 2002
 - Confirmed in rats (Solberg et al. 2006)
and humans (Bolton et al. PLoS Genetics 2014)
- Molecular polymorphisms influencing protein properties
with consequences on adiposity and meat quality
 - Ousova et al. Mol Endocrinol 18:1687, 2004
 - Geverink et al. J Anim Sci 84:204, 2006

QTL cortisol / CBG on Ssr 7 (F2 LWxMS)



Désautés et al. *J Anim Sci* **80**:2276 (2002)

Ousova et al. *Mol Endo* **18**:1687 (2004)

CBG and production traits

Pearson Correlation Coefficients between Carcass Composition Traits, Plasma CBG Capacity, and Total Plasma Control

	CBG	Cortisol (Basal)	Cortisol (Post Stress)
Backfat weight	$r = 0.39$	$r = -0.18$	$r = 0.11$
	$P = 0.014$	$P = 0.28$	$P = 0.50$
% (Ham+ loin)	$r = -0.40$	$r = 0.18$	$r = -0.11$
	$P = 0.012$	$P = 0.27$	$P = 0.52$
% (Back+ leaf fat)	$r = 0.42$	$r = -0.17$	$r = 0.12$
	$P = 0.008$	$P = 0.29$	$P = 0.45$
Estimated muscle content	$r = -0.43$	$r = 0.16$	$r = -0.13$
	$P = 0.0064$	$P = 0.33$	$P = 0.41$

Each cell of the table contains the Pearson correlation (r) coefficient and the probability (P). The number of observations is $n = 39$ in all cases.

Sources of genetic variability in the HPA axis

BIOAVAILABILITY

- Corticosteroid-binding globulin (CBG)
- 11 β -hydroxysteroid dehydrogenases

❖ Corticosteroid-binding globulin (CBG or transcortin)

❖ Hormone metabolism

- Cortisol – cortisone shuttle (hydroxysteroid dehydrogenases 1 & 2)
- Local tissue bioavailability of cortisol may influence adiposity
- Linkage/association between HSD11B1 polymorphism and carcass length and adiposity in pigs
 - Otieno et al. Anim Genet 36:36, 2004

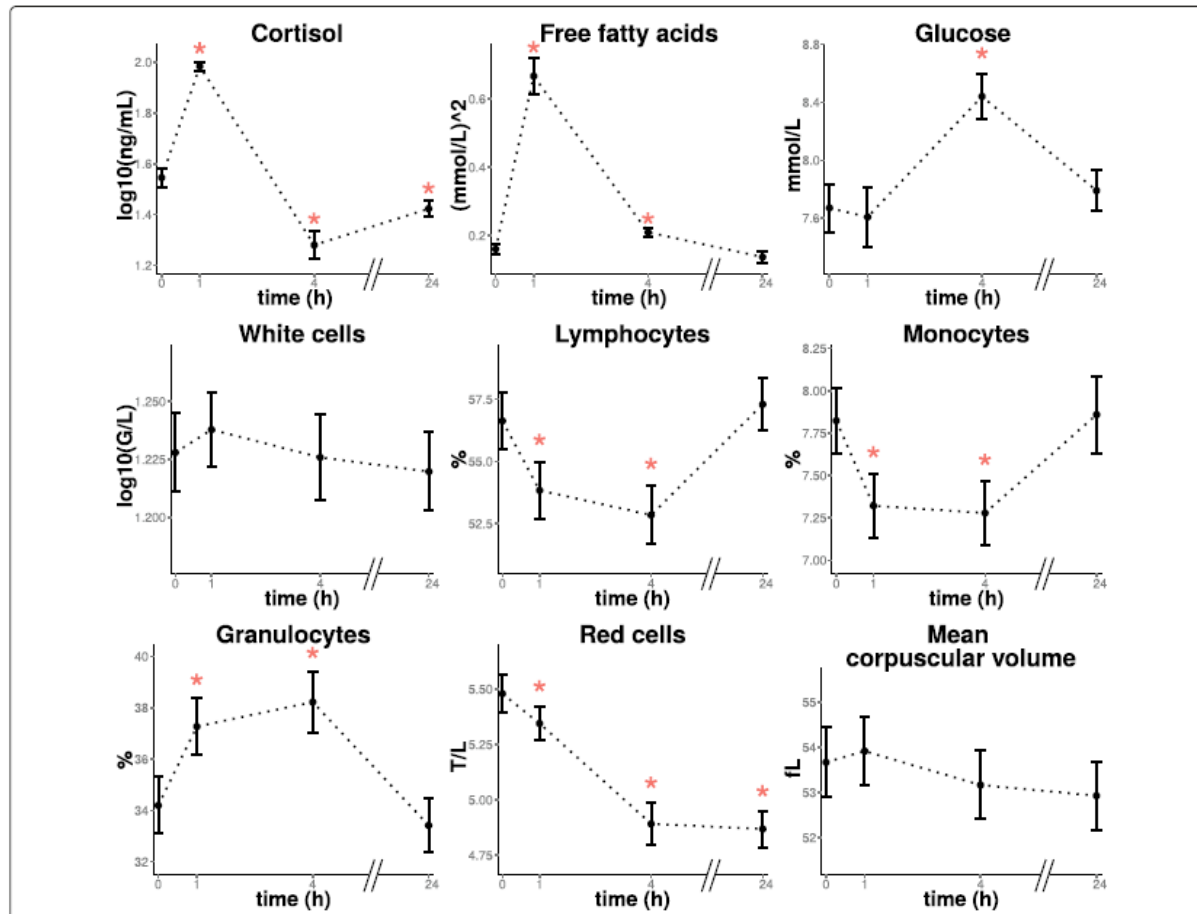
Sources of genetic variability in the HPA axis

ACTION ON TARGET TISSUES

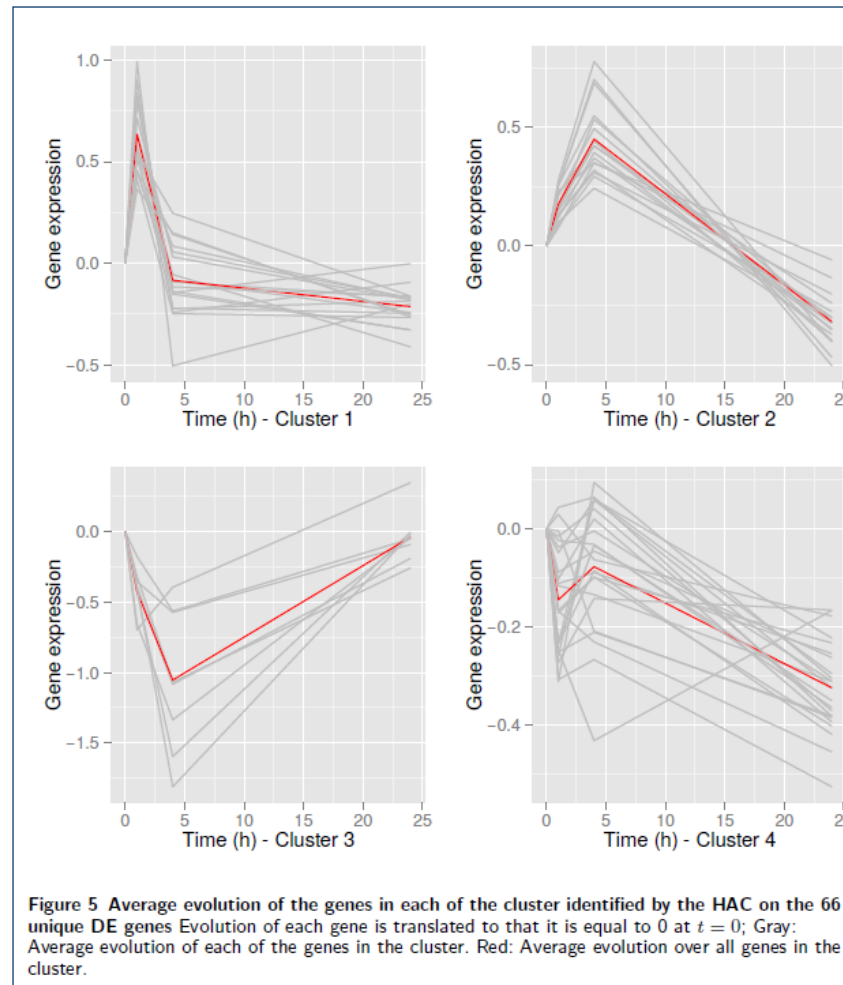
- Receptors or transduction pathways

- ❖ Cortisol receptors are transduction factors
 - Glucocorticoid receptors, GR, gene NR3C1
 - Mineralocorticoid receptors, MR, gene NR3C2
- ❖ Genetic influences and gene polymorphisms
well documented in humans and experimental animals
- ❖ Little information in farm animals
 - Functional differences LW/MS pigs
Perreau et al. Life Sci 17:1501, 1999
 - Association studies (pigs)
Murani et al. BMC Genetics 11:74, 2010; Terenina et al. DAE 44:81, 2013
 - Naturally occurring gain-of-function mutation Ala610Val
Murani et al. 2012, 2016
- ❖ Functional genomic fingerprint of cortisol effects
(e.g. circulating white blood cells)

Response to ACTH – biological measures in peripheral blood

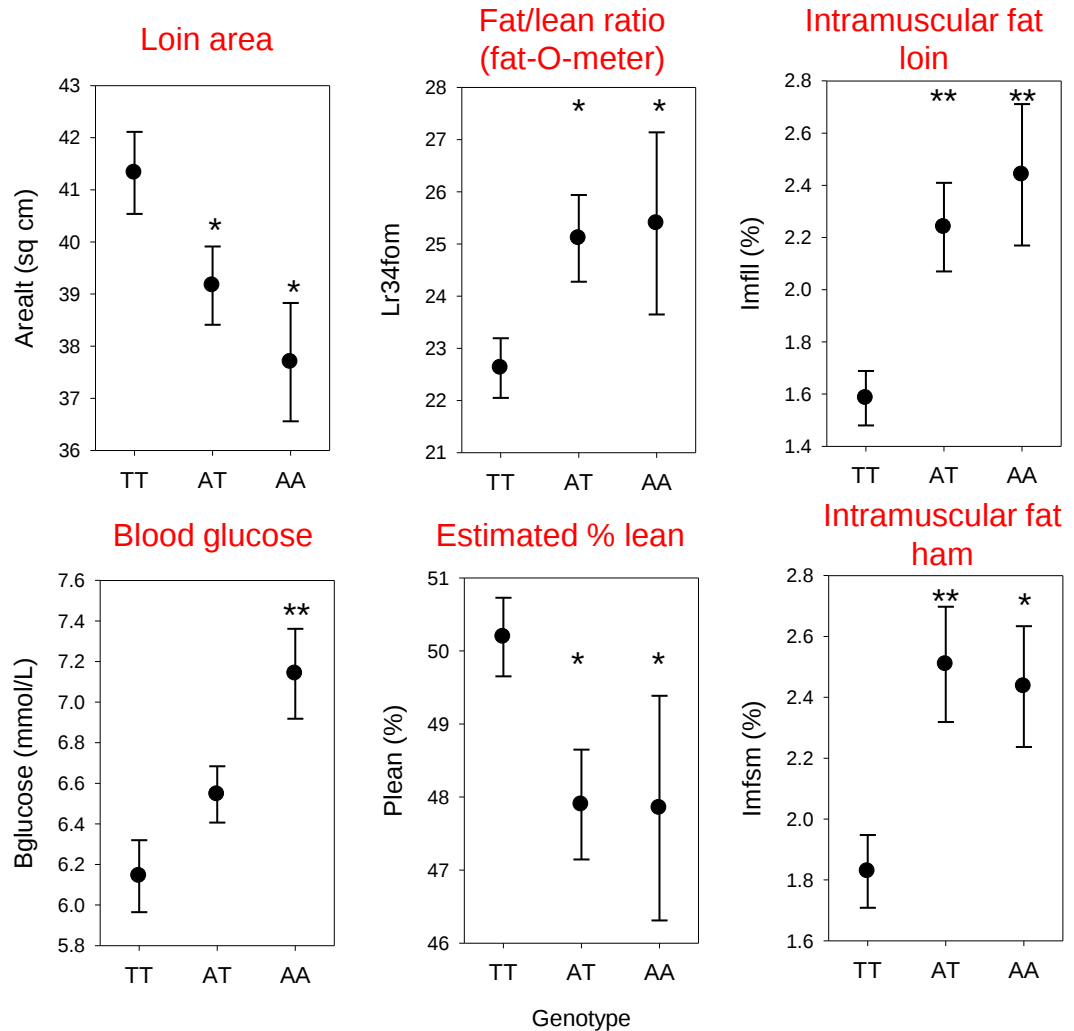


Response to ACTH - differentially expressed genes in circulating leucocytes



Phenotypic effect of the MAOA(-430A>T) SNP in the promoter region of monoamine oxidase A gene

Association study



Female pigs (n = 100) from an advanced intercross line (F9-F11) between Large White and Meishan

TT n = 40
AT n = 41
AT n = 11-13

Terenina et al. DAE 44:81, 2013

Phenotypic effect of the DRD3(-680C>T) SNP in the promoter region of dopamine receptor D3 gene

Association study

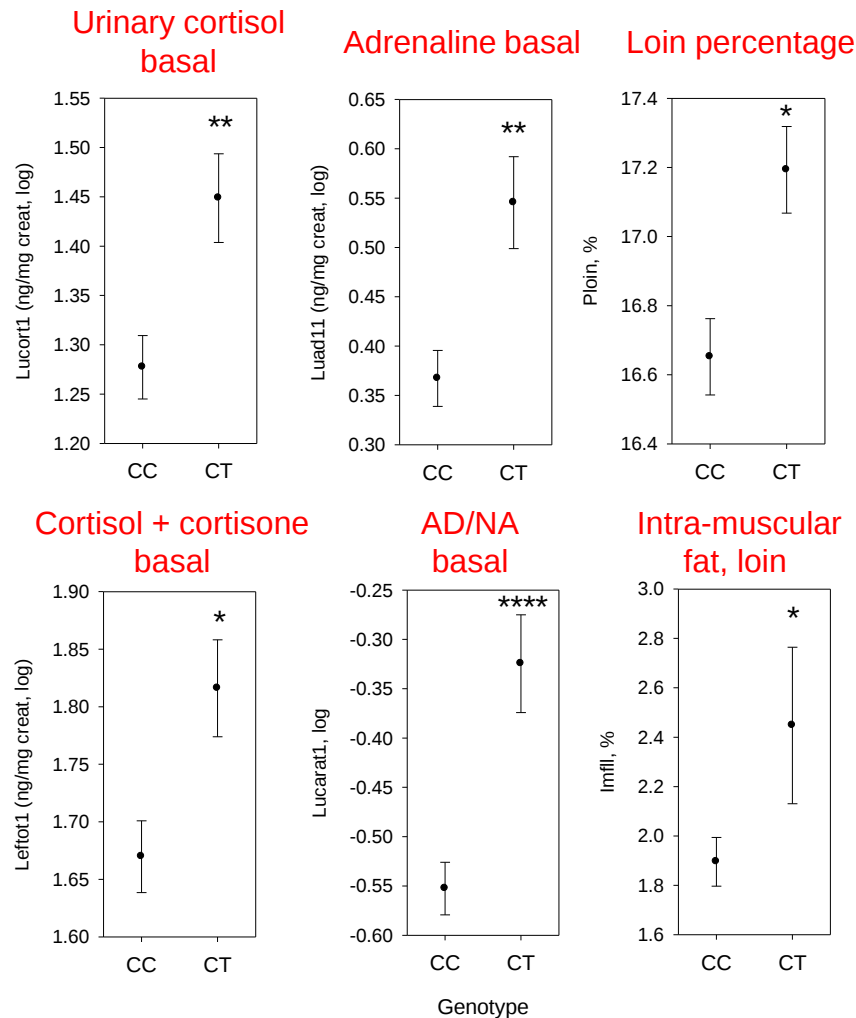
Female pigs (n = 100) from an advanced intercross line (F9-F11) between Large White and Meishan

CC n = 66-68

CT n = 13-20

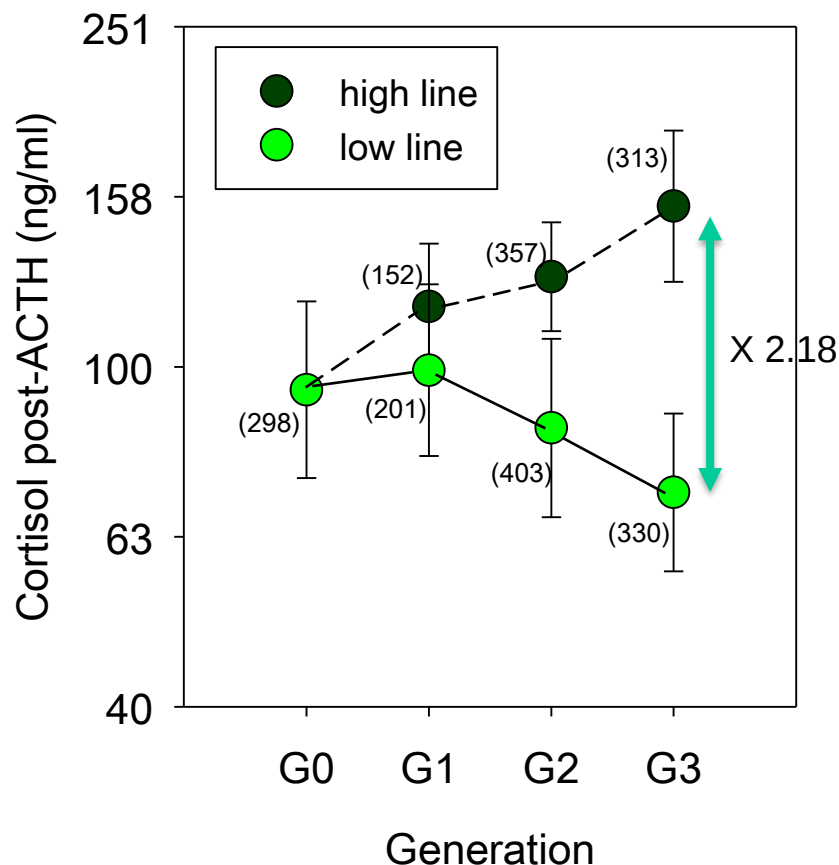
TT n = 3 (not shown)

Terenina et al. DAE 44:81, 2013



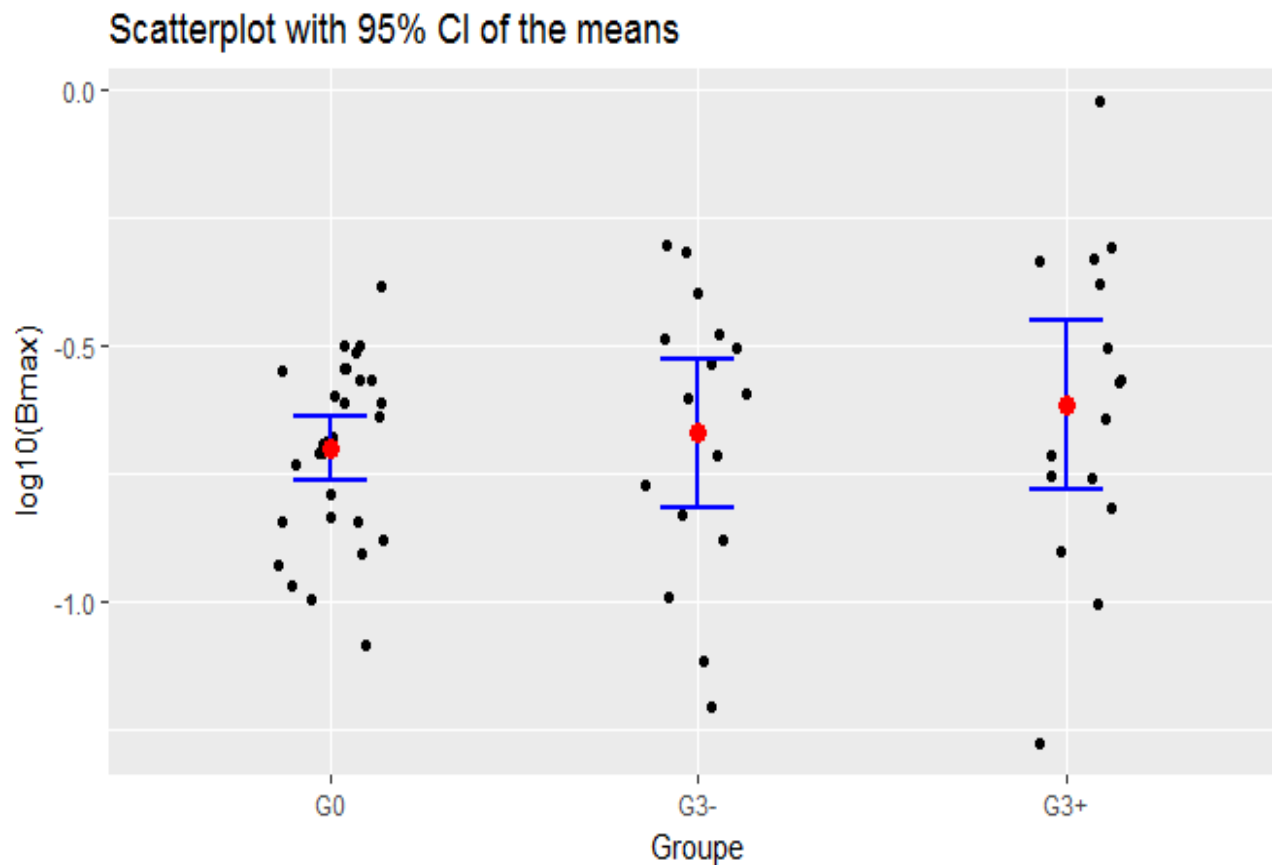
HPA axis and robustness - proof of concept

divergent selection for the response to ACTH in Large White pigs

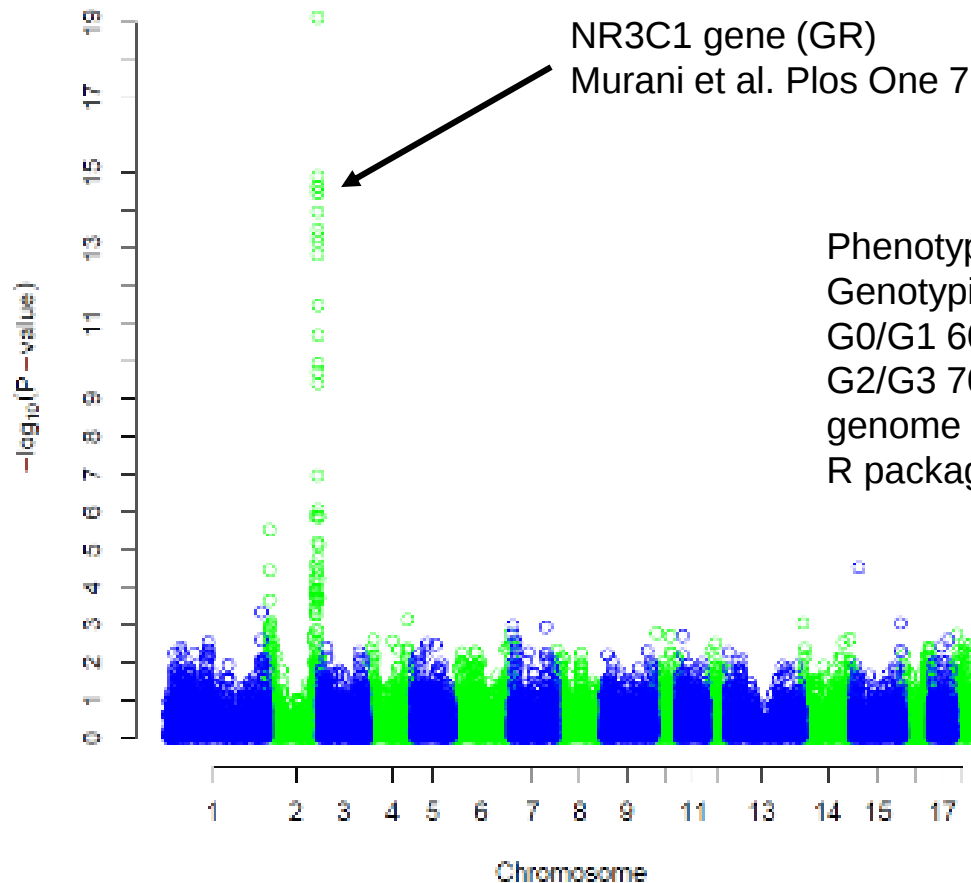


Realized heritability $h^2 = 0.68 \pm 0.12$

CBG (cortisol-binding globulin) in selected lines



Genome-wide association study

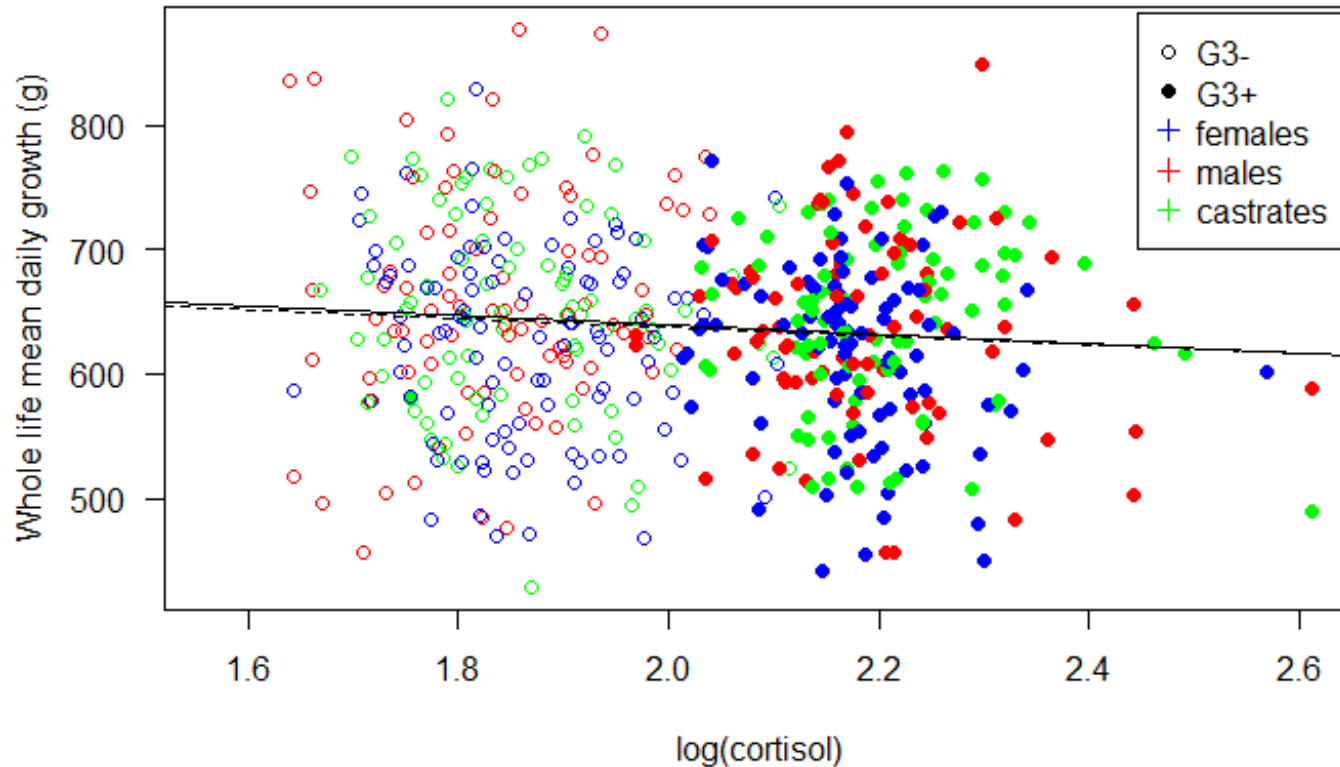


NR3C1 gene (GR)

Murani et al. Plos One 7(9): e45518

Phenotype: cortisol level 1 h post-ACTH
Genotyping with an Illumina Beadchip array
G0/G1 60kSNP array
G2/G3 70kSNP GenSeek array
genome assembly Sscrofa11.1
R package GenABEL

Selected lines G3 – mean daily growth

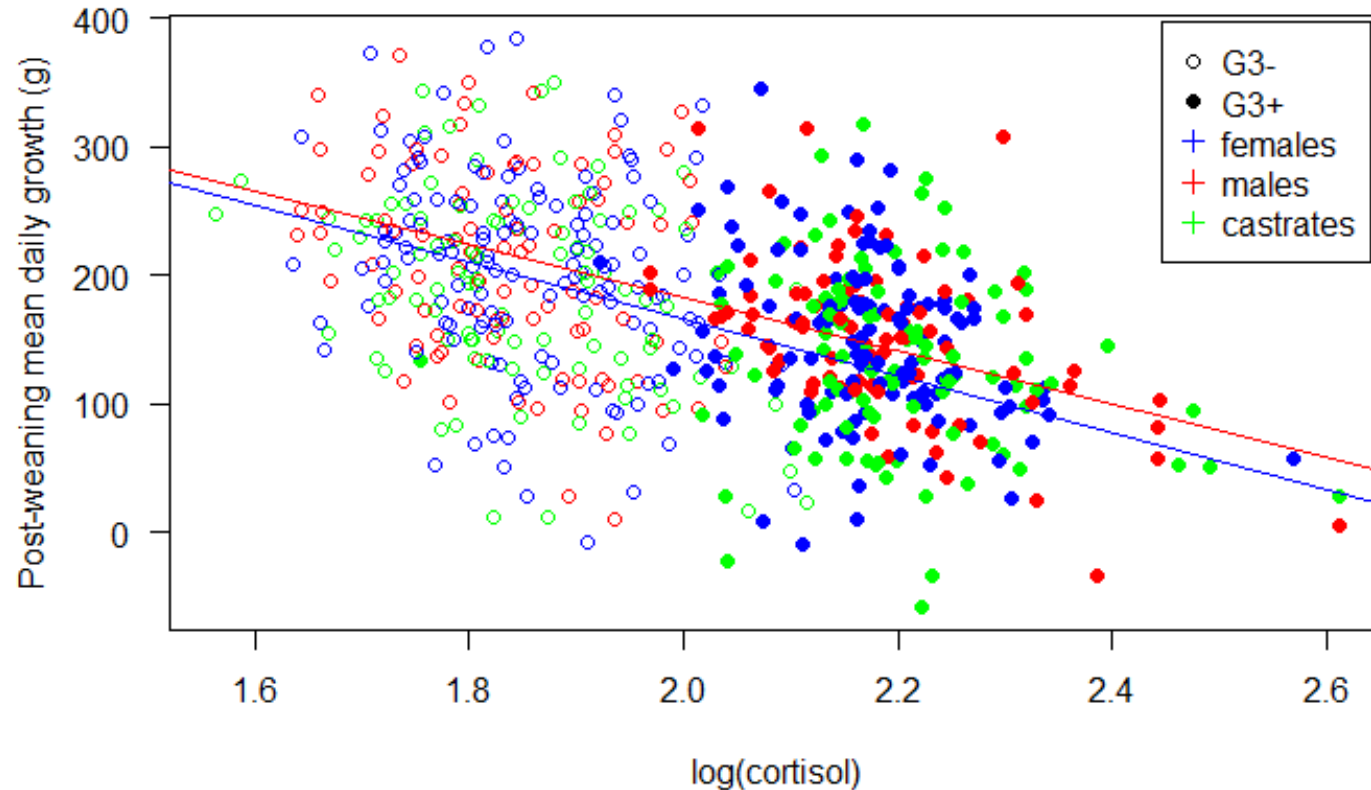


G3+ - G3- = -11 g/day, n.s.

Within groups, - 35 g/day per unit log(cortisol), n.s.

NO effect of selection
NO influence of cortisol

Selected lines G3 – post-weaning growth



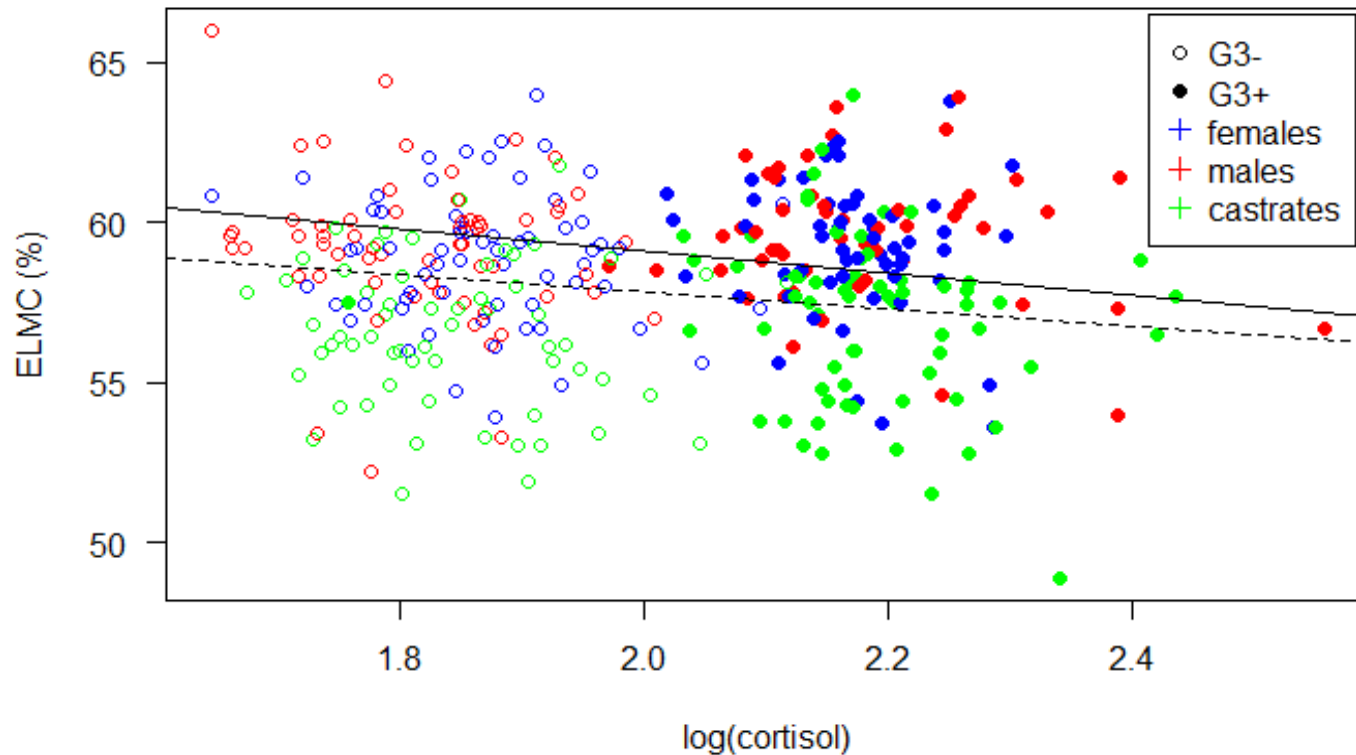
G3+ - G3- = -55 g, $P < 2.10^{-16}$

Within groups, -214 g per unit log(cortisol), $P < 2.10^{-15}$

At the same level of cortisol, G3+ - G3- = +16 g; n.s.

Strong effect of selection
Related to cortisol levels

Selected lines G3 – meat content



G3+ - G3- = 0.49%, $P < 0.05$

Within groups: ELMC -4.12% per unit log(cortisol), $P < 0.002$

At the same level of cortisol, G3+ - G3- = 1.90%, $P < 0.0005$

Small effect of selection
Unrelated to cortisol levels

Sources of genetic variability in the HPA axis

CENTRAL DRIVE

- Inputs to the HPA axis
- CRH, vasopressin and receptors

CORTICOSTEROID HORMONE PRODUCTION

- Response of the adrenal cortex to ACTH

BIOAVAILABILITY

- Corticosteroid-binding globulin (CBG)
- 11β -hydroxysteroid dehydrogenases

ACTION ON TARGET TISSUES

- Receptors or transduction pathways

FUNCTIONAL CONSEQUENCES

Production / Meat quality / Robustness

**SYSTEMS
GENETICS**

FUNDING

INRA, Animal Genetics Division

ANR, Agence Nationale de la Recherche



❖ INRA, Genetics, Physiology and Breeding Systems, Toulouse

- Elena Terenina, Catherine Larzul, Nathalie Villa-Vialaneix
- Valérie Sautron (thesis)

❖ INRA, Genesi, Le Magneraud (experimental unit)

- Yvon Billon

Thank you for your attention



photo David McEnery

Recent review papers

- Ann NY Acad Sci 1220:127, 2011
- Animal 5:651, 2011
- Dom Anim Endocr 43:116, 2012