

Session 77

Trade-offs between health, production and welfare in pigs and poultry
Which tools do we have, and which research is needed?

The future of pig and poultry breeding

Lisanne Verschuren & Pieter Knap

August 2023

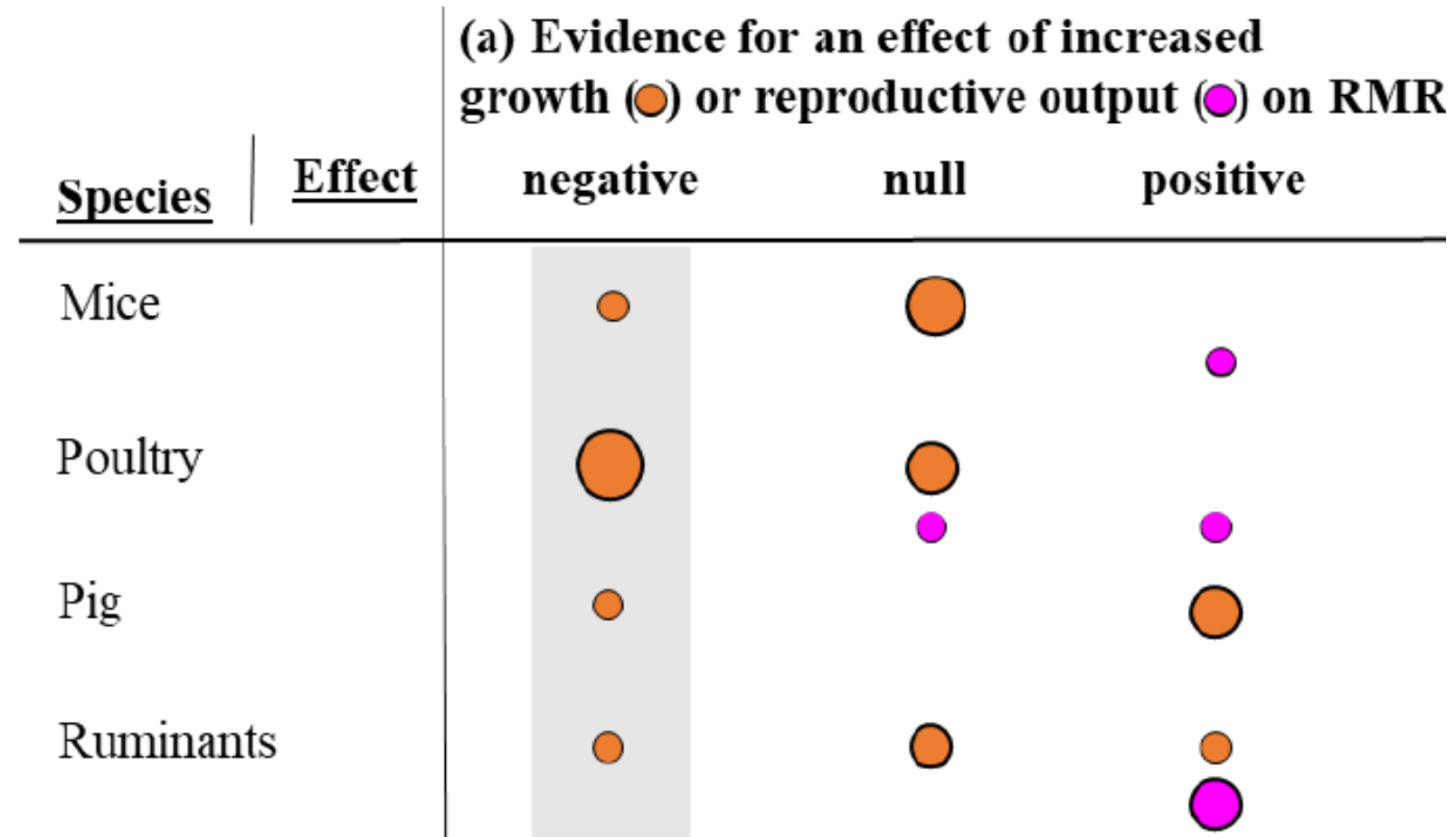


Figure 2. Evidence for association between resting metabolic rate (RMR) and selection for growth or reproductive output when feed intake is not restricted (a), and effects of a genetically reduced RMR on health or reproduction

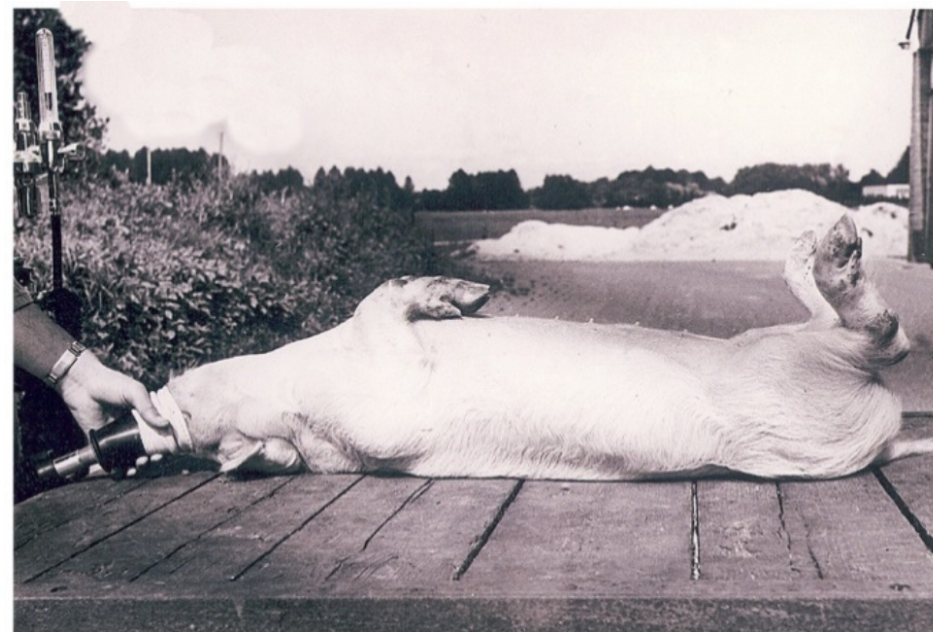
Trade-off systems in breeding: some history

- 1970s: **lean content** ↔ **Porcine Stress Syndrome & Pale Soft Exudative meat**



Figuur 2. Philippe Lampo (Faculteit Diergeneeskunde) voert de halothaantest uit (Uit: *Boerderijrevue*, april 1979).

halothane negative (NN)
or carrier (NP)



halothane positive (PP)

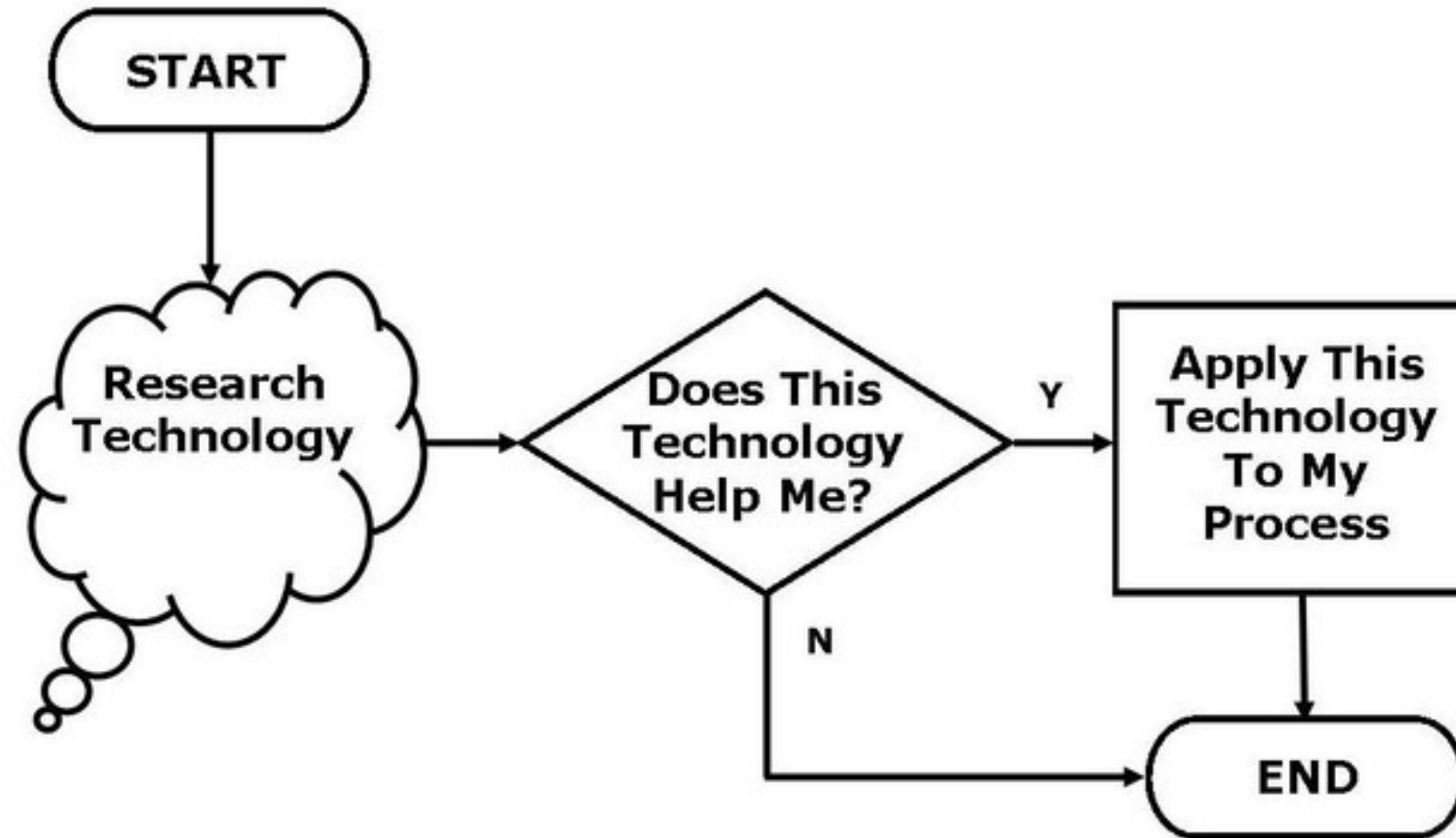


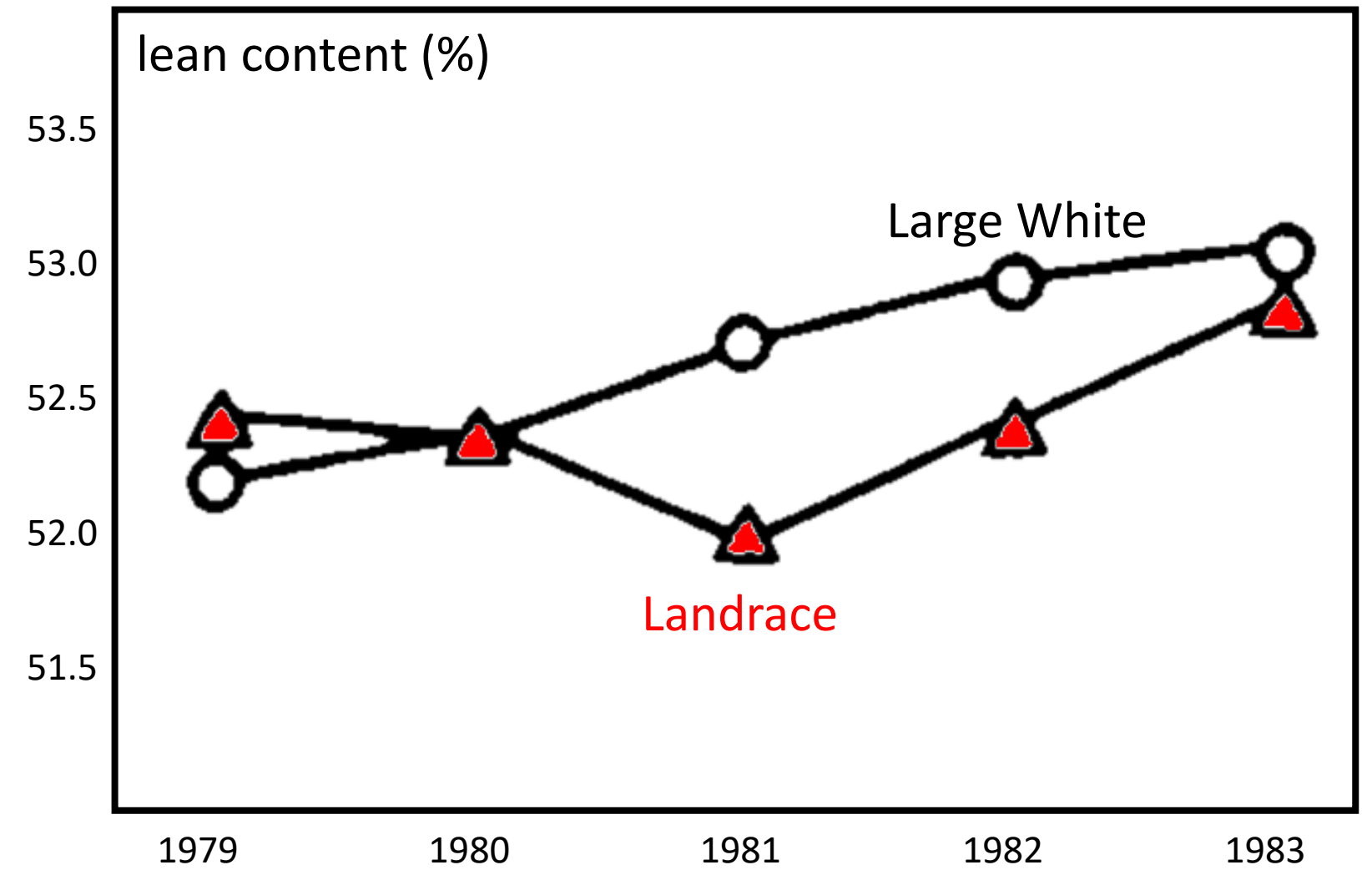
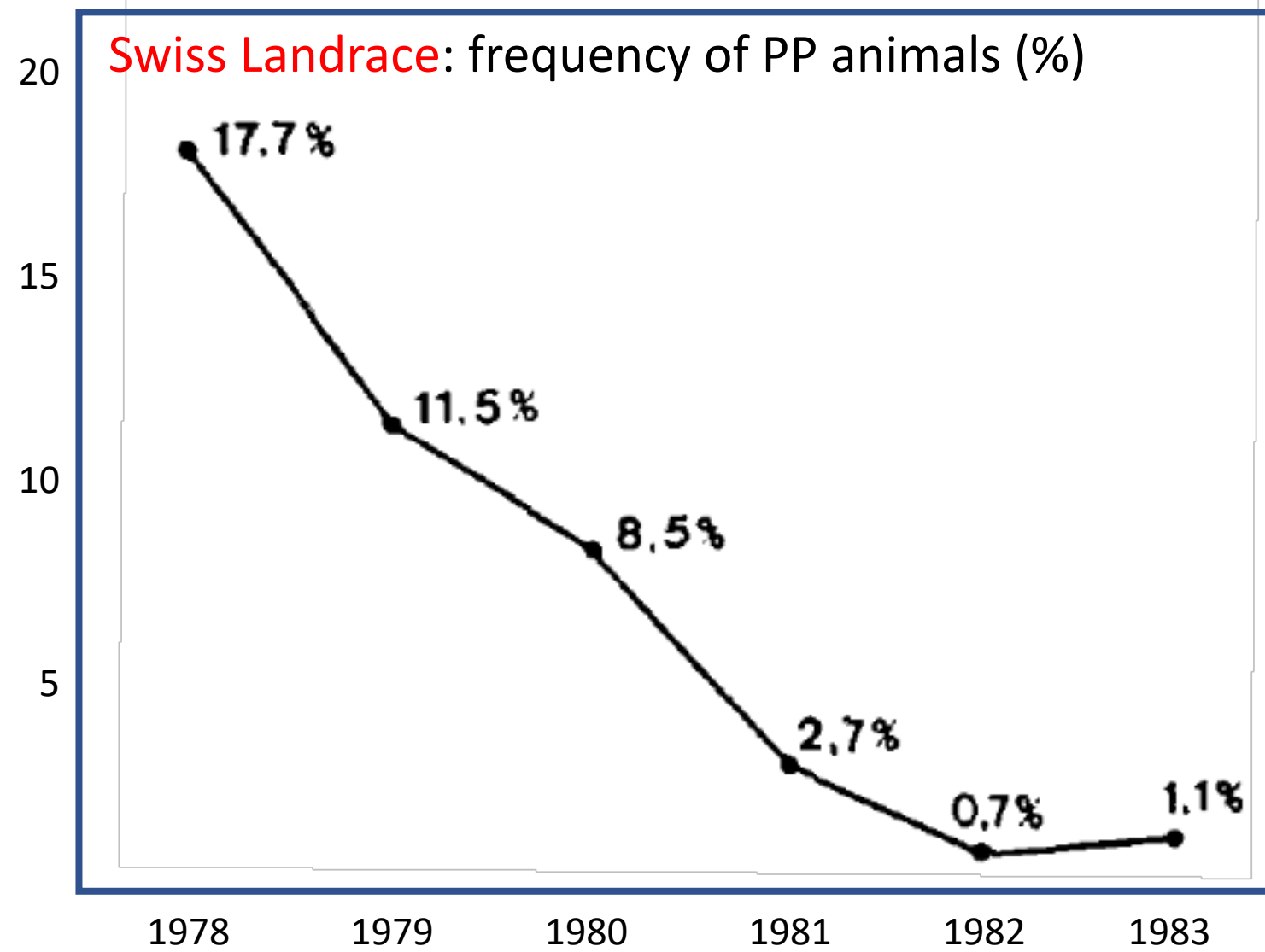
	Number of studies	Mean	Minimum	Maximum
Growth traits (approx. 25-90 kg)				
Growth rate (gm/day)	12	-2	-47	28
Daily food consumption (kg)	9	-0.07	-0.46	0.06
Food conversion ratio (food/liveweight gain)	11	-0.06	-0.30	0.02
Carcase traits (approx. 90 kg)				
Lean (% by weight)	8	2.6	0.9	4.6
Ham (% by weight)	7	0.7	0.3	1.0
Average backfat (mm)	14	-1.0	-4.0	1.0
Killing out%	6	1.0	0.2	2.6
Eye muscle area (cm ²)	7	1.1	-2.7	3.4
Carcase length (mm)	9	-11	-29	1
PSS traits				
Postweaning mortality (and transport losses)%	3	9.8	4.7	17.0
PSE (% of carcasses)	4	46	22	80
Meat colour (% paler than HN)†	14	15	0	50
Meat quality (% worse than HN)†	7	31	16	78
pH 45 min post mortem	11	-0.31	-0.66	0.02
CK activity (log units/litre)	6	0.50	0.06	0.79

Difference PP - NN



Interesting. And what do we do with it?

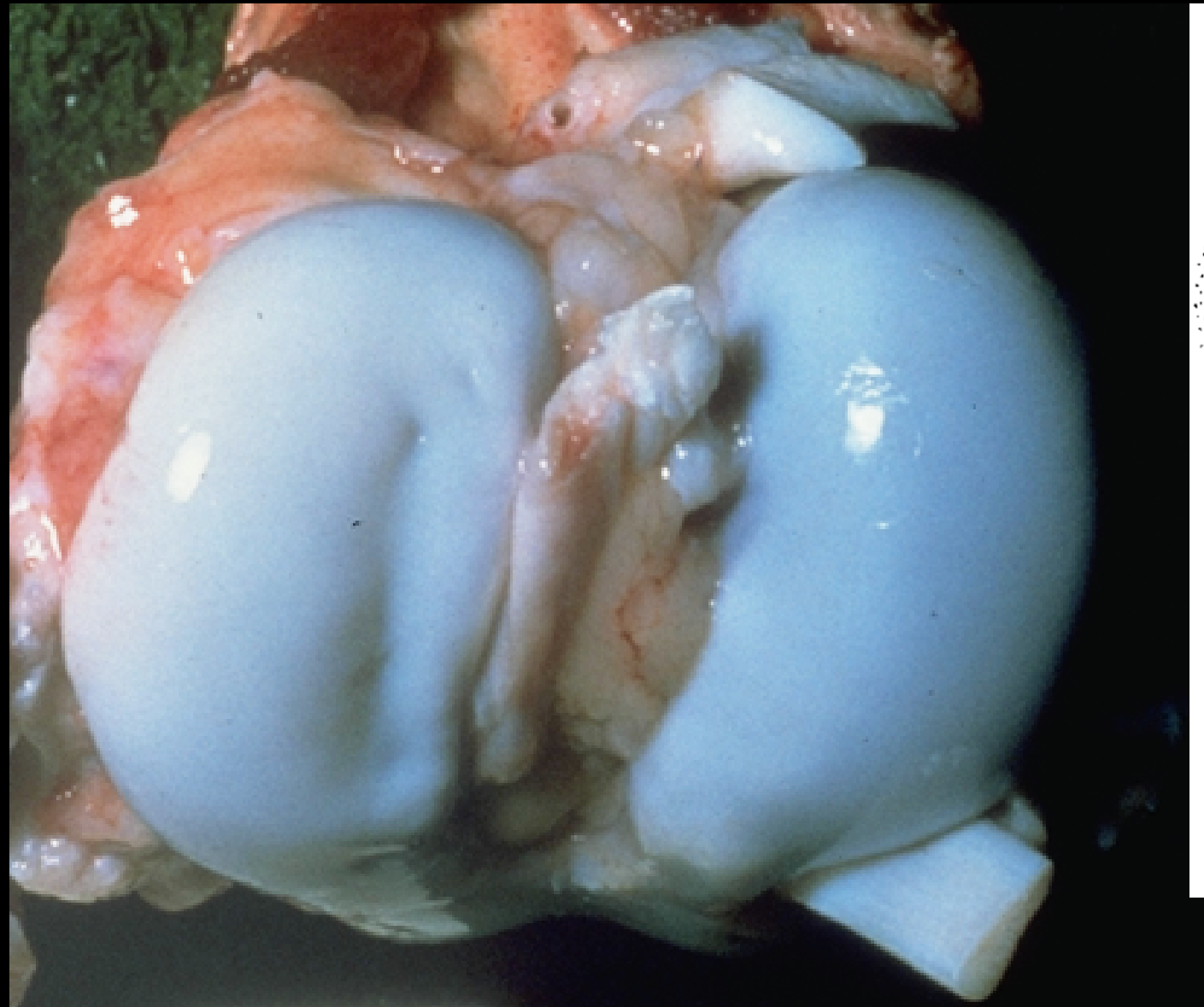




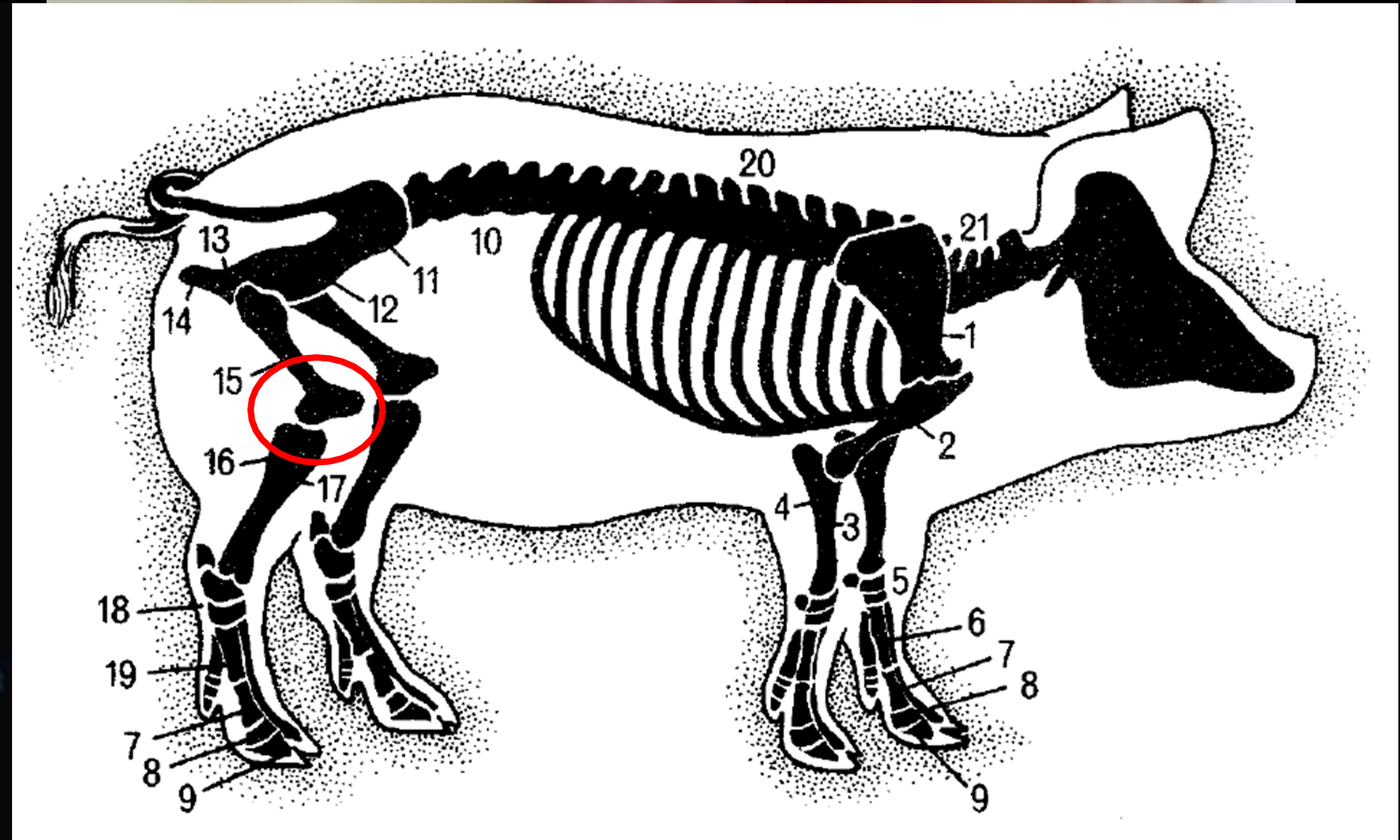
Trade-off systems in breeding: some history

- 1970s: lean content $\leftrightarrow$ PSS & PSE
- 1980s: growth rate $\leftrightarrow$ leg soundness

healthy femoral condyle



osteocondrosis

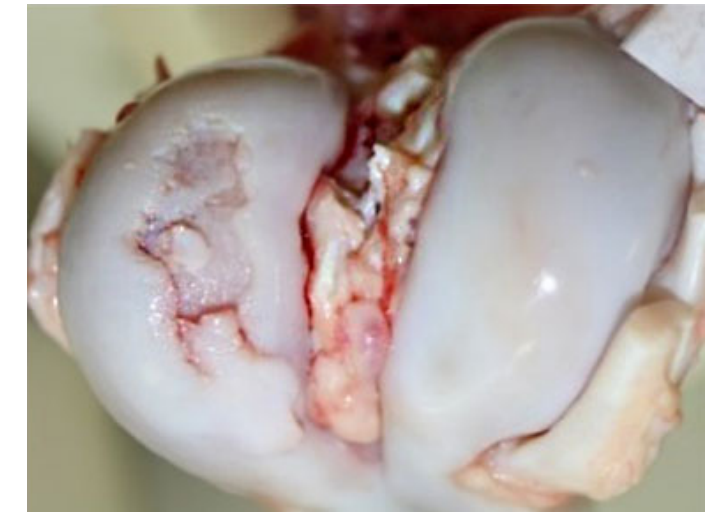


Morfologische onderzoeken bij slachtvarkens laten zien dat er tegenwoordig nauwelijks nog dieren zijn waarbij er geen osteochondropatische veranderingen aantoonbaar zijn (Dämmrich, 1970, 1972; Hermann, 1972; Grøndalen, 1974)

Morphological investigations reveal that currently (= 1976 !) there are hardly any pigs that do not show any osteochondropathic changes.

- An assumed strong connection to selection for a "meaty" type of the past 15 years
- At the same time, housing conditions became more intensive
- Skeletal maturation cannot keep up with fast body growth rate
- Imbalance: body weight $\leftrightarrow$ mechanical load on the bones
- Most where this load is most severe: femoral condyles, ulnar epiphyses (i.e. the joints)

wel aangenomen dat er een sterke samenhang is met het feit dat men de laatste 10 à 15 jaar vooral varkens heeft willen fokken van een duidelijk vlees-type. Ook wordt door sommigen gesteld dat de beengebreen zich meer klinisch openbaren, omdat de huisvestingsomstandigheden veel harder geworden zijn. (Bollwahn, 1966; Bollwahn e.a., 1970; Christensen, 1953; Dämmrich, 1970, 1972; Flock, 1969; Grøndalen, 1974, Hermann, 1969, 1972; Kurzweg e.a., 1972; Sabec, 1961, 1974 en Schilling, 1963). Door de te snelle gewichtstoename zou de skeletrijping hier geen gelijke tred mee kunnen houden en ontstaat er een wanverhouding tussen lichaamsgewicht en de structuur d.w.z. de geschiktheid voor mechanische belasting van het skelet. Men ziet daarom de osteochondropatische veranderingen in het skelet juist vooral op die plaatsen optreden waar de functioneel-mechanische belasting het grootst is (lumbale wervels, mediale condyl van de femur, epiphysairschijven distale ulna etc.)

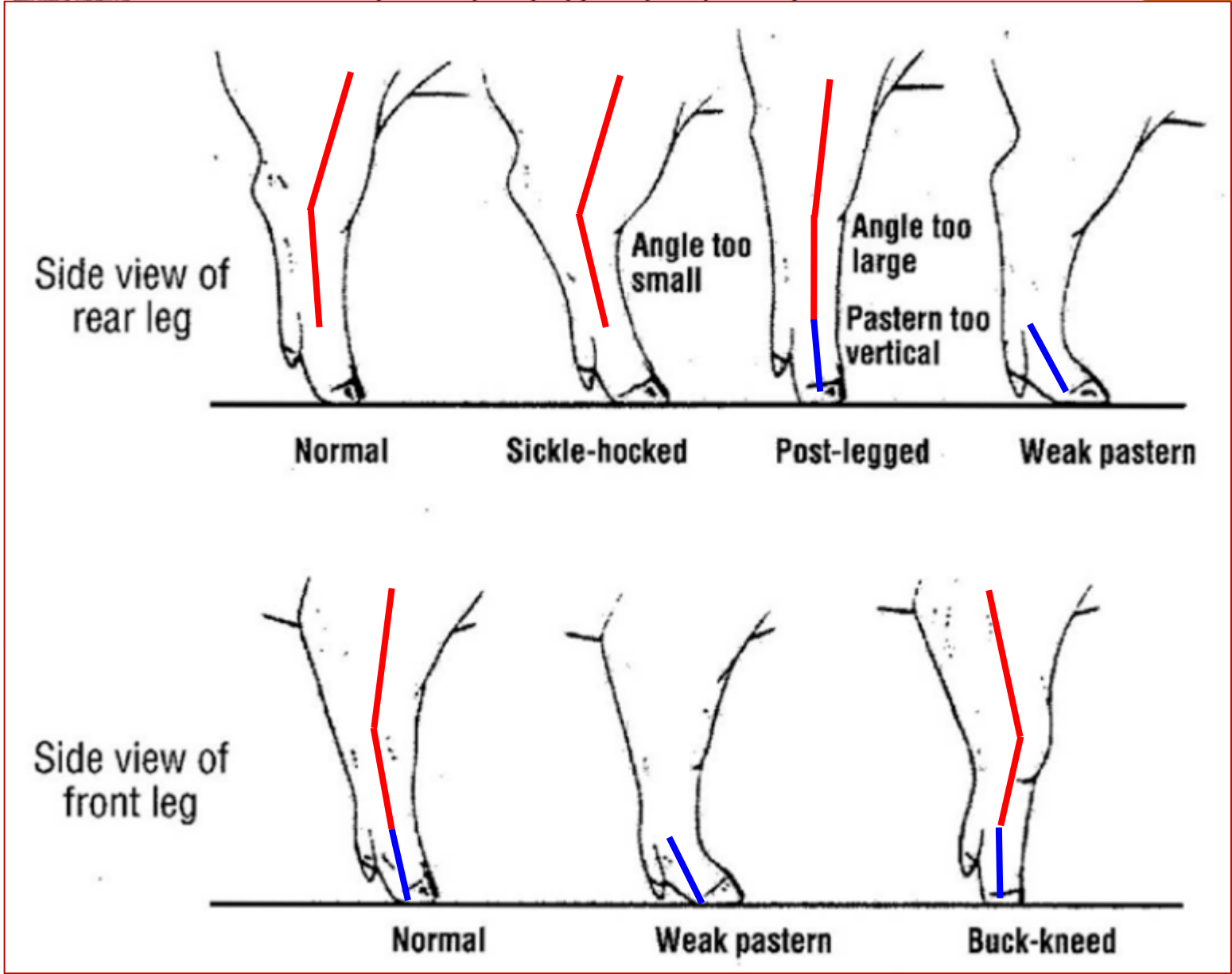


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Morphological investigations reveal that currently there are hardly any pigs that do not show any osteochondropathic changes.

- Followed by 20 years of phenotypic selection
- Visual appraisal of the leg conditions
- Exterior. Effectively scored as *good* or *bad*
- Zero ΔG

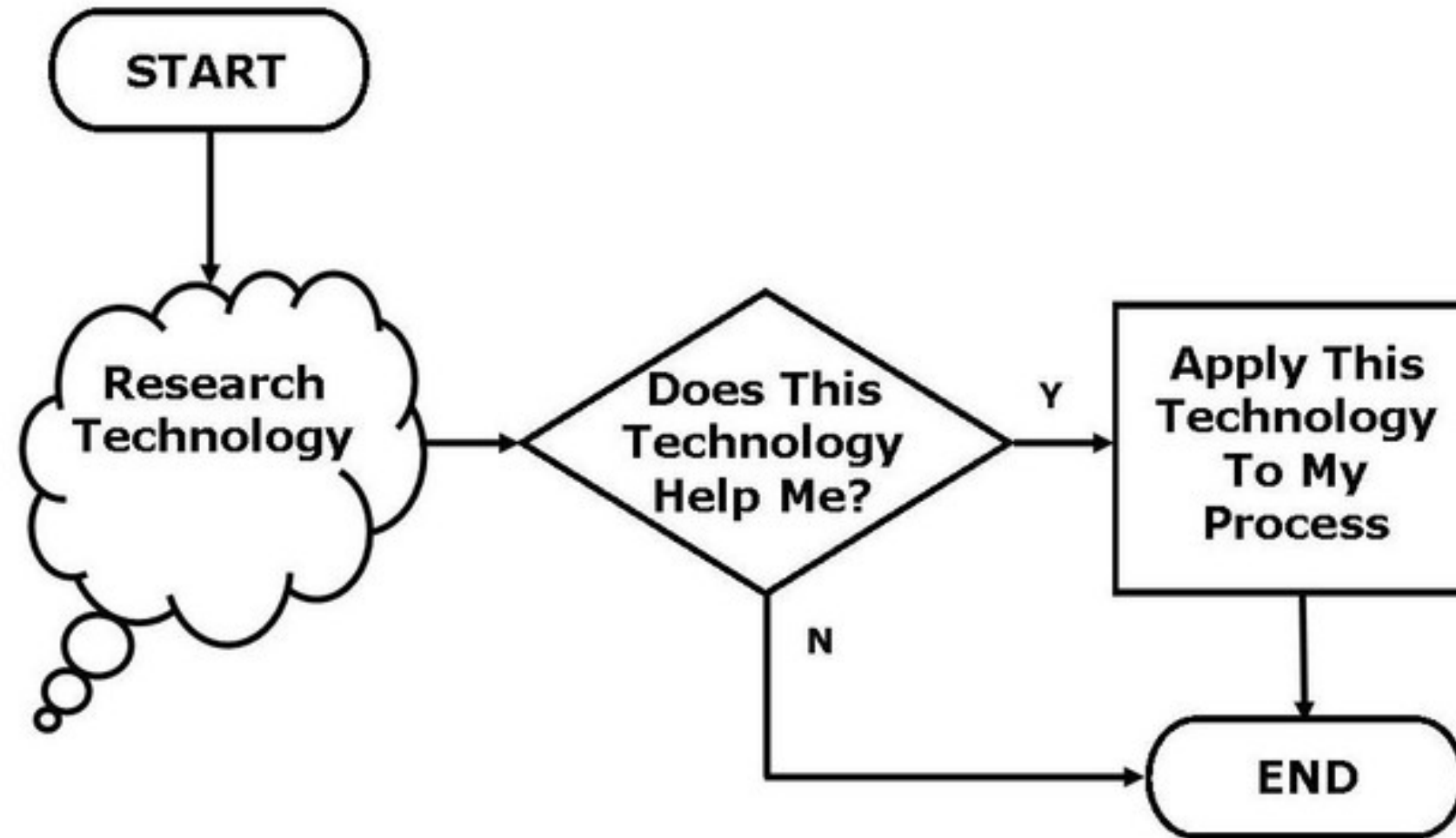
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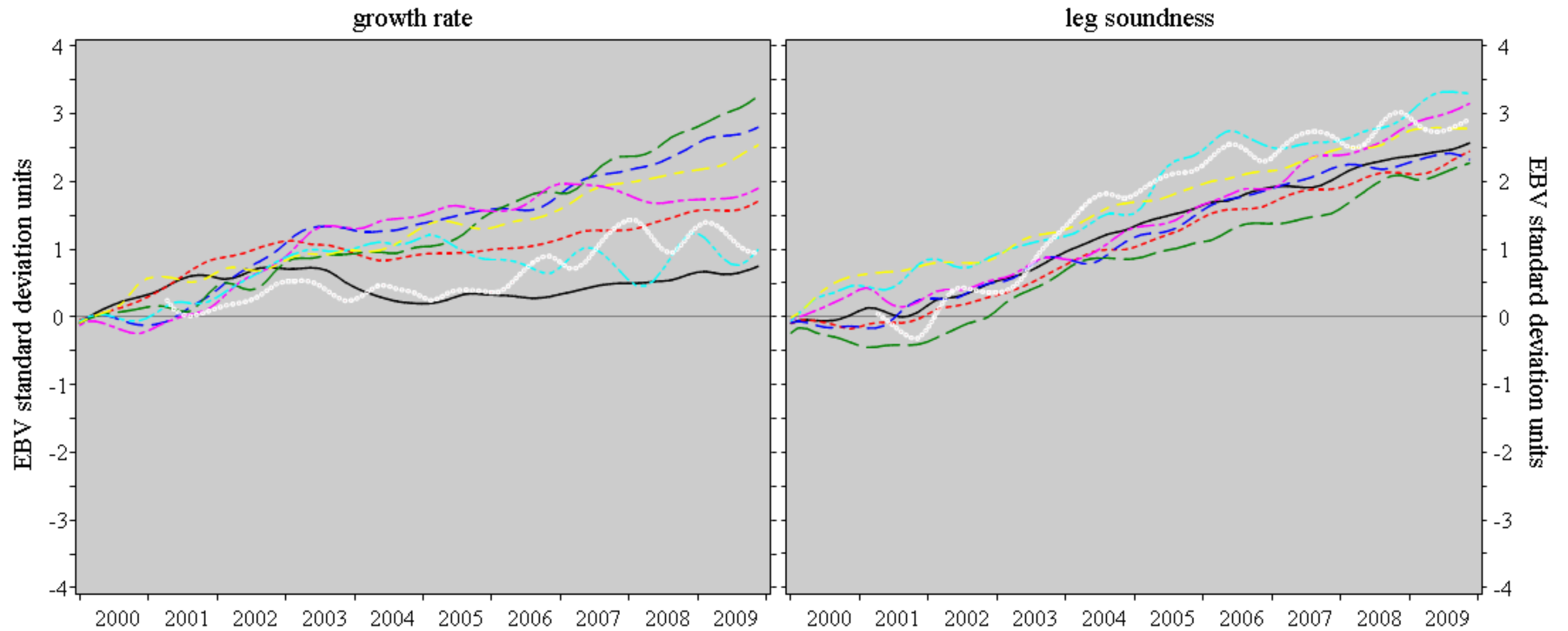


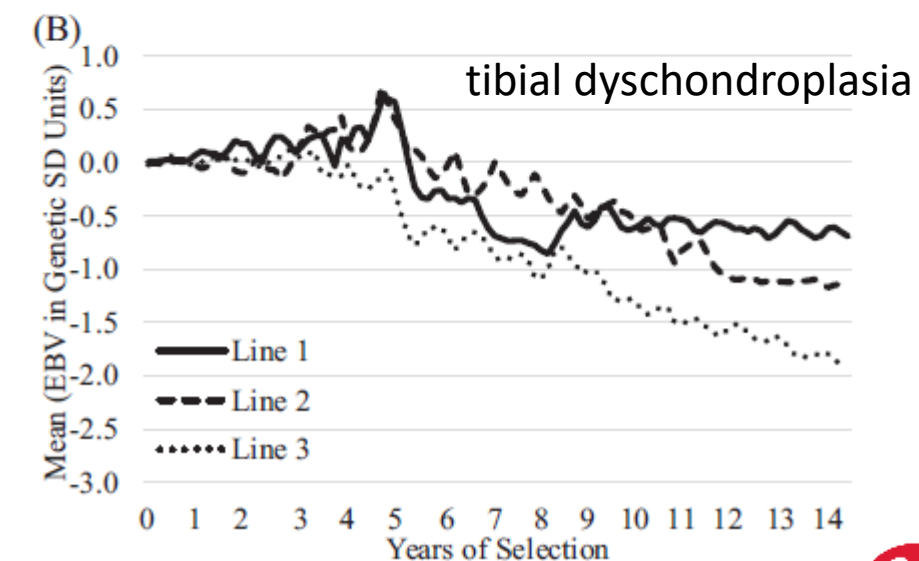
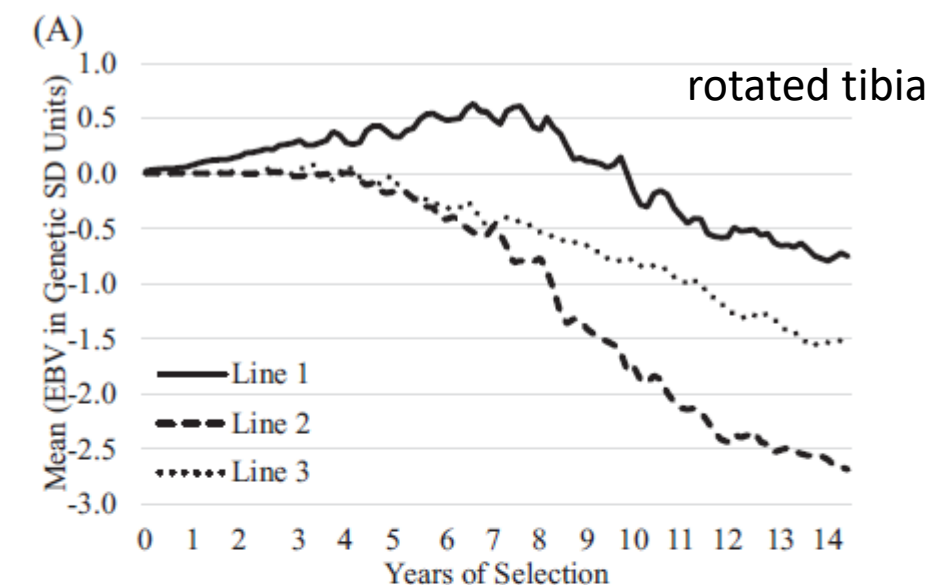
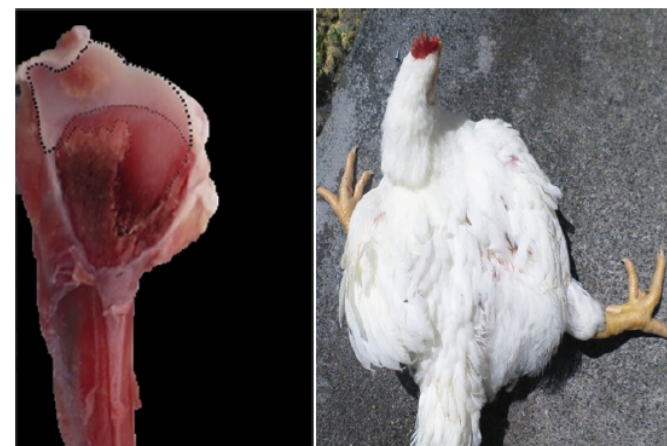
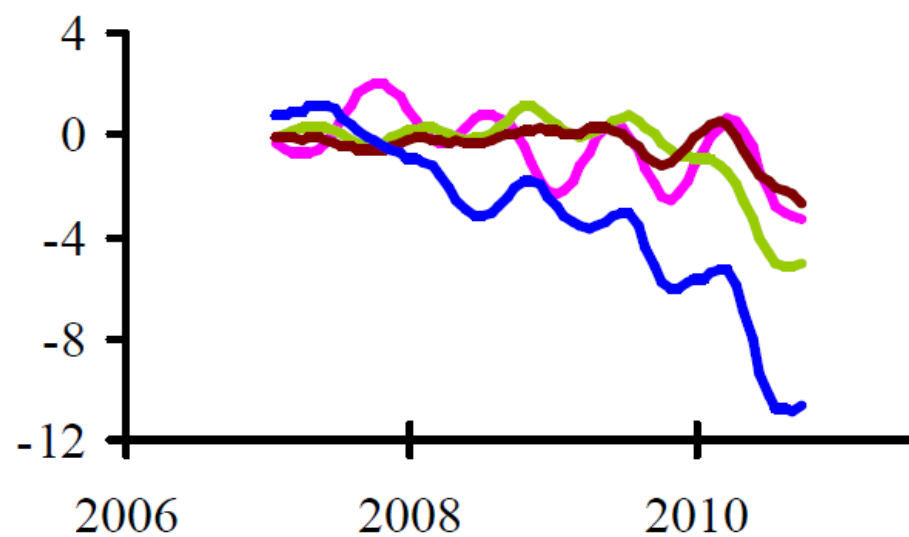
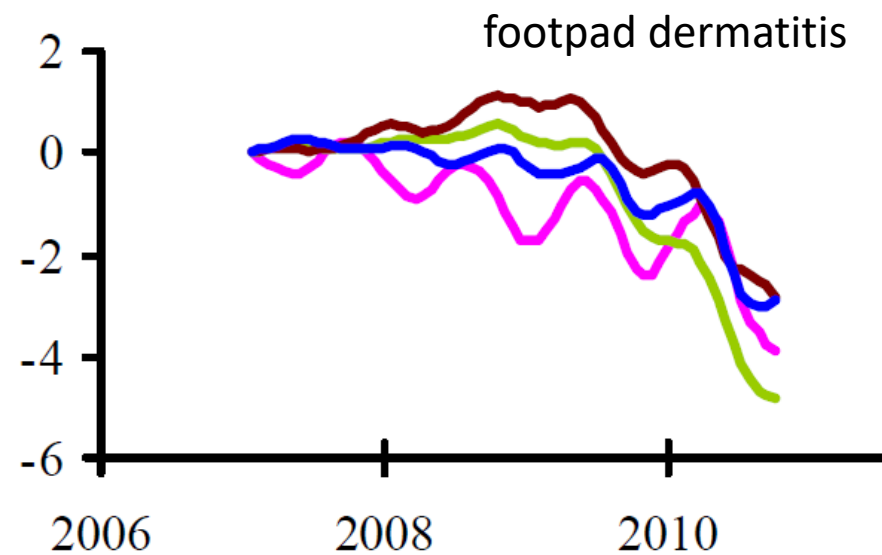
Genetic control of front-leg weakness in Duroc swine. II. Correlated responses in growth rate, backfat and reproduction from five generations of divergent selection ☆

← Phenotyping:
not a narrative, but angles: **numbers**
Can be BLUPped – that works miracles

Interesting. And what do we do with it?

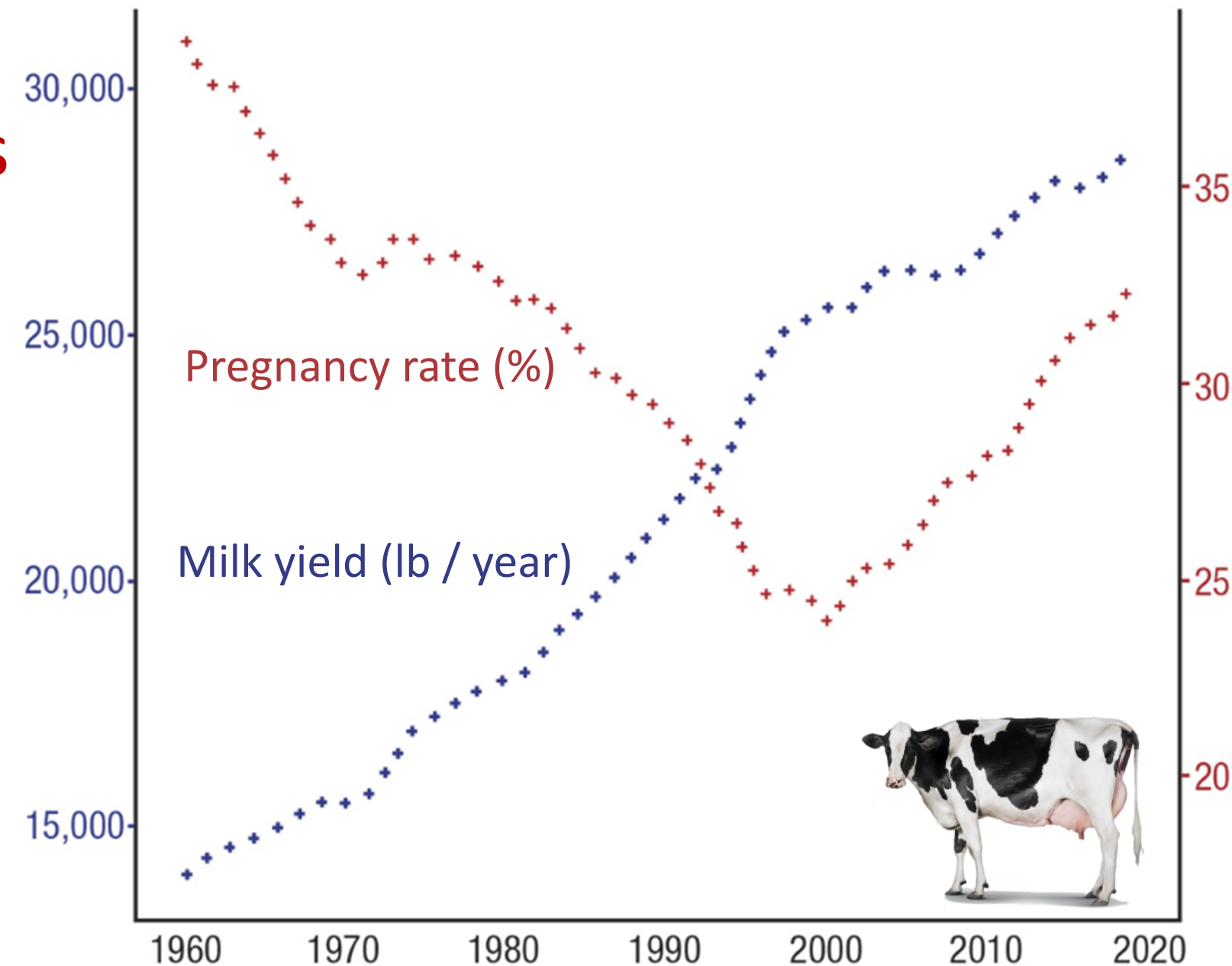






Trade-off systems in breeding: some history

- 1970s: lean content ↔ PSS & PSE
- 1980s: growth rate ↔ leg soundness
- 2000s: milk yield ↔ fertility



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Conclusions

1. Increased production levels

- good for increasing $\Delta\epsilon$
- good for reducing the carbon footprint

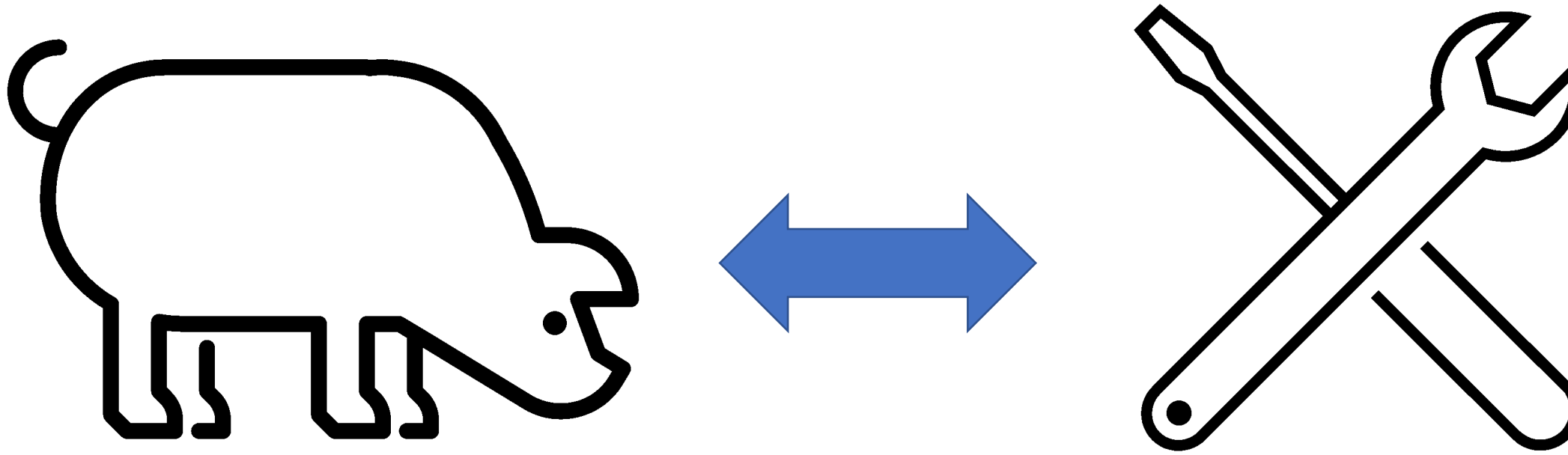
...have often led to trade-offs

- bad for animal welfare

...due to genetic antagonisms

2. Genetic antagonisms can be neutralized:

- select for both traits at the same time
- requires phenotyping & statistics & focus

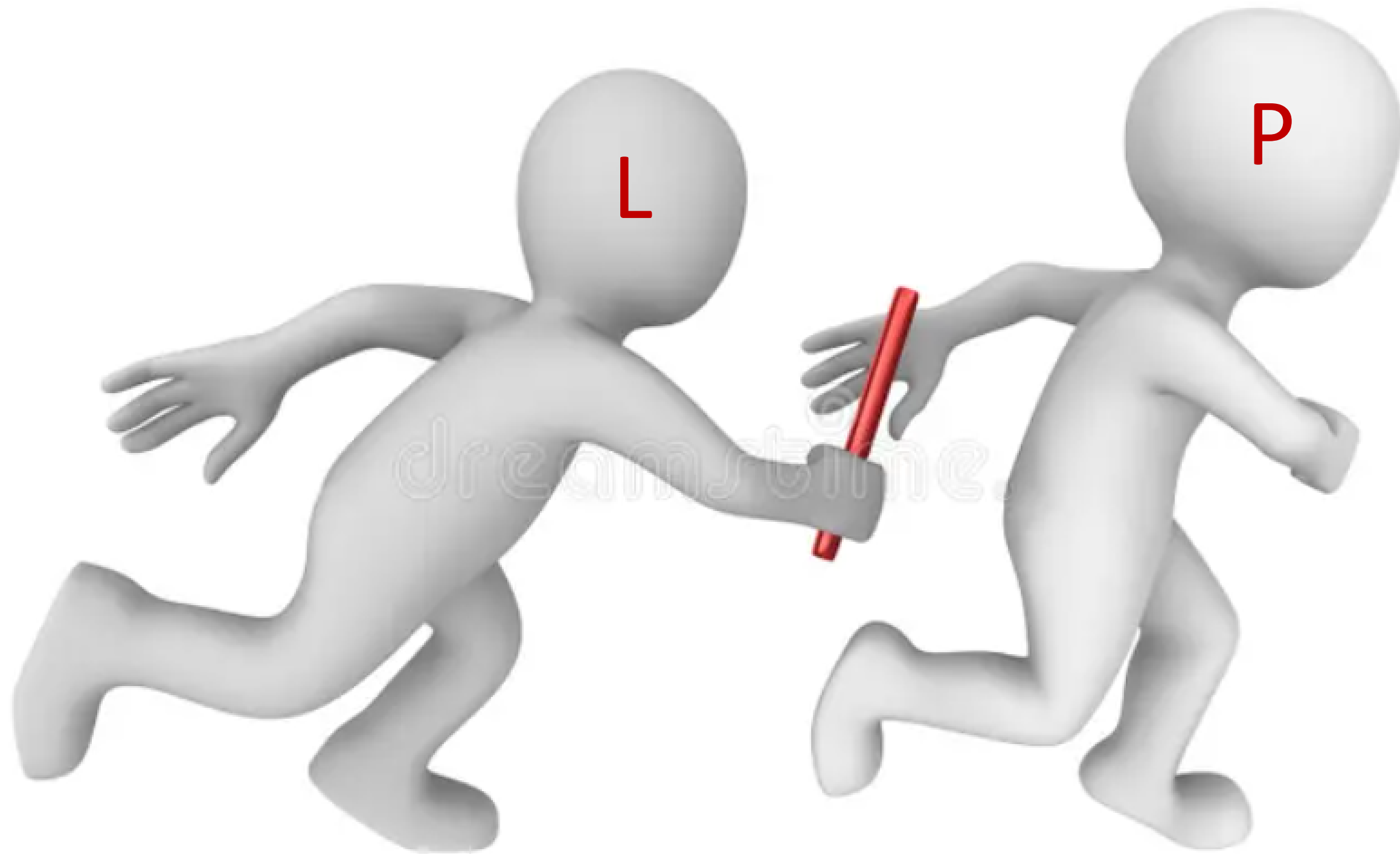


Trends

Simple → Complex

Static → Dynamic

Selection candidate → Group process



Digital phenotyping



OPINION

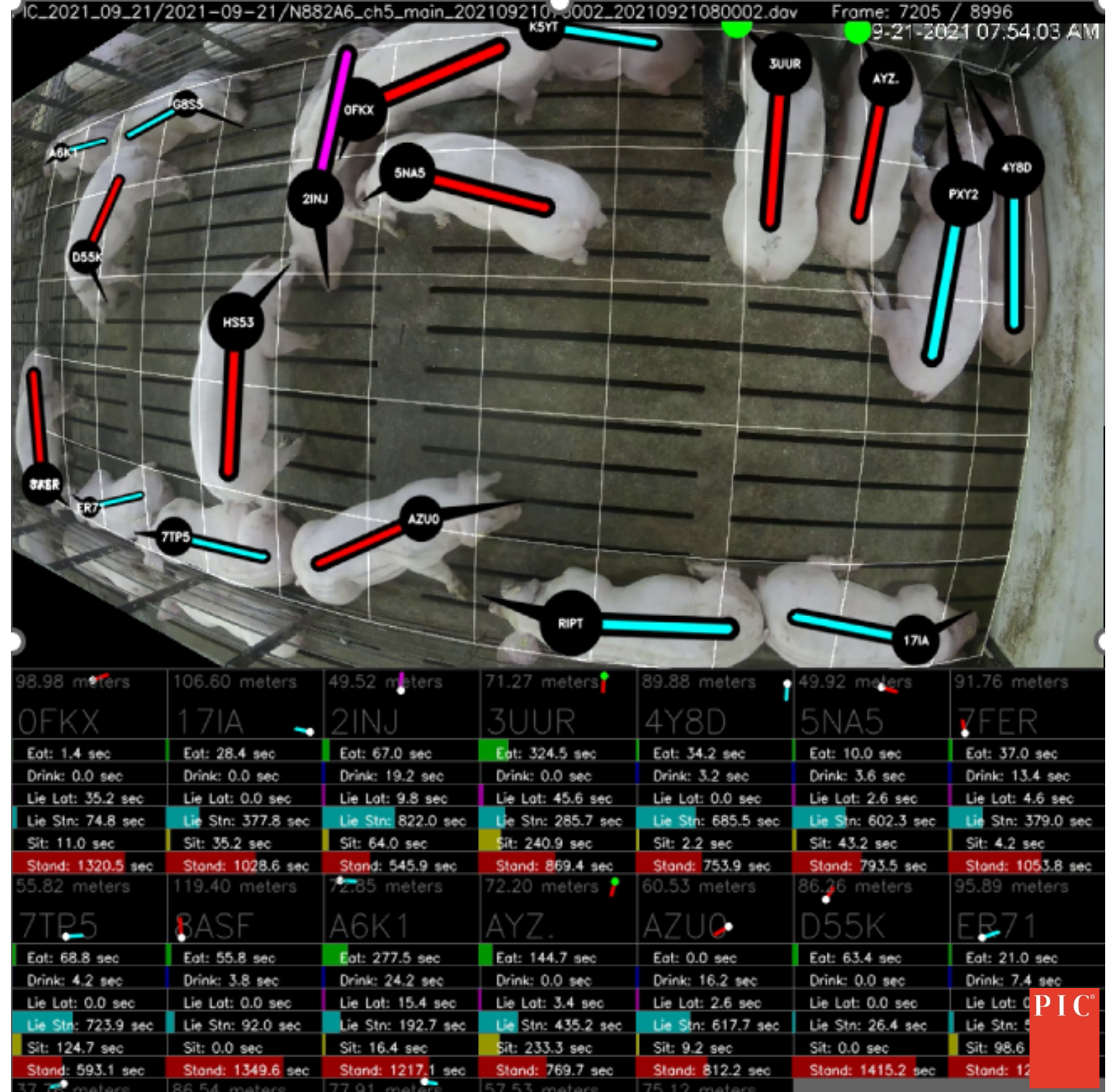
Open Access

Phenomes: the current frontier in animal breeding

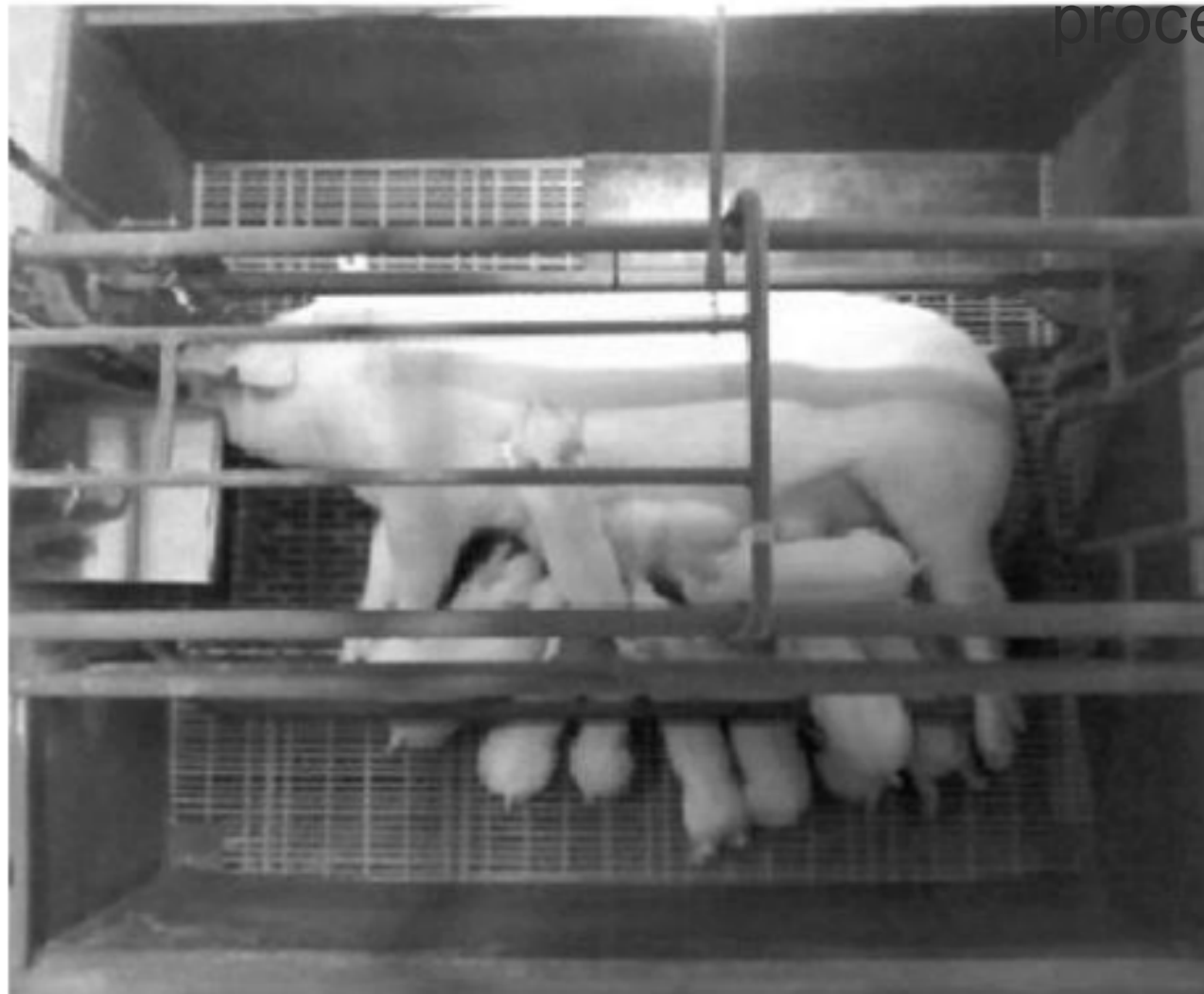


Miguel Pérez-Enciso^{1,2*}  and Juan P. Steibel^{3,4}

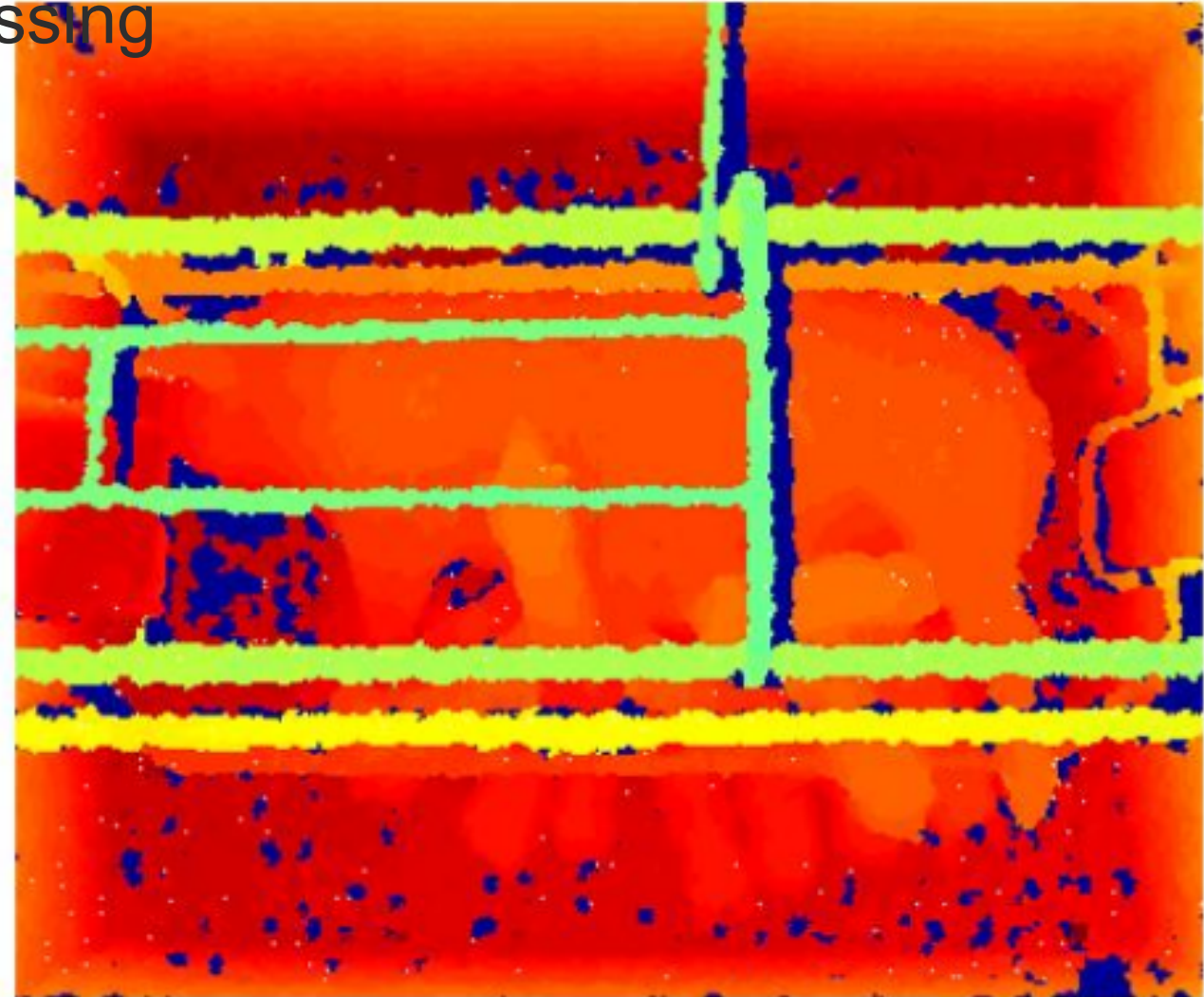
- Phenomics: "the acquisition of high-dimensional phenotypic data on an organism-wide scale" (Houle et al, 2010)
- Big data* challenges ← heterogeneity & rapid change over time of the data, not because of their size
- Two novelties:
 - New traits that could not be recorded before
 - behaviour



Automatic recognition of lactating sow behaviors through depth image processing



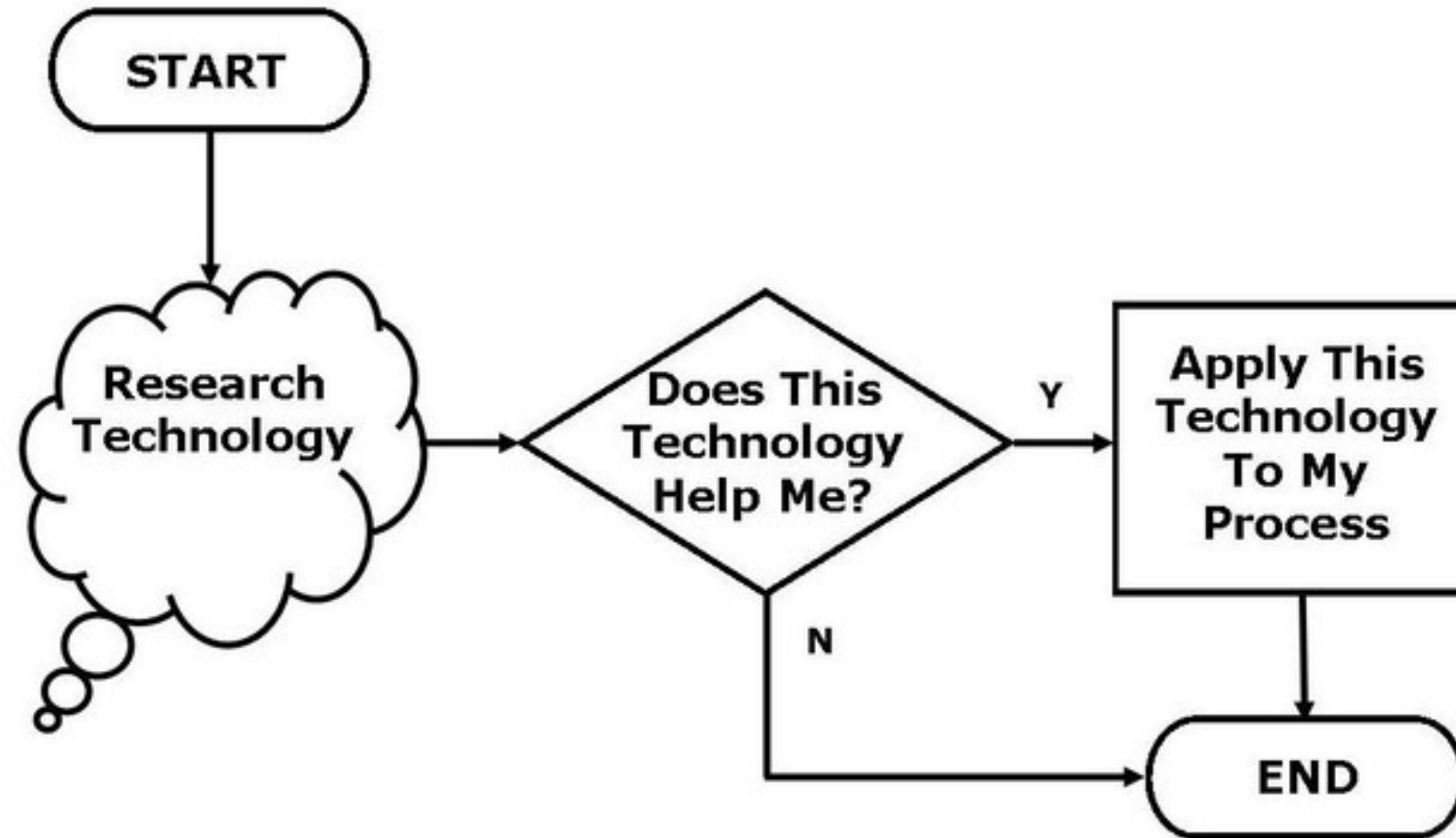
digital image



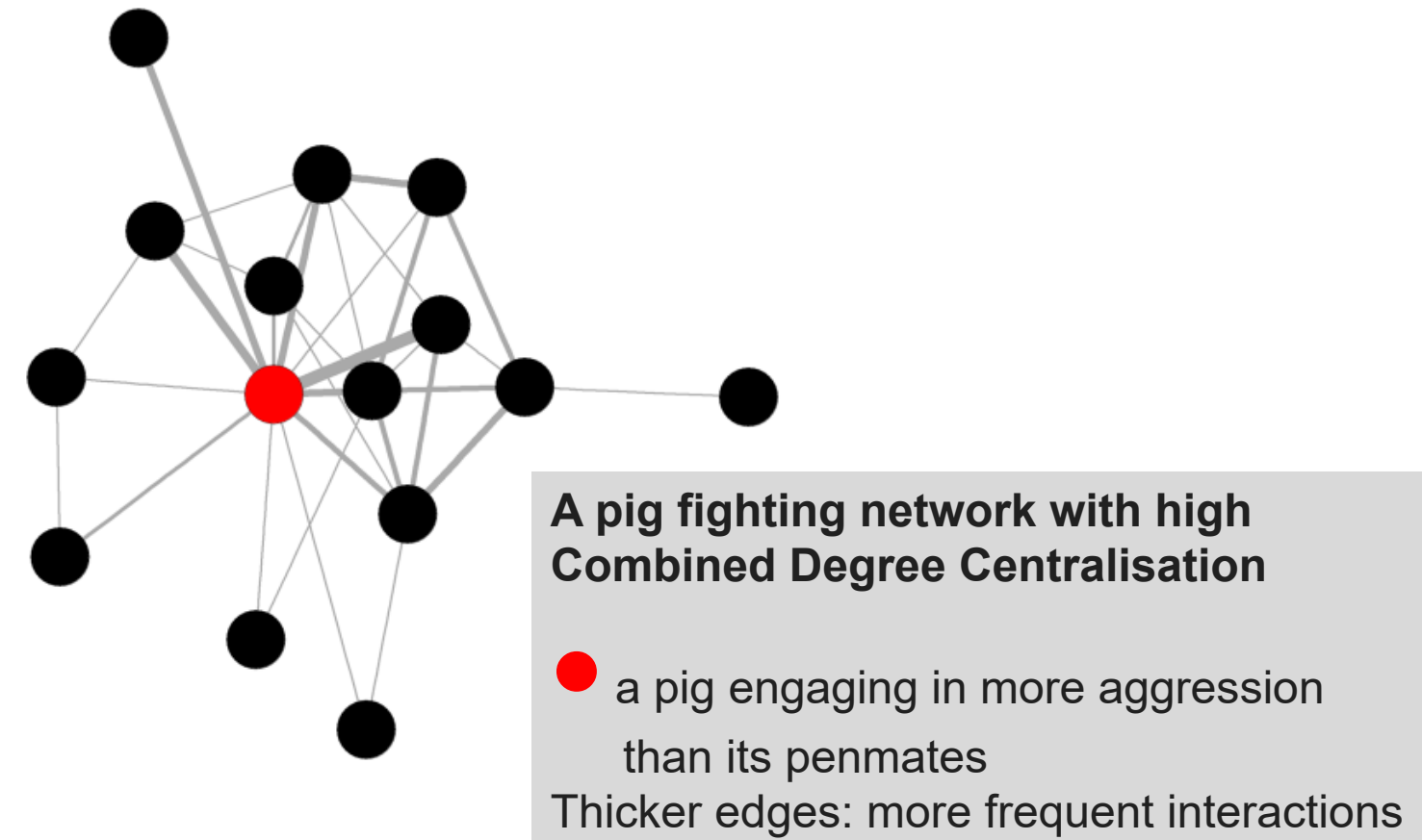
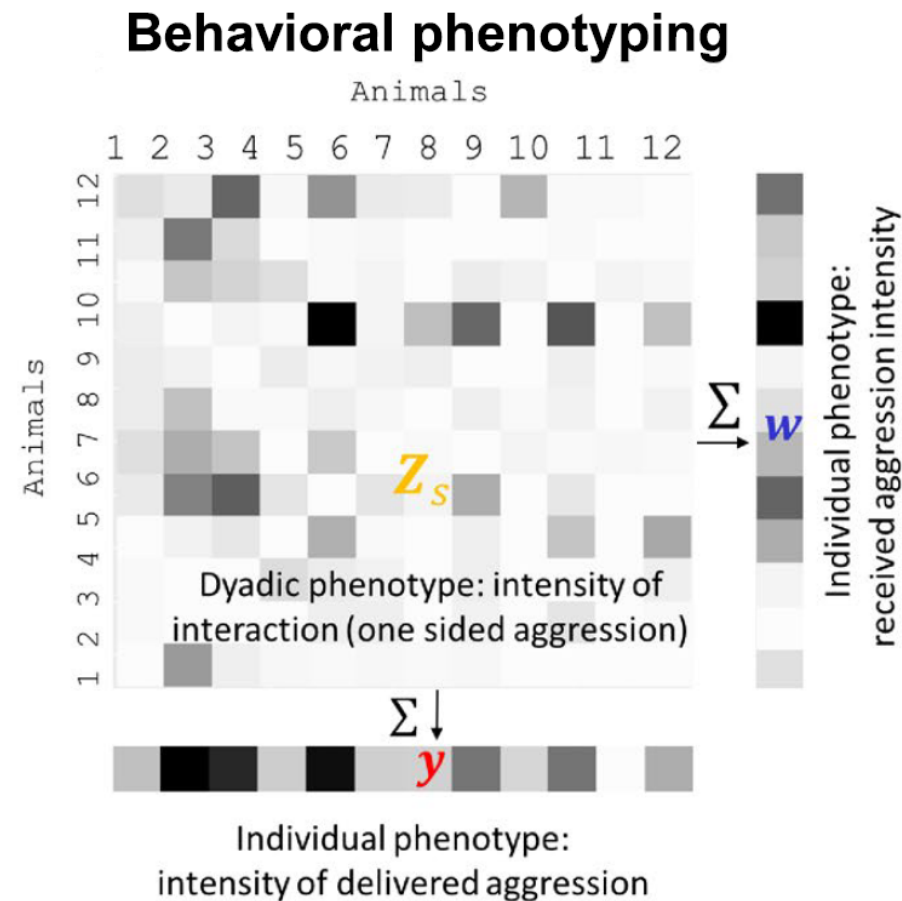
raw depth image

Maternal behaviour, farrowing, savaging, crushing

Interesting. And what do we do with it?

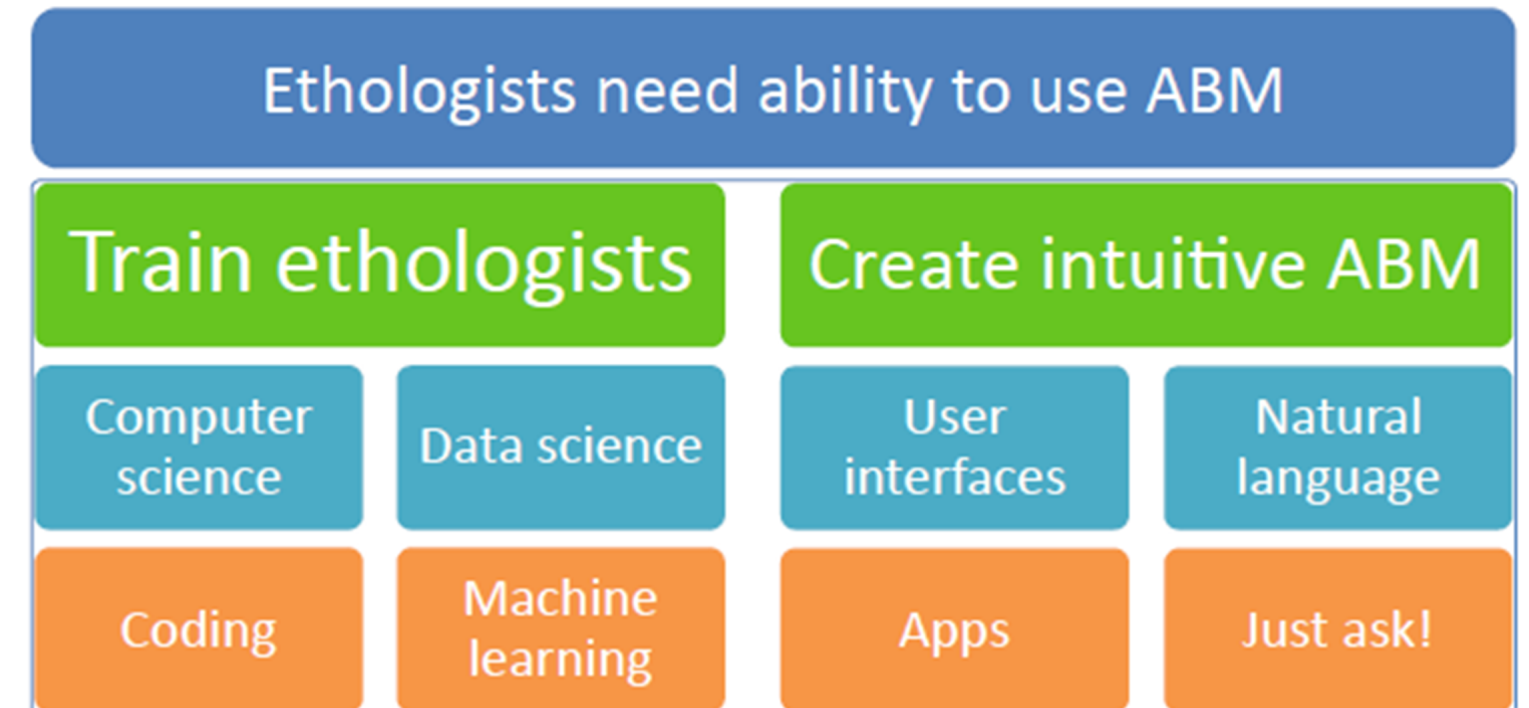
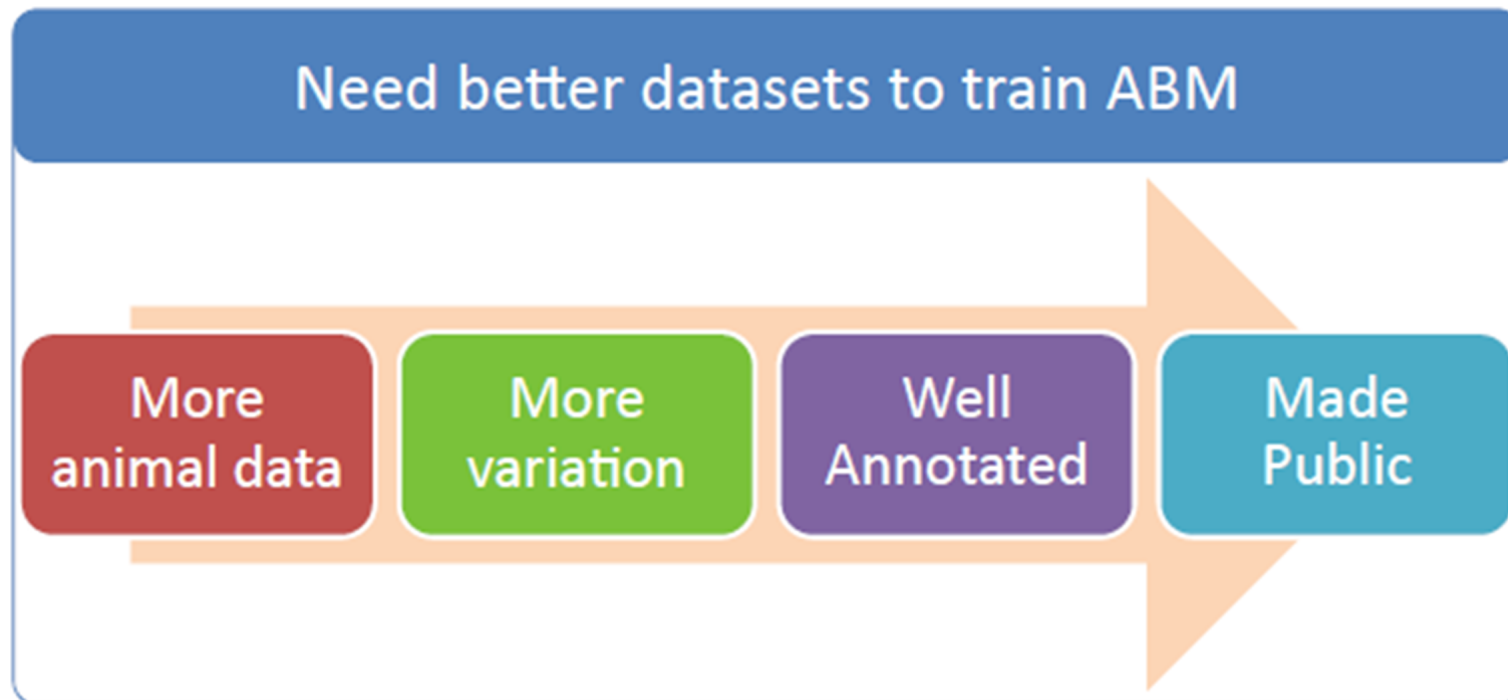


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Automated Behaviour Monitoring



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 - methane emission ← spectroscopy
 - Classical traits can be expanded:
 - more objective → more accurate

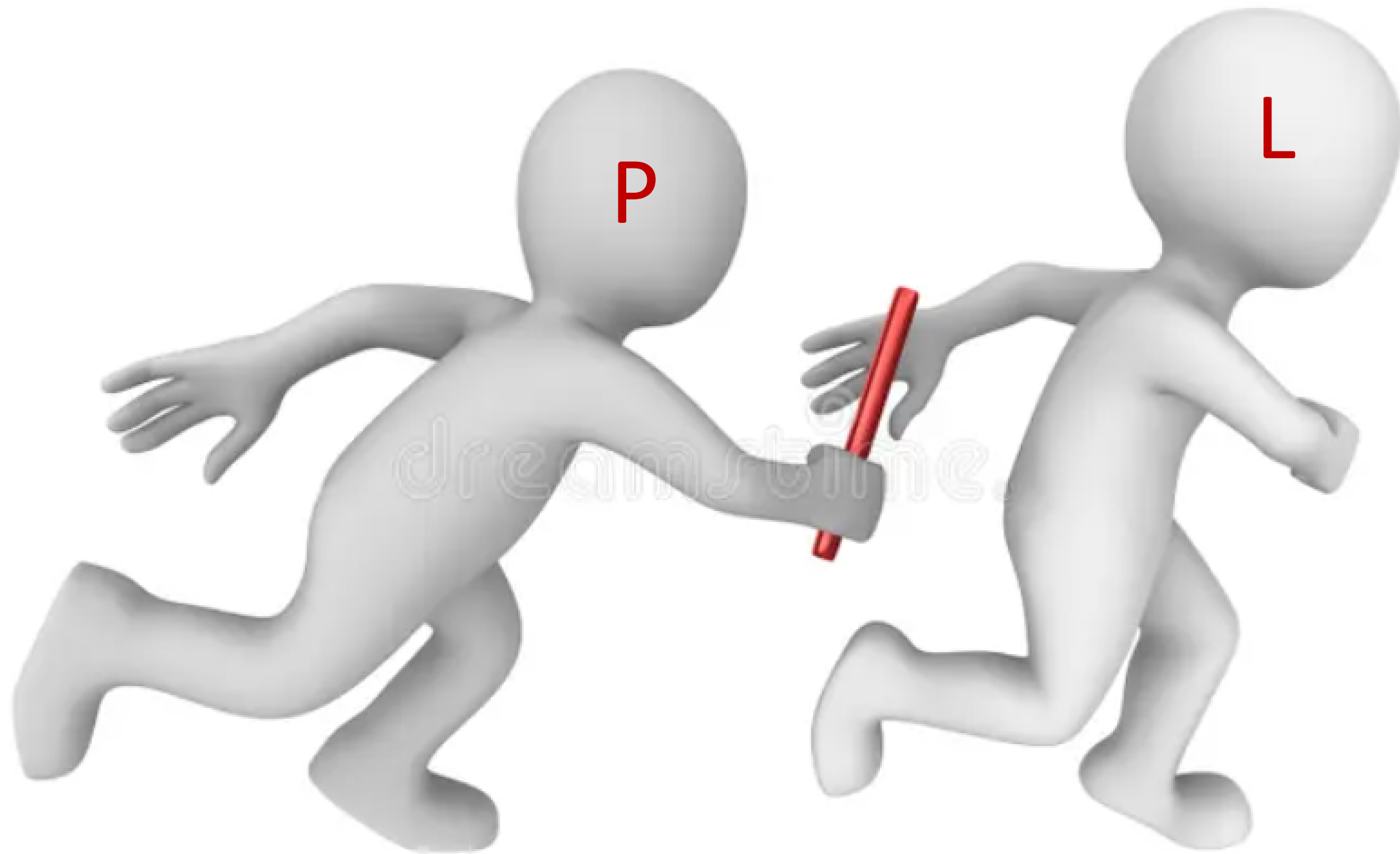


Neural Network vs visual:
heritability goes up
x 2.3 to x 5

	$\frac{h^2(\text{NN})}{h^2(\text{visual})}$	
line	front	rear
A	2.6	3.4
B	2.3	2.7
C	5.0	2.7
D	3.0	2.8
E	2.9	2.0



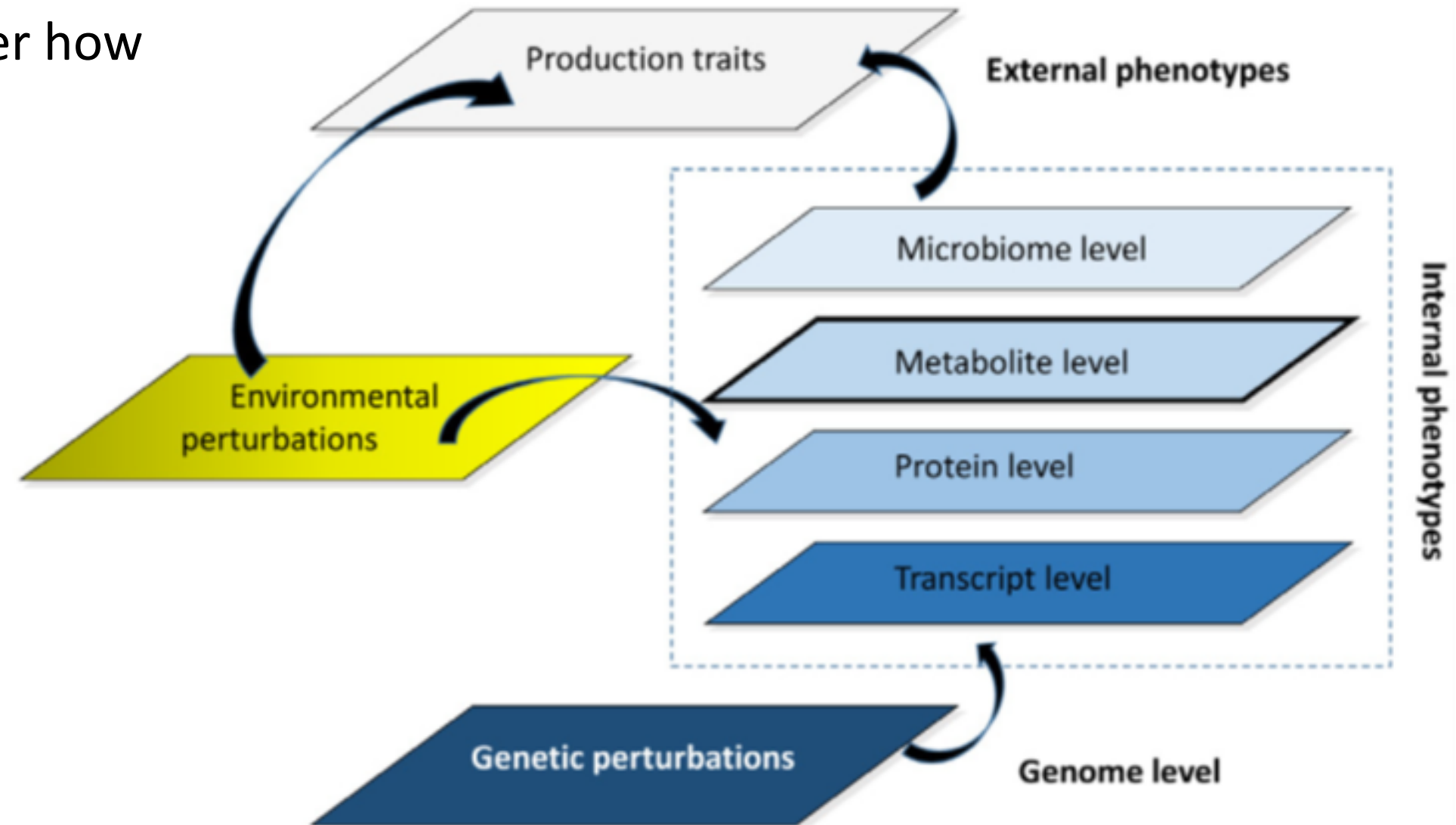
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 - behaviour
 - methane emission ← spectroscopy
- Classical traits can be expanded:
 - more objective → more accurate
 - longitudinal
 - non-invasive
 - many animals
 - commercial conditions



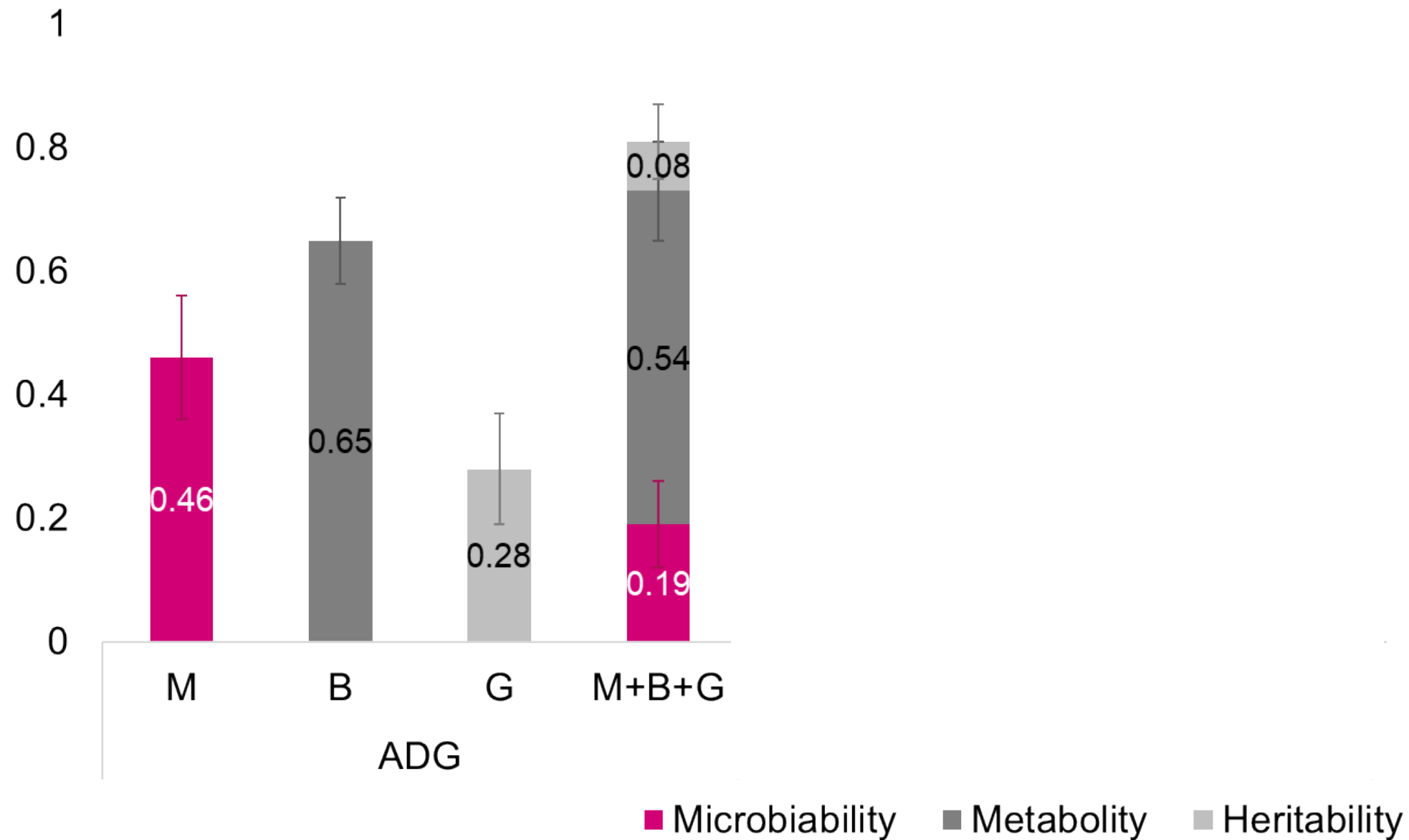
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- **Novel options to explore**

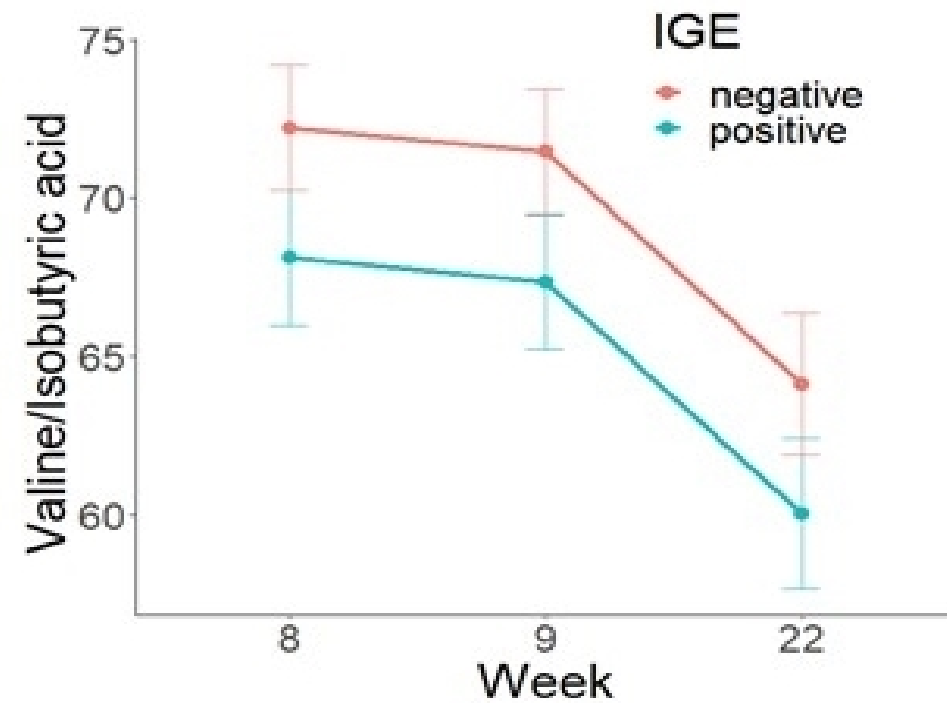
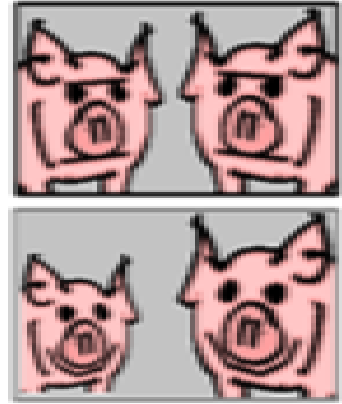
- better biological understanding: discover how
 - selection barriers can be overcome



Phenotypic variance explained

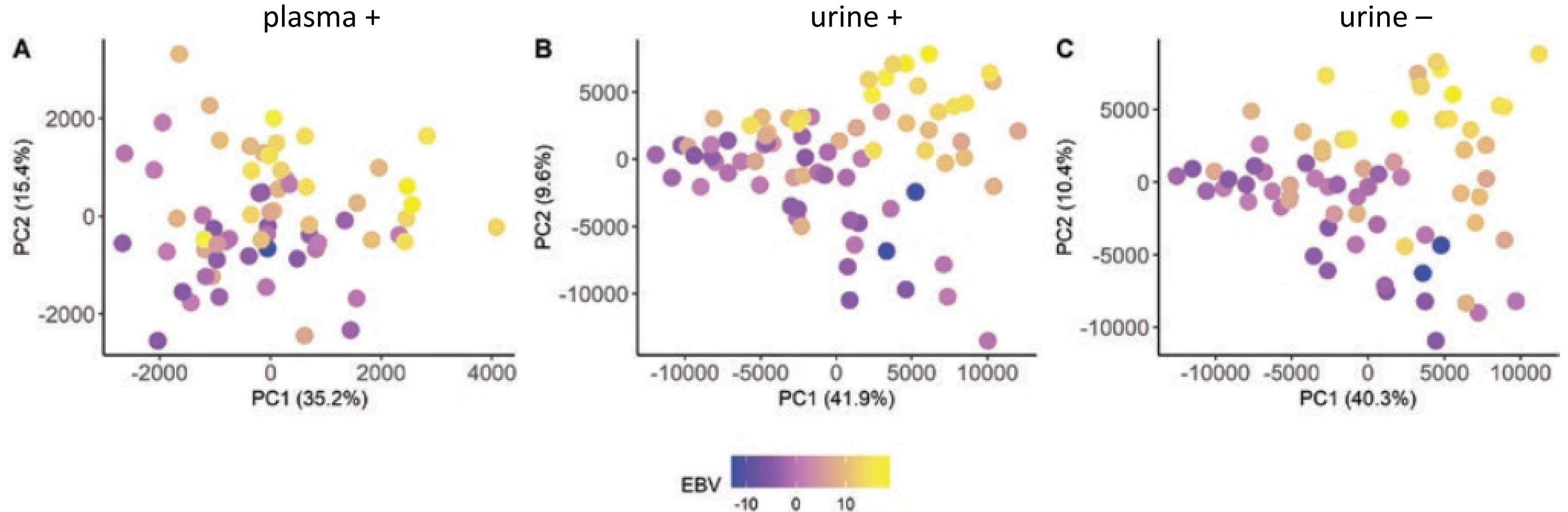


Metabolite levels & Indirect Genetic Effects



Pigs with **negative** or **positive** indirect genetic effects on growth of their penmates
Valine, leucine, isoleucine levels

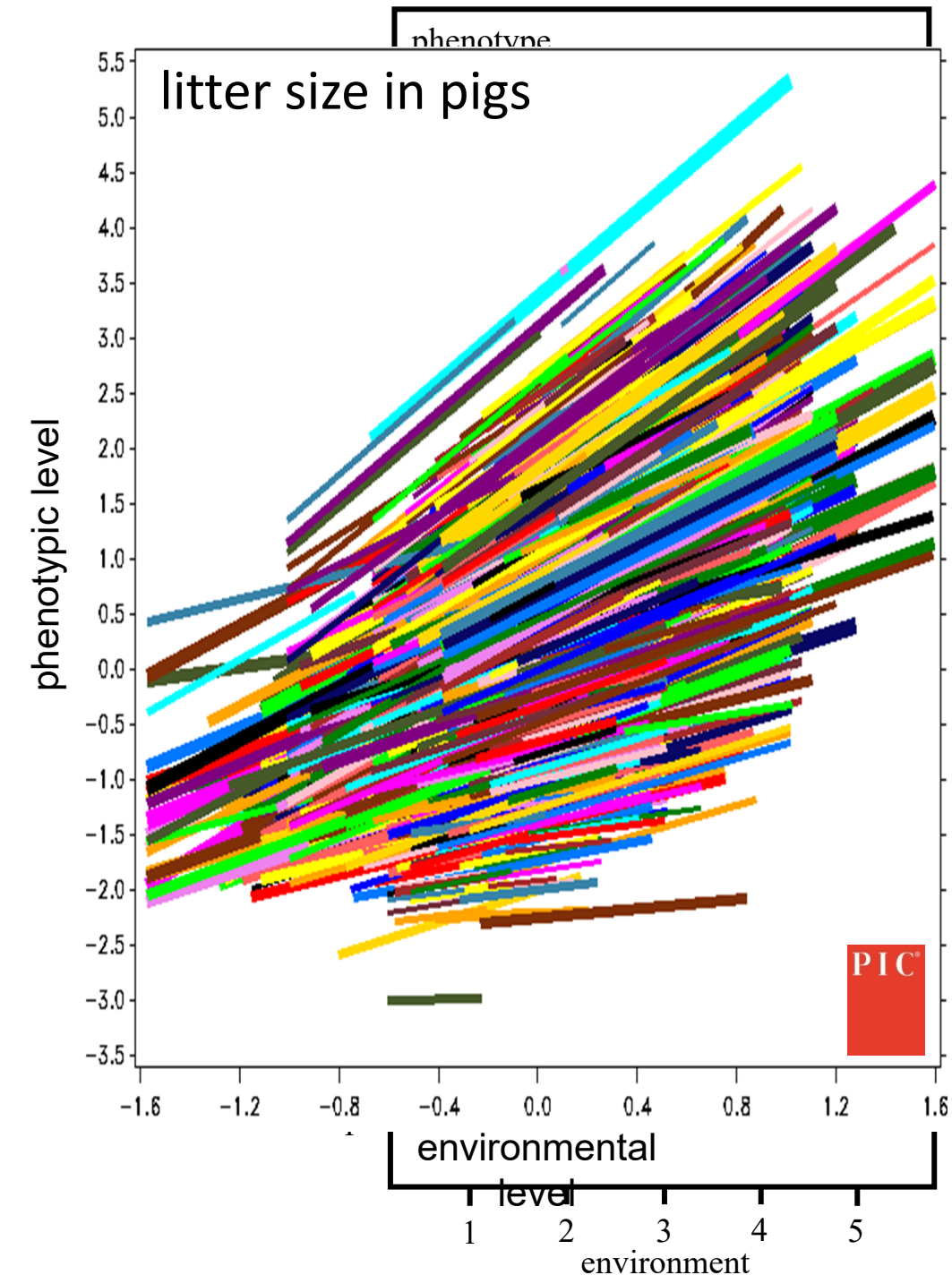
Metabolite levels & nitrogen efficiency



Metabolite ion intensities that discriminate between **low** and **high** EBV for nitrogen retention:
First and second Principal Components

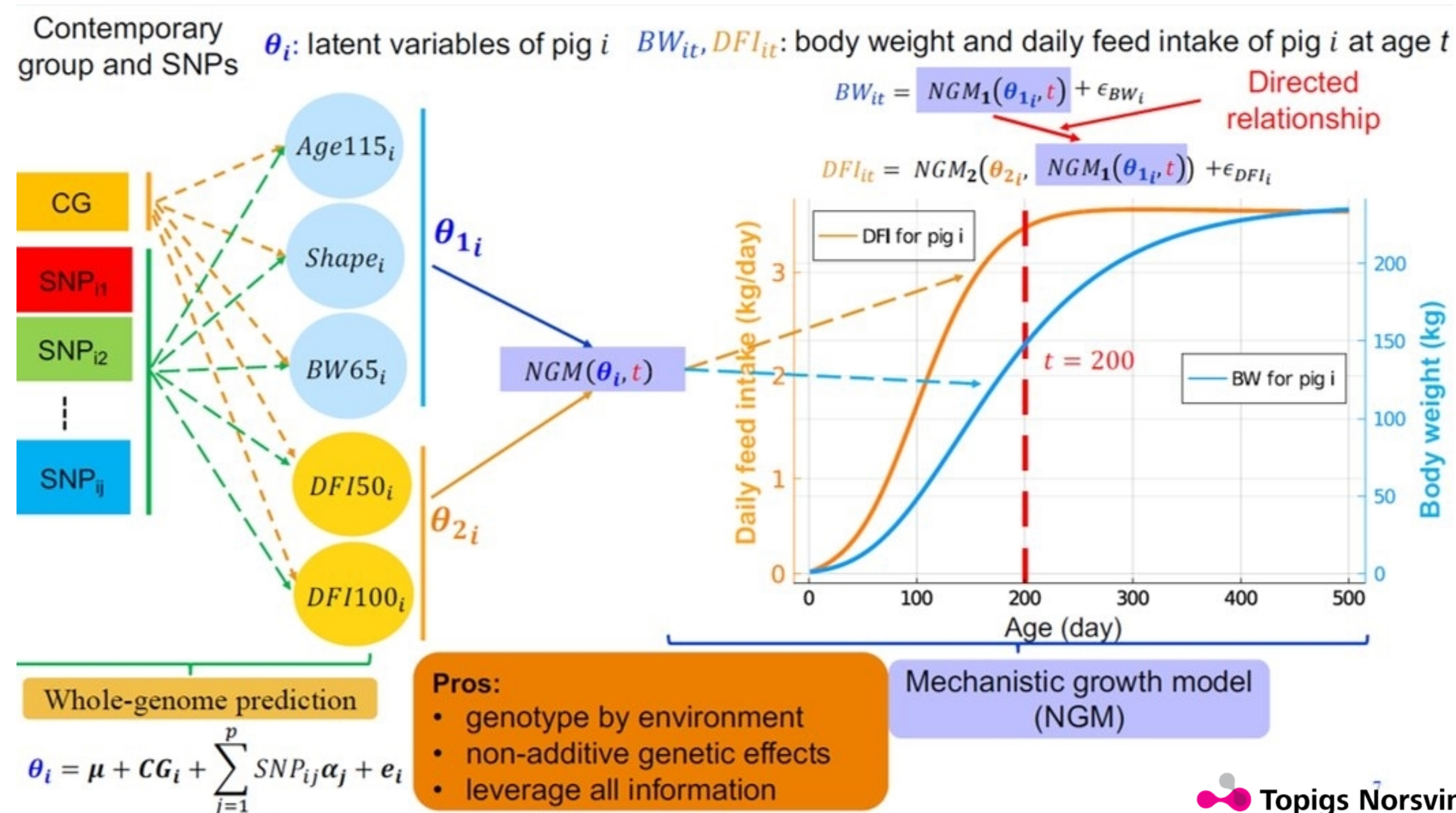
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- Two novelties: new traits, expanded classical traits
- Novel options to explore
 - better biological understanding: discover how
 - selection barriers can be overcome
 - to optimize economic weights dynamically
 - redesign breeding schemes
 - explicitly allow for high-dimension phenomics (~ reaction norms)

Genotype x Environment Interaction
Environmental Sensitivity

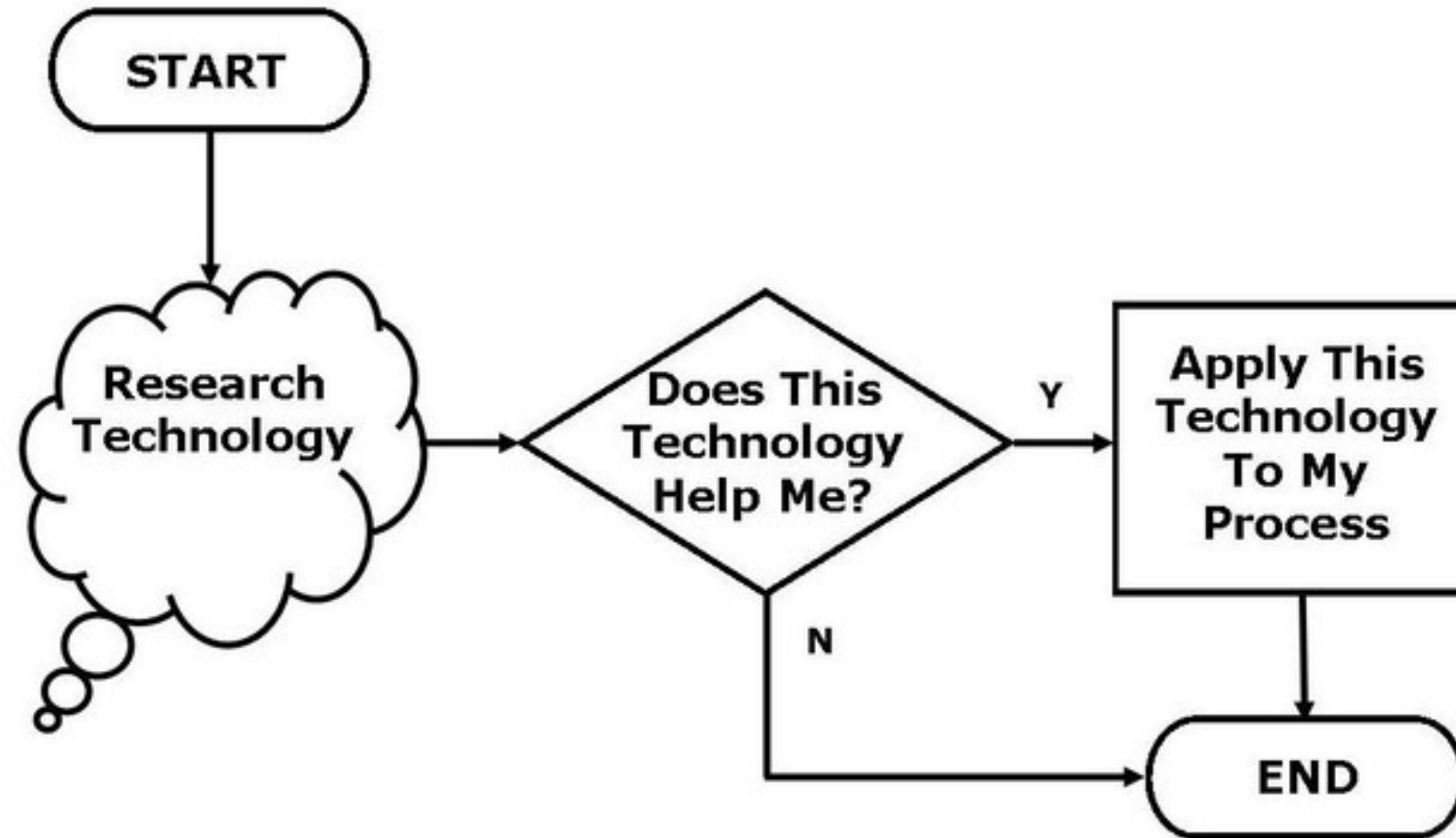


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 - better biological understanding: discover how
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 - to optimize economic weights dynamically
 - redesign breeding schemes
 - explicitly allow for high-dimension phenomics (~ reaction norms)
 - use phenomics data to integrate mechanistic biological models into genetic evaluation

Mechanistic biological models



Interesting. And what do we do with it?



- Phenomics: the acquisition of
- *Big data* challenges ← heterogeneity
- Two novelties: new traits, ex

• Novel options

- better biological understanding or how to optimize eco

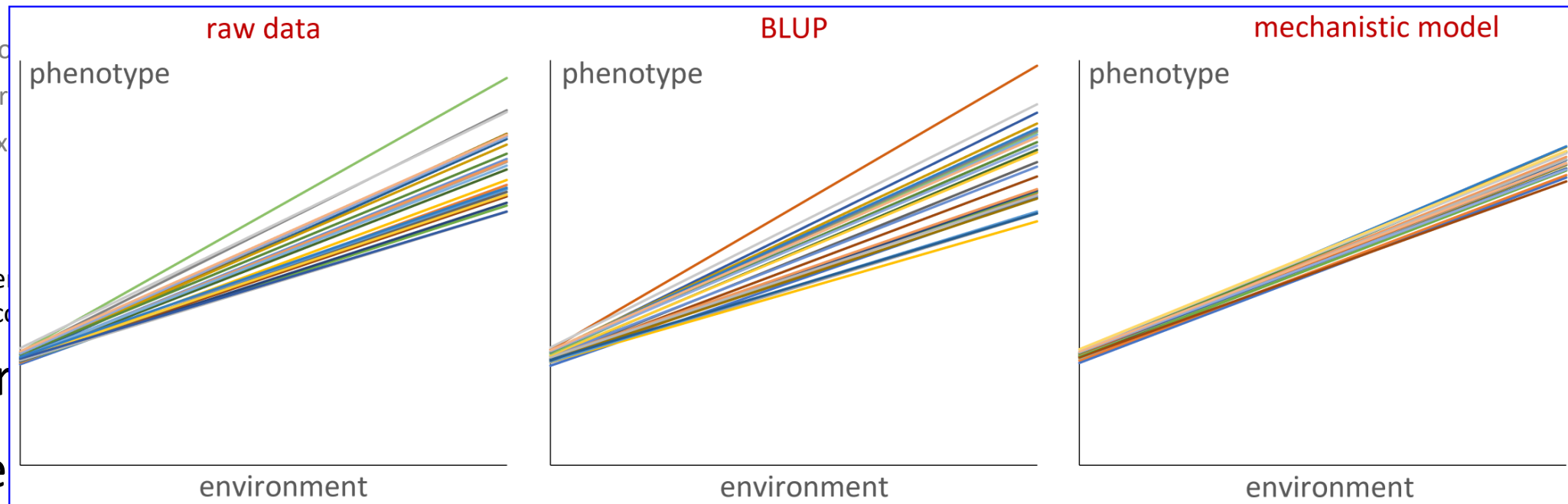
• redesign breeding

• use phenomics

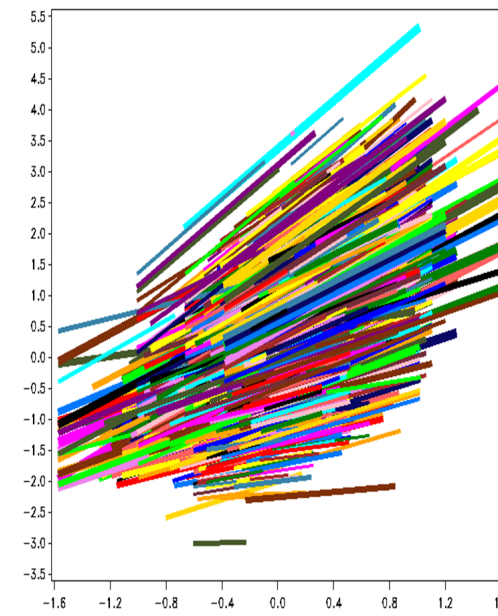
- example: wheat yield per hectare in France, 28 genotypes, 16 locations

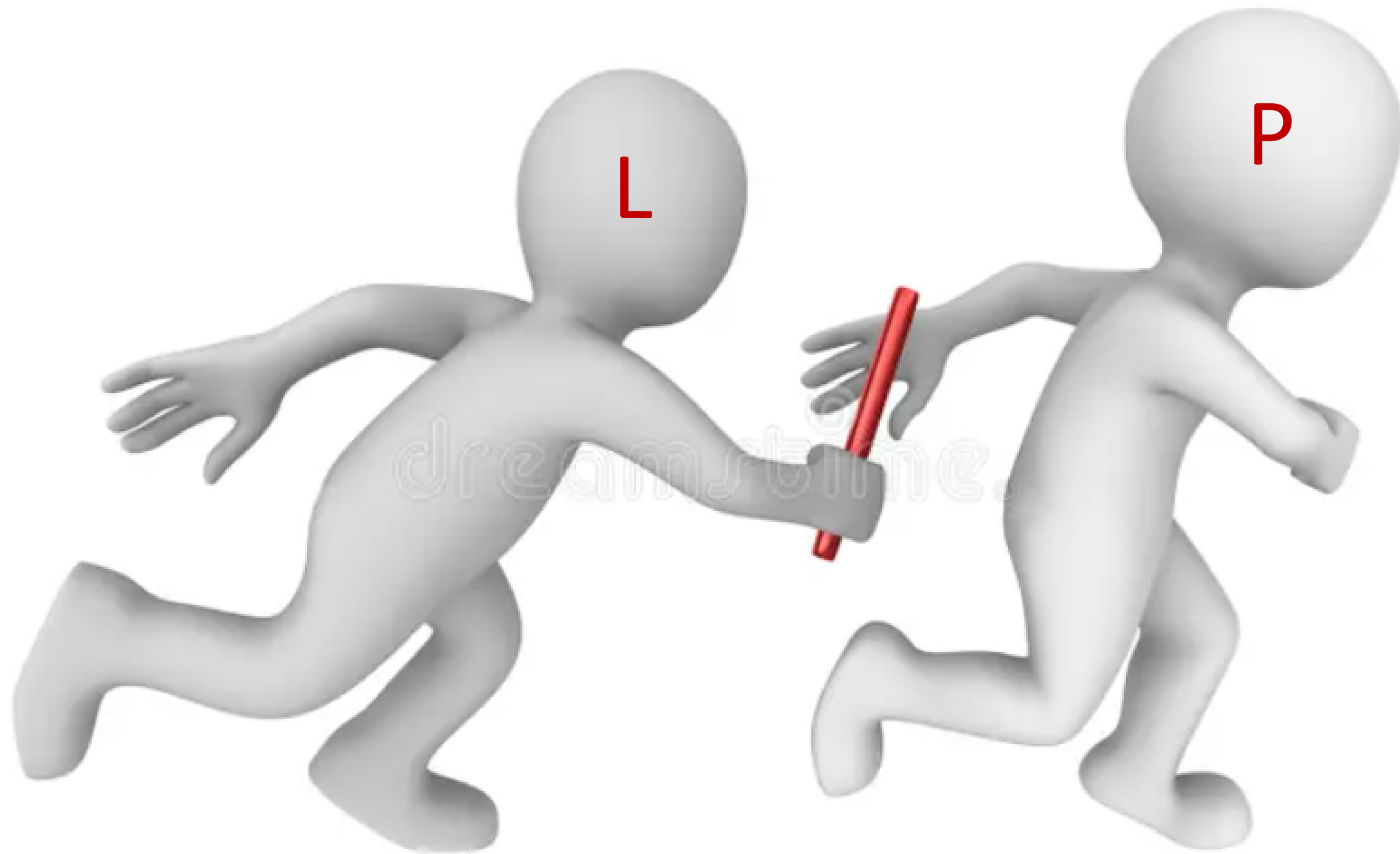
- raw data → BLUP → mechanistic model

- better capture of non-linear effects
- more uniform (and much more accurate) slopes → less evidence for G×E
- a linear model of a non-linear system produces phantom interaction effects
- non-linear model is much more appropriate



reaction norms)
evaluation





- Phenomics: the acquisition of high-dimensional phenotypic data on an organism-wide scale (Houle et al, 2010)
- *Big data* challenges ← heterogeneity & rapid change over time of the data, not because of their size
- Two novelties: new traits, expanded classical traits
- Novel options to explore

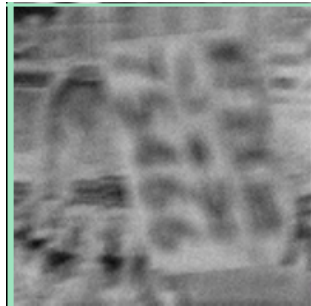
Key factor: animal identification

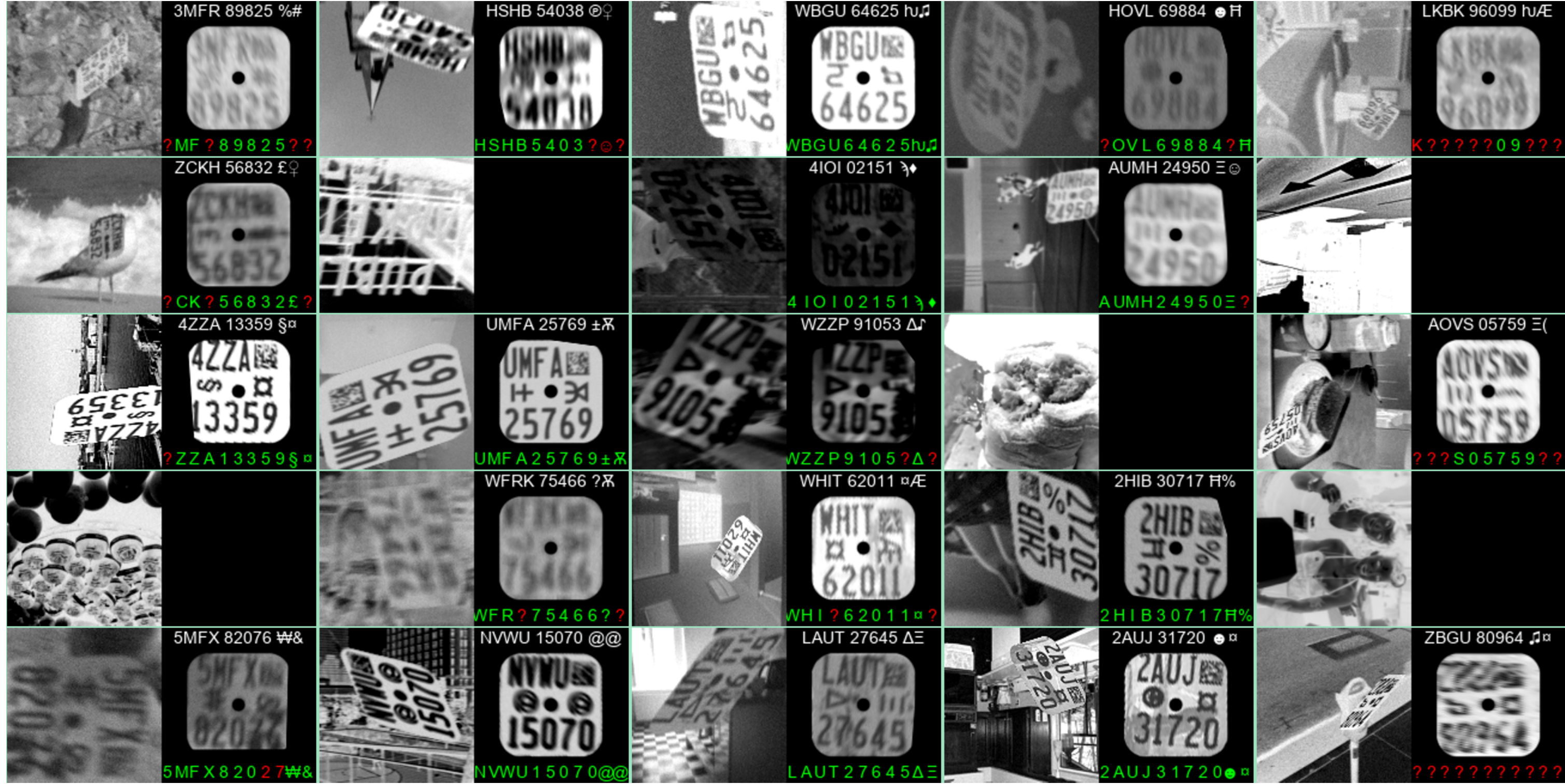
- Necessary, for breeding
- Difficult



LitterMax eartags

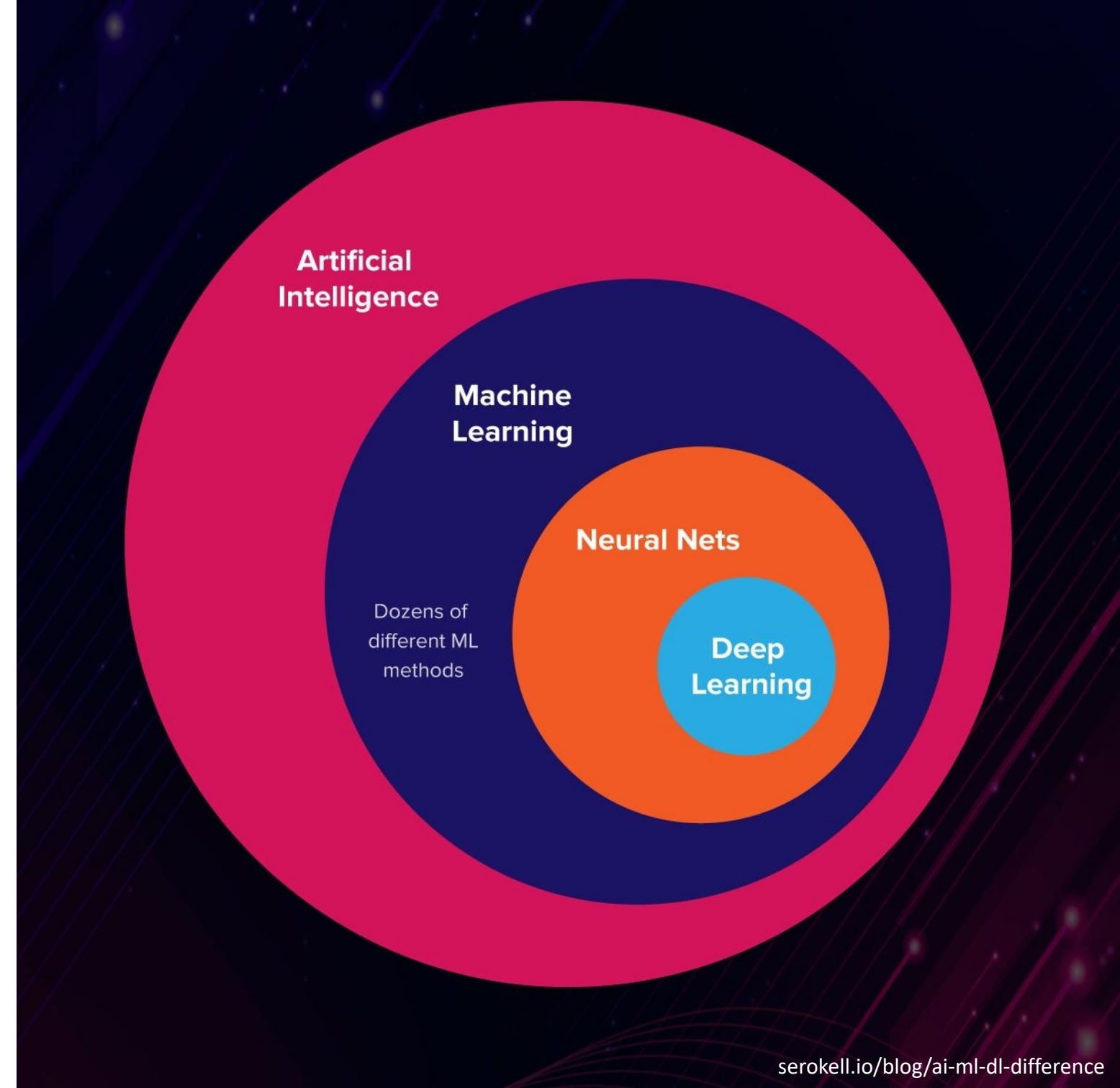
- 2 mio permutations for ID
- QR code
- error trapping codes
- the trick is to make the robot read them correctly





- Phenomics: the acquisition of high-dimensional phenotypic data on an organism-wide scale (Houle et al, 2010)
- *Big data* challenges ← heterogeneity & rapid change over time of the data, not because of their size
- Two novelties: new traits, expanded classical traits
- Key factor: animal identification
 - necessary
 - difficult
- Key factor: internet access
 - challenging in rural areas
- Key factor: heterogeneity of the data
 - non-random missing data
 - imputation: Variational Auto-Encoders, Generational Adversarial Networks
- *There is an enormous range of analytical tools to be developed*

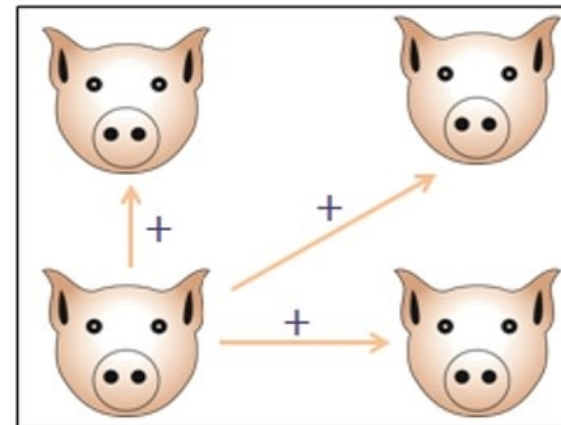
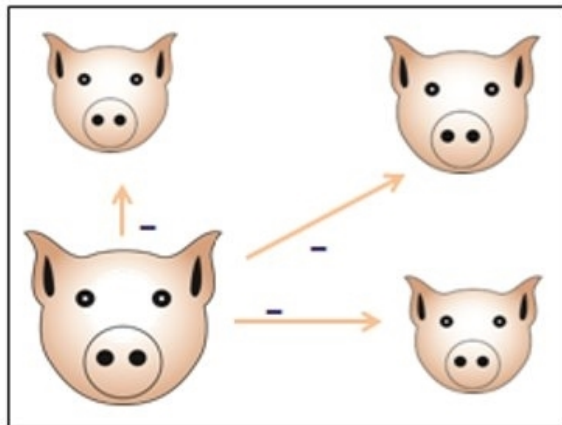
Statistical methods



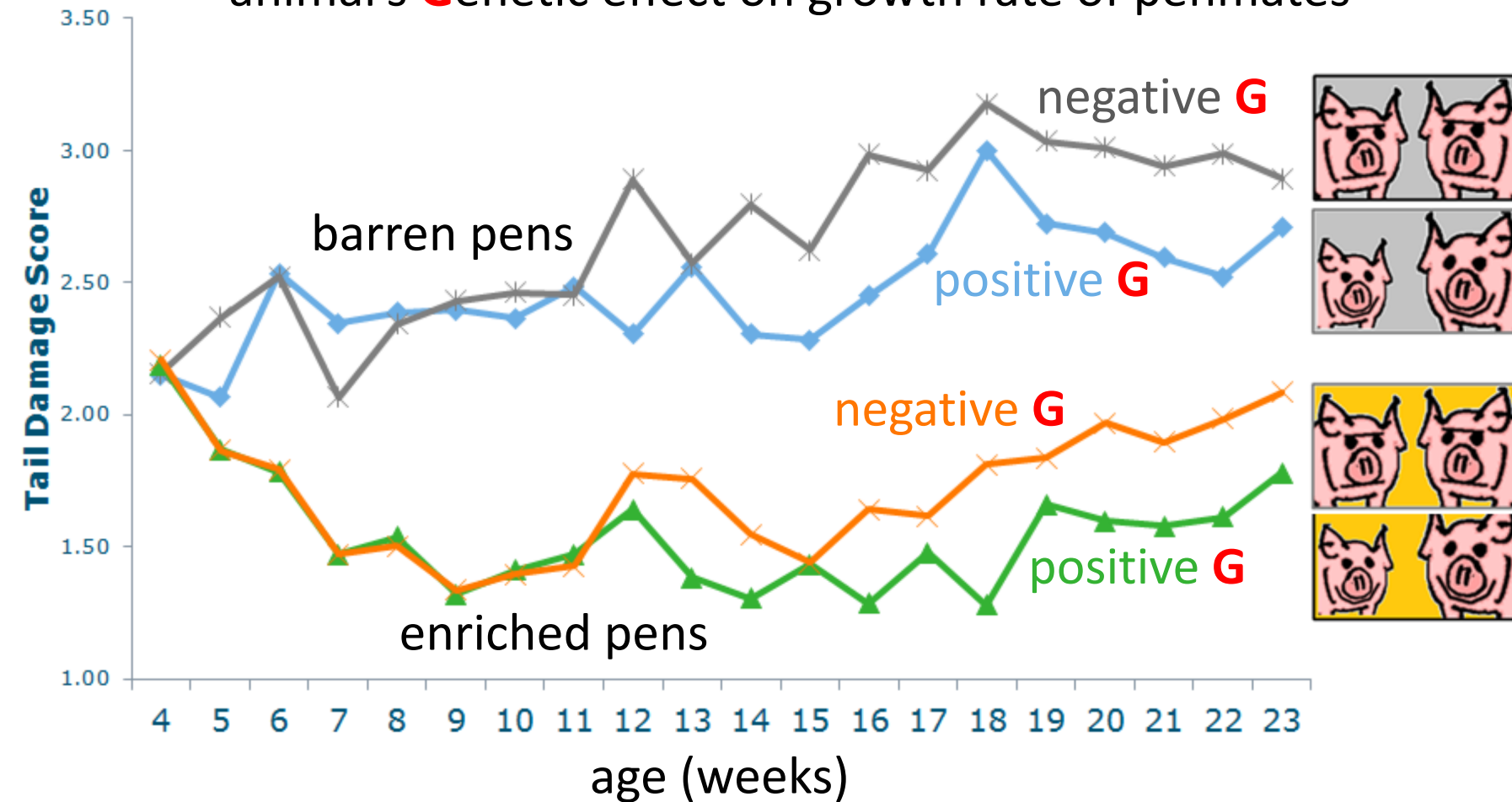
Classic modeling

Multi-trait model with individual behavior and production trait

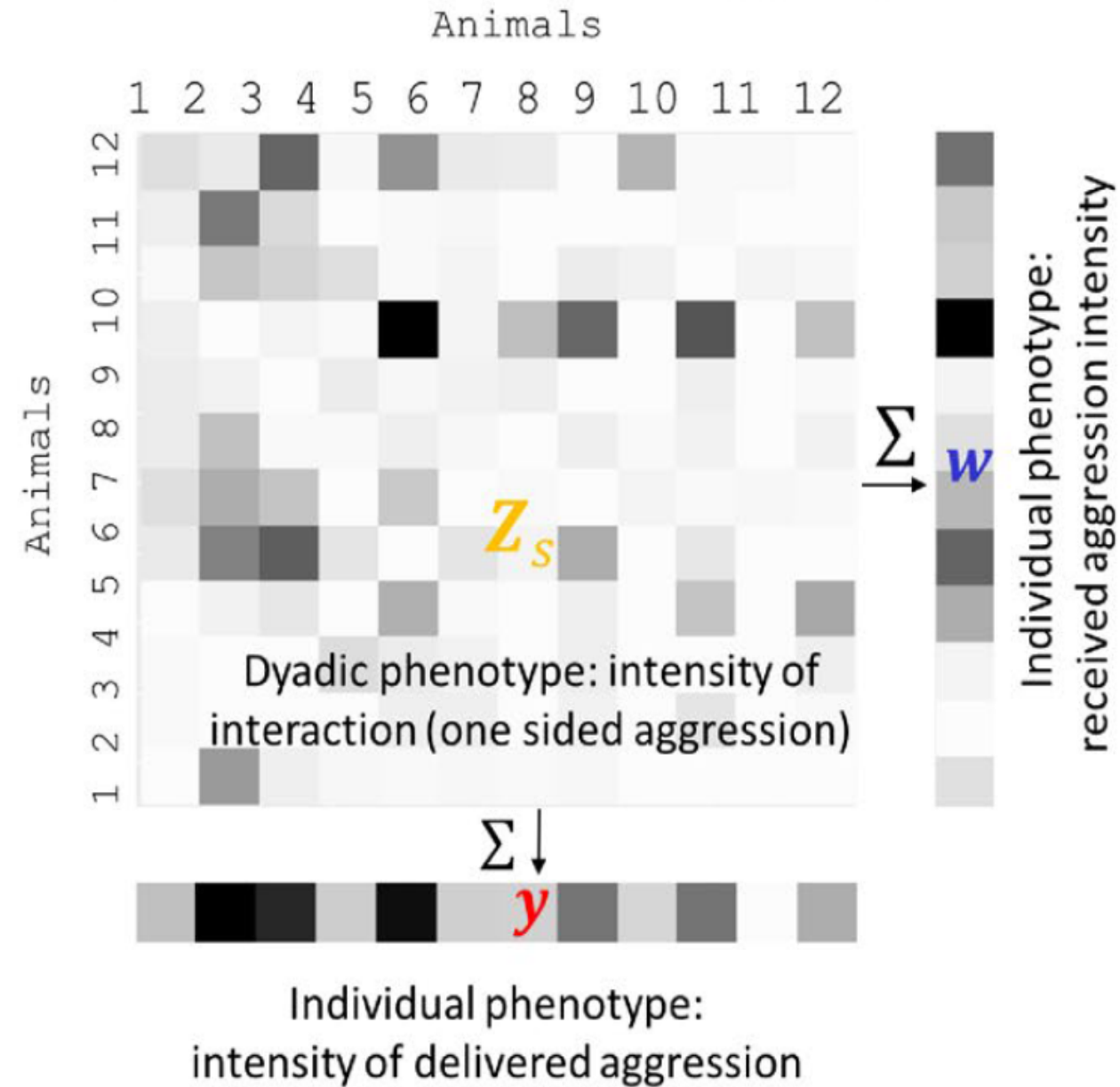
$$\begin{pmatrix} x \\ w \\ y \end{pmatrix} = f(M)$$



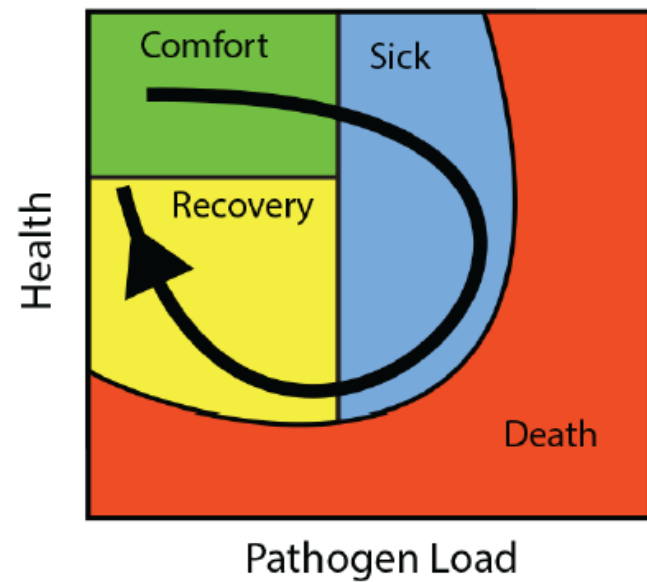
tailbiting:
animal's **G**enetic effect on growth rate of penmates



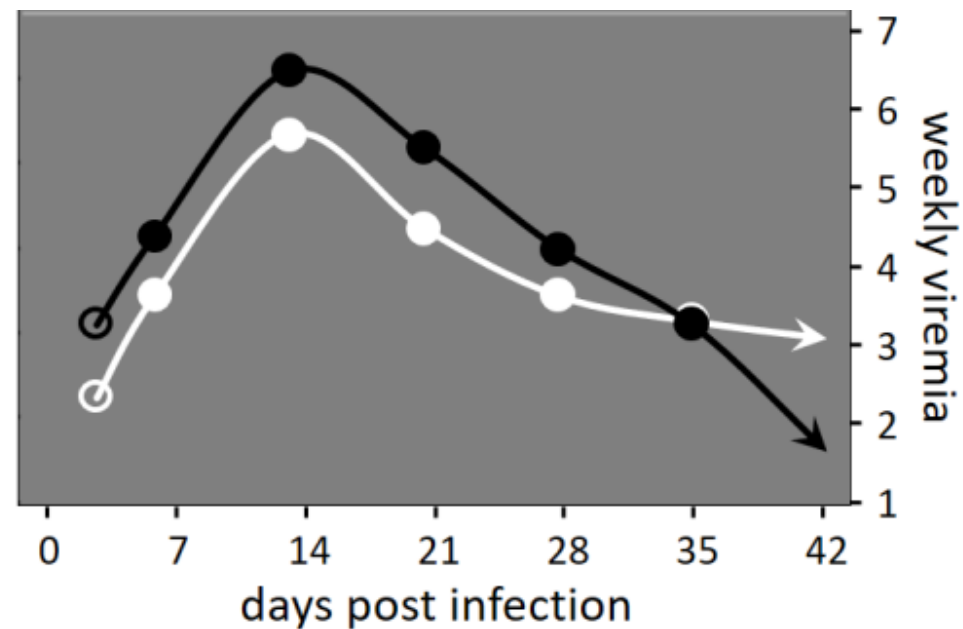
Behavioral phenotyping

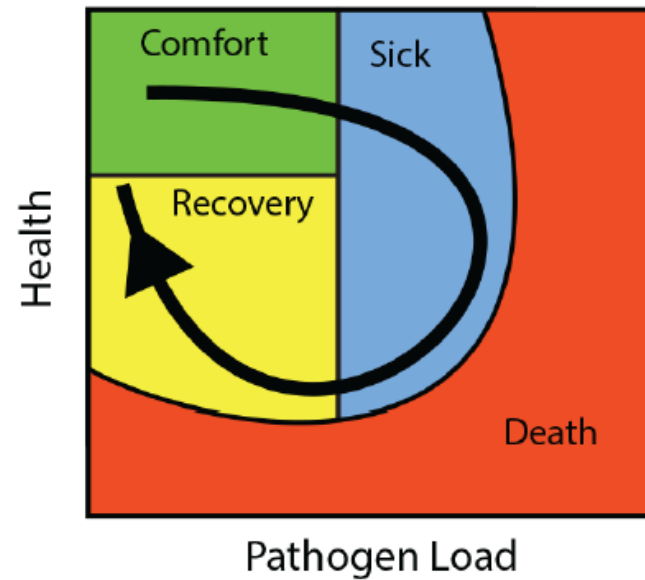


Novel models

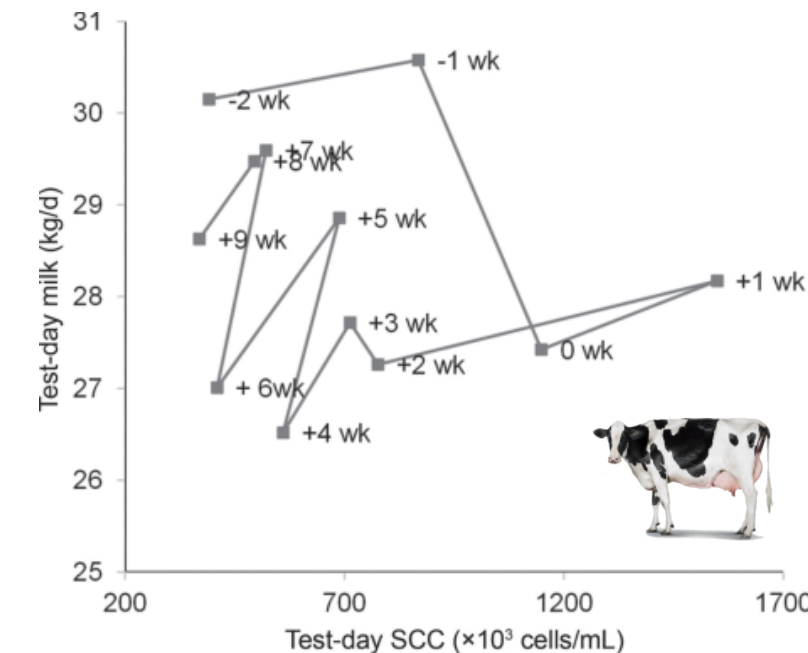
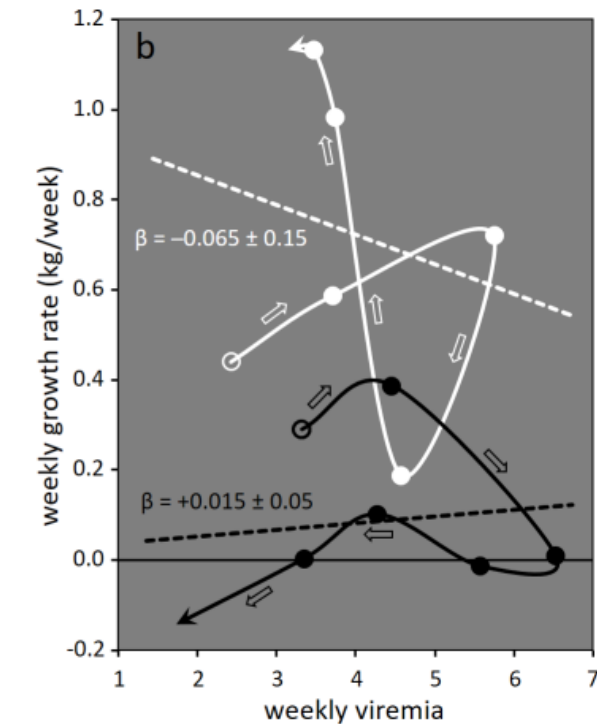


Example: two pigs infected with PRRS virus and followed for 6 weeks





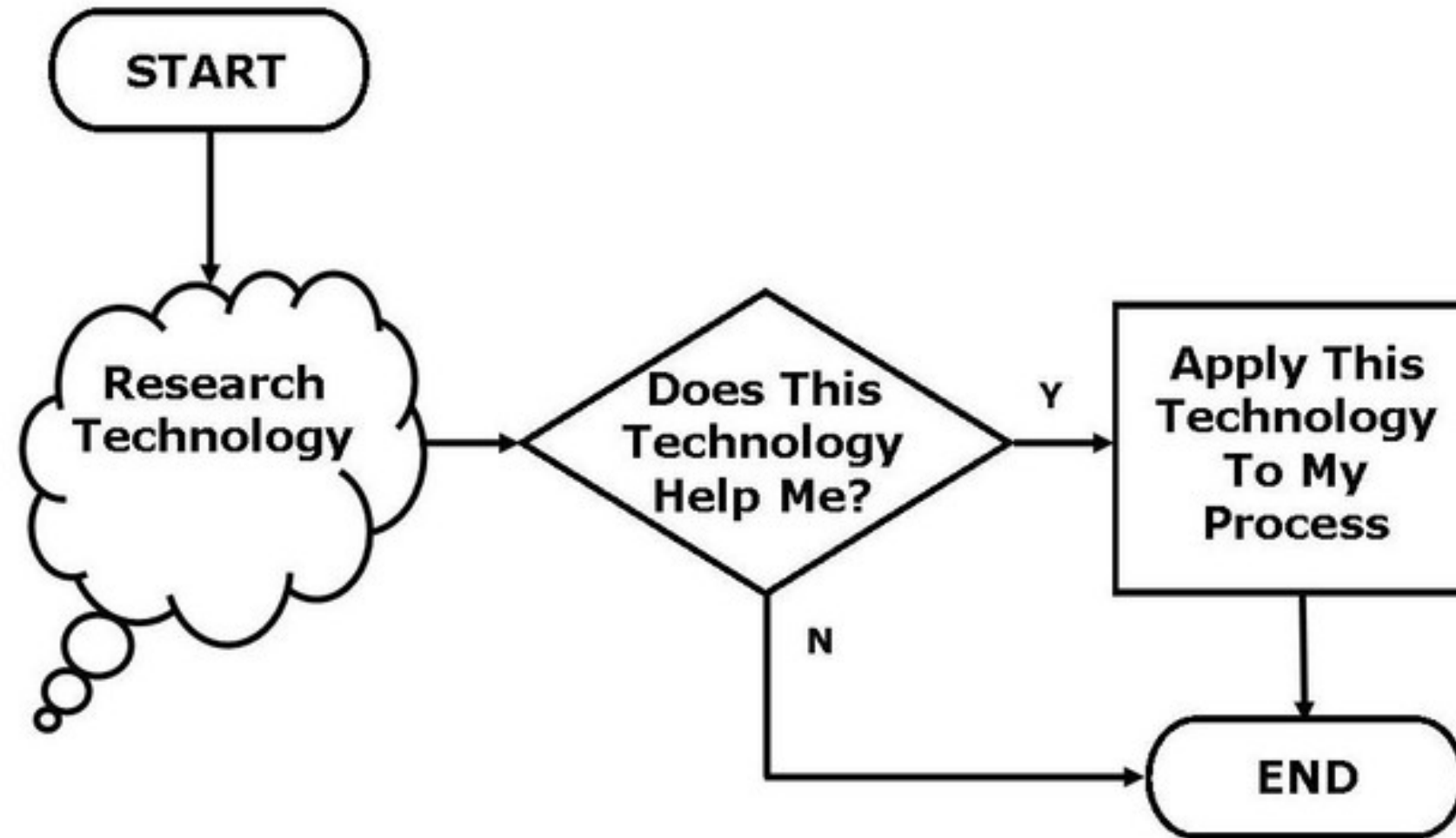
- Stages of infection associated with the strongest loss of health & performance
 - insight into underlying resilience mechanisms and target genes
 - design effective, timely & targeted treatment
- “These patterns cannot be described by mathematical functions”
- Requires novel indicators to
 - capture the trajectory characteristics
 - support routine genetic evaluation

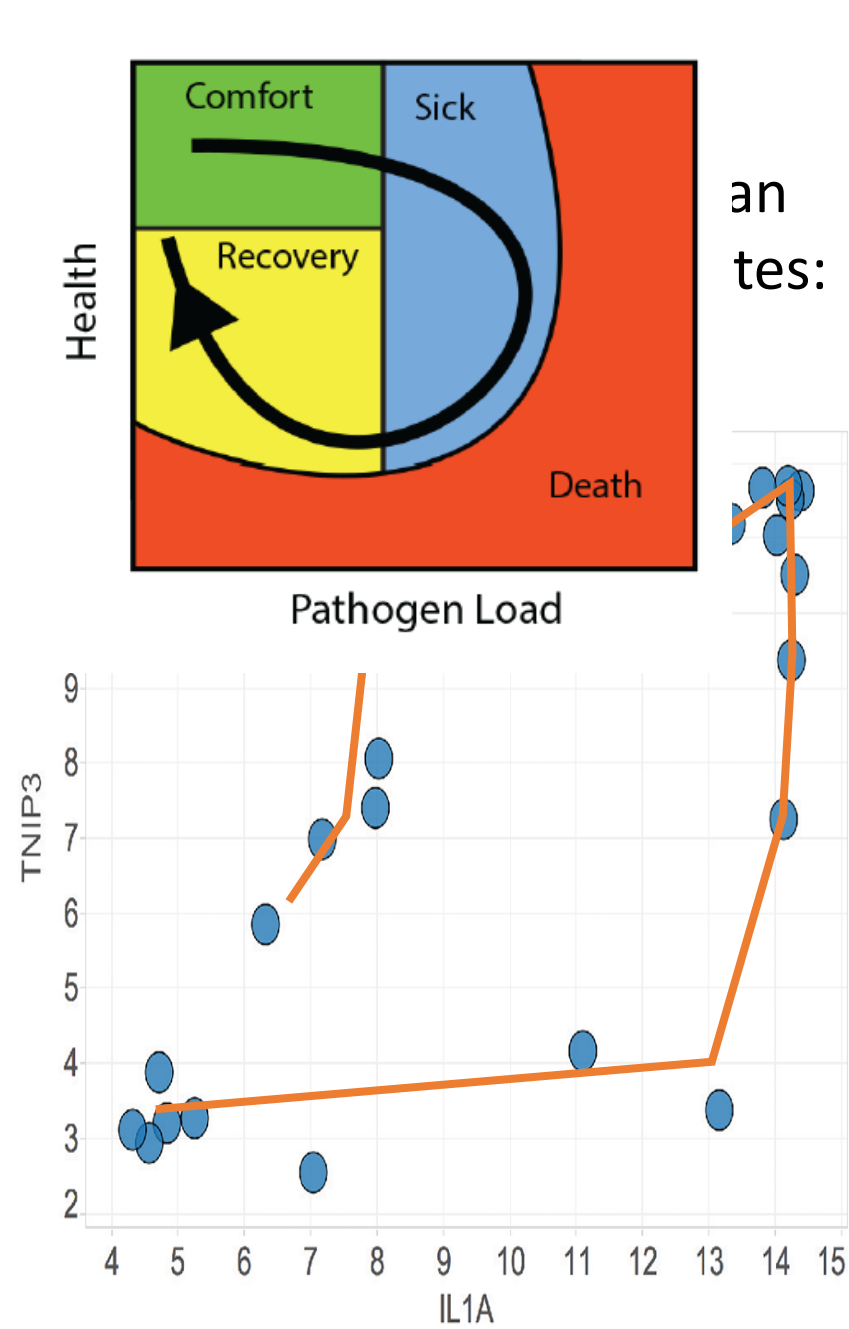


Detilleux (2018) doi.org/10.3168/jds.2017-13976

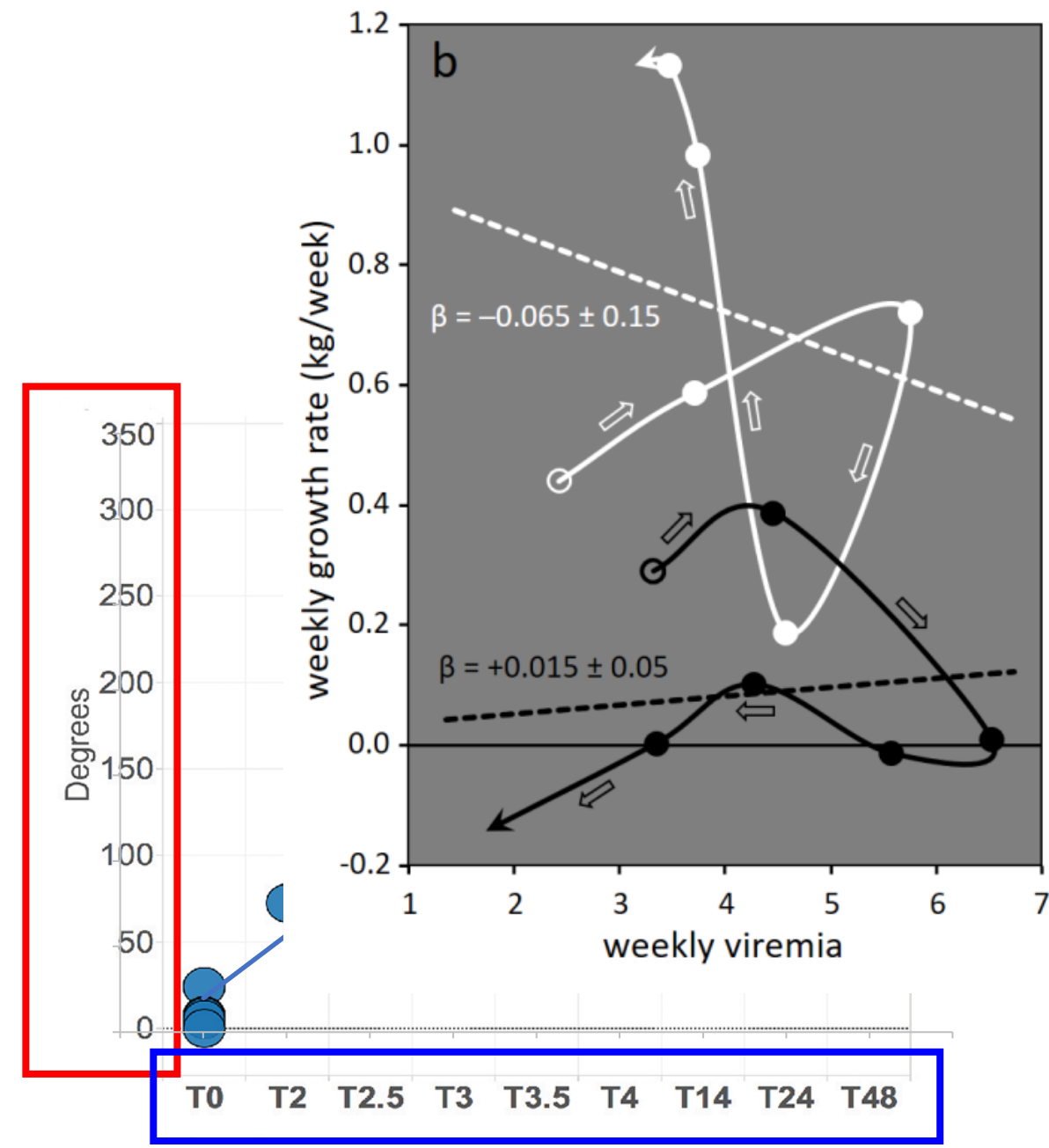
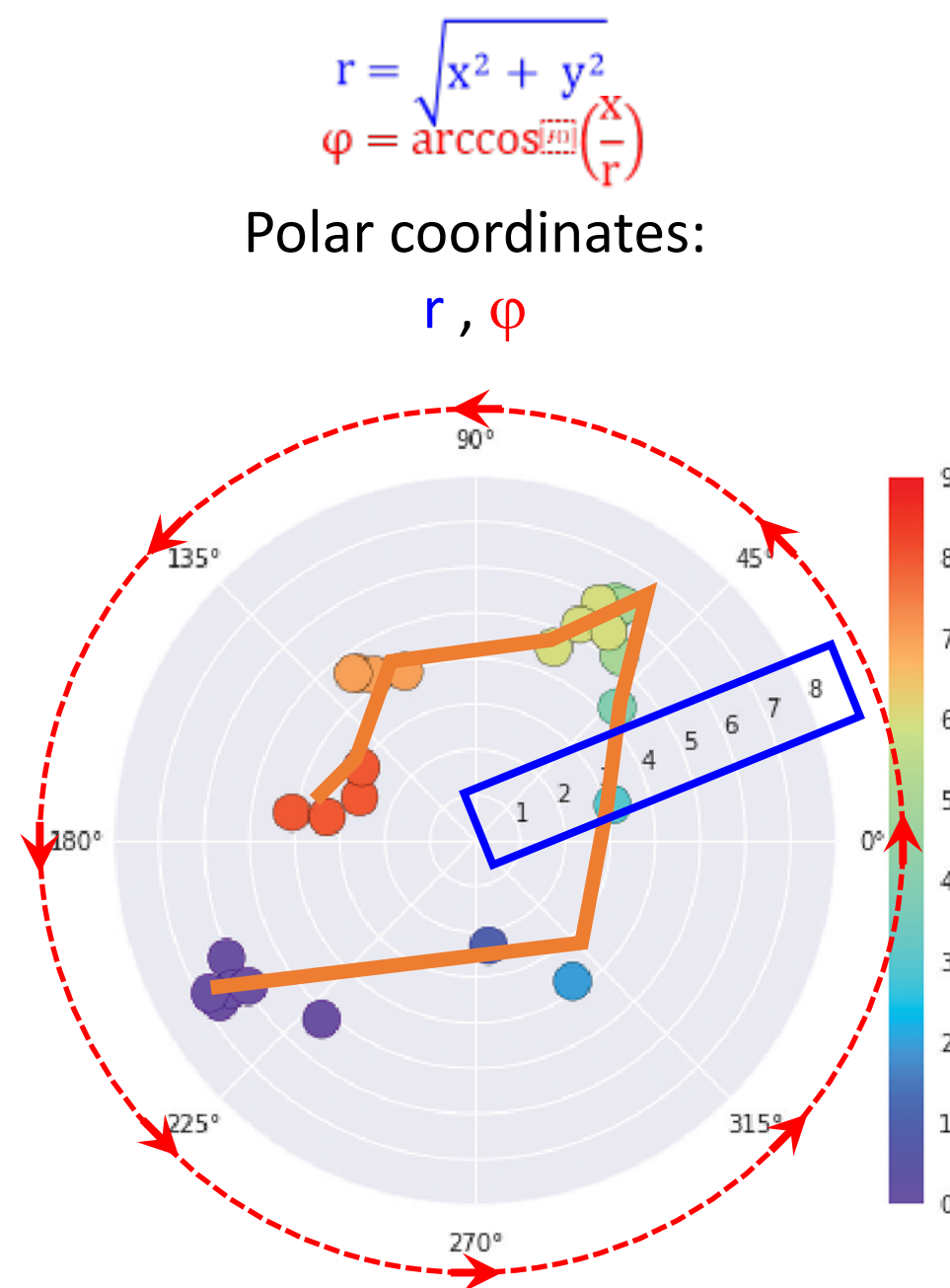
Knap & Doeschl-Wilson (2020) doi.org/10.1186/s12711-020-00580-4

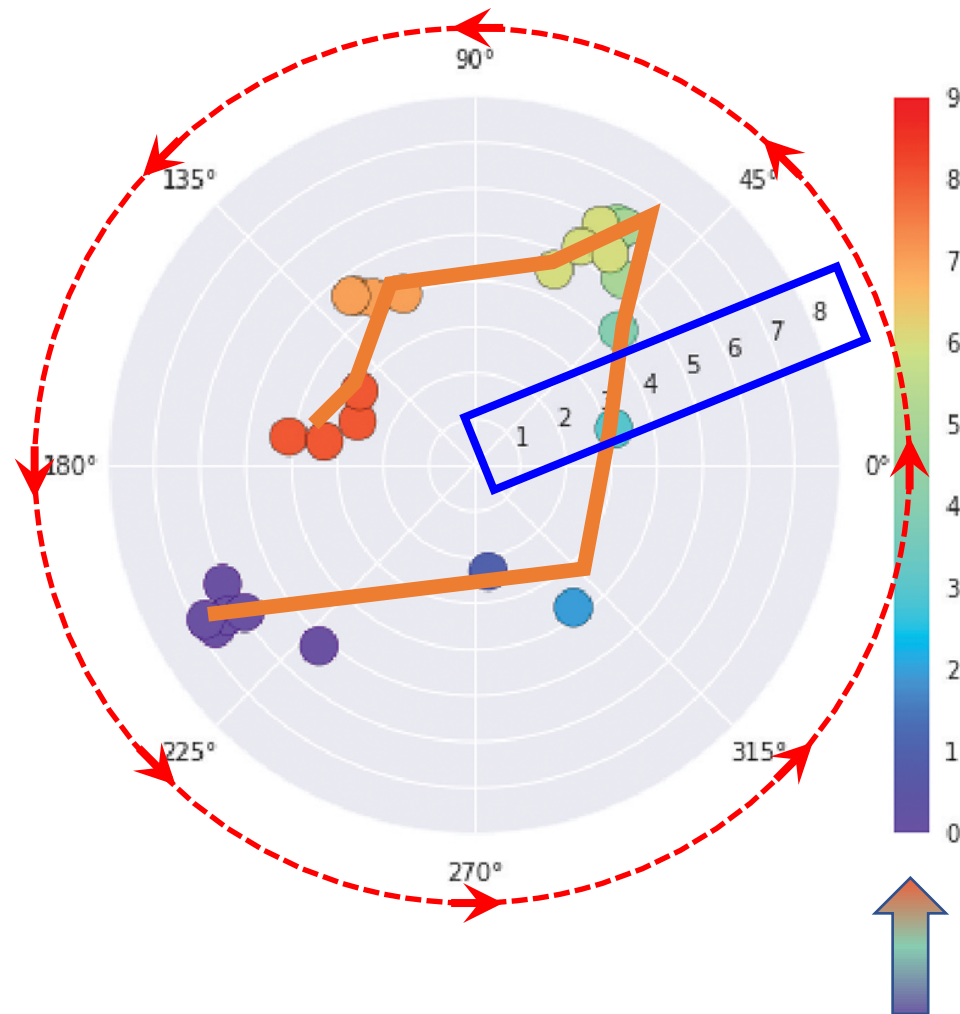
Interesting. And what do we do with it?



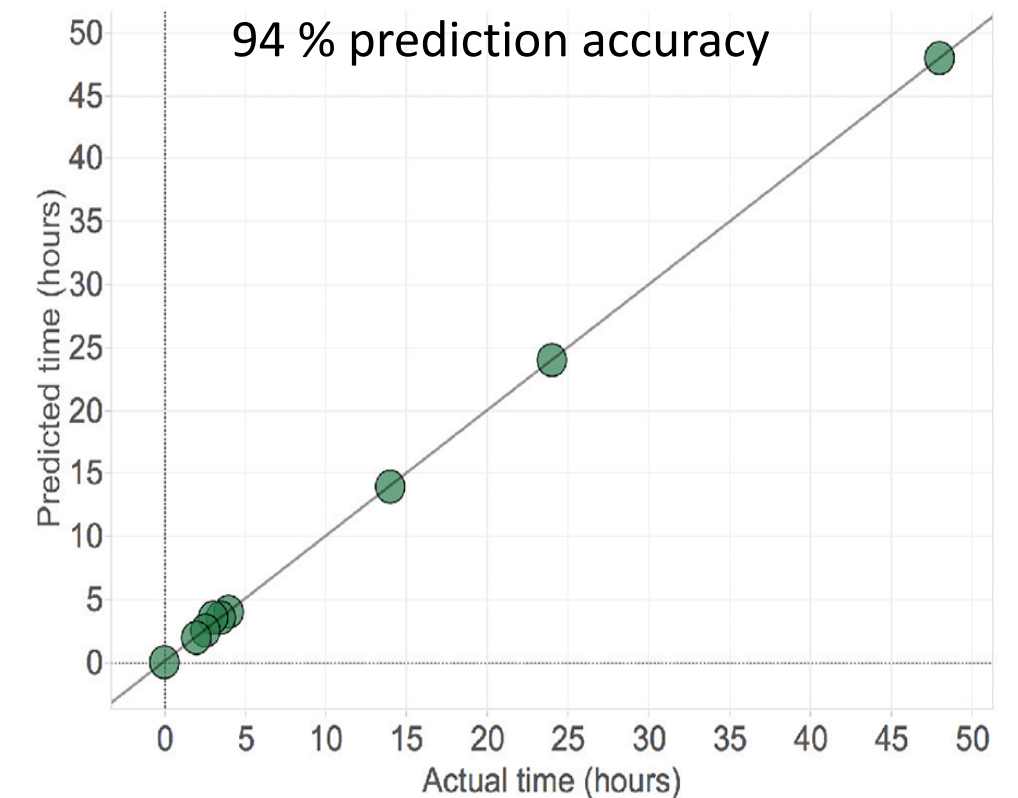
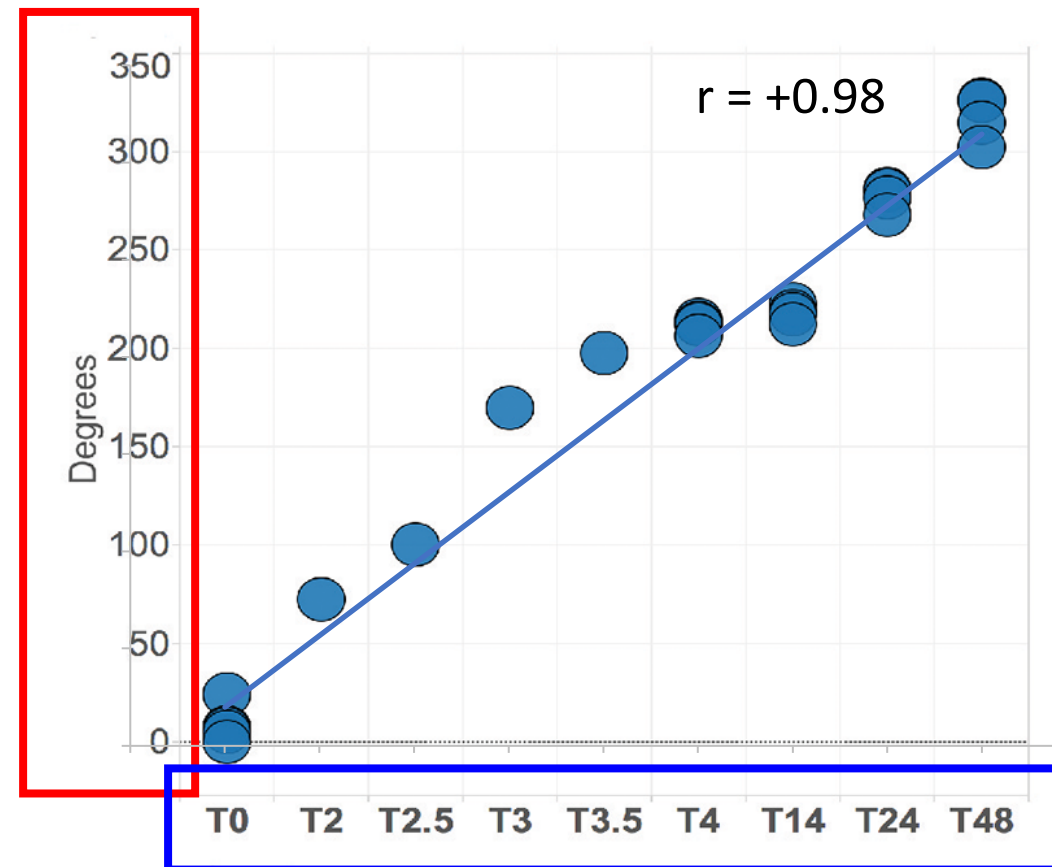


antes:

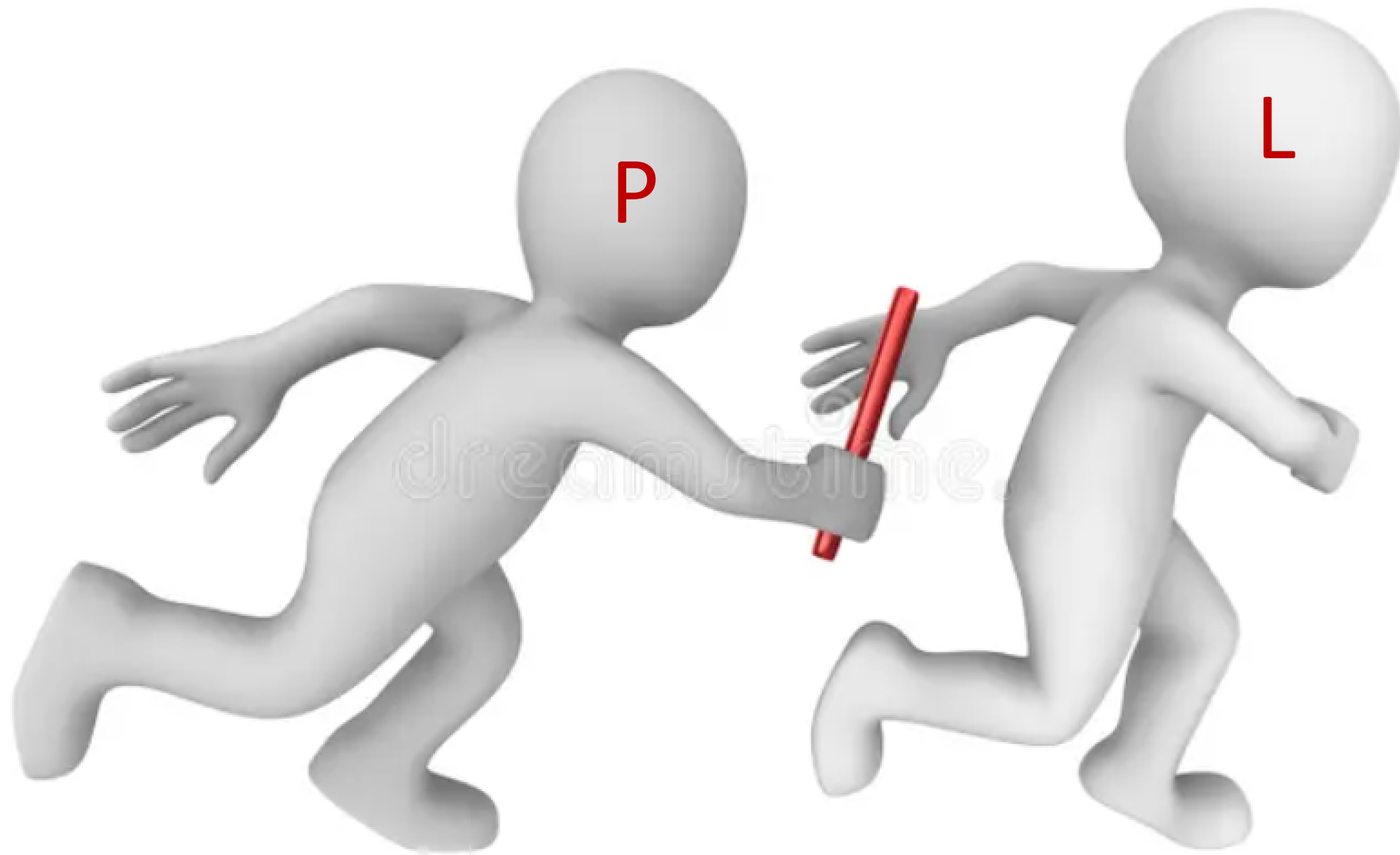


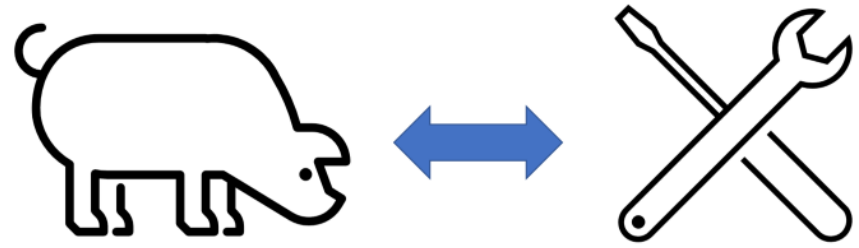


The **effect** is on the timing of an event (the colours of above)



- → a simple linear prediction equation
- crossvalidation:
- predicted **effect** in samples that were not in the training dataset





Conclusions

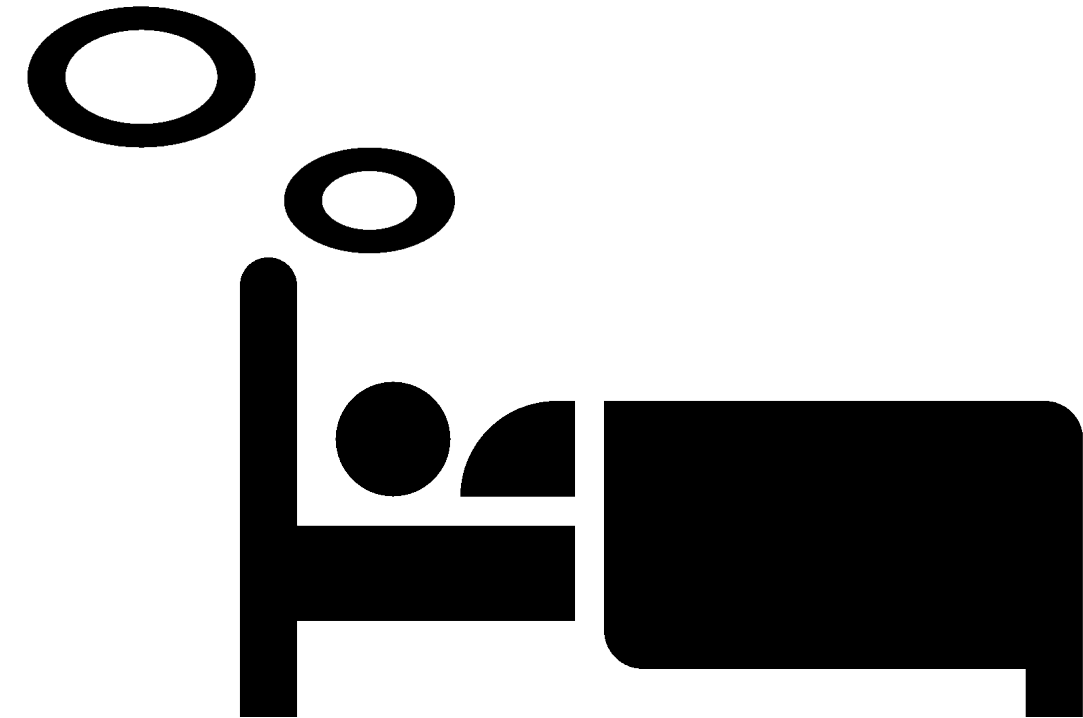
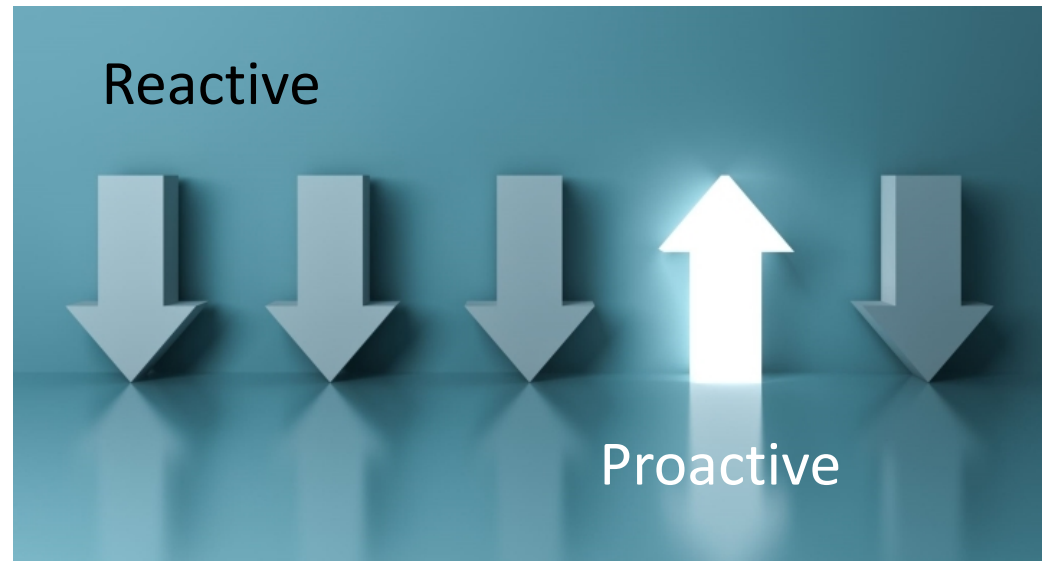
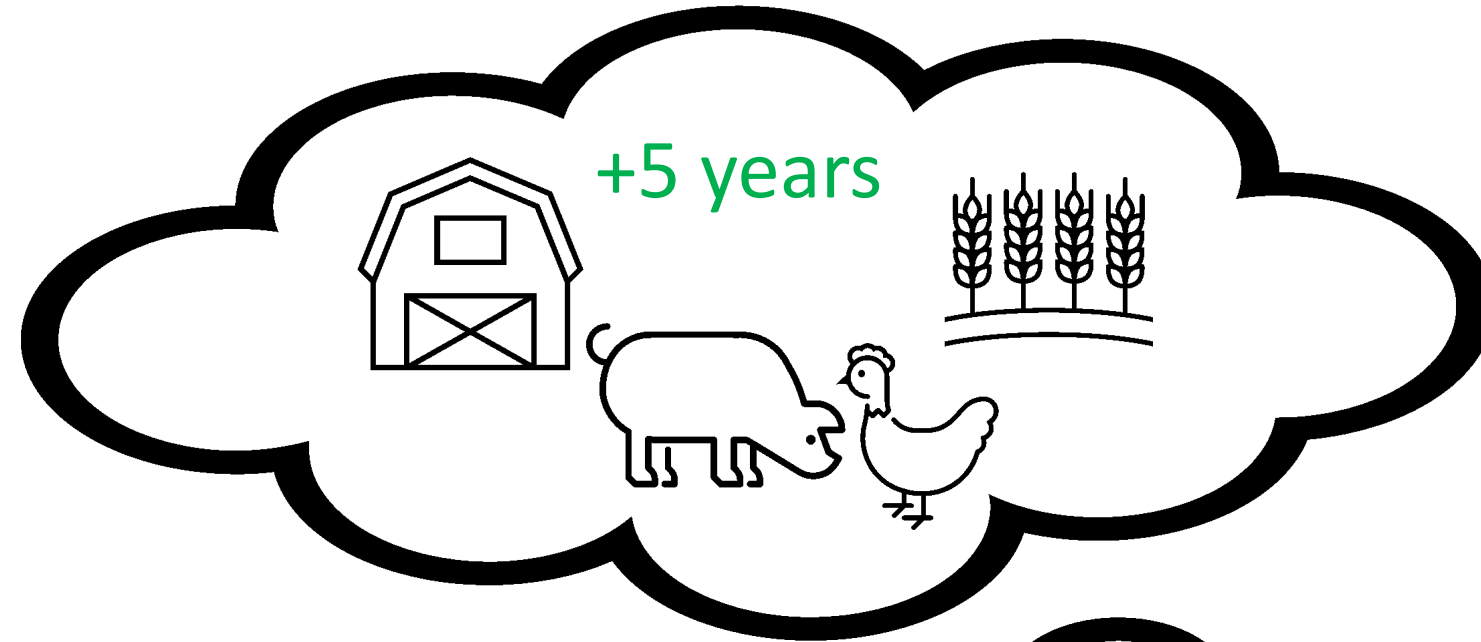
1. Increased production levels have often led to trade-offs due to genetic antagonisms
2. Genetic antagonisms can be neutralized: select for both traits at the same time
3. Rapid increase in availability of tools

Understand the existing phenotype

Create new phenotypes

to simultaneously select for antagonistic traits

We have a dream...



Session 77

Trade-offs between health, production and welfare in pigs and poultry
Which tools do we have, and which research is needed?

The future of pig and poultry breeding

Lisanne Verschuren & Pieter Knap

August 2023