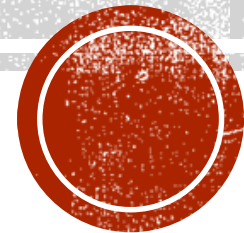


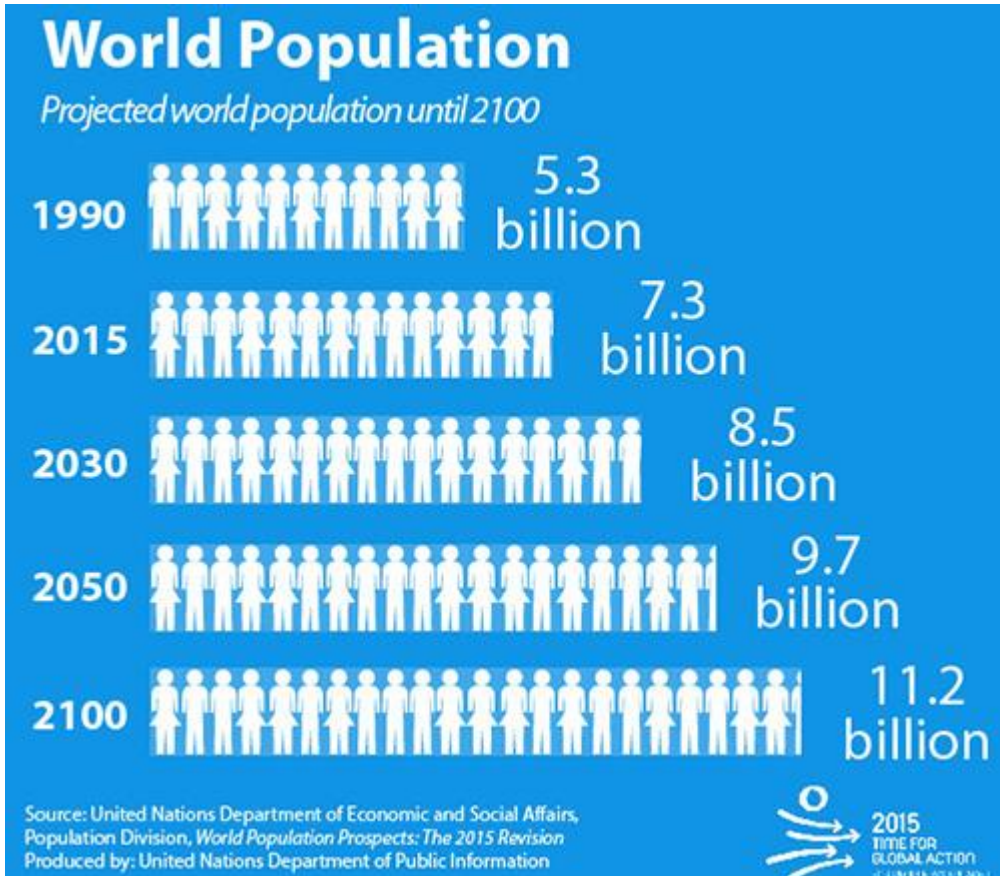
FEEDING ON LARVAE OF BLACK SOLIDER FLIES DOESN'T SUBSTANTIALLY MODIFY CHICKEN CAECAL MICROBIOTA

Moula, N., Cabaraux, J.-F., Dawans, E., Taminiau, B., Detilleux, J.

Veterinary management of animal resources



WE NEED TO FEED THE WORLD



Mostly in developing countries
Food production must increase by 70%

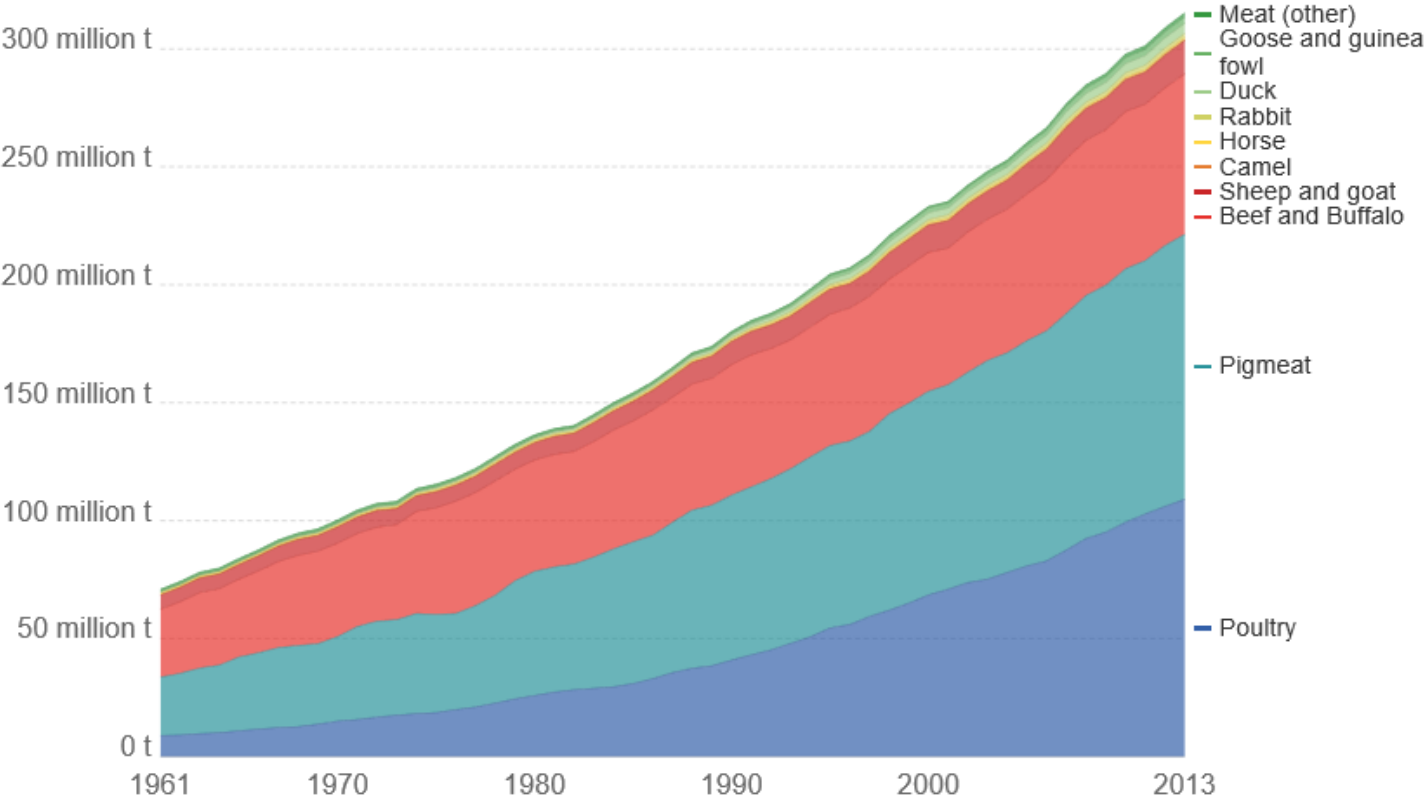
https://www.croatiatraveller.com/southern_dalmatia/Dubrovnik/cruise-schedule.html/



MEAT PRODUCTION AND SUPPLY

Meat production by livestock type

Meat production by commodity or product type, measured in tonnes per year. All data shown relate to total meat production, from both commercial and farm slaughter. Data are given in terms of dressed carcass weight, excluding offal and slaughter fats.



Source: UN Food and Agricultural Organization (FAO)

OurWorldInData.org/meat-and-seafood-production-consumption/ • CC BY-SA



CHICKEN CONSUMPTION



Table 3.10: Food consumption of meat

	1964/66	1974/76	1984/86	1994/96	1997/99	2015	2030
kg per capita, carcass weight equivalent							
World	24.2	27.4	30.7	34.6	36.4	41.3	45.3
Developing countries	10.2	11.4	15.5	22.7	25.5	31.6	36.7
excl. China	11.0	12.1	14.5	17.5	18.2	22.7	28.0
excl. China and Brazil	10.1	11.0	13.1	14.9	15.5	19.8	25.1
Sub-Saharan Africa	9.9	9.6	10.2	9.3	9.4	10.9	13.4
Near East/North Africa	11.9	13.8	20.4	19.7	21.2	28.6	35.0
Latin America and the Caribbean	31.7	35.6	39.7	50.1	53.8	65.3	76.6
excl. Brazil	34.1	37.5	39.6	42.4	45.4	56.4	67.7
South Asia	3.9	3.9	4.4	5.4	5.3	7.6	11.7
East Asia	8.7	10.0	16.9	31.7	37.7	50.0	58.5
excl. China	9.4	10.9	14.7	21.9	22.7	31.0	40.9
Industrial countries	61.5	73.5	80.7	86.2	88.2	95.7	100.1
Transition countries	42.5	60.0	65.8	50.5	46.2	53.8	60.7
<i>Memo item</i>							
World excl. China	28.5	32.6	34.3	34.1	34.2	36.9	40.3
World excl. China and transition countries	26.5	29.0	30.6	32.4	33.0	35.6	39.1
Meat consumption by type (kg per capita, carcass weight equivalent)							
World							
Bovine meat	10.0	11	10.5	9.8	9.8	10.1	10.6
Ovine and caprine meat	1.8	1.8	1.7	1.8	1.8	2.1	2.4
Pig meat	9.1	10.2	12.1	13.7	14.6	15.3	15.1
excl. China	6.7	8.0	11.3	16.0	16.3	20.0	23.7
Poultry meat	3.2	4.6	6.4	9.3	10.2	13.8	17.2
Developing countries							
Bovine meat	4.2	4.3	4.8	5.7	6.1	7.1	8.1
Ovine and caprine meat	1.2	1.1	1.3	1.6	1.7	2.0	2.4
Pig meat	3.6	4.1	6.4	9.6	10.8	12	12.2
excl. China	2.1	2.4	2.8	3.3	3.4	4.0	4.7
Poultry meat	1.2	1.8	2.9	5.8	6.9	10.5	14.0
excl. China and Brazil	1.2	1.9	3.2	4.8	5.2	8.1	11.6

<http://www.fao.org/docrep/005/Y4252E/y4252e05b.htm>



DIETARY PROTEIN

Role: growth, egg production, biological functions, ...

Quantity: ~ 21 % of diet

Quality: methionine and lysine



70-80% of the total costs

Live broilers: Average costs of production at various R\$/\\$ exchange rates, mid-year 2011 (\\$/kg)

	U.S.	Brazil					
	National	Paraná	Goiás	Paraná	Goiás	Paraná	Goiás
	(actual)	(actual)		(hypothetical)		(hypothetical)	
		R\$1.58/\\$1		R\$1.50/\\$1		R\$2.00/\\$1	
Chicks	0.11	0.19	0.26	0.20	0.27	0.15	0.20
Feed	0.69	0.67	0.70	0.70	0.73	0.53	0.55
Grower payment	0.12	0.13	0.14	0.13	0.15	0.10	0.11
All other	0.08	0.07	0.09	0.07	0.09	0.05	0.07
Total	1.01	1.05	1.19	1.10	1.25	0.83	0.94

Exchange rate based on the average of June and July 2011

Brazil - based on conventional grow-out houses

Brazilian grower's payment is constructed based on producer (grow-out farmer) total costs plus water

Insect as a source of protein ?



INSECTS AS A SOURCE OF PROTEIN



<https://www.standardmedia.co.ke/>

- Rapid reproductive rates :
 - 1000 eggs/female
 - egg-adult : 38 days
- Able to convert organic waste
- Self-harvest capability
- Absence of mouthparts in their adult form

Nutrient content of wild black soldier fly (BSF) larvae produced on vegetable and fruit wastes compared to that of fishmeal and soybean.

	Wild BSF larvae	Fishmeal	Soybean
Crude protein (%)	38.98	55.1	47.9
Crude fibre (% w/w)	12.36	0.013	3.6
Ash (% w/w)	14.61	18.2	6.40
Fat (% w/w)	32.62		1.80
Free fatty acids (mh NaOH/g)	42.075	18.7	29.2
Manganese (%)	0.56		
Copper (%)	0.006		
Sodium (%)	3.07		
Iron (%)	0.57	0.82	
Calcium (%)	0.10	1.586	0.35
Potassium (%)	2.27		0.56

<https://www.wageningenacademic.com/>



MICROBIOTA

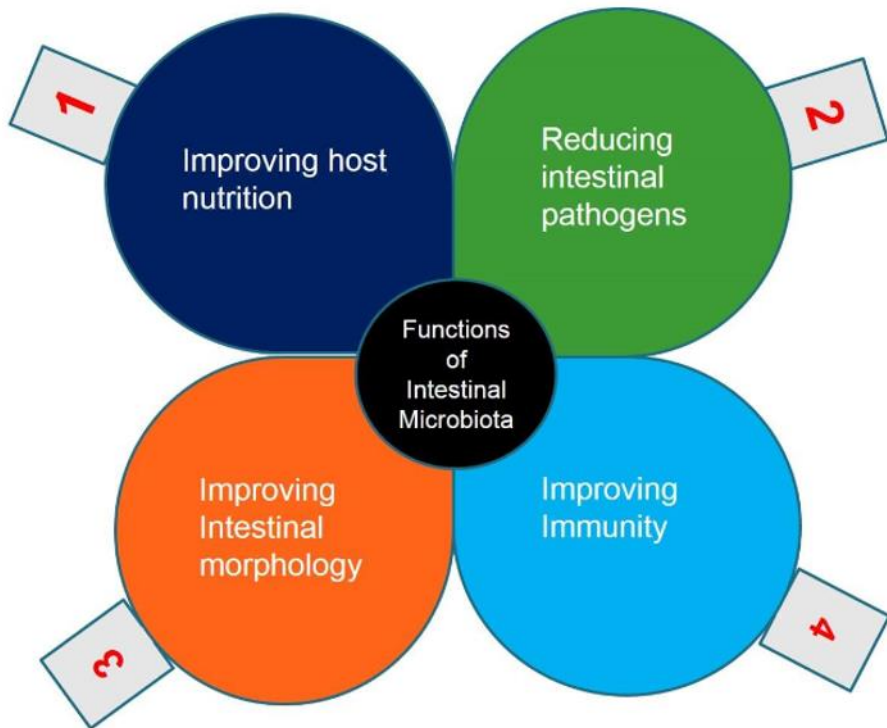
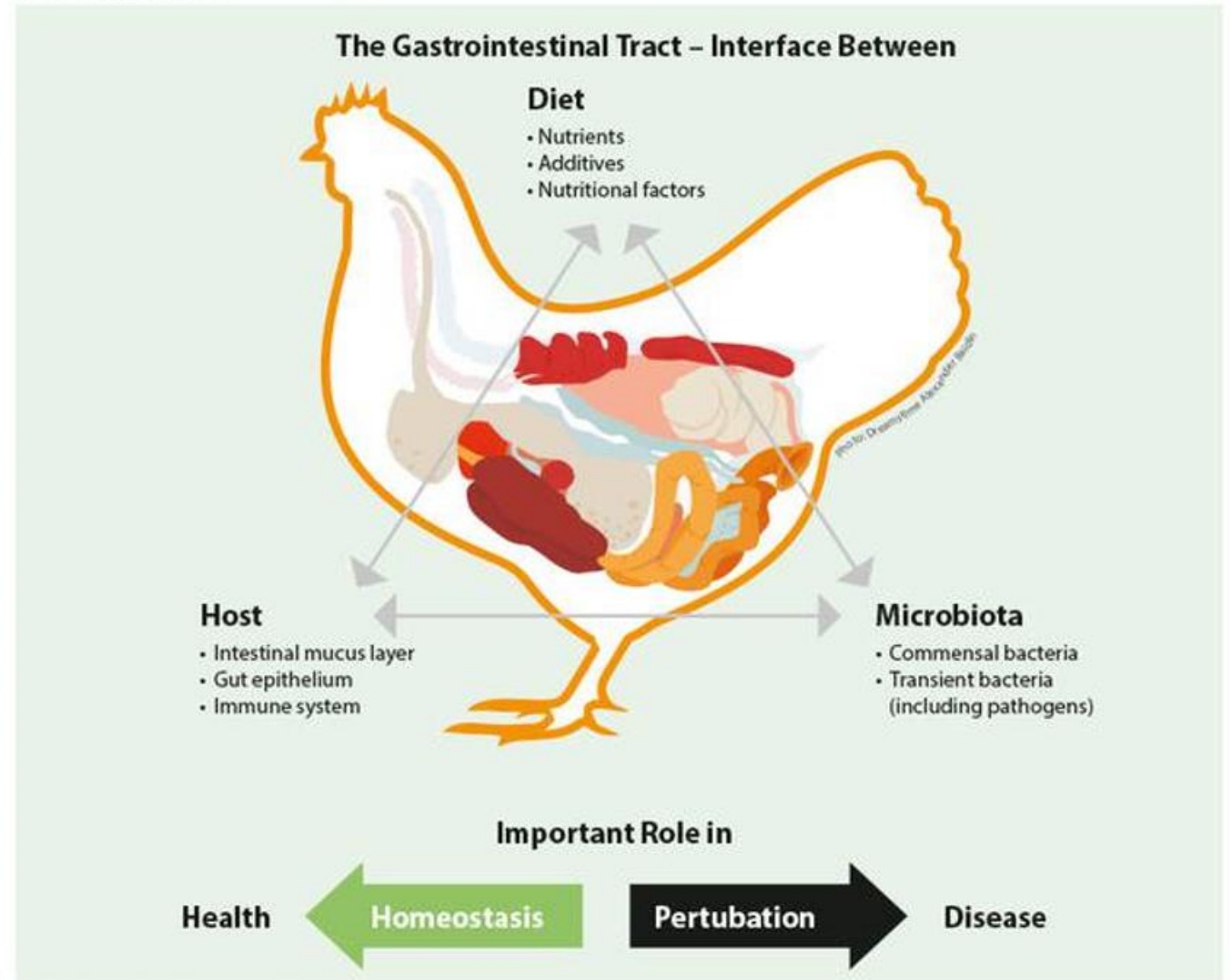


Figure 1. Interplay between diet, host, and gut microbiota - factors influencing the gut ecosystem and contributing to health and disease.

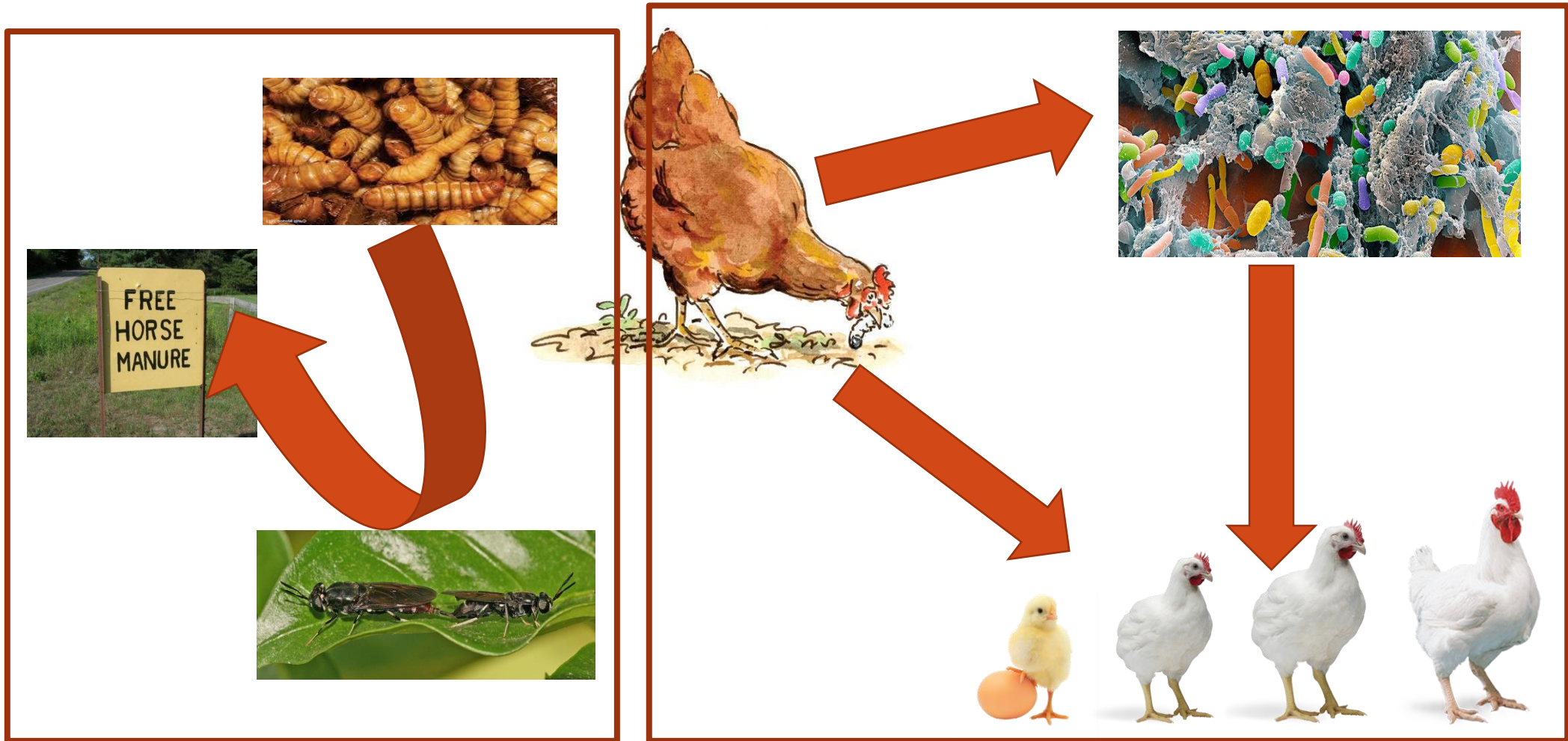


Source: Adapted from Conway, 1994

<http://www.primeanimalhealth.com/microbiota.html>



GOAL OF THE STUDY



Change in caecal microbiota of chicken fed BSF larvae raised on horse manure



I

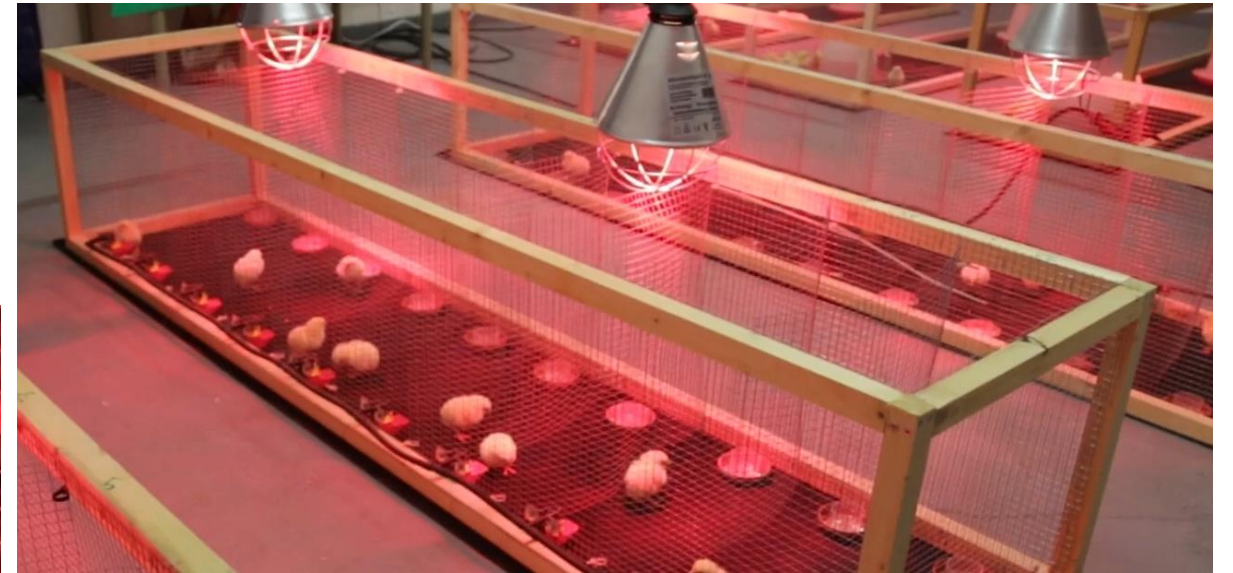


<http://www.fmv-multimedia.ulg.ac.be>



MATERIAL AND METHODS

40 Ross chicks: 0-13 days
2 groups; 2 replicates; 10 birds
Individual cages



MATERIAL AND METHODS



Iso-nitrogenous and iso-caloric	Control	Experimental
Dry matter (g/kg)	149	147
Crude protein (g/kg)	201	201
Crude fat (g/kg)	47	47
Nitrogen-free extract (g/kg)	509	507
Crude fiber (g/kg)	29	31
Ash (g/kg)	65	67
Starch (g/kg)	468	453
Gross energy (MJ/kg)	15.6	15.6

Feed supplied *ad libitum*
Diet as mash
Starter commercial feed
Iso-nitrogenous and iso-caloric



MATERIAL AND METHODS

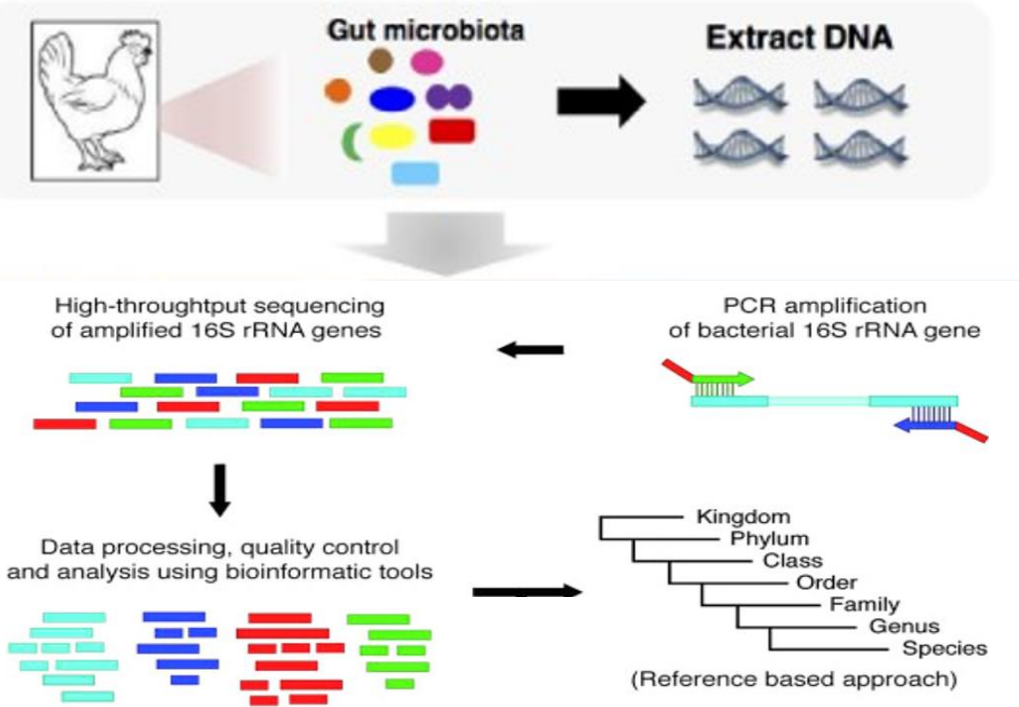


<http://www.fmv-multimedia.ulg.ac.be>



MATERIAL AND METHODS

16s ribosomal RNA sequencing techniques



MOTHUR

Reference alignment and taxonomical assignation



RESULTS (1)

17,616 OTU, 37 families and 5,275 species

Firmicutes: 90.83%

- *Ruminococcaceae* (33.01%)
- *Lachnospiraceae* (46.17%)

Bacteroidetes : 6.93%

- *Bacteroidaceae* (6.65%)

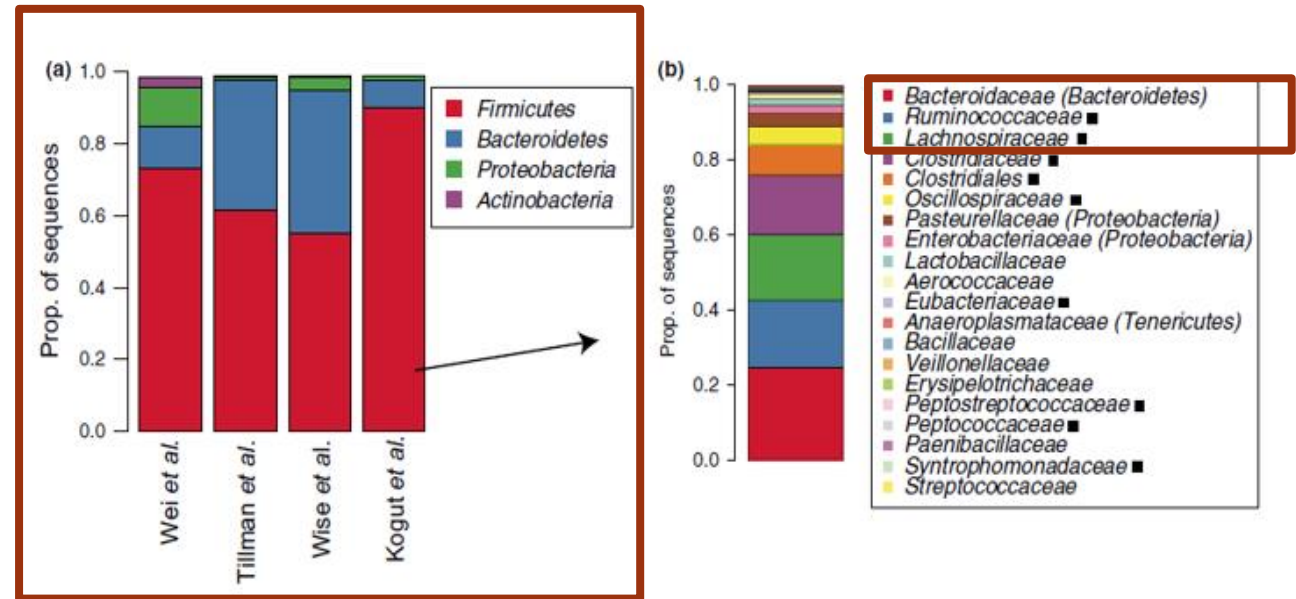


Fig. 1. Relative proportions of bacterial phyla (a) and families (b) found in chicken ceca. Data from Wei *et al.* (2013) represent publicly available sequences retrieved as described. Data from Tillman *et al.* (2011) and Wise & Siragusa (2007) are re-analyzed from data included in (Oakley *et al.*, 2013) representing 8 and 10 birds, respectively. Kogut *et al.* data are unpublished, collected, and analyzed as previously described (Oakley *et al.*, 2012 b, 2013) representing 20 birds and c. 20 000 sequencing reads. Data for each of these three flocks are from 3 weeks posthatch. Sequences from Wei *et al.* were additionally screened by removing sequences with ambiguous base calls, and all sequences were classified against a reference database of type strains from SILVA v115 (Pruesse *et al.*, 2007). Many of the sequences reviewed in (Wei *et al.*, 2013) do not contain metadata regarding bird age, which can have strong effects on community composition and structure. For (b) families belong to the phylum Firmicutes unless otherwise noted; families followed by black squares belong to the Clostridiales.



RESULTS (1)

Firmicutes: 90.83%

- *Ruminococcaceae* (33.01%)
- *Lachnospiraceae* (46.17%)

Bacteroidetes : 6.93%

- *Bacteroidaceae* (6.65%)

Degradation of cellulose and other highly recalcitrant polysaccharides

Degradation of less recalcitrant non-starch polysaccharides and starch

Butyric acid :

- source of energy
- anti-inflammatory properties

production of mucins and antimicrobial peptides

inhibition of the transcription factor NFκB activity

inhibitory activity against *Salmonella* and *Clostridium perfringens*



RESULTS (2)

Phylum	Control group	Experimental group
Actinobacteria	0.02 (0.00 - 0.04)	0.04 (0.00 - 0.06)
Bacteroidetes	7.61 (0.15 - 15.07)	5.70 (0.13 - 11.27)
Firmicutes	89.92 (84.83 - 95.00)	92.39 (86.91 - 97.86)
Proteobacteria	0.78 (0.69 - 0.87)	0.20 (0.17 - 0.24)
Other and unclassified bacteria	1.67 (0.01 - 2.75)	1.67 (0.01 - 2.75)

Means (and 95% confidence intervals) for the relative abundance (%) of bacterial phyla in the caeca of young Ross broilers fed (experimental diet) or not (control diet) larvae of black soldier flies.

Rhodobacteraceae: 0.31% vs 0.02%

Bacillaceae: 0.04% vs 0.00%



RESULTS (2)

Key genera	Species	SD (%)	ID (%)
<i>Bacteroides</i>	<i>Bacteroides coprocola</i>	2.97 ± 1.01	1.20 ± 0.25
	<i>Bacteroides heparinolyticus</i>	2.47 ± 0.85	1.38 ± 0.19
	<i>Bacteroides plebeius</i>	1.00 ± 0.24	3.00 ± 0.58*
	<i>Bacteroides salanitronis</i>	4.37 ± 0.58	0.45 ± 0.15**
<i>Parabacteroides</i>	<i>Parabacteroides merdae</i>	3.69 ± 0.80	0.82 ± 0.51*
<i>Elusimicrobiaceae</i>	<i>Elusimicrobium minutum</i>	0.23 ± 0.06	1.51 ± 0.52*
<i>Clostridiales</i>	<i>Alkaliphilus crotonatoxidans</i>	1.74 ± 0.34	3.04 ± 0.77
	<i>Alkaliphilus transvaalensis</i>	0.12 ± 0.04	1.25 ± 0.32**
	<i>Christensenella minuta</i>	0.66 ± 0.10	3.39 ± 0.35**
	<i>Vallitalea guaymasensis</i>	0.40 ± 0.06	1.03 ± 0.17*
<i>Oscillospira</i>	<i>Flavonifractor plautii</i>	1.36 ± 0.16	4.61 ± 0.71**
	<i>Intestinimonas butyriciproducens</i>	3.13 ± 0.29	4.61 ± 0.61
	<i>Oscillibacter valericigenes</i>	2.61 ± 0.41	2.70 ± 0.09
<i>Ruminococcus</i>	<i>Clostridium leptum</i>	0.53 ± 0.04	1.06 ± 0.20
<i>Veillonellaceae</i>	<i>Unclassified sp</i>	1.25 ± 0.15	0.41 ± 0.07**
<i>Fusobacterium</i>	<i>Unclassified sp</i>	6.02 ± 2.77	0.26 ± 0.08**
<i>Succinivibrionaceae</i>	<i>Succinatimonas hippei</i>	1.61 ± 0.21	0.25 ± 0.08**



SCIENTIFIC REPORTS

OPEN Insect-based diet, a promising nutritional source, modulates gut microbiota composition and SCFAs production in laying hens

14 June 2017
14 November 2017
24 November 2017

Luca Borrelli^{1,4}, Lorena Coretti^{2,3,5}, Ludovico Dipineto^{1,4}, Fulvia Bovera¹, Francesca Menna^{1,4}, Lorenzo Chiariotti^{2,4,6}, Antonio Nizza¹, Francesca Lembo^{5,6} & Alessandro Fioretti^{1,4}



RESULTS (2)

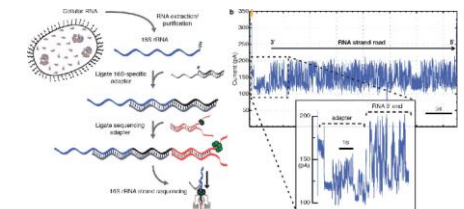
Table 8. Selected microbiota counts in the caecal digesta determined by DAPI staining and fluorescent *in situ* hybridization, log cell number · g⁻¹ of digesta (Experiment 3)

Indices	Treatments ¹				SEM ²	P-value
	NC	SL20	TM20	HI20		
total number of bacteria ³	9.77	9.73	9.56	9.84	0.04	<0.0001
<i>Bacteroides–Prevotella</i> cluster	8.61 ^c	8.81 ^b	8.60 ^c	9.05 ^a	0.09	<0.0001
<i>Clostridium leptum</i> subgroup	8.28	8.32	8.43	8.45	0.07	0.061
<i>Streptococcus</i> spp./ <i>Lactococcus</i> spp.	8.17 ^b	8.33 ^b	8.21 ^b	8.93 ^a	0.20	<0.0001
<i>Clostridium coccoides–Eubacterium rectale</i> cluster	8.22 ^c	8.39 ^{bc}	8.48 ^b	8.87 ^a	0.16	<0.0001
<i>Lactobacillus</i> spp./ <i>Enterococcus</i> spp.	8.27 ^b	8.54 ^{ab}	8.17 ^c	8.60 ^a	0.05	0.007



Table 3. Oligonucleotide probes used for intestinal microbiota analyses using fluorescent *in situ* hybridization (FISH) (Józefiak et al. (2016))

Target	Probe	Sequence (from 5' to 3')
<i>Bacteroides–Prevotella</i> cluster	Bacto303	CCAATGTGGGGACCTT
Enterobacteriaceae	Enter1432	CTTTTGAACCCACT
<i>Lactobacillus</i> spp./ <i>Enterococcus</i> spp.	Lab158	GGTATTAGCAYCTGTTTCCA
<i>Clostridium coccoides–Eubacterium rectale</i> cluster	Erec482	GCTTCTTAGTCARGTACCG
<i>Clostridium leptum</i> subgroup	Clept1240	GTTTTRTCAACGGCAGTC
<i>Streptococcus</i> spp./ <i>Lactococcus</i> spp.	Strc493	GTTAGCCGTCCCTTTCTGG



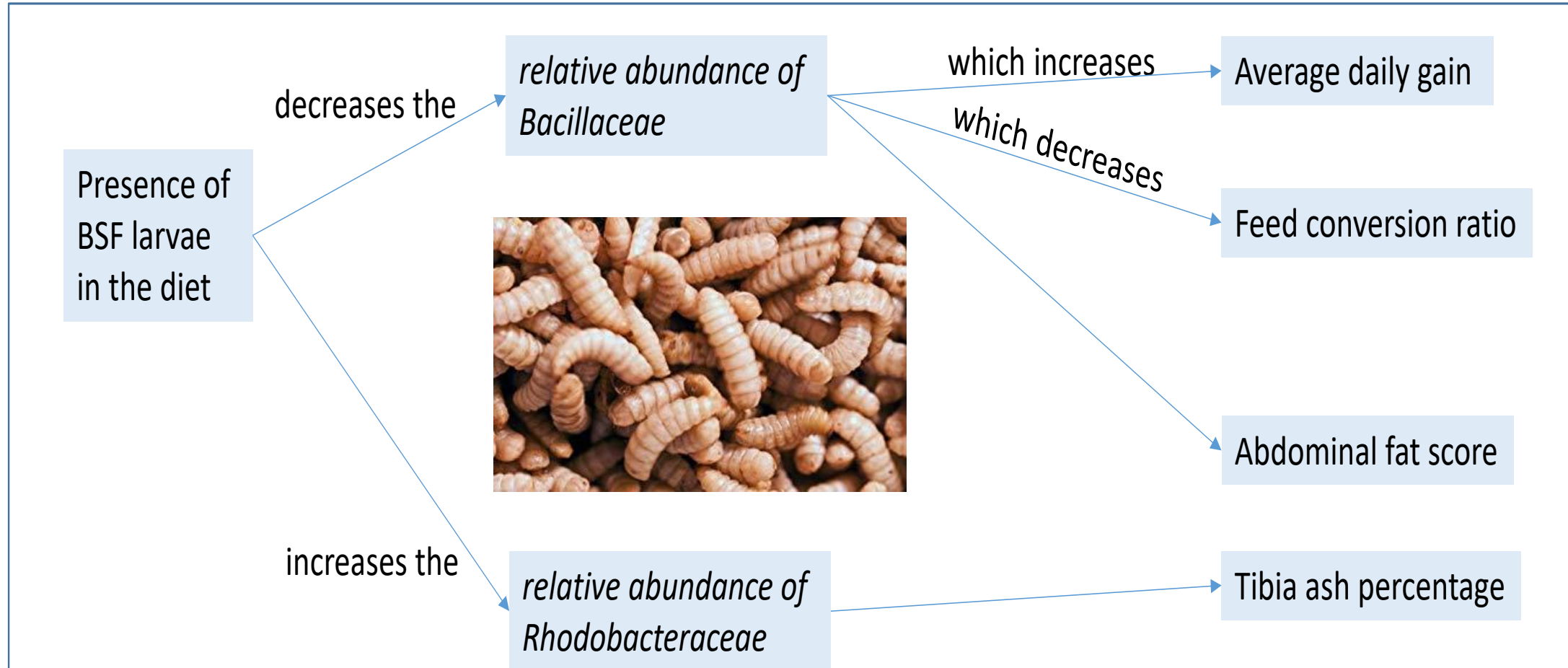
Journal of Animal and Feed Sciences, 27, 2018, 131–139 <https://doi.org/10.22358/jafs/91967/2018>
The Kielanowski Institute of Animal Physiology and Nutrition, Polish Academy of Sciences, Jabłonna

Full-fat insect meals as feed additive – the effect on broiler chicken growth performance and gastrointestinal tract microbiota

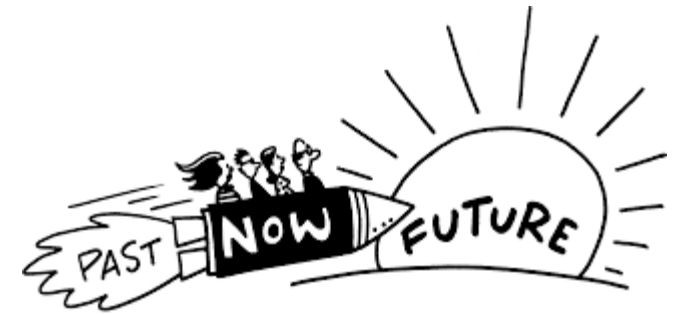
A. Józefiak^{1,6}, B. Kierończyk², M. Rawski^{2,3}, J. Mazurkiewicz^{2,4}, A. Benzertih⁴, P. Gobbi⁴, S. Nogales-Mérida⁴, S. Świątkiewicz² and D. Józefiak²



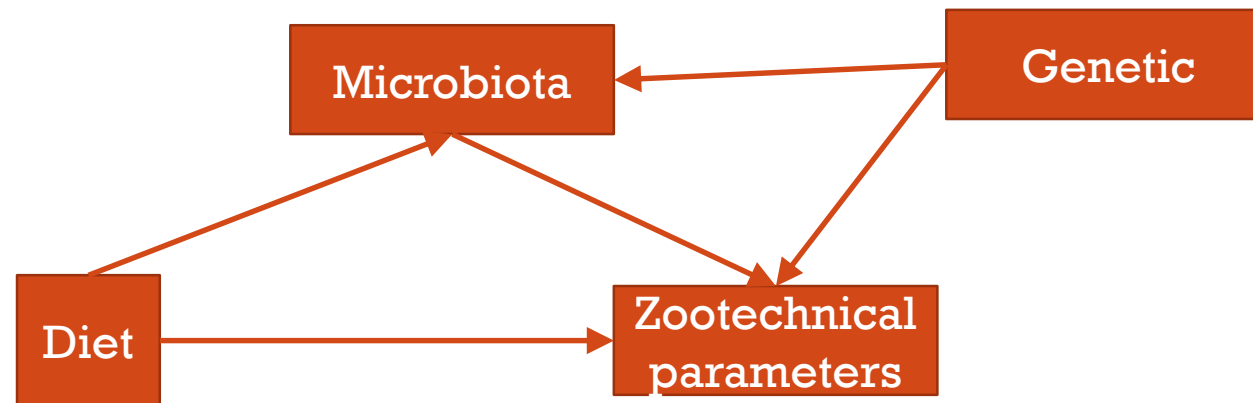
RESULTS (3)



NEXT ?



- Preliminary findings need to be confirmed
 - Higher number of animals
 - Different percentages of BSF in the feed
 - Improved statistical methods
 - Design : non-genetic effects (gender, diet, replication,)
 - Mixed structural equation model



NEXT ?

- Preliminary findings need to be confirmed
 - Higher number of animals
 - Percentage of BSF in the feed
 - Statistical methods (SEM)
- Niger
 - BSF and/or other insects
 - Imported chicken “without head” vs local chicken “bicycle”

