



## Nutritional approach and microbiome in poultry

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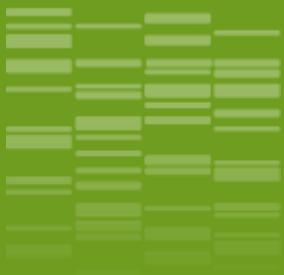
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# Nutritional approach and microbiome in poultry

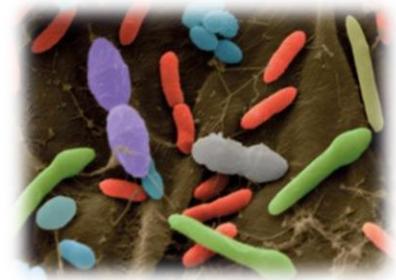


**Irène GABRIEL**

Nutritional Dynamics

UR83 Recherches Avicoles  
INRA Centre Val de Loire  
Nouzilly



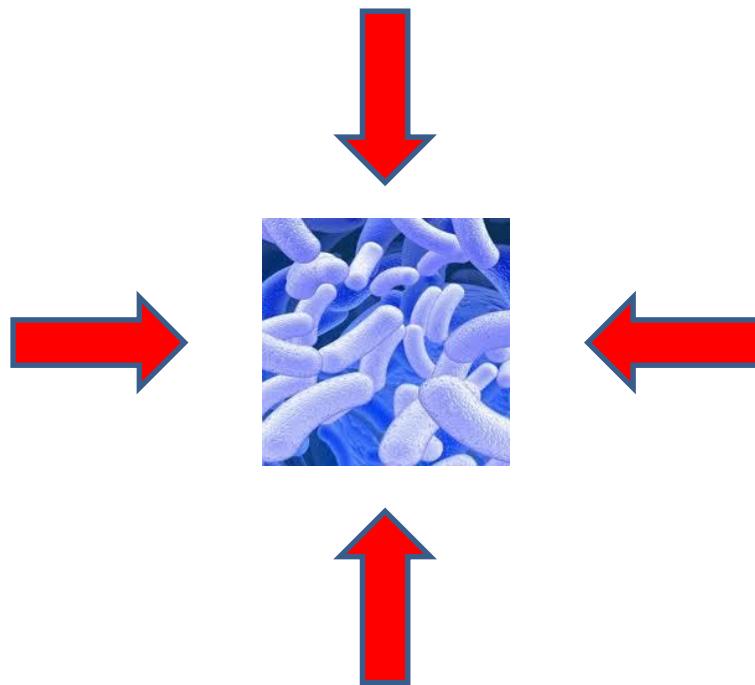


## 1. Factors regulating microbiota

## 2. Dietary factors regulating digestive microbiota



# 1. Factors regulating microbiota



# Factors regulating microbiota (1)

- Environment of first days : first contact with bacteria

Hatcher, eggshell

Human handling

Boxes for transportation

First feed and water

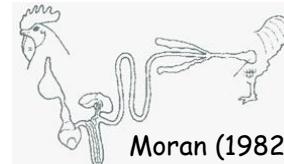
Litter



- Age :

Quantitative increase

Increase in complexity



- Anticoccidial drugs

- Feed additives

- Feed compounds

- Diet structure / Technological treatments

- Water



Gabriel et al 2006, 2014; Oakley et al 2014; Stanley et al 2014; Shaufi et al 2015

# Factors regulating microbiota (2)

- Nervous system



- Stress

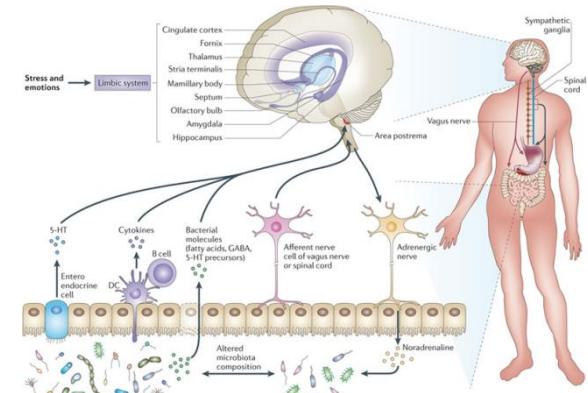
- Rearing environment

- Host genetics



- Maternal nutrition

## Interactions between Microbiota Intestine Brain



Collins et al., 2012

Nature Reviews | Microbiology

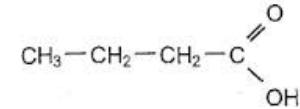
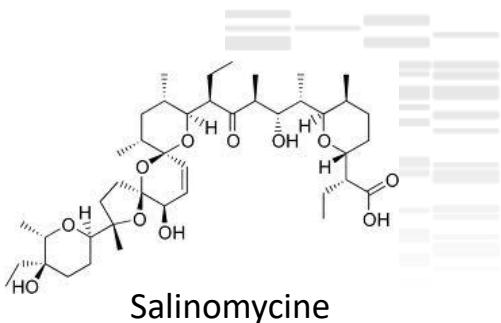


Torok et al. 2011; Guardia et al., 2011; Gabriel et al. 2014; Stanley et al. 2014; Mignon-Grasteau et al., 2015



## 2. Dietary factors regulating digestive microbiota





- 2.1. Anticoccidial drugs
- 2.2. Feed additives
- 2.3. Feed compounds
- 2.4. Diet structure / Technological treatments



# Feed additives

## Additives studied\* in poultry

\* With publications in Web of Science and PubMed

Organic acids

Carbohydrates (Prebiotic, 'Fibre', MOS)

Probiotic and symbiotic

Enzymes : Digestives (Polysaccharidase, Phytase, Protease)

Plant / Plant extracts

Amino acids

Vitamins

Clay

Charcoal

Minerals, trace elements

Bacteriocin

Bacteriophage and their enzymes (Endolysine, ...)

Antimicrobial peptides

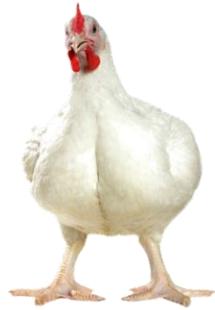
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Seal et al, 2013; Cheng et al, 2014

# Effects of additives reported in literature

Alone or in combination (synergy)

## Results



### Growth performances

Feed intake

Growth

Feed conversion

Mortality

Gut health (whose microbiota)

Immunity

Product quality

### Variable responses

Beneficial effects

No effect

Harmful effects

Effects + on microbiota  
No effect on animal

### According to conditions

Product      Composition / Dose / Mode of application

Animal      Genetic, sexe / Age / Physiological state

Rearing conditions

Diet composition



## Diet composition



→ Indigestible compounds



→ Substrates for the growth of microbiota

Different bacteria have different substrate preferences

→ Diet has a **great potential impact** on the digestive microbiota



# Nutrient composition

Carbohydrates

Proteins

Lipids

Minerals



# Carbohydrates

Water soluble non-starch polysaccharides



Rye / corn

↑ Anaerobe counts

Wagner et Thomas (1978)

Wheat , barley / corn

↑ Bacterial counts

Mathlouthi et al (2002)

Wheat , barley, rye / corn

→ *C. perfringens*

Annett et al (2002), Jia et al (2009)

Barley, oat, wheat / commercial

Change in dominant microbiota (Seq 16S) Ludvigsen et al (2016)

Rk : Environment modify non-dominant microbiota

High fiber diet

DDGS\*, wheat bran / Corn, soybean

\*Dry distillers grains with solubles

Changes in microbiota (Seq 16S) Walugembe et al (2015)

# Protein source and level (1)

Fishmeal / Soybean meal (protein level : 40%)

→ *C. perfringens* Drew et al (2004)



Fishmeal (protein level 33%) / Control diet (protein level : 21.5%)

Changes in microbiota (Seq 16S) Wu et al (2014), Stanley et al (2014)

Fishmeal / Soybean meal (protein level : 23%)

→ *C. perfringens* Drew et al (2004)

Protein level : 40% vs 23% (Fishmeal or Soybean meal)

→ *C. perfringens* Drew et al (2004)

# Protein source and level (2)

## Soybean meal



High concentrations of raffinose family oligosaccharides

→ May stimulate gut fermentation      Zdunczyk et al (2015)

## Rapeseed meal / Soybean meal

Change in SCFA and biogenic amines

Qaisrani et al (2014)



## Sunflower meal

↓ Caecal SCFA (Turkey)

Juskiewicz et al (2010)



# Protein source and level (3)

## Lupin seed meal

Change in microbiota counts

Rubio et al (1998)



## Pea seed meal

Pea (15%) / Control (soybean)

Change in microbiota fingerprint (T-RFLP)

↑SCFA content

Pea level (15 to 30%)

↓ Butyrate in caeca

Konieczka et al (2014)



## Fermented cottonseed meal / Soybean meal

Changes in microbiota counts / Fingerprint

Sun et al (2013)



# Lipids

Dietary fat source : animal / vegetal



Beef tallow / Soybean oil

Changes in microbiota counts

Danicke et al, 1999

Lard and tallow / soybean oil

→ ↑ *C. perfringens*

Knarreborg et al (2002)

Palm kernel fatty acid distillers or lard / Soybean oil

↑ Total organic acid

Jozefiak et al (2014)

Medium chain fatty acids / Soybean oil, animal fat

Changes in microbiota (Microarray) Van der Hoeven-Hangoor et al (2013)

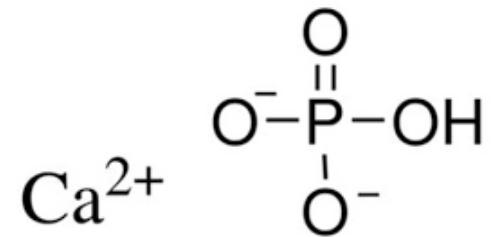
W-3 fatty acids (alpha-linolenic acid) (Flaxseed oil)

Changes in microbiota (Seq 16S) Lee et al (2016)



# Minerals

Ca and digestible P levels



Changes in microbiota counts / SCFA

Ptak et al (2015)

# Diet structure : Particule size

## Whole wheat / ground wheat

↓ *Salmonella Typhimurium*

Bjerrum et al (2005)

↓ *Clostridium perfringens*

Engberg et al (2004)



## Change in microbiota counts

Gabriel et al (2003), Engberg et al (2004), Bjerrum et al (2005), Gabriel et al (2008)

## Coarse grinding / Fine grinding

↓ Branched chain fatty acids

Qaisrani et al (2014)

Change in microbiota counts

Singh et al (2014)

# Technological treatments

## Thermal treatments

Change in bacterial counts and SCFA

Borojeni et al (2014)

## Pelleting

↑ SCFA

Change in microbiota counts

Engberg et al (2002)



↑SCFA

↑Salmonella Typhimurium

Huang et al (2006)

## Extrusion

Change in SCFA

Konieczka et al (2014)

# Regulation of digestive microbiota by diet

## Nutrient composition

Carbohydrates

Proteins

Lipids

Minerals

## Diet structure : Particule size

Whole / ground grains

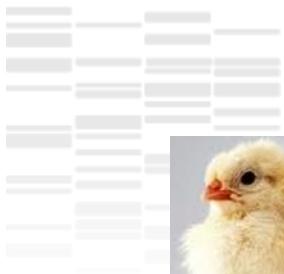
Coarse / Fine grinding

## Technological treatments

Thermal treatments

Pelleting

Extrusion



# Thanks for your attention

