

and may explain the relative importance of *Clostridium*. In the last part of the digestive tract, the caeca, main components may be composed of dietary molecules not hydrolysable by enzymes of the host, as well as endogenous components difficult to hydrolyze. These substrates may be responsible for the development of caecal bacteria, rich in *Clostridium* that may product SCFA that may increase energy extracted from the diet for the host. In summary, in D+ birds, the digestion and maintenance metabolic cost of the digestive tract may be optimized along the different parts of the digestive tract, with action of the host physiology and enzymes in the upper and middle part, and digestive microbiota in the lower part, leading to minimal wastes.

On the contrary in the D- birds, the under-development of their gizzard, due to failure in the gizzard relaxation process, leads to a lower retention time and a higher pH in the proventriculus-gizzard complex, and seems to be responsible for a part of their lower digestive functionality. It may also be responsible for a different selection of digestive microbiota. Indeed, this smaller proventriculus-gizzard complex development may lead to lower efficiency of gizzard bacterial filter, and thus a greater bacterial load passing in the small intestine. Moreover a higher amount of undigested compounds, probably rich in starch gives a high amount of substrate for bacterial growth, and favor *Lactobacillus* development. The low digestibility in the upper part of the digestive tract may be responsible for the more developed intestine, passing through an increase of epithelium area to try to cope with high undigested compounds, and increase muscle layer to move the higher amount of digestive content. The high bacterial load may be responsible for the higher stimulation of innate digestive immune system by increasing the number of goblet cells, suggesting increased mucus production. These changes in microbiota may be responsible for the lower lipid digestibility, and thus AMEn, due in part to bile acid deconjugation. These modifications in the upper and middle part of the digestive tract may lead to modification also in the lower part, with lower fermentative activity in the caeca with different bacterial composition probably due to a change in available substrates, and this may have an effect on energy recovery in this organ. To cope with this low digestive efficiency, D- birds increase their feed intake, contributing to increased amount of digestive content and with consequences for bacterial growth.

Moreover, contrary to D- birds, D+ birds appeared to be more able to adapt to a diet of low quality such as a diet rich on non-starch polysaccharides of wheat as shown by their lower decrease in AMEn than D- birds fed with wheat of high viscosity. They may have developed a mechanism to be less sensitive to deleterious effects of these hydrosoluble polysaccharides on intestinal viscosity.

However further knowledge in physiology of D+ and D- lines is needed to test these mechanistic biological hypotheses. For example, more information is needed on the motricity and regulation of the gastric emptying as this organ

seems to explain an important part of the difference between lines. Moreover, more knowledge is required at the intestinal level. Differences in anatomy of the small intestine have been described, but its digestive functionality is not known, as well as its consequences on digestive efficiency at the end of the small intestine. This knowledge would allow to determine the physiological limiting factors of digestion in order to suggest means to alleviate them. For genetic selection, it may allow to determine how this selection impacts on the phenotype and anticipate any undesirable effects before its introduction in selection schemes. Moreover, it may help to propose a list of candidate genes.

It is also important to know if D+ birds that have been selected by feeding wheat of high viscosity value, are able to adapt to other feedstuffs with other limiting factors in their digestibility. It will also be important to describe these birds in an environment closer to the production environment (on litter with animal groups, not in individual cage), as it would probably affect performances but also microbiota development, among others due to possibility of coprophagy, and then relationships between host, microbiota and digestive performances.

Regarding microbiota, it is important to know when the differences in microbiota composition between D+ and D- lines appear during the development of the birds. Moreover, a more extensive characterization is needed, as minor groups may also largely affect physiology of the host. This characterization may concern as well as the small intestine and the caeca, but also the crop. In addition, the functionality of the microbiota has to be studied, especially regarding to its consequences on the host digestive tract functionality, but also development of digestive immune function that is implied in digestive disease resistance, and consequently on global immune system.

Although relationships have been observed between bacterial groups and AMEn, the results were obtained in specific conditions (wheat diet, cage rearing). Moreover these links do not implied a causality link between these bacterial groups and phenotype such as AMEn, the phenotype being a balance between genotype, environment as diet and rearing conditions, and functionality of the digestive microbiota. Further works are thus needed to determine the involvement of digestive microbiota in digestive efficiency, measured by AMEn.

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