

Chronic stress causes genetic changes in chickens

10 November 2020



Credit: CC0 Public Domain

How can stress in animals be measured? Scientists from Uppsala University and elsewhere have now found that what are known as epigenetic biomarkers could be used to detect long-term exposure to stress in commercially raised chickens. This may, in time, lead to improved conditions in animal rearing. The study has been published in the journal *Frontiers in Genetics*.

Subjected to [chronic stress](#), animals show deterioration in their general state of health and a weakened immune system, which is unfortunate in terms of animal protection. For commercial animal production, it means that [animal products](#) are of a [lower quality](#) and a larger quantity of meat has to be discarded. These repercussions, in turn, adversely affect farmers' finances and consumers' food quality. Nonetheless, there are currently no reliable ways of measuring [long-term stress](#) in animals.

Researchers from Sweden and Brazil have now, in chicken studies, looked for signs of how chronic stress can affect the genes of red blood cells, causing epigenetic changes. In brief, this means

that specific molecule types (methyl groups) attach themselves to different parts of the DNA strand (methylation), depending on how the animal has lived. This may exert long-term effects on gene expression. Genes can, for example, be turned on or off (activated or deactivated).

The chickens studied, males of the popular White Leghorn breed of laying birds, were divided into two groups. One group was raised in a normal commercial environment, housed with other chickens and with good access to food and water. In the other group, the birds were exposed to factors known to induce stress. They were periodically isolated from one another, also with limited access to food and water. The same experiment was performed in both Sweden and Brazil.

"We took [blood samples](#) from the chickens in both the [control group](#) and the stress group after the stress treatment ended. We analyzed the methylation of red blood cells and compared methylation patterns in the two groups," says Fábio Pértile of the University of São Paulo, the first author of the study.

What they then saw was that in the stressed birds, the way in which the [methyl groups](#) had bound to the DNA of the red blood cells was entirely different from how this occurred in the control chickens. Although the scientists were unable to see how long these changes persisted, they were nonetheless an indication that the chickens had been exposed to prolonged stress.

"It's early days, but the results from this study are a step towards being able to identify specific epigenetic biomarkers that are evidence of the stress imposed on commercially raised animals in their living environment. It would be highly useful to have a [diagnostic tool](#) for tracking recurrent stress in production animals. And that could bring about, for example, improved health and protection for

farmed animals; meat and dairy products of higher quality; and reduced use of antibiotics," says Carlos Guerrero-Bosagna, a researcher in environmental toxicology at Uppsala University.

More information: Fábio Pértille et al. Putative Epigenetic Biomarkers of Stress in Red Blood Cells of Chickens Reared Across Different Biomes, *Frontiers in Genetics* (2020). [DOI: 10.3389/fgene.2020.508809](https://doi.org/10.3389/fgene.2020.508809)

Provided by Uppsala University

APA citation: Chronic stress causes genetic changes in chickens (2020, November 10) retrieved 24 February 2021 from <https://phys.org/news/2020-11-chronic-stress-genetic-chickens.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.