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Genetics and Breeding in Sheep and Goats

Guest Editor:

Dr. Shaobin Li

College of Animal Science and Technology/Gansu Key Laboratory of Herbivorous Animal Biotechnology, Gansu Agricultural University, Lanzhou 730070, China

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Message from the Guest Editor

Dear Colleagues,

Domestic sheep and goats are valuable farm animals that provide us with meat, milk, and textile fiber. Most of these traits are quantitative and controlled by multiple genes and environments. Elucidating the genetic and epigenetic mechanisms of these economic traits is critical to understanding how a trait comes into being. As for epigenetics, DNA methylation, histone modification, chromatin remodeling, and non-coding RNA are key factors involved in gene expression regulation. The objective of this issue is to explore genetic and epigenetic factors affecting sheep and goat phenotypes. We focus on not only the economic traits of sheep and goats, but also their resistance and adaptability. We welcome original research articles, brief research reports, and reviews covering (but not limited to) the following topics: 1) Functional gene variation affecting phenotypes; 2) Identification of gene function; 3) Gene regulatory network; 4) Gene expression regulation; 5) Epigenetic regulation (DNA methylation, histone modification, chromatin remodeling and non-coding RNA); 6)Genetic factors affecting the resistance and adaptation of sheep and goats.













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Editor-in-Chief

Prof. Dr. Selvarangan Ponnazhagan

Department of Pathology, The University of Alabama at Birmingham, 1825 University Blvd, SHEL 814, Birmingham, AL 35294-2182, USA

Message from the Editor-in-Chief

Genes are central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fastmoving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised.

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