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# **Breeding of Livestock in the Age of Genomics**

Guest Editor:

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#### Message from the Guest Editor

With the development of SNP-chips (a type of DNA microarray which is used to detect polymorphisms) over 10 years ago, breeding of livestock has undergone a revolution. In the past, genetic evaluations of animals were based only on phenotypes for the traits of interest and relationships among animals. Today, it is possible to derive accurate genetic values for individuals based on genetic markers, even though the trait has not been expressed in the individual, and information on relatives is limited. This Special Issue will consider how genomics has changed animal breeding and attempt to predict future trends.









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1. Institute of Veterinary Medicine and Animal Sciences, Estonian University of Life Sciences, Kreutzwaldi 1, 51014 Tartu, Estonia 2. Curtin University Sustainability Policy (CUSP) Institute, Kent St., Bentley 6102, Australia Animals is an on-line open access journal that was first published in 2011. Animals adheres to rigorous peerreview and editorial processes and publishes only high quality manuscripts that address important issues in the many varied disciplines that involve animals, with a focus on animal science, animal welfare and animal ethics. Animals is covered in the Science Citation Index Expanded (SCIE) in Web of Science, with the latest Impact Factor: 2.7 (2023, ranks 10/80 (Q1) in 'Agriculture, Dairy & Animal Science'; 16/167 (Q1) in 'Veterinary Sciences'), 5-Year Impact Factor: 3.0.

**Message from the Editor-in-Chief** 

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