

Special Issue

Genetic Tools and Techniques in Forensic Science—an In-Depth Look at the Process of Quantification of Forensic Samples

Message from the Guest Editor

The quantification of human DNA extracts from forensic samples plays a key role in the forensic genetics process. In real casework, a trace may not always contain human biological material, rather the biological traces at crime scenes may not be of human origin. Furthermore, the amount of DNA in a sample can vary between tens or hundreds of nanograms. Quantification guarantees maximum efficiency and avoids repeated analyses, overamplified samples or unnecessary examinations. This Special Issue is dedicated to the classic and latest quantitative techniques, studying the status, role, and future of quantification in forensic genetics. We also look forward to receiving other related papers.

Guest Editor

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Deadline for manuscript submissions

closed (15 February 2025)

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Message from the Editor-in-Chief

Genes is central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fast-moving 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. Why not consider *Genes* for your next genetics paper?

Editor-in-Chief

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