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Identification of Human Remains for Forensic and Humanitarian Purposes: From Molecular to Physical Methods

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Message from the Guest Editors

Human remains identification is an important part of a complex multidisciplinary process and a crucial step for contributing to the proper and healthy functioning of a civil society and of justice. Identifying a person in forensic and humanitarian contexts does not just represent a legal necessity but also a fundamental right. The identification process implicates the comparison of information provided by someone who knows the person (antemortem data) with the scientific information obtained by forensic experts during the examination of human remains (postmortem data). Over the past twenty years, we have seen the development of new technologies in diverse scientific fields that have allowed us to obtain results that until recently were not technically feasible and affordable.

This Special Issue is aimed at investigating recent advances in human remains identification. It focuses on collecting reviews and original contributions considering the importance of a multidisciplinary approach to support identification phases of human remains as well as innovative methodological aspects, bioinformatics and statistical tools, and future prospects of forensic/humanitarian research.



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Special Issue



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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?

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