



Non-Coding RNA in Multiple Myeloma

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Message from the Guest Editors

Multiple Myeloma is one of the most common and aggressive hematologic tumors. Despite the great amount of data and with few exception, the Myeloma community is still far from consensus in defining driver events that might be target of personalized therapy.

Major evidences of correlation with coding transcriptome and genomic lesions have emerged involving the small non-coding RNA (ncRNA) fraction, primarily miRNA and snoRNA. The most recent researches have highlighted that also lncRNAs expression levels are modulated in myeloma and their deregulation is candidate for a role in the disease.

This Special Issue will collect works on the latest advances in investigating non-coding RNA involved in myeloma. Topics of interests include:

- Discovery and investigation of novel ncRNAs involved in plasma cell biology
- ncRNAs deregulation in multiple myeloma and plasma cells dyscrasias
- Functional roles of ncRNAs in malignant plasma cells, tumor microenvironment, and serum
- ncRNA transcriptional network and epigenetic mechanisms in myeloma cells
- Pre-clinical studies investigating the role of ncRNA in myeloma

For further reading, please visit the [Special Issue website](https://www.mdpi.com/si/14536):





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Editor-in-Chief

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Message from the Editor-in-Chief

This field finally has a dedicated journal where its broad community can communicate and exchange its latest findings in one centralized place. This field was built stone by stone from the many scientific contributions from extremely diverse horizons, studying gene silencing in plants, position effect variegation in drosophila or quelling in fungi. This field has achieved maturity, but a lot remains to be discovered! Our aim is to publish manuscripts from all horizons that will have a high impact on the development of the field. Let's have fun and wish *Non-Coding RNA* a long and rewarding life!

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