

Special Issue

Osteosarcoma: Current Advances from Molecular and Cellular Mechanisms to Therapy

Message from the Guest Editors

Osteosarcoma is a rare aggressive primary bone cancer. The tumors are highly heterogeneous with complex genomic landscape involving numerous structural and copy number alterations. Alterations in *TP53* are the most frequently somatic changes, while pathogenic germline mutations in *RB1*, *TP53*, *RECQL4*, *BLM*, and *WRN* are associated with an increased risk of osteosarcoma. Tumor heterogeneity and a lack of recurrent driver mutations make it difficult to identify effective molecularly targeted therapies. Recent multi-omics studies have enhanced our understanding of the molecular pathways in osteosarcoma pathogenesis and are opening up new opportunities for biomarker-driven precision therapies based on molecular subtypes or altered genomic or cellular pathways such as PI3K-ATK-mTOR signaling, homologous recombination repair pathway, or therapies based on the immune profile/response of tumors. This Issue will compile recent research on various cellular and molecular processes involved in osteosarcoma growth, progression, and drug sensitivity/resistance, which could lead to the identification of new drug targets and improvements in osteosarcoma treatment.

Guest Editors

Dr. Virinder Kaur Sarhadi

Department of Oral and Maxillofacial Diseases, Helsinki University Hospital and University of Helsinki, 00290 Helsinki, Finland

Dr. Francesca Perut

Biomedical Science and Technologies and Nanobiotechnology Lab, IRCCS Istituto Ortopedico Rizzoli, Via di Barbiano 1/10, 40136 Bologna, Italy

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Editor-in-Chief

Prof. Dr. Maurizio Battino

Department of Odontostomatologic and Specialized Clinical Sciences,
Sez-Biochimica, Faculty of Medicine, Università Politecnica delle
Marche, Via Ranieri 65, 60100 Ancona, Italy

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