Special Issue

In Silico Approaches of Drug-Target Interactions

Message from the Guest Editor

In silico approaches are transforming modern pharmaceutics and biopharmaceutics by providing powerful tools for understanding drug-target interactions and optimizing therapeutics. Using computational techniques such as molecular modeling, docking simulations, molecular dynamics, and virtual screening, these methods predict binding affinities and selectivities between drug candidates and biological targets. Recent advances in machine learning and artificial intelligence have further refined these tools, improving drug design precision. Integrating ADME (Absorption, Distribution, Metabolism, and Excretion) predictions also ensures the selection of candidates with favorable pharmacokinetics.

This Special Issue aims to explore the latest advancements in these in silico methods, focusing on their role in enhancing therapeutic efficacy, safety, and efficiency in biomedical science.

We welcome original research and review articles addressing topics such as molecular modeling, machine learning applications, virtual screening, and the optimization of drug safety and efficacy. We look forward to your contributions to this dynamic field of computational pharmaceutics.

Guest Editor

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Editor-in-Chief

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