Message from the Guest Editor

Dear Colleagues,

Investigations concerning the protein folding led to development of compounds that were named “foldamers”. This definition deals with oligomers that display a secondary structure, mostly evidenced by spectroscopic methods, exploiting non-covalent interactions such as hydrogen bonding and aromatic stacking. Thus, a lot of different structures were obtained, although helices were most frequently observed. Moreover, it is worth mentioning that these compounds display interesting properties in disrupting protein–protein interactions, so that they are largely studied as drug candidates, being suitable for a novel approach to innovative therapeutics.

Prof. Dr. Mario Orena
Guest Editor

Author Benefits

- **Open Access**: free for readers, with publishing fees paid by authors or their institutions.
- **High visibility**: indexed by the Science Citation Index Expanded (Web of Science), MEDLINE (PubMed) and other databases.
- **Rapid publication**: manuscripts are peer-reviewed and a first decision provided to authors approximately 19 days after submission; acceptance to publication is undertaken in 6 days (median values for papers published in this journal in 2016).
- **Sections**: published in twelve topical sections.