



Molecular Recognition in Protein and Peptide Nanotechnology

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Message from the Guest Editors

During the last four decades, a growing body of evidence has been accumulating on the specific interaction and molecular recognition of the complementary peptides, i.e., sense and antisense peptides specified by the complementary DNA and RNA sequences. The interaction of sense and antisense peptides can be described with the set of rules combining physico-chemical properties of amino acids with bidirectional coding and stereochemical interaction. This natural genetic coding algorithm for complementary peptide interactions is neither expensive nor time-consuming, and enables simple and efficient design of interacting protein and peptide motifs. It represents a powerful new tool for the investigation and modeling of natural or de novo designed protein and peptide ligand-acceptor (receptor) systems.

Different applications include:

1. Investigation of ligand and receptor structure, function and evolution
2. Screening of protein and peptide ligands and acceptors (receptors)
3. Development of vaccines, diagnostic tests and biomaterials
4. Label-free technology to track drug candidates
5. Peptidomimetics
6. Investigation of peptide nucleic acids interaction





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Message from the Editor-in-Chief

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