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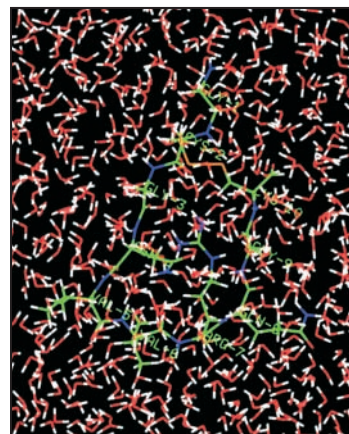
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About the Cover

Molecular dynamics simulation of TMTP1. TMTP1 peptide is solved in a water cubic periodic box. Ten amino acid residues form a cyclic peptide via a disulfide bond between the two cysteines that respectively lie at the head and the tail. The line view of TMTP1 peptide with side chains and the internal hydrogen bonds. For further details, please see Luo et al on page 5494 of this issue.



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