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ADDITION/CORRECTION

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Identification of RNA Pseudoknot-Binding Ligand That Inhibits the −1 Ribosomal Frameshifting of SARS-Coronavirus by Structure-Based Virtual Screening [*Journal of the American Chemical Society* **2011**, *133*, 10094–10100. DOI: 10.1021/ja1098325].

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Page 10096. Figure 1 should be corrected as shown below. In Figure 1a, there were base mispairings in the Stem 3 region. In Figure 1b, "Stem 2" was misspelled.

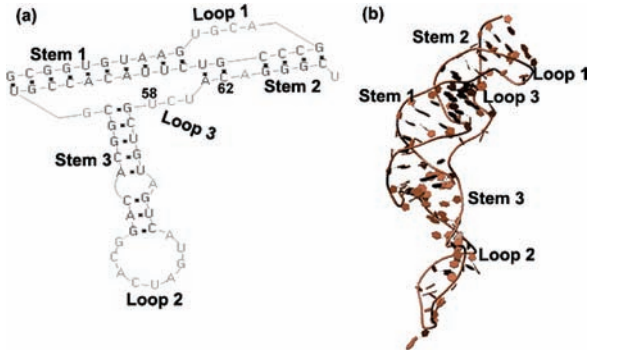

Figure 1. (a) Two-dimensional model of SARS-pseudoknot generated by the PSEUDOVIEWER[19] program. (b) Three-dimensional structural model of the SARS-pseudoknot used in this study. It was optimized by molecular dynamics simulation using the Amber 8.0 program. Brown ribbon renders the phosphate backbone of the RNA pseudoknot.

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