

## Correction to Unfolding the Conformational Behavior of Peptide Dendrimers: Insights from Molecular Dynamics Simulations

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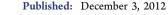
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**P** age 5047. Table 3: The entries presented in Table 3 are incorrect due to a systematic exclusion of some data points

## Table 3. Average Radius of Gyration of the Dendrimers

average $R_{\rm g}~({\rm nm})$
1.439
1.445
1.432
1.418
1.060

in the average radius of gyration calculations. A corrected table is shown. The conclusions of the original article are in no way affected by this correction.





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