

Correction to “An Investigation on the Fundamental Interaction Between Abeta Peptides and the AT-Rich DNA”

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Our original paper stated that the force field used for both protein and DNA is the following:

“All the structures in our simulation setup were represented by the AMBER99SB force field.⁴⁰”

It should have been the following:

“The structures of protein in our simulation setup were represented by the AMBER99SB force field, and DNA was represented by AMBER94.⁴⁰”