


CORRECTION

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Correction to: N⁶-Adenosine methylation on mRNA is recognized by YTH2 domain protein of human malaria parasite *Plasmodium falciparum*

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The original version of this article [1], unfortunately contained a mistake. The presentation of Fig. 2 has been published incorrectly. The correct Fig. 2 is provided.

The original article has been corrected.

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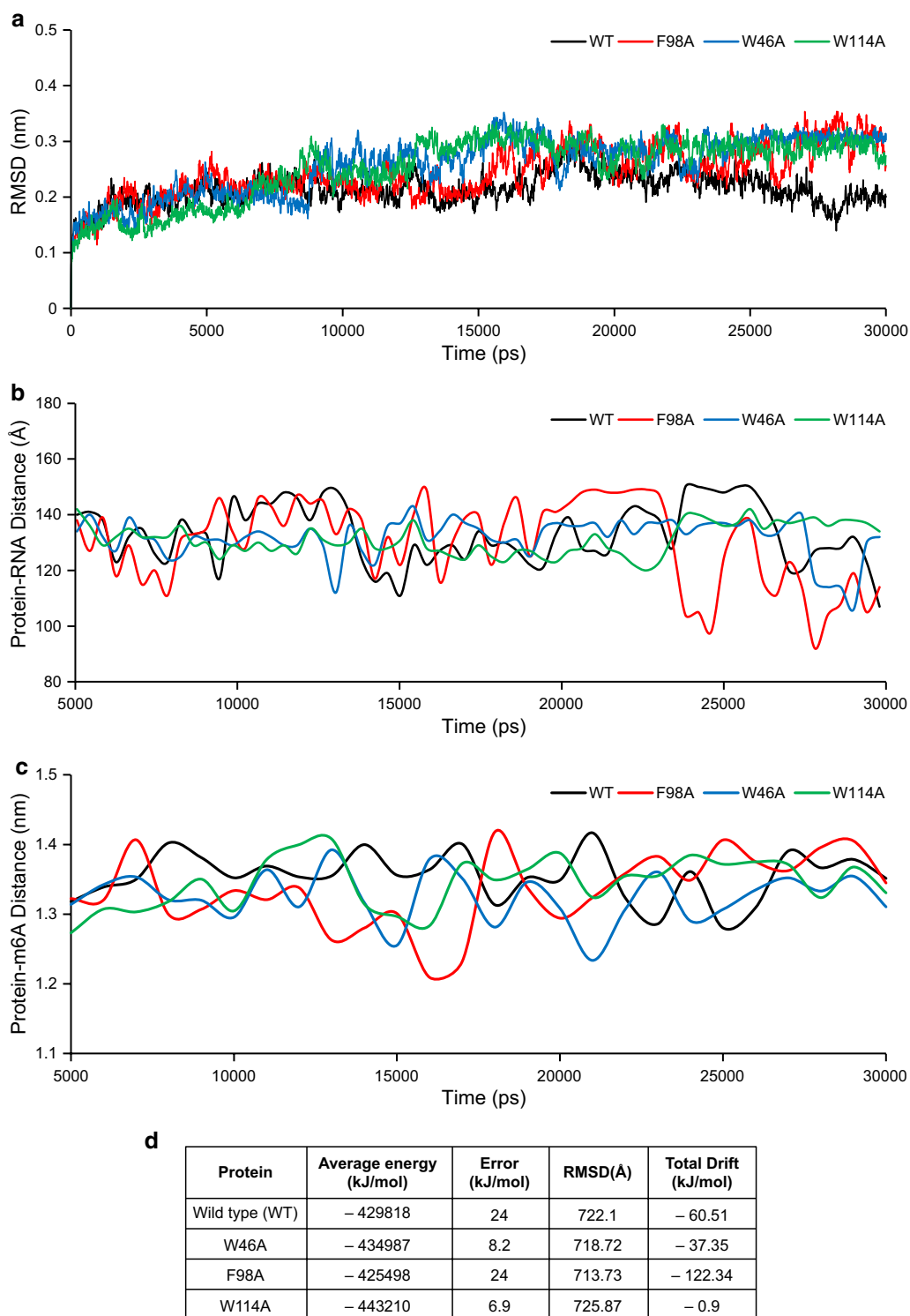


Fig. 2 Molecular dynamics simulation analysis of wild type and mutant PfYTH2 proteins in complex with RNA. **a** Backbone Root Mean Square Deviation (RMSD) profiles of PfYTH2 over molecular dynamics simulations time scale (picoseconds). **b** Spatial distance fluctuations (Angstroms), monitored over the MD simulations timescale (picoseconds) for Wild type and mutant PfYTH proteins in complex with RNA. **c** Distance drifts computed between PfYTH wild type and mutant proteins with m6A residue from RNA (nanometers), monitored over the MD simulations timescale (picoseconds). **d** The table presents the calculated PfYTH (wild type and mutants)-RNA complex potential energy values over the molecular dynamics simulation time scale

Reference

1. Govindaraju G, Kadumuri RV, Sethumadhavan DV, Jabeena CA, Chavali S, Rajavelu A. N6-Adenosine methylation on mRNA is recognized by YTH2 domain protein of human malaria parasite *Plasmodium falciparum*. *Epigenetics Chromatin*. 2020;13:33. <https://doi.org/10.1186/s13072-020-00355-7>.

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