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Author Correction: Artificial Intelligence and Machine learning based prediction of resistant and susceptible mutations in *Mycobacterium tuberculosis*

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In this Article, Figure 7 is a duplication of Figure 10. The correct Figure 7 appears below.
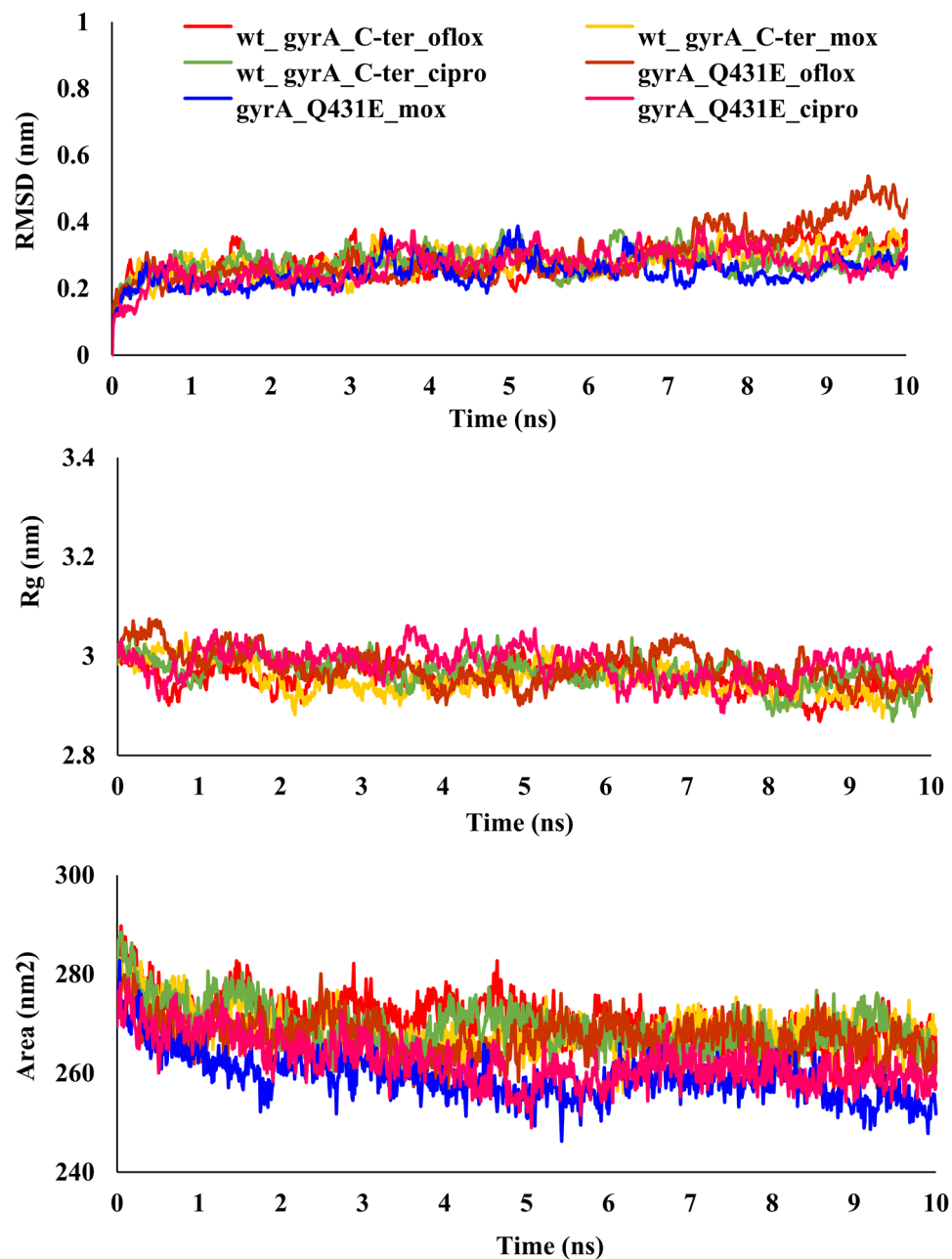
Figure 7. RMSD, Rg and SASA plot for *gyrA* gene, C-terminal protein. For Q431E mutant, the RMSD and Rg were slightly higher than wild type, however SASA was less for mutant protein.