# CORRECTION Open Access

# Correction to: In silico structural homology modelling of EST073 motif coding protein of tea *Camellia sinensis* (L)

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# Correction to: J Genet Eng Biotechnol 18, 32 (2020) https://doi.org/10.1186/s43141-020-00038-6

Following publication of the original article [1], the authors identified an error in the Acknowledgements section.

The updated Acknowledgements section is given below and the changes have been highlighted in **bold typeface**.

### Acknowledgements

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