

Peer Review Report

Review Report on Comparative Genomics of Thermosynechococcaceae and Thermostrictaceae: Insight into Codon Usage Bias

Original Research, Acta Biochim. Pol.

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EVALUATION

Q 1 Please summarize the main findings of the study.

This is an interesting paper with the analysis of the codon usage bias in the light of adaptation to specific conditions. A group of cyanobacteria has been used as a model. The conclusions are interesting and the analyses appear technically sound.

Q 2 Please highlight the limitations and strengths.

The strength of the study is an interesting problem and indication how different codon usage might be related to adaptations of microorganisms. The limitation is a relatively low number of investigated genomes. Nevertheless, if there are all available genomes in databases, this is still fine.

Q 3 Please comment on the methods, results and data interpretation. If there are any objective errors, or if the conclusions are not supported, you should detail your concerns.

The methods are properly chosen and the interpretation of the results are generally correct.

Check List

Q 4 Please provide your detailed review report to the editor and authors (including any comments on the Q4 Check List)

This is an interesting paper with the analysis of the codon usage bias in the light of adaptation to specific conditions. A group of cyanobacteria has been used as a model. The conclusions are interesting and the analyses appear technically sound. Just a few questions:

1. Did the author consider sizes of genomes as possible factors influencing the results of the analysis.
2. Accession numbers of analysed genomes in the database should be provided, not only a link to the general website of the database.

Q 5 Is the English language of sufficient quality?

Yes.

Q 6 Is the quality of the figures and tables satisfactory?

Yes.

Q 7 Does the reference list cover the relevant literature adequately and in an unbiased manner?

Yes.

Q 8 Are the statistical methods valid and correctly applied? (e.g. sample size, choice of test)

Yes.

Q 9 Are the methods sufficiently documented to allow replication studies?

Yes.

Q 10 Are the data underlying the study available in either the article, supplement, or deposited in a repository? (Sequence/expression data, protein/molecule characterizations, annotations, and taxonomy data are required to be deposited in public repositories prior to publication)

No.

Q 11 Does the study adhere to ethical standards including ethics committee approval and consent procedure?

Not Applicable.

Q 12 Have standard biosecurity and institutional safety procedures been adhered to?

Not Applicable.

QUALITY ASSESSMENT

Q 13 Originality

☒ ☒ ☒ ☐ ☐

Q 14 Rigor

☒ ☒ ☒ ☒ ☐

Q 15 Significance to the field

☒ ☒ ☒ ☒ ☐

Q 16 Interest to general audience

☒ ☒ ☒ ☒ ☐

Q 17 Quality of the writing

☒ ☒ ☒ ☒ ☐

Q 18 Overall quality of the study

☒ ☒ ☒ ☒ ☐