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Identification of potential antibiotic targets in the proteome of multi-drug resistant Proteus mirabilis

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Nosheen Ehsan, Sajjad Ahmad, Afifa Navid, Syed Sikander Azam

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Authors: Nosheen Ehsan^{1,a}, Sajjad Ahmad^{1,a}, Afifa Navid¹, Syed Sikander Azam^{1,⊠}

^aThese authors contributed equally to this work

Institutional addresses:

¹Computational Biology Lab, National Center for Bioinformatics, Quaid-i-Azam University, Islamabad, Pakistan.

Email addresses:

Nosheen Ehsan, Email: nosheenehsan23@gmail.com

Sajjad Ahmad, Email: sahmad@bs.qau.edu.pk

Afifa Navid, afifanavid.bi@gmail.com

Syed Sikander Azam, E-mail: syedazam2008@gmail.com, ssazam@qau.edu.pk

Corresponding Author:

[™]Syed Sikander Azam

E-mail: syedazam2008@gmail.com,_ssazam@qau.edu.pk

Phone: 0092-51-906 44130

Computational Biology Lab, National Center for Bioinformatics, Quaid-i-Azam University,

Islamabad 45320, Pakistan

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