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Authors: Reaz Uddin, Faiza Jamil

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Prioritization of Potential Drug Targets against *P. aeruginosa* by Core Proteomic Analysis using Computational Subtractive Genomics and Protein-Protein Interaction Network

Reaz Uddin^{1*} and Faiza Jamil¹

¹Dr. Panjwani Center for Molecular Medicine and Drug Research, International Center for Chemical and Biological Sciences, University of Karachi

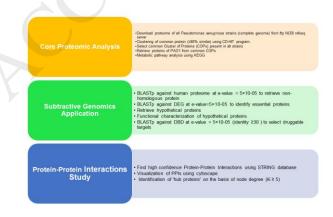
*Correspondence: Dr. Reaz Uddin, Lab 103 PCMD ext. Dr. Panjwani Center for Molecular Medicine and Drug Research, International Center for Chemical and Biological Sciences, University of Karachi, Karachi 75270, Pakistan

*Email: mriazuddin@iccs.edu

Tel: +92-21-34824930

Fax: +92-21-34819018-9

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