EXCELLENT PUBLISHERS

International Journal of Current Microbiology and Applied Sciences (IJCMAS)

ISSN: 2319-7692 (Print) ISSN 2319-7706 (Online)

An International, Monthly, Online, Free Access, Peer Reviewed, Indexed, Scientific Research Journal

CERTIFICATE OF PUBLICATION

This is to certify that the following article reviewed by editorial board and published in

International Journal of Current Microbiology and Applied Sciences (IJCMAS) ISSN: 2319-7692 (Print) ISSN 2319-7706 (Online).

Int.J.Curr.Microbiol.App.Sci.2017.6.(2):1894-1901

http://dx.doi.org/10.20546/ijcmas.2017.602.214

Evaluation of a New Phenotypic Method to Screen for OprD-Deï¬•cient Mutant Strains of Pseudomonas aeruginosa

Omnia M. Ahmed*, A.A Manal and A.G. Samia

Clinical Pathology Department, Faculty of Medicine, Ain shams University, Egypt

Dr. M.Prakash
Editor-in-chief
International Journal of Current Microbiology and Applied Sciences
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