

Nested Multiplex Endpoint Polymerase Chain Reaction: An Alternative Method for Human Papilloma Virus Genotyping in Resource-limited Settings

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In the published article, the authors want to change the sequence of the author byline.¹ This is detailed in the article available at: <https://www.jacmjournal.org/doi/JACM/pdf/10.5005/jacm-11020-0001>.

The authors apologize and deeply regret any inconvenience or challenges resulting from the publication and subsequent corrigendum of this article.

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1. SYMEC Research Group, Mukhopadhyay A, Mathai S, et al. Nested multiplex endpoint polymerase chain reaction: an alternative method for human papilloma virus genotyping in resource-limited settings. *J Acad Clin Microbiol* 2024;26(1):1–6. DOI: 10.5005/jacm-11020-0001