

BIO DATA

Name: Dr. Ajeet Kumar Mohanty, Ph.D
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Date of Birth: 25th Aug, 1982

Education (Post-Graduation onwards & Professional Career)

| Sl No. | Institution Place | Degree | Year | Field of Study |
|--------|--|--|------|------------------------|
| 1. | Utkal University, Bhubaneswar, Odisha | M.Sc. | 2005 | Biotechnology |
| 2. | Goa University, Goa | Ph.D. | 2016 | Mosquito Proteomics |
| 3. | Public Health Foundation of India (PHFI), Delhi | Post Graduate Diploma in Epidemiology (PGDE) | 2018 | Epidemiology |

Research specialization (Major scientific fields of interest)

Entomology, biological vector control agents, vector proteomics, genomics and vector-parasite interaction

Patent:

Patent filed: Temp/E-1/13888/2018-CHE- Novel peptides for developing anti-malarial vaccinesand diagnostic kits.

Awards:

1. “**DGAFMS & Senior Colonel Commandant Award**” for the best published article in the field of Community Medicine for the year 2017 by Armed Forces Medical College (AFMC), Pune, India.
2. Best “**Oral Presentation Award**” during XII Joint Annual Conference of Indian Society for Malaria and Other Communicable Diseases & Indian Association of Epidemiologists” ISMOCD 2017, held from 1st -3rd September, 2017 at Armed Forces Medical College (AFMC), Pune, India.
3. Best“**Poster Presentation Award**” at the National Symposium on “Microbial Diversity and its Applications in Health, Agriculture and Industry” organized by the ICAR Research Complex for Goa, Ela, Old Goa, Goa- from 4-5th March, 2011.

Publications: 25 Research articles, 2 review articles and 3 Book chapters

1. Pereira M, **Mohanty AK**, Garg S, Tyagi S and Kumar A. Characterization of midgut microbiome of *Anopheles stephensi* Liston. Journal of Vector Borne diseases. (Accepted)
2. **Mohanty AK**, de Souza C, Harjai D, Ghavanalkar P, Fernandes M, Almeida A, Walke J, Manoharan SK, Pereira L, Dash R, Mascarenhas A , Gomes E, Thita T, Chery L, Anvikar AR, Kumar A, Valecha N, Rathod P, Patrapuvich R. Optimization of

Plasmodium vivax sporozoite production from *Anopheles stephensi* in South West India. Malaria Journal. 2021 Dec; 20(1):1-3.

3. Dey G, **Mohanty AK**, Sreenivasamurthy SK, Kumar M, Kumar A, Prasad TK. Proteomics dataset of adult *Anopheles stephensi* female brain. Data in brief. 2020 Oct 1; 32:106243.
4. Almeida J, **Mohanty AK**, Dharini N, Hoti SL, Kerkar S, Kumar A. A report on novel mosquito pathogenic Bacillus spp. isolated from a beach in Goa, India. International Journal of Mosquito Research 2020; 7(2): 21-29.
5. Kumar A, Chaturvedi HK, **Mohanty AK**, Sharma SK, Malhotra MS, Pandey A. Surveillance based estimation of burden of malaria in India, 2015–2016. Malaria journal. 2020 Dec; 19 (1):1-2.
6. Almeida J, **Mohanty AK**, Kerkar S, Hoti SL, Kumar A. Current status and future prospects of bacilli-based vector control. Asian Pacific Journal of Tropical Medicine 2020; 13(12): 525-534.
7. **Mohanty AK**, Dey G, Kumar M, Sreenivasamurthy SK, Garg S, Prasad TK, Kumar A. Proteome data of female *Anopheles stephensi* antennae. Data in brief. 2019 Jun 1; 24:103911.
8. Kumar M, **Mohanty AK**, Dey G, Sreenivasamurthy SK, Kumar A, Prasad K. Dataset on fat body proteome of *Anopheles stephensi* Liston. Data in brief. 2019 Feb 1; 22:1068-73.
9. Dey G, **Mohanty AK**, Sreenivasamurthy SK, Kumar M, Prasad TK, Kumar A. Proteome data of *Anopheles stephensi* salivary glands using high-resolution mass spectrometry analysis. Data in brief. 2018 Dec 1; 21:2554-61.
10. Sreenivasamurthy SK, Dey G, Kumar M, **Mohanty AK**, Kumar A, Prasad TK. Quantitative proteome of midgut, malpighian tubules, ovaries and fat body from sugar-fed adult *An. stephensi* mosquitoes. Data in brief. 2018 Oct 1; 20:1861-6.
11. Dey G, **Mohanty AK**, Kumar M, Sreenivasamurthy SK, Patil AH, Prasad TK, Kumar A. Proteome data of *Anopheles stephensi* ovary using high-resolution mass spectrometry. Data in brief. 2018 Oct 1; 20:723-31.
12. **Mohanty AK**, Nina PB, Ballav S, Vernekar S, Parkar S, D'souza M, Zuo W, Gomes E, Chery L, Tuljapurkar S, Valecha N. Susceptibility of wild and colonized *Anopheles stephensi* to *Plasmodium vivax* infection. Malaria journal. 2018 Dec;17(1):1-0.
13. Dey G, **Mohanty AK**, Sreenivasamurthy SK, Kumar M, Prasad TK, Kumar A. Proteome data of *Anopheles stephensi* salivary glands using high-resolution mass spectrometry analysis. Data in brief. 2018 Dec 1; 21:2554-61.
14. **Mohanty AK**, Dey G, Kumar M, Sreenivasamurthy SK, Garg S, Prasad TK, Kumar A. Mapping *Anopheles stephensi* midgut proteome using high-resolution mass spectrometry. Data in brief. 2018 Apr 1; 17:1295-303.
15. Dhawan R, Pillai CR, **Mohanty AK**, Kumar A. Achieving in vitro gametocytogenesis of *Plasmodium falciparum* in optimal conditions: A review.
16. Kumar M, **Mohanty AK**, Sreenivasamurthy SK, Dey G, Advani J, Pinto SM, Kumar A, Prasad TS. Response to blood meal in the fat body of *Anopheles stephensi* using quantitative proteomics: toward new vector control strategies against malaria. Omics: a journal of integrative biology. 2017 Sep 1; 21(9):520-30.
17. Nina PB, **Mohanty AK**, Ballav S, Vernekar S, Bhinge S, D'souza M, Walke J, Manoharan SK, Mascarenhas A, Gomes E, Chery L. Dynamics of *Plasmodium vivax* sporogony in wild *Anopheles stephensi* in a malaria-endemic region of Western India. Malaria Journal. 2017 Dec; 16(1):1-2.
18. Sreenivasamurthy SK, Madugundu AK, Patil AH, Dey G, **Mohanty AK**, Kumar M, Patel K, Wang C, Kumar A, Pandey A, Prasad TS. Mosquito-borne diseases and Omics: tissue-restricted expression and alternative splicing revealed by transcriptome profiling of *Anopheles stephensi*. Omics: a journal of integrative biology. 2017 Aug 1; 21(8):488-97.
19. Dhawan R, **Mohanty AK**, Kumar M, Dey G, Advani J, Prasad TK, Kumar A. Data from salivary gland proteome analysis of female *Aedes aegypti* Linn. Data in brief. 2017 Aug 1; 13:274-7.
20. Prasad TK, **Mohanty AK**, Kumar M, Sreenivasamurthy SK, Dey G, Nirujogi RS, Pinto SM, Madugundu AK, Patil AH, Advani J, Manda SS. Integrating transcriptomic and

- proteomic data for accurate assembly and annotation of genomes. *Genome research*. 2017 Jan 1; 27(1):133-44. (Shared first author)
21. Dhawan R, Kumar M, **Mohanty AK**, Dey G, Advani J, Prasad TK, Kumar A. Mosquito-borne diseases and omics: salivary gland proteome of the female *Aedes aegypti* mosquito. *Omics: a journal of integrative biology*. 2017 Jan 1; 21(1):45-54.
 22. **Mohanty AK**, Garg S, Dhindsa K, Kumar A. Variable region of 16s rRNA is essential for the identification of Group 1 mosquito-pathogenic strains of Lysinibacillus. *Adv Biotech & Micro*. 2(2): 555583.
 23. Kumar A, Hosmani R, Jadhav S, de Sousa T, **Mohanty AK**, Naik M, Shettigar A, Kale S, Valecha N, Chery L, Rathod PK. *Anopheles subpictus* carry human malaria parasites in an urban area of Western India and may facilitate perennial malaria transmission. *Malaria journal*. 2016 Dec; 15(1):1-8.
 24. Chaerkady R, Kelkar DS, Muthusamy B, Kandasamy K, Dwivedi SB, Sahasrabuddhe NA, Kim MS, Renuse S, Pinto SM, Sharma R, Pawar H, Sekhar N R, **Mohanty A K**, Getnet D, Yang Y, Zhong J, Dash A P, MacCallum R M, Delanghe B, Mlambo G, Kumar A, Prasad T S K, Okulate M, Kumar N and Pandey A. A proteogenomic analysis of *Anopheles gambiae* using high-resolution Fourier transform mass spectrometry. *Genome research*. 2011 Nov 1; 21(11):1872-81.
 25. Nagpal BN, Saxena R, Srivastava A, Singh N, Ghosh SK, Sharma SK, Kumar A, Kumar H, Sharma AS, Chand SK, Ojha VP, Mohanty S, **Mohanty AK**, Dasgupta RK, Singh GP and Dash AP. Retrospective study of chikungunya outbreak in urban areas of India. *The Indian journal of medical research*. 2012 Mar; 135(3):351.
 26. Nayak PK, **Mohanty AK**, Gaonkar T, Kumar A, Bhosle SN, Garg S. Rapid identification of polyhydroxy alkanoate accumulating members of *Bacillales* using internal primers for phaC gene of *Bacillus megaterium*. *International Scholarly Research Notices*. 2013;2013.
 27. Sreenivasamurthy SK, Dey G, Ramu M, Kumar M, Gupta MK, **Mohanty AK**, Harsha HC, Sharma P, Kumar N, Pandey A, Kumar A. A compendium of molecules involved in vector-pathogen interactions pertaining to malaria. *Malaria journal*. 2013 Dec; 12(1):1-7.

Book chapters: 3 No.s

| S. No | Title of Chapter | Details |
|-------|--|--|
| 1 | Mohanaty AK , Garg S, Dhindsa K, Kumar H and Kumar A (2012) Phenotypic Characterization of Mosquito Larvicidal Ed. Barbuddhe S.B., Ramesh R., Lysinibacillus Strains Isolated from Paddy Field and Mangrove Vegetation In : microbial diversity and its application, pp. 49-58. | Ed. Barbuddhe S.B., Ramesh R., Singh N.P. New India Publishing Agency, New Delhi |
| 2 | Nayak P, Gaonkar T , Mohanty A , Kumar A, Bhosle S, Garg S (2012) Isolation and Characterization of Polyhydroxyalkanoates Producing Bacteria from Coastal Sand-Dune Ecosystem In : microbial diversity and its application pp. 75-82 | Ed. Barbuddhe S.B., Ramesh R., Singh N.P. New India Publishing Agency, New Delhi |
| 3 | Ajeet Kumar Mohanty , Keshava Prasad, Sandeep Garg and Ashwani Kumar. Proteogenomics of vector mosquitoes: Progress and Prospects: Major tropical Diseases; Public Health Perspective pp.38-50 | Ed. Ashwani Kumar, SavioRodriques and Amit Dias. Broadway Publishing House, Panaji, Goa. |

Popular article

1. Ashwani Kumar, **Ajeet Kumar Mohanty**, T. S. Keshava Prasad. (2017) Mosquito-borne diseases – how can omics help characterize vectors? <https://www.id-hub.com/2017/07/11/mosquito-borne-diseases-can-omics-help-characterize-vectors/> Infectious Diseases Hub.