

## Maria Altaf Satti

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### Education

- 2016-2021 *Ph.D.*, Bioinformatics, **The Graduate University for Advanced Studies, SOKENDAI**, Japan
- 2018-2021 *Ph.D.*, Agricultural, Environmental and Food Science and Technology, **University of Bologna**, Italy (Dual doctoral degree -cotutelle agreement)
- 2016 *M.S.*, Computational Sciences and Engineering, **National University of Sciences and Technology, NUST**, Islamabad, Pakistan
- 2014 *B.S.*, Bioinformatics, **Mohammad Ali Jinnah University, MAJU**, Islamabad, Pakistan

### Research Experience

- 2021 Oct-2022 July *Postdoctoral Research Associate*, **Department of Cellular and Molecular Medicine, Texas A&M**, USA
- Project: “Development of computational method for de novo identification of intronic polyadenylation from standard RNA-seq data, focusing on Cancer dynamics”
- 2018-2021 Worked in collaboration with Paola Mattarelli, Ph.D., Department of Agricultural and Food Sciences, University of Bologna, Italy.  
(Dual doctoral degree -cotutelle agreement signed)
- 2018 (Oct-Dec), 2019 (Sep-Oct) Visiting fellow, Department of Agricultural and Food Sciences, University of Bologna Viale Fanin 44, 40127 Bologna, Italy.  
Principal investigator: Paola Mattarelli, Ph.D.
- Project: Genomic analysis of *Bifidobacterium* novel species
- 2017 Attendee, The 28<sup>th</sup> International conference on Genome Informatics GIW/BIOINFO, Seoul Korea
- 2016 Masters thesis  
Title: “Development of a Computational Method for Identification of Genomic Structural Variants Using Next Generation Sequencing”

2016 Attendee, 2 days hand-on Workshop Next Generation Sequence (NGS) data analysis, COMSATS Institute of Information Technology Islamabad, Pakistan

2014 Bachelors Project  
Title: "Visualization and Mining of Human Microbiota using Computational Techniques"

2013 Assisted, 4<sup>th</sup> One day workshop, Mohammad Ali Jinnah University, Islamabad, Pakistan

**Languages** R, Python, C++, MATLAB, C

### **Work Experience**

June 2013- August 2013 Internship, Genetic Resource Centre, Rawalpindi, Pakistan

2011-2013 Student Assistant Ship, Department of Computer Science, Mohammad Ali Jinnah University, Islamabad, Pakistan

### **Honors and Distinctions**

2021 Morishima Award for outstanding performance during doctoral program in the institute (National institute of Genetics)

2018, 2019 Traveling grant, The Graduate University for Advanced Studies, Japan

2016-Present National Institute of Genetics Japan, MEXT scholarship, 5- year Doctoral program

Fall 2010-Fall 2014 Mohammad Ali Jinnah University, Islamabad, Pakistan, PGC Scholarship

2014 Mohammad Ali Jinnah University, Islamabad, Pakistan, Silver Medal, B.S. Convocation

2014 Mohammad Ali Jinnah University, Islamabad, Pakistan, 2<sup>nd</sup> Prize, Open House

2014 1<sup>st</sup> Prize Poster Presentation, 2<sup>nd</sup> Conference on emerging trends in Bioinformatics and Computational Biology

2010-2013 Mohammad Ali Jinnah University, Islamabad, Pakistan, Chancellor's Honor Roll / Dean's Honor Roll

## Presentations

### Oral Presentations

Comparative genomic analysis of *Bifidobacterium* species isolated from Egyptian fruit bat *Rousettus egyptiacus*, The COMBINE/AYRCOB Symposium, Sydney Australia, 2019

Bifidobacterial occurrence in giant bats: the two new species *Bifidobacterium vespertilionis* and *Bifidobacterium rousettii*, 10<sup>th</sup> National Conference on Research in Parks, Verona Italy, 2019

Comparative analysis of probiotic genus *Bifidobacterium*, Workshop on Commemoration of the centenary of the birth of Prof. Vittorio Scardovi, one of the most relevant Italian microbiologist and taxonomist on the probiotic bacteria, Bologna Italy, 2018

Comparative analysis of probiotic genus *Bifidobacterium* and *Lactobacillus* based on new definition of core genome, The 28<sup>th</sup> International Conference on Genome Informatics GIW/ BIOINFO, Seoul Korea, 2017

### Poster Presentations

Comparative genomic analysis of *Bifidobacterium* species isolated from Egyptian fruit bat *Rousettus egyptiacus*, Joint 30<sup>th</sup> International Conference on Genome Informatics (GIW) & Australian Bioinformatics and Computational Biology Society (ABACUS) Annual Conference, Sydney Australia, 2019

Comparative Analysis of probiotic genus *Bifidobacterium* and *Lactobacillus*, AIST-RIKEN joint exchange meeting for computational biology, Tokyo, Japan, 2018

Insight into the immunomodulatory role of *Bifidobacterium bifidum*, The 12th Annual Meeting of the Japanese Society for Genomic Microbiology, Kyoto, Japan, 2018

Visualization and Mining of Human Microbiota using Computational Techniques, 2nd Conference on emerging trends in Bioinformatics and Computational Biology, Islamabad Pakistan, 2014

## Publications

1. Satti M, Tanizawa Y, Endo A, Arita M. Comparative analysis of probiotic bacteria based on a new definition of core genome. Journal of Bioinformatics and Computational Biology; 2018.
2. Modesto M, Watanabe K, Arita M, Satti M, et al. *Bifidobacterium jacchi* sp. nov., isolated from the faeces of a baby common marmoset (*Callithrix jacchus*). International journal of systematic and evolutionary microbiology; 2019.
3. Modesto M\*, Satti M\*, Watanabe K, Puglisi E, Morelli L, et al. Characterization of *Bifidobacterium* species in faeces of the Egyptian fruit bat: Description of *B.*

- vespertilionis* sp. nov. and *B. rousetti* sp. nov. Systematic and applied microbiology: 2019.
4. Modesto M, Satti M, Watanabe K, Sciavilla P, Felis GE, et al. *Alloscardovia theropitheci* sp. nov., isolated from the faeces of gelada baboon, the bleeding heart monkey (*Theropithecus gelada*). International Journal of Systematic and Evolutionary Microbiolo
  5. Modesto M\*, Satti M\*, Watanabe K, Scarafile D, Huang CH, et al. Phylogenetic characterization of two novel species of the genus *Bifidobacterium*: *Bifidobacterium saimiriisciurei* sp. nov. and *Bifidobacterium platyrrhinorum* sp. nov. Systematic and applied mi
  6. Modesto M, Satti M, Watanabe K, Huang CH, Liou JS, et al. Bifidobacteria in two-toed sloths (*Choloepus didactylus*): phylogenetic characterization of the novel taxon *Bifidobacterium choloepi* sp. nov. International Journal of Systematic and Evolutionary Mic
  7. Satti M, Modesto M, Endo A, Kawashima T, Mattarelli P, Arita M. Host-Diet Effect on the Metabolism of *Bifidobacterium*. Genes; 2021.

\*co-first authors

## References

### 1. Masanori Arita

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National Institute of Genetics, Yata 1111, Mishima, 411-8540 Japan  
Email: [arita@nig.ac.jp](mailto:arita@nig.ac.jp)

### 2. Paola Mattarelli

Assistant Professor  
Department of Agricultural and Food Sciences  
University of Bologna, Viale Fanin 44, 40127 Bologna, Italy  
Email: [paola.mattarelli@unibo.it](mailto:paola.mattarelli@unibo.it)

### 3. Sahar Fazal

Professor, Head of Biosciences Department  
Capital University of Science & Technology, Islamabad, Pakistan  
Email: [sahar@cust.edu.pk](mailto:sahar@cust.edu.pk)