Curriculum Vitae

Nanami (Masuyama) Yamaguchi

Ph.D. Student

Institute for Advanced Biosciences, Keio University

Systems Biology Program, Graduate School of Media and Governance, Keio University

E-mail: nananannsutann@gmail.com

Education

March 2020 Master of Media and Governance with honor, Systems Biology Program, Keio University, Tokyo, Japan

Supervisors: Drs. Nozomu Yachie and Masaru Tomita

2019 SFC STUDENT AWARD

March 2018 Bachelor of Arts in Environment and Information Studies with honor, Keio University, Tokyo, Japan.

Supervisors: Drs. Nozomu Yachie and Masaru Tomita

Graduation Thesis Award

Research Experience

October 2015-March 2018

Visiting student @Yachie laboratory, the University of Tokyo, Tokyo, Japan

Work Experience

February 2021 –

Part-time researcher @Yachie laboratory, University of British Columbia, , British Columbia, Canada

April 2018 – Current

Part-time researcher @Yachie laboratory, the University of Tokyo, Tokyo, Japan

Awards/Fellowships

April 2021– JSPS Fellowship (DC2), Japan Society for the Promotion of Science (JSPS)

April 2015–Today TTCK Research Fellowship, Keio University
March 2020 SFC STUDENT AWARD, Keio University

June 2019 Best Poster Award, The 13th International Workshop on Advanced Genomics, Tokyo, Japan, June 2019.

March 2018 SFC, Graduation Thesis Award, SFC, Keio University

Publications

Peer-reviewed articles

- 1- Rina C. Sakata[†], Soh Ishiguro[†], Hideto Mori[†], Mamoru Tanaka, Kenji Tatsuno, Hiroki Ueda, Shogo Yamamoto, Motoaki Seki, Nanami Masuyama, Keiji Nishida, Hiroshi Nishimasu, Kazuharu Arakawa, Akihiko Kondo, Osamu Nureki, Masaru Tomita, Hiroyuki Aburatani, and Nozomu Yachie. Base editors for simultaneous introduction of C-to-T and A-to-G mutations. *Nature Biotechnology*, 38, pp865–869, 2020.
- 2- Hirofumi Nishizono, Mohamed Darwish, Hideki Uosaki, <u>Nanami Masuyama</u>, Motoaki Seki, Hiroyuki Abe, Nozomu Yachie and Ryohei Yasuda. Use of Freeze-thawed Embryos for High-efficiency Production of Genetically Modified Mice. *Journal of Visualized Experiments* 158, e60808, 2020.
- 3- Yumi Murai, Takahiro Masuda, Yasuhide Onuma, Daniel Evans-Yamamoto, Nao Takeuchi, Hideto Mori, Nanami Masuyama, Soh Ishiguro, Nozomu Yachie, Kazuharu Arakawa. Complete Genome Sequence of Bacillus sp. Strain KH172YL63, Isolated from Deep-Sea Sediment. *Microbiology Resource Announcements*, Accepted on April 2, 2020.
- 4- Daniel Evans-Yamamoto, Nao Takeuchi, Takahiro Masuda, Yumi Murai, Yasuhide Onuma, Hideto Mori, <u>Nanami Masuyama</u>, Soh Ishiguro, Nozomu Yachie, and Kazuharu Arakawa. Complete genome sequence of Psychrobacter sp. strain KH172YL61 isolated from deep-sea sediments in the Nankai Trough, Japan. *Microbiology Resource Announcements*, 8, e00326-19, 2019.
- 5- Nanami Masuyama, Hideto Mori and Nozomu Yachie, DNA barcodes evolve for high-resolution cell lineage tracing. *Current Opinion in Chemical Biology* 52, 63-71, 2019.
- 6- Nozomu Yachie, Robotic Biology Consortium (including Nanami Masuyama), and Toru Natsume. Robotic crowd biology with Maholo LabDroids. *Nature Biotechnology*, 35, pp310-312, 2017.

Review articles (in Japanese)

1- Soh Ishiguro, <u>Nanami Masuyama</u>, Nozomu Yachie. DNA barcode tricks to tackle combinatorial explosion of experiments in omic measurements. *Journal of Japanese Biochemical Society*, 89:538-545, 2017.

Talks

1- Nanami Masuyama, Hideto Mori, Soh Ishiguro, Osamu Masui, Hirohumi Nishizono, Mikiko Negishi, Rina Sakata, Arman Adel, Motoaki Seki, Keiji Nishida, Akihiko Kondo, Masaru Tomita, Hiroyuki Aburatani, Haruhiko Koseki and Nozomu Yachie. *The 20th International Conference on System Biology*, Okinawa, Japan, October 2019.

Poster Presentations

- 1- Nanami Masuyama, Hideto Mori, Soh Ishiguro, Osamu Masui, Hirohumi Nishizono, Mikiko Negishi, Rina Sakata, Arman Adel, Motoaki Seki, Keiji Nishida, Akihiko Kondo, Masaru Tomita, Hiroyuki Aburatani, Haruhiko Koseki and Nozomu Yachie. *The 13th International Workshop on Advanced Genomics, Tokyo, Japan, June 2019.*
- 2- <u>Nanami Masuyama</u>, Hideto Mori, Soh Ishiguro, Osamu Masui, Hirohumi Nishizono, Mikiko Negishi, Rina Sakata, Arman Adel, Motoaki Seki, Keiji Nishida, Akihiko Kondo, Masaru Tomita, Hiroyuki Aburatani, Haruhiko Koseki and Nozomu Yachie. *The 4th Annual Meeting of The Japanese Society for Genome Editing*, *Tokyo*, *Japan*, *May 2019*.
- 3- Nanami Masuyama, Soh Ishiguro, Hideto Mori, Osamu Masui, Motoaki Seki, Kentaro Kakoi, Ayaka Matsuo, Rina Sakata, Keiji Nishid, Akihiko Kondo, Masaru Tomita, Hiroyuki Aburatani, Haruhiko Koseki and Nozomu Yachie. *The Molecular Biology Society of Japan, Kobe, Japan, December 2017.*
- 4- <u>Nanami Masuyama</u>, Soh Ishiguro, Hideto Mori, Osamu Masui, Motoaki Seki, Kentaro Kakoi, Ayaka Matsuo, Rina Sakata, Keiji Nishid, Akihiko Kondo, Masaru Tomita, Hiroyuki Aburatani, Haruhiko Koseki and Nozomu Yachie. From Genetic Networks to a Cellular Wiring Diagram, Tokyo, Japan, April 2017.

Research Funding

- 1- Doctoral Program Support Program, Keio University (April 2020–March 2021)
- 2- Taikichiro Mori Memorial Research Grant, Taikichiro Mori Memorial Research Fund (April 2020–March 2021)
- 3- Taikichiro Mori Memorial Research Grant, Taikichiro Mori Memorial Research Fund (April 2019–March 2020)
- 4- Taikichiro Mori Memorial Research Grant, Taikichiro Mori Memorial Research Fund (April 2018-March 2019)
- 5- Yamagishi Student Project Support Program. (June 2017-February 2018)
- 6- Yamagishi Student Project Support Program. (June 2016-February 2017)

Patents submitted

1- Nanami Yamaguchi, Hideto Mori, Nozomu Yachie. FRACTAL Assembly (Japan), Application number: 2020-53077.