Sungsik Kong | Curriculum Vitæ

Department of Electric and Computer Engineering – University of California San Diego 3180 Voigt Dr. Franklin Antonio Hall, La Jolla, CA., USA, 92093

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Summary

- I am a computational evolutionary biologist with trainings in biology, statistics, and bioinformatics
- My research focuses on phylogenetic theory, phylogenetic networks, and reticulate evolution
- I have international education and work experiences in South Korea, Australia, Canada, and USA
- I love adventure and travelled in Asia for a year on foot, by bicycle, and motorbike

Professional Experience

Wisconsin Institute for Discovery

University of Wisconsin-Madison, WI., USA

Mentor: Prof. Dr. Claudia Solís-Lemus

University of California San Diego, CA., USA

Mentor: Prof. Dr. Siavash Mirarab

2023-Presnet

Research Associate

2023

2018-2023

2013-2015

Postdoctoral Scholar

Education

Ph.D. in Evolution, Ecology, and Organismal Biology

Department of Electrical and Computer Engineering

The Ohio State University, Columbus, OH., USA

Dissertation: Towards the network of life: Phylogenetic networks as a tool to understand complex

evolutionary histories

Advisor: Prof. Dr. Laura Kubatko

M.Sc. in Ecology and Evolutionary Biology

University of Toronto, Toronto, ON., Canada

Thesis: On the use of Median-Joining networks: A philosophical and empirical evaluation of its suitability

in evolutionary biology

Advisor: Prof. Dr. Robert Murphy

B.Sc. in Zoology and Ecology and Evolutionary Biology

University of Toronto, Toronto, ON., Canada

2006-2012

Publications

Total number: 16

h-index: 7 [Google scholar]

[‡]Co-correspondence; [†]Equal contribution

Selected peer-reviewed journal articles.

- [16] Kong S (2022) Digest: Frequent hybridization in *Darevskia* rarely leads to the evolution of asexuality. *Evolution* 76:9 2216–2217 doi:10.1111/evo.14587
- [15] Kong S[†], Pons JC[†], Kubatko LS, and Wicke K. (2022) Classes of phylogenetic networks and their mathematical and biological significance. *Journal of Mathematical Biology* 84:47 doi:10.1007/s00285-022-01746-y
- [14] Kong S and Kubatko LS. (2021) Comparative performance of popular methods for hybrid detection using genomic data. *Systematic Biology* 70:5 891–907 doi:10.1093/sysbio/syaa092
- [13] Sánchez-Pacheco SJ[‡], Kong S[‡], Pulido-Santacruz P, Murphy RW, and Kubatko L. (2020) Median-joining network analysis of SARS-CoV-2 genomes is neither phylogenetic nor evolutionary. *Proc. Natl. Acad. Sci. U.S.A* 117:23 12518–12519 doi:10.1073/pnas.2007062117
- [12] Kong S[†], Sánchez-Pacheco S[†], and Murphy RW. (2016) On the use of Median-Joining Networks in evolutionary biology. *Cladistics* 32: 691–699 doi:10.1111/cla.12147

Collaborative peer-reviewed journal articles.....

- [11] Glon H, Häussermann V, Brewin P, Brickle P, Kong S, Smith M, and Daly M. (2023) There and back again: The unexpected journeys of *Metridium* de Blainville, 1824 between the old oceans and throughout the modern world. *Biological Bulletin*
- [10] Nguyen HQ, Ho P-T, Kong S, Bae Y, Pham TH, and Jang Y. (2022) A time-calibrated mitogenomic phylogeny suggests that Korean *Hyalessa fuscata* is a bridge between Chinese and Japanese *H. maculaticollis*. *Journal of Genetics* 102:9 doi:10.1007/s12041-022-01405-7
- [9] Nguyen HQ, Kim E, Bae Y, Chae S, Ji S, Heo J, Kong S, Nguyen TK, Pham TH, and Jang Y. (2022) An effective method for accurate life-stage delimitation of cicadas. *Journal of Asia-Pacific Entomology* 101952 doi:10.1016/j.aspen.2022.101952
- [8] Glon MG, Broe MB, Crandall KA, Daly M, Kong S, Thoma RF, and Freudenstein JV. (2022) Anchored hybrid enrichment resolves the *Lacunicambarus* (Decapoda: Cambaridae) phylogeny. *Journal of Crustacean Biology* 42:1 ruab073 doi:10.1093/jcbiol/ruab073
- [7] Kim M, Nguyen HQ, Yi Y, Ahn J, Kim YE, Jang S, Kim S, Kong S, and Borzée A. (2020) Policy recommendation on whaling, trade and watching of cetaceans in the Republic of Korea. *Biodiversity Journal* 11: 255–258 doi:10.31396/Biodiv.Jour.2020.11.1.255.258

- [6] Weiringa JG, Boot MR, Dantas-Queiroz MV, Duckett D, Fonseca EM, Glon H, Hamilton N, Kong S, Lanna FM, Mattingly KZ, Parsons DJ, Smith ML, Stone BW, Thompson C, Zuo L, and Carstens BC. (2020) Does habitat stability structure intraspecific genetic diversity? It's complicated. . . Frontiers in Biogeography e45377 doi:10.21425/F5FBG45377
- [5] Borzée A, Purevdorj Z, Kim YI, Kong S, Choe M, Yi Y, Kim K, Kim A, and Jang Y. (2020) Breeding preferences in the Treefrogs *Dryophytes japonicus* (Hylidae) in Mongolia, the northern limit of their range. *Journal of Natural History* 53: 2685-2698 doi:10.1080/00222933.2019.1704458
- [4] Bae Y, Kong S, Yi Y, Jang Y, and Borzée A. (2019) Additional new type of threat to *Hynobius* salamander eggs: predation by loaches (*Misgurnus* sp.) in agricultural wetlands. *Animal Biology* 69:4 451–461 doi:10.1163/15707563-20191070
- [3] Groffen J, Kong S, Jang Y, and Borzée A. (2019) The invasive American bullfrog (*Lithobates catesbeianus*) in the Republic of Korea: history and recommendation for population control. *Management of Biological Invasions* 10:3 517–535 doi:10.3391/mbi.2019.10.3.08
- [2] Borzée A, Kong S, Didinger C, Nguyen HQ, and Jang Y. (2018) A ring-species or a ring of species? Phylogenetic relationship between two treefrog species around the Yellow Sea: *Dryophytes suweonensis* and *D. immaculatus*. The Herpetological Journal 28: 160–170
- [1] Kim K, Kong S, Kim YI, Borzée A, Bae Y, and Jang Y. (2018) Japanese hard ticks (*Ixodes nipponensis*) parasitizing on the endangered leopard cat (*Prionaliurus bengalensis euptilura*) in the Republic of Korea. *Animal Systematics, Evolution and Diversity* 34:1 23–26 doi:10.5635/ASED.2018.34.1.032

Policy recommendations.

[1] Borzée A, Nguyen HQ, Ahn J, Kim M, Kim S, Jang S, Kong S, Kim YE, and Yi Y. (2017) Policy recommendation on the negative impact of infrastructure development on the population of endangered Long-Tailed Goral in Oseck-Kkeutchong. Society for Conservation Biology Korean Chapter

Submitted

- [3] Kong S, Swofford DL, and Kubatko LS. (In Review) Inference of phylogenetic networks from sequence data using composite likelihood. Preprint available on *bioRxiv* at 10.1101/2022.11.14.516468v1
- [2] Kong S, Sánchez-Pacheco SJ, and Murphy RW. (In Revision) Topological incongruence between median-joining networks and Bayesian inference phylogenies. Preprint available on *bioRxiv* at 10.1101/2022.10.01.510440
- [1] Fu M, Eimes JA, Kong S, Lamichhaney S, and Waldman B. (In Review) Chytrid-resistant genotypes arise from multiple mutations.

Presentations *Virtual presentation Invited talks..... [11] Algorithms and Complexity in Phylogenetics* 2022 Title: Inferring phylogenetic networks from sequence data using composite likelihood Video recording available on YouTube [link] [10] Indiana University, Bloomington, IN., USA 2022 Dr. Matthew Hahn laboratory Title: Towards the Network of Life: Phylogenetic networks as a tool to understand complex evolutionary histories [9] University of California San Diego, CA., USA* 2022 Drs. Siavash Mir Arabbaygi and Vineet Bafna laboratory Title: Towards the Network of Life: Phylogenetic networks as a tool to understand complex evolutionary histories [8] Evolution Meeting, Cleveland, OH., USA 2022 Society of Systematic Biologists (SSB) Ernst Mayr award competition Title: Inferring phylogenetic networks using composite likelihood directly from sequence data [7] Louisiana State University, Baton Rouge, LA., USA 2022 Dr. Jeremy Brown laboratory and Department of Biological Sciences Title: Inferring phylogenetic networks using composite likelihood directly from sequence data 2021 [6] The Korean Association of Biological Sciences, Seoul, South Korea* The Ecological Society of Korea Section Title: Comparative performance of popular methods for hybrid detection using genomic data [5] Workshop in Mathematical and Computational Biology* 2021 Bioinformatics Section Title: Comparative performance of popular methods for hybrid detection using genomic data [4] Misericordia University, Dallas, PA., USA* 2021 Biology Seminar Series Title: How my gap year changed my perspectives [3] The Stone Laboratory, Gibraltar Island, OH., USA 2019 EEOB 1930 Introduction to Biological Studies-Aquatic Biology Title: Phylogenetic networks and behavioral evolution of Primates [2] Ewha Womans University, Seoul, South Korea 2016

Eco-science/Eco-creative Graduate Seminar Title: On the use of Median-Joining network

[1]	Ewha Womans University, Seoul, South Korea Laboratory of Behavior and Ecology Weekly Seminar Title: The evolution of infanticide and male takeover in primates in a macroevolutionary framework	2013
R	egular talks	
	Evolution Meeting* Title: Comparative performance of popular methods for hybrid detection using genomic data	2021
[5]	The Ohio State University, Columbus, OH., USA* EEOB Annual Darwin Series Title: Comparative performance of popular methods for hybrid detection using genomic data	2021
[4]	The Ohio State University, Columbus, OH., USA Evolution & Ecology Club Title: Which tool to use to detect hybridization?	2020
[3]	SSB Standalone Meeting, Gainesville, FL., USA Title: Comparison and evaluation on the accuracy of popular hybrid detection methods using simulate	2020 d data
[2]	The Ohio State University, Columbus, OH., USA Evolution & Ecology Club Title: The giving tree: how the branching pattern of a tree is used in evolutionary biology	2019
[1]	Ewha Womans University, Seoul, South Korea Animal Communication Symposium Title: Illustrating hybridization	2017
Р	oster presentations.	
[5]	Intelligent Systems for Molecular Biology (ISMB), The International Society for Computational Biology (ISCB), Madison, WI., USA Best Poster Award Winner Title: Inference of phylogenetic networks from sequence data using composite likelihood	2022
[4]	Evolution Meeting, Providence, RI., USA Title: Performance and accuracy of Median-Joining Networks and comparisons with Bayesian phylogenesis.	2019 enies
[3]	The Korean Association of Biological Sciences, Yong-In, South Korea Title: Performance and accuracy of Median-Joining Networks and comparisons with Bayesian phylogenesis of the Company	2017 enies
[2]	Korean-American Scientists and Engineers Association (KSEA) Young Generation Technical and Leadership Conference (YGTLC), Houston, TX., U Title: Matrilineal relationships of green pitvipers with the description of a new species of the genus <i>Trime</i> (Squamata: Viperidae: Crotalinae) from the central highlands of Vietnam	
[1]	University of Toronto, Toronto, ON., Canada Department of Ecology and Evolutionary Biology undergraduate research Fair Title: The evolution of infanticide and male takeover in primates in a macro- evolutionary framework	2012

Academic Experience

Academic visit	
University of Wisconsin-Madison, WI., USA Wisconsin Institute for Discovery	2022 <i>Visiting Ph.D. student</i>
Mentor: Prof. Dr. Claudia Solís-Lemus	
Louisiana State University, Baton Rouge, LA., USA	2022
Department of Biological Sciences Mentor: Prof. Dr. Jeremy Brown	Visiting Ph.D. student
Research experience.	
Ewha Womans University, Seoul, South Korea Division of Eco-science	2016–2017 <i>Researcher</i>
Mentor: Prof. Dr. Yikweon Jang	Kesearcher
Ewha Womans University, Seoul, South Korea	2013
Division of Eco-science Mentor: Prof. Dr. Jae Chun Choe	Student researcher
University of Toronto, ON., Canada	2011–2013
Faculty of Forestry Mentor: Prof. Dr. Sandy Smith and Eric Davies	Work-Study student Biodiversity technician
Royal Ontario Museum, Toronto, ON., Canada	2012
Centre for Biodiversity and Conservation Biology Mentor: Prof. Dr. Robert Murphy	Student researcher
University of Toronto, ON., Canada	2011–2012
Department of Ecology and Evolutionary Biology Mentor: Prof. Dr. Deborah McLennan	Student researcher
University of Toronto, ON., Canada	2010
Cities Centre, Department of Geography	Student data collector
Fieldwork experience.	
South Korea, Mongolia, and ChinaIndonesia, Laos, and South Korea	2017 2016
Greater Toronto Area (GTA)	2010–2013

Teaching and Mentoring Experience

The Ohio State University.	
Teaching assistant	
o EEOB 3310 Evolution	(Spring, Fall) 2021
	(Spring, Summer, Fall) 2020
- FEOD 0500 H	(Spring, Fall) 2019
• EEOB 2520 Human Physiology	(Summer) 2021
 EEOB 1930 Introduction to Biological Studies-Aquatic Biology BIOL 1113 Energy Transformation and Development 	(Summer) 2019 (Fall) 2018
University of Toronto	
Teaching assistant	
 BIO 120 Adaptation and Biodiversity 	(Summer) 2015
	(Summer) 2014
	(Fall) 2013
 BIO 220 From Genomics to Ecosystems in a Changing World 	(Spring) 2015
 EEB 362 Macroevolution 	(Fall) 2014
o <i>EEB 386</i> Avian Biology	(Spring) 2014
Miscellaneous.	
o Daruma Eco-Farm, Bang Phra, Si Racha District, Chon Buri, Thailanc	2016
 Cheong-Shim International English Camp, Ga-Pyeong, South Korea 	2009
(Co-)supervision of undergraduate students.	
Alanna Kong	2021–2022
Undergraduate student at University of Toronto	
University of Toronto Woodsworth Mentorship Program	
Devansh Pandey	2021–2022
Undergraduate student at Indian Institute of Technology Kharagpur	
Currently a graduate student at University of Texas at Austin	
Científico Latino	2020 2021
McKenna Molnar Color	2020-2021
Graduate student at The Ohio State University	
EEOB Graduate Mentoring Program, The Ohio State University, OH., USA • Shelby Jansens and Kimberly Wiles	2020
Co-supervised with Dr. Laura Kubatko	2020
Mathematical Biosciences Institute Research Experiences for Undergraduates	nrogram
Title: Detecting shifts in the evolutionary process in SARS-CoV-2 genomes	program
Tulio Soares and Ellen Southam	2014-2015
Co-supervised with Dr. Pedro Bernardo	
University of Toronto and Royal Ontario Museum	
Shreeram Senthivasan	2014
Co-supervised with Drs. Pedro Bernardo, Santiago Sánchez-Pacheco, and Ro	obert Murphy
University of Toronto and Royal Ontario Museum	

Conference and Workshop Participation

*Virtual event	
 Participation with a role. Technical Moderator ISMB, ISCB, Madison, WI., USA Symposium Organizer (co-organized with Md Rejuan Haque) Evolution Meeting, Cleveland, OH., USA Symposium title: Towards the network of life: Phylogenetic networks as a tool to understand complex evolutionary histories. Chair/Moderator in multiple Faux-Live Sessions, Evolution Meeting* 	2022 2022 2022
Participant. • Workshop in Mathematical and Computational Biology* • Annual Conference of the Korean Association of Biological Sciences, South Korea* • Workshop in Mathematical and Computational Biology*	2022 2021 2021
 SMRT Leiden 2021 – Young Investigator Virtual Conference, hosted by Pacific Biosciences and Leiden University Medical Center* Virtual Asilomar 2021, American Society of Naturalists* Future of Systematics Workshop at SSB Standalone Meeting, funded by NSF, hosted by Dr. Emily Sessa, University of Florida, Gainesville, FL., USA 	2021 2021 2020
 Evolution Meeting, Providence, RI., USA Midwest Phylogenetics Workshop, Itasca Biological Station, funded by NSF, hosted by Dr. Emma Goldberg, University of Minnesota, MN., USA Annual Conference of the Korean Association of Biological Sciences, Yong-In, South Korea 	2019 2019 2017
 Society for Conservation Biology (SCB) Asia Section Workshop on Conservation policy, funded by SCB, Bangkok, Thailand World Congress of Herpetology, Hangzhou and Tonglu, China Phylogenomics Symposium and Software School hosted by Dr. Tandy Warnow at SSB Standalone Meeting, University of Michigan, Ann Arbor, MI., USA 	2017 2016 2015
Diversity and Inclusion Activities	
 Participation with a role. Moderator International Mixer, Evolution Meeting* Moderator Decolonizing Evolution Workshop, Evolution Meeting* Moderator Strategies for Responding to Harassment and Bullying: Improving Workplace Climate, Evolution Community Resources for Early Career Researchers 2020 (ECR²)* Moderator Fireside ECR Chat, ECR²* 	2021 2021 2020 2020
 Participant. International Mixer, Evolution Meeting, Cleveland, OH., USA Transformative Justice within Academia: A radical institutional reflection, The Centre for Criminology and Sociolegal Studies Graduate Student Conference, University of Toronto* 	2022

Grants and Awards

All in US Dollars unless stated. †Canadian Dollar; ‡Approximately converted from South Korean Won Total amount of funding received up to date \approx US\$50,000

 Highlights. The Presidential Fellowship, The Ohio State University [\$36,500] SSB Ad Hoc Funds [\$3,000] SSE Graduate Research Excellence Grant – R.C. Lewontin Award competition [\$1,800] University of Toronto Graduate Fellowship [Stipend with tuition waived for 18 months] 2 E.M. Walker Graduate Award in Biodiversity, University of Toronto [†\$2,615] 	2022 2021 2020 2013–2015 2014
Other grants. • EvolCompGen Best Poster Award, ISMB, ISCB [\$100] • EEOB Graduate Development Grant, The Ohio State University [Funded]	2022
 EEOB Graduate Evolution & Ecology Student Organization (GEES) Career Development Grant, The Ohio State University [\$200] Council of Graduate Students (CGS) Career Development Grant, 	2022
The Ohio State University [\$350] • EEOB Darwin Talk Society Membership Grant, The Ohio State University [\$70]	2022 2021
 EEOB Graduate Development Grant, The Ohio State University [Funded] Funding for participation in Future of Systematics Workshop, 	2021
SSB Standalone Meeting (NSF) [\$500]	2020
• Funding for participation in Midwest Phylogenetics Workshop (NSF) [Funded]	2019
EEOB Graduate Development Grant, The Ohio State University [Funded]Major contribution to successful grant application to National Research Foundation of K	2019 (orea
PI: Dr. Yikweon Jang, Ewha Womans University, Seoul, South Korea [‡\$500,000]	2017
• University of Toronto Undergraduate Bursaries [$\dagger \approx $3,000$]	2010–2012
Outreach and Service	
 Awardee Graduate School Awards Ceremony, The Ohio State University 	2022
 Invited Panelist EEOB 8894: EEOB Graduate Seminar, The Ohio State University Invited Guest Translational Data Analytics Institute Kickoff Event, 	2020
College of Arts and Sciences, The Ohio State University	2020
 Museum of Biological Diversity's Open House, The Ohio State University 	2019
	2020–2021
·	2019–2020
o Communication Officer Society for Conservation Biology (SCB), South Korean Chapter 2	2016–2017
 Kuang Si Butterfly Park, Ban Thapene, Luang Prabang District, Laos 	2016, 2022

Peer-Review Activities

- Systematic Biology
- Molecular Biology and Evolution
- Genome Biology and Evolution
- Scientific Reports
- Molecular Phylogenetics and Evolution
- Infection, Genetics and Evolution

Professional Memberships

- Society of Systematic Biologists
- Society for Study of Evolution
- o International Society for Computational Biology

Skills

Programming UNIX, R, Python, Julia, HTML, JavaScript
Typesetting Languages English (Fluent), Korean (Native), Japanese (Intermediate)

References

[1] Prof. Dr. Laura S. Kubatko (Ph.D. advisor)

Department of Statistics and Department of Evolution, Ecology, and Organismal Biology The Ohio State University 219 Cockins Hall, Columbus, OH., 43210, USA lkubatko@stat.osu.edu +1 614-247-8846

[2] Prof. Dr. Robert W. Murphy (M.Sc. advisor)

Department of Natural History-Royal Ontario Museum And Professor of Ecology and Evolutionary Biology University of Toronto 100 Queen's Park Toronto, ON., M3S 2C6, Canada bob.murphy@utoronto.ca +1 416-586-8099

[3] Prof. Dr. Deborah A. Mclennan (B.Sc. advisor)

Department of Ecology and Evolutionary Biology University of Toronto, 25 Harbord St. Toronto, ON., M5S 3G5, Canada deborah.mclennan@utoronto.ca +1 416-978-8538

[4] Prof. Dr. Bryan C. Carstens

Department of Evolution, Ecology, and Organismal Biology And Tetrapod Division, Museum of Biological Diversity The Ohio State University, 318 W. 12th Avenue, Columbus, OH., 43210, USA carstens.12@osu.edu +1 614-292-6587

[5] Prof. Dr. Jeremy M. Brown

Department of Biological Sciences Louisiana State University A243 Life Science Annex, Baton Rouge, LA., 70803, USA jembrown@lsu.edu +1 225-578-1745