#### **KEY SKILLS & EXPERIENCE**

# Joha Park, Ph.D.

- RNA Biology, Spatial Omics, Genomics, Gene Editing
- Multi-omics: (sc)RNA-seq, ATAC-seq, WGS/WES, eQTLs
- AI/ML: Biological LLM, Variant effect prediction, Genotoxicity
- Image processing: 3D confocal/light-sheet microscopy
- Tools: PyTorch, Scikit-learn/image, Nextflow, Docker, AWS

Intellia Therapeutics, Inc. 40 Erie St, Cambridge MA 02139, United States email: johapark@gmail.com webpage: johapark.github.io

Computational biologist with expertise in RNA biology, spatial omics, genomics, and gene editing. Skilled in developing computational frameworks for analyzing large scale sequencing and imaging data by employing advanced bioinformatics tools and AI/ML approaches. Demonstrated track record of conducting high-quality research, which has been published in esteemed journals.

#### **CURRENT POSITION**

#### 03/2023 - present Senior Computational Scientist

Intellia Therapeutics

**Computational Sciences** 

**Genomic Innovation** 

Intellia Therapeutics, Inc., Cambridge, MA, US (Remote)

- Develop platforms for CRISPR-Cas9 off-target discovery and build AI/ML prediction models
- Predict genetic variant effect for de-risking genotoxicity of gene therapy
- Build scalable, reproducible, and streamlined computational analysis pipelines
- Analyze various custom DNA/RNA sequencing datasets

#### PROFESSIONAL EXPERIENCE

#### 01/2020 - 03/2023 Postdoctoral Associate/Fellow

Massachusetts Institute of

Technology

3 yrs 3 mos The Picower Institute for Learning and Memory Institute for Medical Engineering and Science

Department of Chemical Engineering

Massachusetts Institute of Technology, Cambridge, MA, US

### Roles

- Develop hydrogel-based tissue clearing and expansion methods for spatial omics
- Build a scalable computational framework for 3D light-sheet microscopy
- Generate a single-cell mouse brain reference atlas

#### 03/2019 - 12/2019 Postdoctoral Researcher

Seoul National University

10 months Center for RNA Research

Institute for Basic Science

Seoul National University, Seoul, Korea

#### Roles

- Lead projects on mRNA poly(A)-tail length regulations
- Perform conventional and custom NGS library prep & data analysis
- Carry out wet and dry lab experiments for RNA biology

#### **EDUCATION AND TRAINING**

2012 – 2019 Ph.D. in Biological Sciences (RNA Biology & Bioinformatics)

RNA Biology Lab (Advisor: V. Narry Kim)

Dissertation: "Poly(A) length regulation: deadenylases and the poly(A) barricade"

2015 – 2018 Expert Research Personnel (Compulsory Military Service)

Republic of Korea Army

2008 – 2012 B.S. in Biological Sciences (cum laude)

Seoul National University

#### **PUBLICATIONS**

\*: co-first author, #: co-corresponding author

- 1 J. Park\*, M. Kim\*, H. Yi\*, K. Baeg\*, Y. Choi, Y.-s. Lee, J. Lim, V. N. Kim (2023) "Short poly(A) tails are protected from deadenylation by the LARP1-PABP complex" Nat. Struct. Mol. Biol., 30(3):330–338
- 2 J. Park\*, S. Khan\*, D. H. Yun\*, T. Ku, K. L. Villa, J. E. Lee, Q. Zhang, G. Feng, J. Park, E. Nedivi#, K. Chung# (2021) "Epitope-preserving magnified analysis of proteome (eMAP)" Sci. Adv. 7(16):eabf6589
- 3 Y. Kim\*#, J. Park\*, S. Kim\*, M. Kim, M.-G. Kang, C. Kwak, M. Kang, B. Kim, H.-W. Rhee, and V. N. Kim# (2018) "PKR senses nuclear and mitochondrial signals by interacting with endogenous double-stranded RNAs" *Mol. Cell*, 71(6):1051–1063.e6
  - Selected as "Research Highlights" in Nat. Chem. Biol., 14(11):989
- 4 H. Yi\*, J. Park\*, M. Ha, J. Lim, H. Chang and V. N. Kim (2018) "PABP cooperates with the CCR4-NOT complex to promote mRNA deadenylation and block precocious decay" *Mol. Cell*, 70(6):1081–1088.e5
  - Previewed in *Mol. Cell*, 70(6):987–988
- 5 T. A. Nguyen\*#, J. Park\*, T. L. Dang, Y.-G Choi, and V. N. Kim# (2018) "Microprocessor depends on hemin to recognize the apical loop of primary microRNA" Nucleic Acids Res., 46(11):5726–5736
- 6 K. T. You, **J. Park**, and V. N. Kim (2015) "Role of the small subunit processome in the maintenance of pluripotent stem cells" *Genes Dev.*, 29(19):2004–2009
- 7 T. A. Nguyen, M. H. Jo, Y.-G. Choi, J. Park, S. C. Kwon, S. Hohng, V. N. Kim# and J.-S. Woo# (2015) "Functional anatomy of the human Microprocessor" *Cell*, 161(6):1374–1387
- 8 Y.-K. Kim, G. Wee, J. Park, J. Kim, D. Baek, J.-S. Kim#, and V. N. Kim# (2013) "TALEN-based knockout library for human microRNAs" Nat. Struct. Mol. Biol., 20(12):1458–1464

Links: Google Scholar, PubMed

#### **HONORS AND AWARDS**

- 2022 2023 Picower Postdoctoral Fellows, The Picower Institute for Learning and Memory, MIT
  - 2021 AKN Outstanding Research Awards, Association of Korean Neuroscientists
  - 2019 Conference Scholarship, Keystone Symposia: Small Regulatory RNAs
  - 2019 Best Dissertation Awards, School of Biological Sciences, SNU

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2008 – 2012 Presidential Science Scholarship, Korea Science and Engineering Foundation

2014 Keystone Symposia: RNA Silencing — poster

#### **INTERNATIONAL CONFERENCES**

2019	Cell Symposia: Regulatory RNAs — <b>short talk</b> "Poly(A) barricade as a rate-limiting factor that shapes poly(A) tail length"
2018	${\it CSHL Meeting: Regulatory \& Noncoding RNAs-short talk} \\ {\it ``PKR senses nuclear and mitochondrial signals by interacting with endogenous dsRNAs''} \\$
2017	CSHL Meeting: Eukaryotic mRNA Processing — <b>short talk</b> "Genome-wide evaluation of the role and specificity of deadenylases"
2016	EMBL Symposium: The Complex Life of mRNA — poster
2016	IMBA: 11th Microsymposium on Small RNAs — poster

#### **INVITED TALKS**

08/2023	Special Seminar – Graduate School of Medical Science and Engineering, KAIST
11/2022	Special Seminar – Yonsei University College of Medicine
08/2022	Young Global Leader Symposium – Dept. of Chemical and Biomolecular Engineering, KAIST
07/2022	Department Seminar – Department of Pharmacology, SNU College of Medicine
05/2022	Young Investigator Seminar Series – Department of Biomedical Engineering, UNIST

#### **TEACHING EXPERIENCE**

2022	Kaufman Teaching Certificate Program, Massachusetts Institute of Technology
2014	Teaching Assistant, Advanced RNA Biology Lab, School of Biological Sciences, SNU
2013	Teaching Assistant, Biology Lab 2, School of Biological Sciences, SNU
2012	Teaching Assistant, Biology Lab 1, School of Biological Sciences, SNU
2011	Tutor, Basic Course for Biology, School of Biological Sciences, SNU

# **TECHNICAL EXPERTISE**

## **Computational biology**

- General skills
  - Unix/Linux, AWS cloud computing environment
  - Programming languages: Python, R, Rust
  - Python scientific computing libraries: NumPy, SciPy, Pandas, etc.
  - Python machine learning frameworks: Scikit-learn, PyTorch
  - Building a reproducible analysis pipeline: Nextflow, Snakemake, Docker, Singularity
  - Efficient data visualization: Matplotlib, Seaborn, Plotly

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- Next-generation sequencing (NGS)
  - (sc)RNA-seq, ATAC-seq, Amp-seq, CLIP-seq, TAIL-seq, etc.
  - Statistical analysis for genome-wide, transcriptomic studies
  - Tools: Samtools, Picard, Scanpy, Seurat, STAR, Bedtools, etc.
- Image processing
  - Concurrent programming for large-scale images
  - 2D/3D rigid and non-rigid biomedical image registration
  - Convolutional Neural Networks
  - Tools: PyTorch, Zarr, Dask, Scikit-image, OpenCV, ImageJ/Fiji, Imaris, etc.

#### **Wet experiments**

- Routine experiments
  - Molecular cloning: Gibson, Golden Gate assembly
  - Mammalian, bacterial cell culture
  - DNA and siRNA transfection
  - Quantitative RT-PCR
  - Immunoprecipitation
  - Western blotting
- RNA biology
- Custom sequencing library preparation
- Custom high-resolution poly(A)-tail length assay
- Tet-On inducible reporter cell line generation
- RNA co-immunoprecipitation
- Lentiviral transduction
- In vitro transcription
- Northern blotting
- Luciferase reporter assay
- Tissue clearing and expansion
  - SHIELD clearing
  - Expansion microscopy: eMAP