

# Kejun “Albert” Ying

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*Studying aging at the intersection of biology and AI*

## Education

### Harvard University

Cambridge, MA

*Ph.D., Biological Science in Public Health*

2019 – Expected May 2025

- Advisor: Dr. Vadim Gladyshev, Harvard Medical School, Brigham and Women’s Hospital
- Dissertation Advisory Committee: Dr. Brendan Manning, Dr. David Sinclair, Dr. Shamil Sunyaev
- Focused on understanding the mechanism of aging through multi-omic modeling & causal inference

### Harvard University

Cambridge, MA

*M.S., Computational Science Engineering*

2022 – 2024

- Secondary field during Ph.D. study

### University of California, Berkeley

Berkeley, CA

*Visiting Student, Integrative Biology*

2017 – 2018

### Sun Yat-Sen University

Guangzhou, China

*B.S., Life Science*

2015 – 2019

- Thesis: Screening for the Interactome of hTERC based on Molecular Fluorescence Complementation System in Living Cells
- Yat-Sen Honor School Program (Top 0.5%)
- National college admissions exam (Top 0.6%)

## Grants

**Using causal aging biomarkers and protein design to develop novel anti-aging interventions** NIH/NIA

F99/Koo, *Transition to Aging Research for Predoctoral Students*

2024 – 2028

- Award Document Number: FAGO88431A (PI)
- Received a *perfect* Impact Score of **10**

## Publications

<sup>†</sup> Corresponding author; <sup>\*</sup> Co-first author; <sup>+</sup> Contributed as consortium author

**Ying, K.**, Castro, J. P., Shindyapina, A. V., Tyshkovskiy, A., Moqri, M., Goeminne, L. J. E., Milman, S., Zhang, Z. D., Barzilai, N., & Gladyshev, V. N. (2024). Depletion of loss-of-function germline mutations in centenarians reveals longevity genes. **Nature Communications**, in press.

Lyu, YX.<sup>\*</sup>, Fu, Q.<sup>\*</sup>, Wilczok, D.<sup>\*</sup>, **Ying, K.**<sup>\*</sup>, King, A., ..., Bakula, D. (2024). Longevity biotechnology: Bridging AI, biomarkers, geroscience and clinical applications for healthy longevity. **Aging**, in press.

**Biomarkers of Aging Consortium**<sup>+</sup>, Herzog, C. M. S., Goeminne, L. J. E., Poganik, J. R., Barzilai, N., Belsky, D. W., Betts-LaCroix, J., Chen, B. H., Chen, M., Cohen, A. A., Cummings, S. R., Fedichev, P. O., Ferrucci, L., Fleming, A., Fortney, K., Furman, D., Gorbunova, V., Higgins-Chen, A., Hood, L., Horvath, S., ... Gladyshev,

V. N. (2024). Challenges and recommendations for the translation of biomarkers of aging. **Nature Aging**, 1–12. <https://doi.org/10.1038/s43587-024-00683-3>

Castro, J. P., Shindyapina, A. V., Barbieri, A., **Ying, K.**, Strelkova, O. S., Paulo, J. A., Tyshkovskiy, A., Meinl, R., Kerepesi, C., Petrashen, A. P., Mariotti, M., Meer, M. V., Hu, Y., Karamyshev, A., Losyev, G., Gallhardo, M., Logarinho, E., Indzhukulian, A. A., Gygi, S. P., Sedivy, J. M., Manis, J. P., & Gladyshev, V. N. (2024). Age-associated clonal B cells drive B cell lymphoma in mice. **Nature Aging**, 4(8), 1–15. <https://doi.org/10.1038/s43587-024-00671-7>

Moqri, M., Cipriano, A., Simpson, D. J., Rasouli, S., Murty, T., de Jong, T. A., Nachun, D., de Sena Brandine, G., **Ying, K.**, Tarkhov, A., Aberg, K. A., van den Oord, E., Zhou, W., Smith, A., Mackall, C., Gladyshev, V. N., Horvath, S., Snyder, M. P., & Sebastiano, V. (2024). PRC2-AgeIndex as a universal biomarker of aging and rejuvenation. **Nature Communications**, 15(1), 5956. <https://doi.org/10.1038/s41467-024-50098-2>

Tarkhov, A. E., Lindstrom-Vautrin, T., Zhang, S., **Ying, K.**, Moqri, M., Zhang, B., Tyshkovskiy, A., Levy, O., & Gladyshev, V. N. (2024). Nature of epigenetic aging from a single-cell perspective. **Nature Aging**, 1–17. <https://doi.org/10.1038/s43587-023-00555-2>

Moqri, M., Herzog, C., Poganik, J. R., **Ying, K.**, Justice, J. N., Belsky, D. W., Higgins-Chen, A. T., Chen, B. H., Cohen, A. A., Fuellen, G., Hägg, S., Marioni, R. E., Widschwendter, M., Fortney, K., Fedichev, P. O., Zhavoronkov, A., Barzilai, N., Lasky-Su, J., Kiel, D. P., ... Ferrucci, L. (2024). Validation of biomarkers of aging. **Nature Medicine**, 1–13. <https://doi.org/10.1038/s41591-023-02784-9>

Griffin, P. T., Kane, A. E., Trapp, A., Li, J., Arnold, M., Poganik, J. R., Conway, R. J., McNamara, M. S., Meer, M. V., Hoffman, N., Amorim, J. A., Tian, X., MacArthur, M. R., Mitchell, S. J., Mueller, A. L., Carmody, C., Vera, D. L., Kerepesi, C., **Ying, K.**, ... Sinclair, D. A. (2024). TIME-seq reduces time and cost of DNA methylation measurement for epigenetic clock construction. **Nature Aging**, 1–14. <https://doi.org/10.1038/s43587-023-00555-2>

**Ying, K.**, Liu, H., Tarkhov, A. E., Sadler, M. C., Lu, A. T., Moqri, M., Horvath, S., Kutalik, Z., Shen, X., & Gladyshev, V. N. (2024). Causality-enriched epigenetic age uncouples damage and adaptation. **Nature Aging (Featured on the February Cover)**, 1–16. <https://doi.org/10.1038/s43587-023-00557-0>

Moqri, M., Herzog, C., Poganik, J. R., **Biomarkers of Aging Consortium**<sup>+</sup>, Justice, J., Belsky, D. W., Higgins-Chen, A., Moskalev, A., Fuellen, G., Cohen, A. A., Bautmans, I., Widschwendter, M., Ding, J., Fleming, A., Mannick, J., Han, J.-D. J., Zhavoronkov, A., Barzilai, N., Kaeblerlein, M., ... Gladyshev, V. N. (2023). Biomarkers of aging for the identification and evaluation of longevity interventions. **Cell**, 186(18), 3758–3775. <https://doi.org/10.1016/j.cell.2023.08.003>

Lieberman, N., Rothi, M. H., Gerashchenko, M. V., Zorbas, C., Boulias, K., MacWhinnie, F. G., **Ying, A. K.**, Flood Taylor, A., Al Haddad, J., Shibuya, H., Roach, L., Dong, A., Dellacona, S., Lafontaine, D. L. J., Gladyshev, V. N., & Greer, E. L. (2023). 18S rRNA methyltransferases DIMT1 and BUD23 drive intergenerational hormesis. **Molecular Cell**, 83(18), 3268–3282.e7. <https://doi.org/10.1016/j.molcel.2023.08.014>

Bitto, A., Grillo, A. S., Ito, T. K., Stanaway, I. B., Nguyen, B. M. G., **Ying, K.**, Tung, H., Smith, K., Tran, N., Velikanje, G., Urfer, S. R., Snyder, J. M., Barton, J., Sharma, A., Kayser, E.-B., Wang, L., Smith, D. L., Thompson, J. W., DuBois, L., ... Kaeblerlein, M. (2023). Acarbose suppresses symptoms of mitochondrial disease in a mouse model of Leigh syndrome. **Nature Metabolism**, 5(6), 955–967. <https://doi.org/10.1038/s42255-023-00815-w>

Emmrich, S., Trapp, A., Tolibzoda Zakusilo, F., Straight, M. E., **Ying, A. K.**, Tyshkovskiy, A., Mariotti, M., Gray, S., Zhang, Z., Drage, M. G., Takasugi, M., Klusmann, J.-H., Gladyshev, V. N., Seluanov, A., & Gorbunova, V. (2022). Characterization of naked mole-rat hematopoiesis reveals unique stem and progenitor cell patterns and neotenic traits. **The EMBO Journal**, 41(15), e109694. <https://doi.org/10.15252/embj.2021109694>

Yang, Z., Macdonald-Dunlop, E., Chen, J., Zhai, R., Li, T., Richmond, A., Klarić, L., Pirastu, N., Ning, Z., Zheng, C., Wang, Y., Huang, T., He, Y., Guo, H., **Ying, K.**, Gustafsson, S., Prins, B., Ramisch, A., Dermitzakis, E. T., ... Shen, X. (2022). Genetic Landscape of the ACE2 Coronavirus Receptor. **Circulation**, 145(18), 1398–1411. <https://doi.org/10.1161/CIRCULATIONAHA.121.057888>

**Ying, K.**, Zhai, R., Pyrkov, T. V., Shindyapina, A. V., Mariotti, M., Fedichev, P. O., Shen, X., & Gladyshev, V. N. (2021). Genetic and phenotypic analysis of the causal relationship between aging and COVID-19. **Communications Medicine**, 1(1), 35. <https://doi.org/10.1038/s43856-021-00033-z>

Li, T., Ning, Z., Yang, Z., Zhai, R., Zheng, C., Xu, W., Wang, Y., **Ying, K.**, Chen, Y., & Shen, X. (2021). Total genetic contribution assessment across the human genome. **Nature Communications**, 12(1), 2845. <https://doi.org/10.1038/s41467-021-23124-w>

Zhu, J., Xu, M., Liu, Y., Zhuang, L., **Ying, K.**, Liu, F., Liu, D., Ma, W., & Songyang, Z. (2019). Phosphorylation of PLIN3 by AMPK promotes dispersion of lipid droplets during starvation. **Protein & Cell**, 10(5), 382–387. <https://doi.org/10.1007/s13238-018-0593-9>

## Preprints

Galkin, F., Naumov, V., Pushkov, S., Sidorenko, D., Urban, A., Zagirova, D., Alawi, K. M., Aliper, A., Gumerov, R., Kalashnikov, A., Mukba, S., Pogorelskaya, A., Ren, F., Shneyderman, A., Tang, Q., Xiao, D., Tyshkovskiy, A., **Ying, K.**, Gladyshev, V. N., & Zhavoronkov, A. (2024). Precious3GPT: Multimodal Multi-Species Multi-Omics Multi-Tissue Transformer for Aging Research and Drug Discovery. **bioRxiv**. <https://doi.org/10.1101/2024.07.25.605062>

**Ying, K.**, Paulson, S., Eames, A., Tyshkovskiy, A., Li, S., Perez-Guevara, M., Emamifar, M., Martínez, M. C., Kwon, D., Kosheleva, A., Snyder, M. P., Gobel, D., Herzog, C., Poganik, J. R., Biomarker of Aging Consortium, Moqri, M., & Gladyshev, V. N. (2024). *A Unified Framework for Systematic Curation and Evaluation of Aging Biomarkers*. **bioRxiv**. <https://doi.org/10.1101/2023.12.02.569722>

Tyshkovskiy, A., Kholdina, D., **Ying, K.**, Davitadze, M., Molière, A., Tongu, Y., Kasahara, T., Kats, L. M., Vladimirova, A., Moldakozhayev, A., Liu, H., Zhang, B., Khasanova, U., Moqri, M., Van Raamsdonk, J. M., Harrison, D. E., Strong, R., Abe, T., Dmitriev, S. E., & Gladyshev, V. N. (2024). Transcriptomic Hallmarks of Mortality Reveal Universal and Specific Mechanisms of Aging, Chronic Disease, and Rejuvenation. **bioRxiv**. <https://doi.org/10.1101/2024.07.04.601982>

Rothi, M. H., Sarkar, G. C., Al Haddad, J., Mitchell, W., **Ying, K.**, Pohl, N., Sotomayor-Mena, R. G., Natale, J., Dellacono, S., Gladyshev, V. N., & Greer, E. L. (2024). The 18S rRNA Methyltransferase DIMT-1 Regulates Lifespan in the Germline Later in Life. **bioRxiv**. <https://doi.org/10.1101/2024.05.15.570935>

Moqri, M., Poganik, J. R., Herzog, C., **Ying, K.**, Chen, Q., Emamifar, M., Tyshkovskiy, A., Eames, A. W., Mur, J., Matei-Dediu, B., Goeminne, L., Mitchell, W., McCartney, D. L., Marioni, R. L., Lasky-Su, J. A., Snyder, M., & Gladyshev, V. N. (2024). Integrative epigenetics and transcriptomics identify aging genes in human blood. **bioRxiv**. <https://doi.org/10.1101/2024.05.30.596713>

Goeminne, L. J. E., Eames, A., Tyshkovskiy, A., Argentieri, M. A., **Ying, K.**, Moqri, M., & Gladyshev, V. N. (2024). Plasma-based organ-specific aging and mortality models unveil diseases as accelerated aging of organismal systems. **medRxiv**. <https://doi.org/10.1101/2024.04.08.24305469>

**Ying, K.**, Tyshkovskiy, A., Trapp, A., Liu, H., Moqri, M., Kerepesi, C., & Gladyshev, V. N. (2023). *ClockBase : A comprehensive platform for biological age profiling in human and mouse*. **bioRxiv**. <https://doi.org/10.1101/2023.02.28.530532>

Zhang, B., Tarkhov, A. E., Ratzan, W., **Ying, K.**, Moqri, M., Poganik, J. R., Barre, B., Trapp, A., Zoller, J. A., Haghani, A., Horvath, S., Peshkin, L., & Gladyshev, V. N. (2022). *Epigenetic profiling and incidence of disrupted development point to gastrulation as aging ground zero in *Xenopus laevis**. **bioRxiv**. <https://doi.org/10.1101/2022.08.02.502559>

Castro, J. P., Shindyapina, A. V., Barbieri, A., **Ying, K.**, Strelkova, O. S., Paulo, J. A., Tyshkovskiy, A., Meinl, R., Kerepesi, C., Petrashen, A. P., Mariotti, M., Meer, M., Hu, Y., Karamyshev, A., Losyev, G., Indzhikulian, A. A., Gygi, S. P., Sedivy, J. M., Manis, J. P., & Gladyshev, V. N. (2021). *Integrative analyses uncover mechanisms by which aging drives B cell lymphoma*. **bioRxiv**. <https://doi.org/10.1101/2021.02.23.432500>

## Patents

V. N. Gladyshev, **K. Ying**, “High-dimensional measurement of biological age” (2024). *Provisional Patent Application*

V. N. Gladyshev, **K. Ying**, “Mapping CpG sites to quantify aging traits” (2024). *WO2024039905A2*

## Software and Database

**Biolearn** (2024) <https://bio-learn.github.io/>

**ClockBase** (2023) <https://www.clockbase.org/>

## Presentations

### ORAL PRESENTATIONS

**Biomarkers of Aging Symposium** Boston, MA  
*Standardization of aging biomarkers and BoA challenge* 2024

**Harvard GRIP Presentations** Boston, MA  
*Causal Aging Biomarker empowers Unbiased Anti-Aging Therapy Screening* 2024

**4th TimePie Longevity Forum** Shanghai, China  
*Causal Aging Biomarker as a Tool for Unbiased Anti-Aging Therapy Screening* 2023

**Global Congress on Aesthetic and Anti-Aging (GCAA2023)** Singapore  
*Causal Aging Biomarker as a Tool for Unbiased Anti-Aging Therapy Screening* 2023

**10th Aging Research and Drug Discovery conference (ARDD2023)** Copenhagen, Denmark  
*Causal Epigenetic Age Uncouples Damage and Adaptation* 2023

**AGE 2023 51st Annual Meeting** Oklahoma City, OK  
*Causal Epigenetic Age Uncouples Damage and Adaptation* 2023

**Broad Institute MPG Retreat** Cambridge, MA  
*Causal Epigenetic Age Uncouples Damage and Adaptation* 2023

**Harvard GRIP Presentations** Boston, MA  
*Causal Epigenetic Age Uncouples Damage and Adaptation* 2022

**Targeting Metabesity 2022**, ‘Honorable Mention’ Virtual Conference  
*Causal Epigenetic Age Uncouples Damage and Adaptation* 2022

**GSA 2021 Annual Scientific Meeting** Virtual Conference  
*Genetic and phenotypic evidence for causal relationships between aging and COVID-19* 2021

## POSTER PRESENTATIONS

### CHSL Mechanisms of Aging Meeting

Cold Spring Harbor, NY

- A unified framework for systematic curation and evaluation of aging biomarkers* 2024  
*Causal epigenetic age and transcriptomic clock facilitate systemic anti-aging therapy screening* 2024  
*Depletion of loss-of-function germline mutations in centenarians reveals novel longevity genes* 2024  
*High-dimensional representations of biological aging in functional modules* 2024

### Biomarker of Aging Symposium

Novato, CA

- Causal Aging Biomarker as a Tool for Unbiased Anti-Aging Therapy Screening* 2023

### Gordon Research Conference, Systems Aging

Maine, MA

- Causal Epigenetic Age Uncouples Damage and Adaptation* 2022

## INVITED TALKS

### BioAge Seminar, hosted by Dr. Robert Hughes & Dr. Paul Timmers

Boston, MA

- Ageome: Biological age with higher-dimensionality* 2024

### MRC Integrative Epidemiology Unit Seminar

Bristol, UK

- Epigenetic Clocks and Mendelian Randomization* 2024

### NIA EL Projects Joint Meeting, National Institute on Aging

Online Webinar

- Aging Clocks* 2024

### Biomarkers of Aging Challenge, Foresight Institute

Online Webinar

- Update Webinar with Foresight* 2024

### Everything Epigenetics, podcast hosted by Hannah Went

Online Podcast

- Causal Epigenetic Age Uncouples Damage and Adaptation* 2024

### Chinese University of Hong Kong, hosted by Dr. Xin Wang

Hong Kong, China

- Causal Aging Biomarker as a Tool for Systemic Anti-Aging Therapy Screening* 2024

### Everything Epigenetics, podcast hosted by Hannah Went

Online Podcast

- Causal Epigenetic Age Uncouples Damage and Adaptation* 2023

### Chinese University of Hong Kong, hosted by Dr. Xin Wang

Hong Kong, China

- Causal Aging Biomarker as a Tool for Systemic Anti-Aging Therapy Screening* 2023

### Peking University, hosted by Dr. Jingdong Han

Beijing, China

- Causal Aging Biomarker and ClockBase* 2023

### Chinese Academy of Sciences, hosted by Dr. Xuming Zhou

Beijing, China

- Causal Epigenetic Age Uncouples Damage and Adaptation* 2022

### Foresight Institute, hosted by Allison Duettmann

Online Seminar

- Genetic Variation, Aging & Relationship to COVID-19 | Joris Deelen, Albert Ying* 2020

## Research Experience

### Harvard Medical School, Brigham and Women's Hospital

Boston, MA

**Biological Aging** Graduate Researcher, Vadim Gladyshev's Lab

2020 – Present

<b>Harvard Medical School, Boston Children's Hospital</b> <b>RNA Modifications</b> <i>Rotation Student, Eric Greer's Lab</i>	Boston, MA 2020
<b>Harvard Medical School</b> <b>IPSC Reprogramming &amp; DNA methylation</b> <i>Rotation Student, David Sinclair's Lab</i>	Boston, MA 2019L
<b>Harvard T. H. Chan School of Public Health</b> <b>mTORC1</b> <i>Rotation Student, Brendan Manning's Lab</i>	Boston, MA 2019
<b>Sun Yat-Sen University</b> <b>Telomere &amp; Telomerase</b> <i>Undergraduate Researcher, Zhou Songyang's Lab</i>	Guangzhou, China 2018 – 2019
<b>University of Edinburgh</b> <b>Population genetics</b> <i>Undergraduate Researcher, Xia Shen's Lab</i>	Edinburgh, UK 2018
<b>University of Washington</b> <b>Acarbose &amp; Rapamycin</b> <i>Undergraduate Researcher, Matt Kaeberlein's Lab</i>	Seattle, WA 2018
<b>Buck Institute for Research on Aging</b> <b>Senolytics</b> <i>Undergraduate Researcher, Judith Campisi's Lab</i>	Novato, CA 2018
<b>University of California, Berkeley</b> <b>SIRT7</b> <i>Undergraduate Researcher, Danica Chen's Lab</i>	Berkeley, CA 2017
<b>Sun Yat-Sen University</b> <b>Telomere &amp; DNA Methylation</b> <i>Undergraduate Researcher, Yikang Rong's Lab</i>	Guangzhou, China 2015 – 2017

## Honors

<b>Best Poster Award</b> , Inaugural Biomarker of Aging Symposium	2023
<b>Best Poster Award</b> , Gordon Research Conference, Systems Aging	2022
<b>Hackathon Winner</b> , Longevity Hackathon, VitaDAO	2021
<b>Yan-Sen Honor School Program</b> , Sun Yat-Sen University	2016 – 2019
<b>Yan-Sen Scholarship</b> , Sun Yat-Sen University	2016 – 2019

## Professional Experience

### SERVICE & LEADERSHIP

<b>President</b> , Harvard Interdisciplinary Discussion on Disease and Health	2024 – Present
<b>Organizer</b> , Biomarker of Aging Challenge	2024 – Present
<b>Organizing Committee Member</b> , Biomarker of Aging Symposium 2024	2024
<b>Organizing Committee Member</b> , Biomarker of Aging Symposium 2023	2023

### TEACHING & MENTORING

<b>Mentor</b> , Yuanpei Young Scholars Program	2023 – 2024
<b>Instructor</b> , Harvard Public Health Symposium For Young Generation	2023

## STUDENTS SUPERVISED

**Predoctoral Students:** Ali Doga Yucel, Siyuan Li, Hanna Liu, Han Weng

## JOURNALS REVIEWED

*Nature Aging, Nature Communications, BMC Nephrology, Lipids in Health and Disease, Clinical Proteomics, Evidence-Based Complementary and Alternative Medicine, Scientific Report*

## References

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**Dr. Vadim Gladyshev**, Dissertation Advisor vgladyshev@bwh.harvard.edu  
Professor of Medicine, Harvard Medical School

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Professor of Human Genetics, UCLA

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Professor of Genetics, Harvard Medical School

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Professor of Pathology, University of Washington