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 - Expected May 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/K00, Transition to Aging Research for Predoctoral Students 2024 - 2028

- Award Document Number: FAG088431A (PI)
- Received a *perfect* Impact Score of **10**

Publications

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., Galhardo, M., Logarinho, E., Indzhykulian, 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**, in press.

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**, in press.

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, I–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

Liberman, 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., ... Kaeberlein, 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**, *t*(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

Bitto, A., Tung, H., Ying, K., Smith, D. L., Kayser, E.-B., Morgan, P. G., Sedensky, M. M., & Kaeberlein, M. (2019). AGING AND MITOCHONDRIAL DISEASE: SHARED MECHANISMS AND THERAPIES? Innovation in Aging, 3(Supplement_1), S395–S395. https://doi.org/10.1093/geroni/igz038.1459

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

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., Paulson, S., Perez-Guevara, M., Emamifar, M., Martinez, M. C., Kwon, D., Poganik, J. R., Moqri, M., & Gladyshev, V. N. (2023). *Biolearn, an open-source library for biomarkers of aging.* bioRxiv. https://doi.org/10.1101/2023.12.02.569722

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., Indzhykulian, 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, Mapping CpG sites to quantify aging traits (2024). WO2024039905A2

Software and Database

Biolearn (2024)

https://bio-learn.github.io/

ClockBase (2023)

<www.clockbase.com>

Presentations

ORAL PRESENTATIONS

Harvard GRIP Presentations

Boston, MA

Causal Aging Biomarker enpowers 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 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 **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 2.02.2 Virtual Conference GSA 2021 Annual Scientific Meeting Genetic and phenotypic evidence for causal relationships between aging and COVID-19 2.O2.I Poster Presentations 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 2.02.2 INVITED TALKS 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 Online Podcast Everything Epigenetics, podcast hosted by Hannah Went Causal Epigenetic Age Uncouples Damage and Adaptation 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 Everything Epigenetics, podcast hosted by Hannah Went Online Podcast Causal Epigenetic Age Uncouples Damage and Adaptation 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 Graduate Researcher, Vadim Gladyshev's Lab 2020 - Present Harvard Medical School, Boston Children's Hospital Boston, MA Rotation Student, Eric Greer's Lab 2020 Harvard Medical School Boston, MA Rotation Student, David Sinclair's Lab 2019 Harvard T. H. Chan School of Public Health Boston, MA Rotation Student, Brendan Manning's Lab 2019 Sun Yat-Sen University Guangzhou, China Undergraduate Researcher, Zhou Songyang's Lab 2018 - 2019 University of Edinburgh Edinburgh, UK Undergraduate Researcher, Xia Shen's Lab 2018 University of Washington Seattle, WA Undergraduate Researcher, Matt Kaeberlein's Lab **Buck Institute for Research on Aging** Novato, CA Undergraduate Researcher, Judith Campisi's Lab 2018 University of California, Berkeley Berkeley, CA Undergraduate Researcher, Danica Chen's Lab Sun Yat-Sen University Guangzhou, China 2015 - 2017 Undergraduate Researcher, Yikang Rong's Lab Honors Best Poster Award, Inaugural Biomarker of Aging Symposium 2023 Student Spotlight, Harvard Chan School of Public Health 2023 Best Poster Award, Gordon Research Conference, Systems Aging 2022 Hackathon Winner, Longevity Hackathon, VitaDAO 202I Yan-Sen Honor School Program, Sun Yat-Sen University 2016 - 2019

Yan-Sen Scholarship, Sun Yat-Sen University

2016 - 2019

Professional Experience

SERVICE & LEADERSHIP

President, Harvard Interdisciplinary Discussion on Disease and Health

2024 - Present

Advisory Committee Member, Massachusetts Community Health & Healthy Aging Funds

2024 - Present

Organizing Committee Member, Biomarker of Aging Symposium

2023

TEACHING & MENTORING

Mentor, Yuanpei Young Scholars Program

2023 - 2024

Instructor, 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

Dr. Vadim Gladyshev, Dissertation Advisor

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Professor of Medicine, Harvard Medical School

Dr. Steve Horvath, Collaborator

Professor of Human Genetics, UCLA

shorvath@mednet.ucla.edu

Dr. David Sinclair, Dissertation Advisory Committee

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Professor of Genetics, Harvard Medical School

Dr. Matt Kaeberlein, Advisor

kaeber@uw.edu

Professor of Pathology, University of Washington