YUAN HE, Ph.D.

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EDUCATION

Postdoc	Life Sciences Division Lawrence Berkeley National Laboratory – Berkeley, California, USA	2014
Ph.D.	Interdepartmental Biological Sciences Graduate Program Northwestern University – Evanston, Illinois, USA	2008
B.S.	Department of Chemical Engineering Beijing Tech. and Bus. University – Beijing, China	2003
RESEARCH		
Associate Pr Northwestern	ofessor University, Department of Molecular Biosciences	2021 - present
Assistant Pr Northwestern	ofessor University, Department of Molecular Biosciences	2015 - 2021
Princi	ntist rkeley National Laboratory oal Investigator: Eva Nogales arch Project: <i>Near-atomic resolution visualization of human transcription in</i>	2013 - 2014 itiation
Princi	l Fellow rkeley National Laboratory oal Investigator: Eva Nogales arch Project: <i>Structural studies of human transcription by cryo-electron mic</i>	2008 - 2013 rroscopy
Advis	a te Student University, Department of Molecular Biosciences or: Ishwar Radhakrishnan arch Project: <i>Structure-function analysis of the mammalian Sin3 corepress</i>	2003 - 2008 or complex
Tsinghua Uni Advise	ate Research versity, Department of Biological Sciences and Biotechnology or: Zengyi Chang arch Project: <i>Biochemical analysis of hemoglobin o of Mycobacterium tube</i>	2002 - 2003 rculosis

RESEARCH INTERESTS

Transcription regulation, DNA repair, structural biology

HONORS & AWARDS

Burton Medal Microscopy Society of America	2022
Rising Star Award Fusion: 5 th DNA repair/replication structures & cancer conference	2022
NIGMS Director's Featured Research Advance National Institutes of Health	2013
Spot Award for outstanding contributions to the study of transcription Lawrence Berkeley National Laboratory	2013

PEER REVIEWED PUBLICATIONS

Note: Underlined names indicate He Lab graduate student co-authors and asterisked names indicate He Lab postdoc co-authors

(27) Koo, C.W., Tucci, F.J., **He, Y.** and Rosenzweig A.C. (2022) Recovery of particulate methane monooxygenase structure and activity in a lipid bilayer. *Science*, 375:1287-91.

(26) <u>Reyes, A.A.</u>, Fishbain, S. and He, Y. (2022) Structural and functional analysis of the SET3 histone deacetylase complex. *Acta Crystallogr F Struct Biol Commun.*, 78:113-118.

(25) <u>Chen, S.</u>, Lees-Miller, J.P., **He, Y.** and Lees-Miller, S.P. (2021) Structural insights into the role of DNA-PK as a master regulator in NHEJ. *Genome Instability & Disease*, 2, 195-210.

(24) <u>Chen, S.</u>, Lee, L., Naila, T., Fishbain, S., Wang, A., Tomkinson, A.E., Lees-Miller, S.P. and **He, Y.** (2021) Structural basis of Long-range to Short-range synaptic transition in NHEJ. *Nature*, 593, 294-8.

(23) <u>Reyes, A.A.</u>, Marcum, R.D.* and **He**, **Y**. (2021) Structure and function of ATP-dependent chromatin remodeling complexes. *Journal of Molecular Biology*, doi: 10.1016/j.jmb.2021.166929

(22) <u>Abdella, R., Talyzina, A.</u>, Inouye, C., Tjian, R. and **He, Y.** (2021) Structure of the human Mediatorbound transcription pre-initiation complex. *Science*, 372:52-56.

(21) Mashtalir, N., Suzuki, H., Farrell, D.P., Sankar, A., Luo, J., D'Avino, A.R., Filipovski, M., Yang, Y., Valencia, A.M., Pierre, R., Onikubo, T., Roeder, R.G., Han, Y.*, **He, Y.**, Ranish, J.A., DiMaio, F., Walz, T., Kadoch, C. (2020) A structural model of the endogenous human mSWI/SNF (BAF) complex informs disease mechanisms. *Cell*, 183(3):802-817.e24

(20) Marcum, R.D.*, <u>Reyes, A.A.</u> and **He, Y**. (2020) Structural insights into the evolutionarily conserved BAF chromatin remodeling complex. *Biology*, 9(7):E146

(19) Han, Y.*, <u>Reyes, A.A.</u>, Malik, S. and **He, Y.** (2020) Cryo-EM structure of SWI/SNF complex bound to a nucleosome. *Nature*, 579, 452-5.

(18) <u>Abdella, R.</u>, Aggarwal, M., Okura, T., Lamb R.A. and **He, Y.** (2020) Structure of a paramyxovirus polymerase complex reveals a unique methyltransferase-CTD conformation. *Proc Natl Acad Sci USA* 117, 4931-4941.

(17) Yan, C., Dodd, T., **He, Y.**, Tainer, J.A., Tsutakawa, S.E., Ivanov, I. (2019) Transcription preinitiation complex structure and dynamics provide insights into genetic diseases. *Nature Structural and Molecular Biology*, 26, 397-06.

(16) Han, Y.*, Yan, C., Fishbain, S., Ivanov, I. and **He, Y.** (2018) Structural visualization of RNA polymerase III transcription machineries. *Cell Discovery* 4, 40.

(15) Jackobel, A.J., Han, Y.*, **He, Y.** and Knutson, B.A. (2018) Breaking the mold: structures of the RNA polymerase I transcription complex reveal a new path for initiation. *Transcription* 9 (4) 255-261.

(14) Han, Y.*, Yan, C., Nguyen, T.H.D., Jackobel, A.J., Ivanov, I., Knutson, B.A. and **He, Y.** (2017) Structural mechanism of ATP-independent transcription initiation by RNA polymerase I. *Elife* doi: 10.7554/elife.27414.

(13) Nogales, E., Louder, R.K., **He, Y.** (2017) Structural insights into the eukaryotic transcription initiation machinery. *Annual Review in Biophysics* 46, 59-83.

(12) Nogales, E., Louder, R.K., **He, Y.** (2016) Cryo-EM in the study of challenging systems: the human transcription pre-initiation complex. *Current opinion in structural biology* 40, 120-127.

(11) Han, Y., **He**, **Y**. (2016) Eukaryotic transcription initiation machinery visualized at molecular level. *Transcription* 7(5), 203-208.

(10) **He, Y.**, Yan, C., Fang, J., Inouye, C., Tjian, R., Ivanov, I., and Nogales, E. (2016) Near atomic resolution visualization of human transcription promoter opening. *Nature*, 533, 359-65.

(9) Louder, R.K., **He, Y.**, Lopez-Blanco, J.R., Fang, J., Chacon, P., and Nogales, E. (2016) Structure of promoter-bound TFIID and model of human pre-initiation complex assembly. *Nature*, 531, 604-9.

(8) Zhang, E., **He, Y.**, Grob, P., Fong, Y., Nogales, E., and Tjian, R. (2015) Architecture of the human XPC DNA repair and stem cell coactivator complex. *Proc Natl Acad Sci USA* 112(48):14817-22.

(7) **He, Y.**, Fang, J., Taatjes, D.J., and Nogales, E. (2013) Structural visualization of key steps in human transcription initiation. *Nature* 495, 481-6.

(6) Xie, T., **He, Y.**, Korkeamaki, H., Zhang, Y., Imhoff, R., Lohi, O., Radhakrishnan, I. (2011) Structure of the 30 kDa Sin3-Associated Protein (SAP30) in Complex with the Mammalian Sin3A Corepressor and its Role in Nucleic Acid Binding. *J. Biol. Chem.* 286, 27814–24.

(5) **He, Y.**, Imhoff, R., Sahu, A. and Radhakishnan, I. (2009). Solution structure of a novel zinc finger motif in the SAP30 polypeptide of the Sin3 corepressor complex and its potential role in nucleic acid recognition. *Nucleic Acids Res.* 37, 2142–2152.

(4) **He, Y.**, and Radhakishnan, I. (2008). Solution NMR studies of apo-mSin3A and mSin3B reveal that the PAH1 and PAH2 domains are structurally independent. *Protein Sci.* 17, 171-175.

(3) He, Y., Hicke, L., and Radhakishnan, I. (2007). Structural basis for ubiquitin recognition by SH3 domains. *J. Mol. Biol.* 373, 190-196.

(2) Stamenova, S.D., French, M.E., **He, Y.**, Francis, S.A., Kramer, Z.B., and Hicke, L. (2007). Ubiquitin binds to and regulates a subset of SH3 domains. *Mol. Cell* 25, 273-284.

(1) Liu, C., **He, Y.**, and Chang, Z. (2004). Truncated hemoglobin *o* of *Mycobacterium tuberculosis*: the oligomeric state change and the interaction with membrane components. *Biochem. Biophys. Res. Commun.* 316, 1163-1172.

Link to published work in MyBibliogrphy: https://www.ncbi.nlm.nih.gov/myncbi/yuan.he.1/bibliography/public/

INVITED TALK

Department of Chemical Physiology and Biochemistry, OHSU, Portland, OR	2022
NNF Symposium, Rethinking Transcription Factors, Copenhagen, Danmark	2022
Van Andel Institute Symposium on Epigenetics, Grand Rapids, MI	2022
Microscopy and Microanalysis, Portland, OR	2022
The 18 th SCBA International Symposium, Boston, MA	2022
6 th Telluride Workshop on Chromatin Structure and Dynamics, Telluride, CO	2022
Molecular Biology, Memorial Sloan Kettering Cancer Center, New York, NY	2022
Fusion: 5 th DNA repair/replication structures & cancer conference, Cancun, Mexico	2022
Department of Biology, Johns Hopkins University, Baltimore, MD	2022
Social DNAing Webinar, Columbia University, NY	2022
Department of Biochemistry and Molecular Biology, Southern Illinois University, Carbondale, IL	2022
Department of Chemical Pathology, The Chinese University of Hong Kong, China	2022
	2022
Gordon Research Conference: "Mammalian DNA Repair", Ventura, CA	
Functional consequences of genetic variants seminar series, Emory University, Atlanta, GA	2021
Department of Molecular Biology, Max Planck Institute for Biophysical Chemistry	2021
Department of Pharmacology, Case Western Reserve University, Cleveland, OH	2021
Department of Biochemistry, Vanderbilt University, Nashville, TN	2021
Division of Life Science, The Hong Kong University of Science and Technology, China	2021
Advanced Science Research Center, City College of New York, NY	2021
Department of Biochemistry, University of Colorado Boulder, CO	2021
Department of Biophysics and Biophysical Chemistry, Johns Hopkins University, Baltimore, MD	2021
Fragile Nucleosome webinar	2021
•	2021
Cold Spring Harbor Lab Meeting: "Mechanisms of Eukaryotic Transcription", New York	
Department of Molecular and Cellular Oncology, MD Anderson Cancer Center, Houston, TX	2021
Department of Biological Chemistry, University of Michigan, Ann Arbor, MI	2021
Department of BCMP, Harvard Medical School, Boston, MA	2020
Department of Biochemistry and Molecular Biophysics, Washington University in St. Louis, MO	2020
Department of Biochemistry and Molecular Biology, University of Chicago, Chicago, IL	2020
17 th Annual CBC Symposium: "Epigenetics and Disease", Chicago, IL	2020
The American Society for Biochemistry and Molecular Biology Conference, Snowbird, UT	2020
Keystone Symposium: "Cryo-EM for Health Science", Stockholm, Sweden	2020
Keystone Symposium: "Gene Regulation: From Mechanisms to Disease", Keystone, CO	2020
Van Andel Institute Symposium on Structural Biology, Grand Rapids, MI	2019
	2013
Eppley Institute for research in Cancer, University of Nebraska Medical Center, Omaha, NE	
2018 Purdue Cryo-EM Symposium, Purdue University, Lafayette, IN	2018
26 th Conference on Intelligent Systems for Molecular Biology, Chicago, IL	2018
American Crystallography Association Transactions Symposium, New Orleans, LA	2017
Department of Biochemistry and Molecular Genetics, University of Illinois at Chicago, Chicago, IL	2017
Department of Chemistry, University of Illinois at Chicago, Chicago, IL	2017
Third Coast Workshop on Biological Cryo-EM, Chicago, IL	2017
Department of Biochemistry and Molecular Genetics, Northwestern University, Chicago, IL	2017
Division of Life Science, The Hong Kong University of Science and Technology, China	2016
Department of Chemistry, Georgia State University, Atlanta, GA	2016
Fusion: Dynamic DNA and RNA Structures in Damage Responses & Cancer, Cancun, Mexico	2016
FASEB Science Research Conferences: "Machines on Genes", Snowmass, Colorado	2014
Plexxikon, Berkeley, CA	2014
UCSF ACSS "Great People and Science" Seminar, San Francisco, California	2013

CONFERENCE POSTER PRESENTATION

HHMI Scientific Meeting, Ashburn, VA	2013
Cold Spring Harbor Lab Meeting: "Mechanisms of Eukaryotic Transcription", New York	2013
Gordon Research Conference: "Three Dimensional Electron Microscopy", New London, NH	2013
The American Society for Biochemistry and Molecular Biology Meeting, Snowbird, UT	2012
Annual Structural Cell Biology of DNA Repair Machines Workshop, Berkeley, CA 2	2008 - 2013
Cold Spring Harbor Lab Meeting: "Mechanisms of Eukaryotic Transcription", New York	2007
Keystone Symposium: "Frontiers of NMR in Molecular Biology", Banff, Canada	2005

TEACHING AND MENTORSHIP

Classroom Teaching: Instructor, Biophysics (BIOL SCI 363) overall instructor evaluation: 5.71 out of 6	Spring 2022
overall course rating: 5.5 out of 6 Instructor, Biophysics (BIOL SCI 363) overall instructor evaluation: 5.5 out of 6	Spring 2021
overall course rating: 5.17 out of 6 Instructor, Biophysics (BIOL SCI 363) overall instructor evaluation: 5.78 out of 6	Spring 2020
overall course rating: 5.6 out of 6 Instructor, Biophysics (BIOL SCI 363) overall instructor evaluation: 5.36 out of 6 overall course rating: 5.2 out of 6	Spring 2019
Co-Instructor, Biochemistry (IGP 401)	Fall 2019
Co-Instructor, Biophysical Methods for Macromolecular Analysis (IBIS 409)	Fall 2019
Special Topics, Frontiers on Transcription Regulation (IBIS 455) overall instructor evaluation: 4.67 out of 6 overall course rating: 4.42 out of 6	Winter 2019
Co-Instructor, Biochemistry (IGP 401)	Fall 2018
Instructor, Biophysics (BIOL SCI 363) overall instructor evaluation: 5.0 out of 6 overall course rating: 4.9 out of 6	Spring 2018
Co-Instructor, Biochemistry (IGP 401)	Fall 2017
Co-Instructor, Biophysical Methods for Macromolecular Analysis (IBIS 409)	Fall 2017
Instructor, Biophysics (BIOL SCI 363) overall instructor evaluation: 4.93 out of 6 overall course rating: 4.93 out of 6	Spring 2017
Co-Instructor, Biophysical Methods for Macromolecular Analysis (IBIS 409)	Winter 2016
Graduate Students:	
Weifeng Lu	2022 - present
Jinkang Qing	2021 - present
Alex Vogt	2021 - present
Weifeng Lu	2020 - 2021
Anna Talyzina	2020 - present
Siyu Chen	2017 - 2022

Alexis Reyes Luis Schachner (as secondary mentor for this CLP trainee) Ryan Abdella

Graduate Rotation Students:

Young Jun Kim Weifeng Lu Ryan McKeown Tonatiuh Ocampo Jinkang Qing Alex Vogt Elias Guan Frank Tucci **Reyvin Reyes David Bushhouse** Anna Talyzina Alexander Lee Ziyu Zhao Joy Nyaanga Amy Neely Bethany Sump Siyu Chen Kevin Gallagher Alexis Reyes Christopher Koo Ryan Abdella

Undergraduate Student Researchers:

Ethan Less Andrew Kearney Adele Javal Robert Luca Maria Fernandez Davila Pastor Annie Wang (Undergraduate Summer Research Grant Recipient) Eugenie Bang (Undergraduate Summer Research Grant Recipient) Sara Malik (Undergraduate Summer Research Grant Recipient) Srijit Paul (BIOL SCI 398)

Postdoctoral fellows:

Noah Bradley Moustafa Saleh Jingfei Xu Avinash Patel Ryan Marcum Yan Han

Research Technicians:

Oliver Vickman Susan Fishbain

Fall 2022 Spring 2022 Winter 2022 Fall 2021 Summer 2021 Spring 2021 Winter 2021 Spring 2020 Spring 2020 Winter 2020 Fall 2019 Summer 2019 Winter 2019 Fall 2018 Fall 2017 Spring 2017 Fall 2016 Spring 2016 Winter 2016 Fall 2015 Spring 2015

Fall 2022 - present Fall 2022 - present Summer 2022 Summer 2021 - present Summer 2021 Spring 2019 – Summer 2021 Spring 2018 – Summer 2020 Spring 2018 - Winter2020 Spring 2017

Fall 2022 – present Summer 2022 – present Spring 2022 – present Fall 2020 – present Fall 2019 – Fall 2020 Summer 2015 – Fall 2019

Summer 2021 – Winter 2021 Winter 2015 - present

2016 - 2022 2016 - 2021 2015 - 2021

Qualifying Exams and Thesis Defense Committe	es:
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Co-organizer of the IBiS Graduate Program Annual Retreat

Poster Judge for the Annual Biophysics Symposium, Evanston, IL

Poster Judge for the Annual Biophysics Symposium, Evanston, IL

IBiS Graduate Program Admissions Committee

QSB Master's Program Admissions Committee

IBiS Graduate Program Admissions Committee

Structural Biology Core Facility Advisory Committee

Department of Molecular Biosciences Vision Committee

MSTP Admissions Committee

Brooke Angell, Qualifying Exam and Thesis Committee	2021 - present
Alexandra Karagiaridi, Qualifying Exam and Thesis Committee	2021 - present 2020 - present 2020 - present 2020 - present 2019 - present 2018 - 2022 2018 - present 2017 - present
Elijah Taylor, Qualifying Exam and Thesis Committee	
Frank Tucci, Qualifying Exam and Thesis Committee	
Bin Zheng, Qualifying Exam and Thesis Committee	
Jake VanBelzen, Qualifying Exam and Thesis Committee	
Luyi Cheng, Qualifying Exam and Thesis Committee	
Michael Schamber, Qualifying Exam and Thesis Committee	
Katherine Berman, Qualifying Exam and Thesis Committee	
Peter Doubleday, Thesis Committee	2017 - 2021
Ronald Biggs, Qualifying Exam and Thesis Committee	2016 - 2021
Christopher Koo, Qualifying Exam and Thesis Committee	2016 - present
Katarzyna Soczek, Thesis Committee and Thesis Defense Committee	2016 - 2019
Arnold Huang, Masters Defense Committee	2015 - 2016
Amy Nilles, Masters Defense Committee	2015 - 2016
Other Teaching Related Activities:	
IBIS 519 Ethics Refresher	Spring 2022
IBIS 423 Ethics in Biological Research, session on peer review process	2017 - present
Guest lecture in IBIS Structural Biology Workshop	2015 - present
SERVICE	
Departmental/Graduate Program/University Service:	
Molecular Biosciences faculty search committee	2022 - 2023
Weinberg College of Arts and Sciences ad hoc tenure committee	2022 - 2023
Molecular Biosciences DEI committee	2021 - present
Associate Director of Structural Biology Facility	2021 - present
IBiS Graduate Program Admissions Committee	2021 - 2022
IBiS Graduate Program 1 st year student advisor	2020 - present
HHMI Gilliam Fellowship Northwestern Nominating Committee	2020
Poster Judge for the Annual Biophysics Symposium, Evanston, IL	2020
IBiS curriculum Committee	2019 - present
PBS curriculum Committee	2019 - present
Molecular Biophysics Training Program Steering Committee	2019 - present
Judge for the Northwestern University Undergraduate Research and Arts Exposition	2019
Judge for the Chicago Area Undergraduate Research Symposium	2019
Poster Judge for the Annual Biophysics Symposium, Evanston, IL	2019
IBiS Graduate Program Admissions Committee	2018 - 2019
Poster Judge for the Annual Biophysics Symposium, Evanston, IL	2018

2018

2017

2016

2017 - present

2017 - present

2016 - present

2016 - present 2016 - 2017

2017 - 2018

Manuscripts reviewed for the following journals:

Nature, Science, Nature Structure & Molecular Biology, Nature Communication, Elife, Cell Reports, Molecular Cell, Proc Natl Acad Sci USA, Journal of Structural Biology, Structure, Biochemistry, Transcription, Epigenetics & Chromatin, Journal of Nanomaterials

2015

RESEARCH SUPORT

Ongoing Research Support:

R01 (He) Structure and Mechanism of Eukaryotic Transcription Regulation Agency: NIGMS The goal of this study is to dissect the mechanism of eukaryotic transcription regulation by both RNA polymerase II and III using the single particle cryo-EM approach. Total Direct Costs: \$1,024,500 Time Period: 04/01/2022 – 03/31/2026

R01 (He)

Structure and Mechanism of Non-Homologous End Joining Agency: NIGMS The goal of this study is to dissect the mechanism of non-homologous end joining using the single particle cryo-EM approach. Total Direct Costs: \$ 1,170,400 Time Period: 02/11/2020 – 01/31/2025

P01 CA092584 (Tainer) (He, subaward PI) Structural Cell Biology of DNA Repair Machines (SBDR) Agency: LBNL (NIH/NCI, Originating Sponsor) SBDR will identify fundamental structural principles for repair proteins, and provide the framework for understanding DNA repair. The He lab will carry out cryo-EM studies of double strand break repair complexes. Total Direct Costs: \$225,000

Total Direct Costs: \$225,000 Time Period: 09/01/2021 – 08/31/2026

U54CA231638 (Kadoch, Shilatifard) (He, subaward) The Center for Synovial Sarcoma Biology and Therapeutics Agency: DANA-FARBER CANCER INST (NIH/NCI, Originating Sponsor)

The overarching goal for the center is to develop and execute a comprehensive, multidisciplinary set of approaches rooted in protein biochemistry and structural biology to define the mechanistic underpinnings of synovial sarcoma and unmask opportunities for therapeutic development. The He lab will carry out cryo-EM studies of human SWI/SNF complex in both wild type and cancer relevant mutant forms.

Total Direct Costs: \$34,929 Time Period: 09/10/2021 – 06/30/2023

Completed Research Support:

PSOC Pilot project award (He)

Molecular mechanism of the SWI/SNF complex in regulating chromatin structure and gene transcription Agency: NCI

The goal of this study is to dissect the mechanism of chromatin remodeling by SWI/SNF using the single particle cryo-EM approach.

Time Period: 05/01/2019 - 04/30/2020

IRG-15-173-21 (He) Structural Basis of Human Transcription Promoter Proximal Pausing Agency: American Cancer Society The goal of this study is to dissect the mechanism of transcription promoter proximal pausing using the single particle cryo-EM approach. Time Period: 01/01/2016 – 12/31/2017

C-064 Catalyst Award (He) Toward the Visualization of Spliceosomal Intermediates Agency: Chicago Biomedical Consortium The goal of this study is to determine structures of spliceosome complexes trapped at various splicing intermediate states using single particle cryo-EM approach. Time Period: 02/01/2016 – 01/31/2018

Cornew Innovation Award (He, Co-PI) Toward elucidating RNA co-transcriptional folding pathways from RNA polymerase II Agency: The Chemistry of Life Processes Institute at Northwestern University The goal of this study is to dissect the mechanism of RNA polymerase II directed mRNA cotranscriptional folding using the single particle cryo-EM approach. Time Period: 01/01/2017 – 12/31/2017

PROFESSIONAL MEMBERSHIPS

Member, American Society for Biochemistry and Molecular Biology Member, Biophysical Society Member, Robert H. Lurie Comprehensive Cancer Center at Northwestern University Member, The Chemistry of Life Processes Institute at Northwestern University

Date of CV preparation: 12/9/2022