James P. Lingford

Website: jameslingford.com

Last updated: July 2024

Education

02/2009 - II/2012 University of Western Australia

B.Sc. (Hons) in Genetics

Research positions

OI/2023 - Present Monash Biomedicine Discovery Institute, VIC

Research Officer – Greening Lab

The structural biology of enzymes that extract energy from air

08/2016 - 08/2021 Walter and Eliza Hall Institute of Medical Research, VIC

Research Assistant - Goddard-Borger Lab

The structural biology of mucosal proteins and sulfoglycolysis

08/2015 Children's Medical Research Institute, NSW

Visiting Scientist - Host: Dr. Scott B. Cohen

Learning how to run a direct telomerase activity assay

02/2014 - 08/2016 Harry Perkins Institute of Medical Research, WA

Research Assistant - Rackham Lab

Designing PPR proteins for programmable nucleic acid binding

Honours & awards

2024 Chemistry Biology Interface Horizon Prize: Rita and John Cornforth Award – Royal

Society of Chemistry. For developing an understanding of sulfosugar metabolism and

the discovery of new enzymes and pathways of sulfur recycling. [html]

2012 Honours stipend scholarship – ARC Centre for Plant Energy Biology, The University of

Western Australia.

Publications

18.

JOURNAL ARTICLES

Leung PM[†], Grinter R, Tudor-Matthew E, **Lingford JP**, Jimenez L, Lee H-C, Milton M, Hanchapola I, Tanuwidjaya E, Kropp A, Peach HA, Carere CR, Stott MB, Schittenhelm RB, Greening C[†]. Trace gas oxidation sustains energy needs of a thermophilic archaeon at suboptimal temperatures. *Nature Communications*. **2024**. [html]

Arumapperuma T, Lee M, Sharma M, Zhang Y, Snow AJD, **Lingford JP**, Goddard-Borger ED[†], Davies GJ[†], Williams SJ[†]. Capture-and-release of a sulfoquinovose-binding protein on sulfoquinovose-modified agarose. *Organic & Biomolecular Chemistry*. **2024**. [html] [pdf]

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- Sharma M, Kaur A, Soler NM, **Lingford JP**, Epa R, Goddard-Borger ED, Davies GJ[†], Williams SJ[†]. Defining the molecular architecture, metal dependence, and distribution of metal-dependent class II sulfofructose-I-phosphate aldolases. *Journal of Biological Chemistry*. **2023**. [html]
- Snow AJD, Sharma M, **Lingford JP**, Zhang Y, Mui JWY, Epa R, Goddard-Borger ED, Williams SJ, Davies GJ[†]. The sulfoquinovosyl glycerol binding protein SmoF binds and accommodates plant sulfolipids. *Current Research in Structural Biology*. **2022**. [html]
- Sharma M, Lingford JP, Petricevic M, Snow AJD, Zhang Y, Järvå MA, Mui JWY, Scott NE, Saunders EC, Mao R, Epa R, da Silva BM, Pires DEV, Ascher DB, McConville MJ, Davies GJ, Williams SJ, Goddard-Borger ED. Oxidative desulfurization pathway for complete catabolism of sulfoquinovose by bacteria. *Proceedings of the National Academy of Sciences*. 2022. [html]
- Mao R, Xi S, Shah S, Roy MJ, John A, **Lingford JP**, Gäde G, Scott NE, Goddard-Borger ED[†]. Synthesis of C-mannosylated glycopeptides enabled by Ni-catalyzed photoreductive cross-coupling reactions. *JACS*. **2021**. [html] [pdf]
- Sharma M, Abayakoon P, Jin Y, Epa R, **Lingford JP**, Shimada T, Nakano M, Mui J, Ishihama A, Goddard-Borger ED[†], Davies GJ[†], Williams SJ[†]. The Molecular Basis of Sulfosugar Selectivity in Sulfoglycolysis. *ACS Central Science*. **2021**. [html]
- Li J, Epa R, Lingford JP, Scott NE, Skonesczny D, Sharma M, Snow A, Goddard-Borger ED, Davies GJ, McConville MJ, Williams SJ[†]. A sulfoglycolytic Entner-Doudoroff pathway in *Rhizobium leguminosarum* bv. *trifolii* SRDI565. *Applied and Environmental Microbiology*. 2020. [html]
- Järvå MA, Lingford JP, John A, Scott NE, Goddard-Borger ED[†]. Trefoil factors share a lectin activity that defines their role in mucus. *Nature Communications*. 2020. [html]
- Järvå MA*, Dramicanin M*, **Lingford JP**, Mao R, John A, Jarman K, Grinter RW, Goddard-Borger ED[†]. Structural basis of substrate recognition and catalysis by fucosyltransferase 8. *Journal of Biological Chemistry*. **2020**. [html]
- Sharma M, Abayakoon P, **Lingford JP**, Jin Y, Epa R, Goddard-Borger ED[†], Davies GJ[†], Williams SJ[†]. Dynamic structural changes accompany the production of dihydrox-ypropanesulfonate by sulfolactaldehyde reductase. *ACS Catalysis*. **2020**. [html] [pdf]
- Zhang Y*, Mui J*, Arumaperuma T, Lingford JP, Goddard-Borger ED, White J, Williams SJ†. Concise synthesis of sulfoquinovose and sulfoquinovosyl diacylglycerides, and development of a fluorogenic substrate for sulfoquinovosidases. *Organic & Biomolecular Chemistry*. 2020. [html] [pdf]
- Abayakoon P, Ruwan E, Petricevic M, Christopher C, Mui J, van der Peet P, Zhang Y, Lingford JP, White J, Goddard-Borger ED, Williams SJ[†]. Comprehensive synthesis of substrates, intermediates and products of the sulfoglycolytic Embden-Meyerhoff-Parnas pathway. *The Journal of Organic Chemistry*. 2019. [html] [pdf]
- Abayakoon P*, Jin Y*, **Lingford JP**, Petricevic M, John A, Ryan E, Wai-Ying Mui J, Pires DEV, Ascher DB, Davies GJ[†], Goddard-Borger ED[†], Williams SJ[†]. Structural and biochemical insights into the function and evolution of sulfoquinovosidases. *ACS*

Central Science. 2018. [html]

Spåhr H*, Chia T*, **Lingford JP**, Siira SJ, Cohen SB, Filipovska A, Rackham O[†]. Modular ssDNA binding and inhibition of telomerase activity by designer PPR proteins. *Nature Communications*. **2018**. [html]

Abayakoon P, **Lingford JP**, Jin Y, Bengt C, Davies GJ, Yao S[†], Goddard-Borger ED[†], Williams SJ[†]. Discovery and characterization of a sulfoquinovose mutarotase using kinetic analysis at equilibrium by exchange spectroscopy. *Biochemical Journal*. **2018**. [html]

Lopaticki S*, Yang ASP*, John A, Scott NE, Lingford JP, O'Neill MT, Erickson SM, McKenzie NC, Jennison C, Whitehead LW, Douglas DN, Kneteman NM, Goddard-Borger ED[†], Boddey JA[†]. Protein O-fucosylation in Plasmodium falciparum ensures efficient infection of mosquito and vertebrate hosts. *Nature Communications*. 2017. [html]

Coquille S*, Filipovska A*, Chia T, Rajappa L, **Lingford JP**, Razif MF, Thore S, Rackham O[†]. An artificial PPR scaffold for programmable RNA recognition. *Nature Communications*. **2014**. [html]

PREPRINTS

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06/2024

02/2019

02/2018

08/2019

08/2017

Appler KE, **Lingford JP**, Gong X, Panagiotou K, Leão P, Langwig M, Greening C, Ettema TJG, De Anda V, Baker BJ[†]. Oxygen metabolism in descendants of the archaealeukaryotic ancestor. *bioRxiv*. **2024**. [html]

Kropp A*, Gillett GL*, Venugopal H, Gonzálvez MA, **Lingford JP**, Barlow CK, Zhang J, Greening C[†], Grinter R[†]. Quinone extraction drives atmospheric carbon monoxide oxidation in bacteria. *bioRxiv*. **2024**. [html]

ARTICLES IN NEWS MEDIA

Samorodnitsky D, Bird K, Carlson J, **Lingford J**, Phillips J, Sear R, Townsend C. Journals that published Richard Lynn's racist 'research' articles should retract them. *STAT*. [html]

Conference poster presentations

Structural, mechanistic, and evolutionary insights into sulfoglycolysis. *44th Lorne Conference on Protein Structure and Function*, VIC. [html]

Sulfoquinovosidases as the gatekeepers to sulfoglycolysis: insight into structure, function, mechanism, and evolution. *43rd Lorne Conference on Protein Structure and Function*, VIC.

Seminar presentations

James Lingford – Curriculum Vitae

The structural biology of cysteine rich mucin domains. *ACRF Chemical Biology Divisional Seminar* — *Walter and Eliza Hall Institute of Medical Research*, VIC.

What controls mucus viscoelasticity? ACRF Chemical Biology Divisional Seminar — Walter and Eliza Hall Institute of Medical Research, VIC.

^{*} denotes equal contribution | † denotes corresponding author

Student mentorship & supervision

2021	– Sean Smyth, Masters student.
2018 – 2019	– Runyu Mao, Рн.D. candidate.
2018	– Hanna Mayerhofer, Masters student.
2018	– Keyu Su, InSPIRE program student.
2016	– Dana Tabara, Honours student.
2016	– Kimberley Callaghan, Honours student.
2015	 Peter Alfrich, Honours student.

2015

- Undergraduate laboratory demonstrator, Course: UWA GENE3330 2012

Service

2024 -	- Organising committee member of the Monash Structural Biology Symposium. [html]
2017 – 2021	– Safety Officer, Walter and Eliza Hall Institute.
2017 – 2021	– Fire Warden, Walter and Eliza Hall Institute.
2014 – 2016	– Safety Committee member representing the Rackham lab.

Public outreach

2019	– Tour guide: Public Discovery Tour, Walter and Eliza Hall Institute.
2018	– Tour guide: SEAMS Discovery Tour, Walter and Eliza Hall Institute.

Skills

STRUCTURAL
Biology

Cryo-EM training at the Monash Ramaciotti Centre for Cryo-Electron Microscopy, VIC.

Cryo-EM three-day training workshop at the ARC CCeMMP facility, VIC.

MX2 Beamline training at the Australian Synchrotron, VIC. Crystallography basics: crystallisation trials and fishing crystals.

BIOPHYSICS

Surface plasmon resonance (SPR).

Isothermal titration calorimetry (ITC).

Mass photometry.

PROTEIN Віоснемізтку Recombinant protein expression and purification.

Membrane protein purification, including ultracentrifugation of membrane fractions.

Eukaryotic recombinant protein expression in Sf 21 insect cells.

Anaerobic protein expression in *E. coli*.

Affinity chromatography.

Size exclusion chromatography (SEC). Ion exchange chromatography (IEX).

ÄKTA Pure System: handling, method design, and basic maintenance.

Direct telomerase activity assay with radioisotopes.

Michaelis-Menten kinetic analysis with spectrophotometric plate readers.

Blue native polyacrylamide electrophoresis (BN-PAGE).

Electrophoretic mobility shift assays (EMSA).

Protein quantification: BCA assay, Bradford assay, and nanodrop.

SDS-PAGE and silver staining.

Gas chromatography.

Molecular Recombinant construct design for protein expression and purification.

Biology Primer design and PCR.

Traditional molecular cloning and Gibson assembly cloning.

Plasmid minipreps and maxi-preps.

Site-directed mutagenesis: overlap extension PCR and QuickChange PCR.

Preparation of competent *E. coli* cells.

Cellular Working in a laminar flow hood with correct ascetic technique. Biology

Passaging, monitoring, and counting mammalian cell lines.

Transfecting CHO cells (non-lentiviral work).

Сомрита-Working with a Linux command-line interface. TIONAL Programming and scripting basics: Bash, Python, R. BIOLOGY Data analysis basics: Python (JupyterLab), R.

High Performance Computing: ssh, sbatch scripting, Singularity containers.

Workflow tools: git, conda, Vim/Neovim, Markdown, LATEX.

Protein structure modelling and analysis: AlphaFold2, FoldSeek, UCSF ChimeraX.

Figure design: UCSF ChimeraX, Inkscape, GIMP image software.

Lab Ordering lab reagents and supplies.

MANAGEMENT Making and autoclaving liquid media and buffers.

Calibrating pH meters.

Adhering to PC2 safety precautions.

Other information

LinkedIn linkedin.com/in/jameslingford

Scholar goo.gl/xdjuRr

Orcid Id 0000-0003-1980-3949 **GITHUB** github.com/jlingford