

Interdisciplinary Pathways In Medicine-from Neurodegenerative to Oncology Disease Research And Drug Discovery



FRANK LEE, PH.D. Scientific Associate, **Krembil Research Institute**



MARK A REED, PH.D. Chief Science Officer, **Treventis Corporation**



LIVE

Thursday, March 21 2024 8:00 AM (PDT) | 11:00 (EDT) 3:00 (GMT)



ERIC GIFFORD, PH.D. **Business Development Consultant** Collborative Drug Discovery



Have a question to ask our panel?

Open the ZOOM Q&A and type in your question during the webinar





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Dr. Mark A. Reed

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- 2018-present Staff Scientist II and Director, Centre for Medicinal Chemistry and Drug Discovery (CMCDD), Krembil Research Institute, UHN, Toronto
- 2023-present Assistant Professor Department of Chemistry, University of Toronto
- 2019-present Assistant Professor Department of Pharmacology and Toxicology and Department of Chemistry, University of Toronto
 - 2008–present Co-founder and CSO, Treventis Corporation
 - 2003–2008 Senior Scientist, Medicinal Chemistry, **ICOS Corporation**, Seattle, WA and **Schering Plough Research Institute**, Cambridge, MA
 - 2000–2003 Postdoctoral Fellow, Queen's University, Canada. (**Organic** chemistry Aromatic Lithiation)
- 1996–2000 Ph.D., University of Sussex, UK**. (Organic Chemistry Total Synthesis)**





Dr. C. Frank Lee

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- 2020-present Scientific Associate II, Centre for Medicinal Chemistry and Drug Discovery (CMCDD), Krembil Research Institute, University Health Network (UHN), Canada
 - 2018–2020 Postdoctoral Fellow, **Centre for Medicinal Chemistry and Drug Discovery (CMCDD)**, Krembil Research Institute, University Health Network (UHN), Canada
- 2013–2018 Ph.D., University of Toronto, Canada (Organic Chemistry Organoboron methodology & Macrocyclic peptides)
- 2009–2013 B.Sc.H., Queen's University, Canada (Organic chemistry-Aromatic Lithiation, Total Synthesis)





Targeting Protein Misfolding Disorders: Discovery of small oligomerization inhibitors for the treatment of neurodegeneration





oronto General Foronto Western Princess Margaret Centre for Medicinal Chemistry and Drug Discovery (CMCDD): Early-stage translation research within Canada largest research hospital

Proteinopathies: Intrinsically Disordered Proteins (IDPs)



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Common Conformational Morphology – CCM



- IDPs
 - No X-ray structures; solution NMR only
 - SBDD not possible
- HTS difficult due protein/ assay nature
- Technology: In silico model of protein misfolding, the Common Conformational Morphology (CCM)
 - An *in silico* model based on epitope commonality between multiple misfolding amyloidogenic proteins (e.g, A Syn, Aβ, tau).
 - Used to build "surrogate crystal structures" of incipient oligomers.



Target: Common Conformational Morphology – CCM





Alzheimer's Disease







Intracellular neurofibrillary tangles Tau – 441 amino acids, Intrinsically disordered protein

Extracellular plaques

 $A\beta$ - 40/42 amino acids, Intrinsically disordered protein

Discovery of a Dual Aβ/tau Oligomer Inhibitor



In Vitro Pharmacology

Biotin-tau 4R2N ELISA Biotin-Aβ42 ELISA SDS PAGE (cell free)

Counter Screens Tublin assembly DLS IDP selectivity MoA

TRESI-HDX/Native-MS



LeVine, H. Anal. Biochem, 356, 265-72, (2006)

ADMET: Property Based Lead Optimization

- BBB score, LogD/ Kinetic Sol. - MDR1-MDCK, Caco2 (Permeability; PgP ER) - M/R/H Hepatocytes/ microsomal clearance) Rat PK (cassette/ discrete) Kp,uu (Total and free drug brain exposure)



Gupta, M. et al., JMedChem, 62, 9824-36, (2019).

Target Engagement (PK-PD)

tau: rTg4510 murine; quant. otau in cortex; FRET- based biosensor/HTRF <u>Aβ42:</u> APP/PS1 murine; oAβ42 ISF microdialysis; ELISA



Cirrito, J.R. et al., J. Neurosci 23, 8844, (2003)

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Aβ Oligomer Target Engagement



- TRV101 in TgAPP.PS1 (n=8): 1000 Kda MWCO Microdialysis/ELISA (ISF)
- APP/PS1 mice, 12 months old
- P.O. / Q.D. in the same animals
 - 30mg/kg on day 1
 - 100mg/kg on day 2
 - 300mg/kg day 3
- Interstitial fluid sampled via microdialysis probe every 2-3 hours
- Aβ Oligomers analyzed by ELISA
- Study conducted by John Cirrito University of Washington St Louis

tau Oligomer Target Engagement





Current Status





CCM Discovery Engine: Enables identification of druggable sites on IDPs for virtual screening against multiple targets



Target Engagement: Dose dependent reduction in vivo of $\text{OA}\beta$ and Otau



PCC identification (Aβ/tau): ~1500 NCEs in lead series, high CNS exposure (10-20X fold over PoC compounds); heterocycles developed for de-risking and supporting robust IP portfolio. PCC selected - IND enabling studies underway



Translational : Multiple chronic model studies underway and biomarker discovery



RESEARCH INSTITUTES





Krembil Research Institute arthritis | vision | neuroscience



McEwen Stem Cell Institute cell-based regenerative therapies



KITE Research Institute rehabilitation science



The Institute for Education Research *health care education*



Princess Margaret Cancer Centre cancer



Toronto General Hospital Research Institute cardiovascular | respiratory and critical care | metabolism | infection and immunity | communities of health

RESEARCH AT



Centre for Medicinal Chemistry & Drug Discovery (CMCDD)

- Established at Krembil in 2018 to identify druggable targets from novel disease biology
- 10 medicinal chemists
- 30 collaborations at UHN and academic institutes across Canada
- Oncology, Neurology, Ophthalmology, Arthritis and Metabolic diseases
- Foster UHN institute collaborations based on therapeutic targets

Portfolio

	Disease Indication/PI	Target Class	Pre-clinical Validation	Stage
Neurology/ Ophthalmology	Rett Syndrome and dementia - Neuroinflammation	Ion Channel	Genetic	Hit to lead
	Depression and dementia - Neuroinflammation	enzyme	Pharmacological	Lead Optimization
	Glaucoma and dementia – Neuroinflammation	unknown	Genetic/ Pharmacological	Receptor Identification
	Stroke, Spinal cord injury: Axonal Regeneration	Kinase	Genetic/ Pharmacological	Hit Finding
	BBB Integrity	Kinase	Genetic/ Pharmacological	Hit Finding
	Depression	Receptor	Genetic/ pharmacological	In Silico HTS
Oncology	AML	Kinase	Genetic/ Pharmacological	Hit to lead
	Radiation Fibrosis	Unknown	Genetic/ Pharmacological	Target ID
	Ovarian Cancer	Kinase	Genetic/ Pharmacological	Hit to lead
	BRCA1/2-mutant breast and ovarian cancer	E3 ligases	Genetic	Hit ID
	Pancreatic cancer	Receptor	Genetic	Assay development
	AML	Stress granule	Genetic	Structure Based Design
	Prostate Caner	transporter	Genetic	Hit to lead
Arthritis	Cartilage Degeneration	Transcription factor	Genetic	Hit finding
	Fibrosis/ myofibroblast activation	enzyme	Genetic/ Pharmacological	Hit ID
	Spondyloarthritis	Chemokine	Genetic/ Pharmacological	Hit to lead
Metabolic	NASH	enzyme	Genetic/ Pharmacological	Hit ID



The Princess Margaret Cancer Foundation 🔮 UHN

Toronto General & Western Hospital Foundation **QUHN**







Neuroinflammation: New Therapeutic Targets



Lower Proinflammatory Phenotype: TRPM2 Ion Channel (Dr. James Eubanks)



Dr. James Eubanks

- Senior Scientist, Krembil Research Institute, UHN
- Research Division Head, Krembil Research Institute, UHN

Enhance Neuroprotection: <u>Target Unknown</u> (Dr. Jeremy Sivak)



Dr. Jeremy Sivak

- Senior Scientist & Glaucoma Research Chair, Krembil Research Institute, UHN and Associate Professor UofT School of Medicine
- 5+ years at Novartis leading multi-diciplenary ophthmalic drug discovery team

TRPM2 Ion Channel in Neuroinflammation

- Member of Melastatin sub-group of TRP channel
- Redox sensing, non-selective cation channel that allows Ca²⁺ influx and efflux
- Functions as a surveillance receptor in the brain, where it monitors local environments for signs of "stress"
- Highly expressed in microglia
- Under stress, TRPM2 activates and allows influx of Ca²⁺ → pro-inflammatory cytokines.
- <u>TRPM2</u> inhibition rescue AβO-induced microglia activation and hippocampal LTP



Raghunatha P, Vosoughi A, Kauppinen TM, Jackson MF. Microglial NMDA receptors drive pro-inflammatory responses via PARP-1/TRMP2 signaling. Glia. 2020;68(7):1421-1434. doi:10.1002/glia.23790

CNS-Penetrant TRPM2 Inhibitor Development



Glaucoma: A Neurodegenerative Disease

- Top cause of permanent blindness and growing (>80 million worldwide)
- Associated with many risk factors:
 - Intraocular pressure (IOP) the only current modifiable risk factor
 - o IOP not associated with all cases
- Neurodegenerative disease of retinal ganglion cells (RGCs)
 - No direct treatment
 - Improved imaging has made clinical trials practical
 - $\circ \quad \text{Accessible to local delivery} \\$
- Goal: target glaucoma through
 neuroprotection

Livne-Bar, I.; Wei, J.; Liu, H.-H.; Alqawlaq, S.; Won, G.-J.; Tuccitto, A.; Gronert, K.; Flanagan, J. G.; Sivak, J. M. J. Clin. Investig. 2017, 127, 4403-4414; and references therein Nucci, C.; Martucci, A.; Giannini, C.; Morrone, L. A.; Bagetta, G.; Mancino, R. Eye 2018, 32, 938-945; and references therein.



GLAUCOMA

DRAINAGE CANAL BLOCKED TOO MUCH FLUID STAYS IN THE EYE THIS INCREASES PRESSURE Projected Glaucoma Cases (U.S.)



*National Eve Institute



HIGH PRESSURE

MAGES OPTIC NERVE

Disc Center (-0.18,-0.36) mm

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LXB₄ Protects RGCs from Glaucoma Injury



- Sivak group discovered novel neuroprotective role of lipoxins
- Lipoxins (LXA₄ and LXB₄): special pro-resolving lipid mediators (SPMs)
- Potent effectors of inflammation resolution
 - $\,\circ\,$ Not well studied in retina or CNS
 - o LXA₄ better understood and has an established receptor (GPCR)
 - $\circ~$ LXB4 less studied, and its receptor is unknown
 - \circ LXB₄ (EC₅₀ = 31nM) is ~20x more potent than LXA₄ (EC₅₀ = 631 nM)
- LXB₄ treatment is efficacious in a 15-week chronic glaucoma model

LXB₄ target deconvolution strategies: Photo Affinity Labelling

(PALMS) (Evotec)

Tritiation in Cell Microarray (Retrogenix)

Livne-Bar, I.; Wei, J.; Liu, H.-H.; Alqawlaq, S.; Won, G.-J.; Tuccitto, A.; Gronert, K.; Flanagan, J. G.; Sivak, J. M. J. Clin. Investig. 2017, 127, 4403-4414; and references therein

Synthesis of LXB₄ & Aromatic Analogs



A Stereocontrolled Total Synthesis of Lipoxin B4 and its Biological Activity as a Pro-Resolving Lipid Mediator of Neuroinflammation

C. Frank Lee,^[a] Carla E. Brown,^[a] Alexander J. Nielsen,^[a] Changmo Kim,^[b, c, i] Izhar Livne-Bar,^[b, c] Philip J. Parsons,^[d] Christophe Boldron,^[e] François Autelitano,^[e] Donald F. Weaver,^[f, g, h] Jeremy M. Sivak,^[b, c, i] and Mark A. Reed*^[a, j]

Lee, C. F.; Brown, C. E.; Nielsen, A. J.; Kim, C.; Livne-Bar, I.; Parsons, P. J.; Boldron, C.; Autelitano, F.; Weaver, D. F.; Sivak, J. M.; Reed, M. A. Chem. Eur. J. 2022, 28, e202200360. Brown, C. E.; Lee, C. F.; Nielson, A. J.; Sivak, J. M.; Reed, M. A. Unpublished results; patent pending.

LXB₄ Target Deconvolution



Portfolio

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Metabolic	NASH	enzyme	Genetic/ Pharmacological	Hit ID





KREMBIL FOUNDATION





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Acknowledgements

TREVENTIS

Senior Management Christopher Barden Mark Reed Marcia Taylor Donald Weaver Chemistry Leela Anagani Scott Banfield **Christopher Barden** Kunal Keskar Erhu Lu Ross Mancini Autumn Meek Shengguo Sun Thomas Wood Fan Wu Arun Yadav Elena Diez Cecilia **Carlos** Zepeda

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RETROGENIX



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Conseil de recherches en sciences naturelles et en génie du Canada



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Collaborators TGH/UofT Dr. Mamatha Bhat Dr. Moumita Baru Dr. Dan Winer











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Questions?





Inventory Keep track of samples, biologicals and compounds







Visualization Plot datasets and mine them



Deep Learning Computer aided design

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