

WELLCOME-WOLFSON **INSTITUTE FOR EXPERIMENTAL MEDICINE** **Queen's University Belfast IENGAGE Program**



OBJECTIVES

6-week course comprised of:

- Daily seminars delivered by world-leading experts in the areas of immunobiology and microbes, vision and vascular medicine and respiratory medicine.
- Weekly roundtables hosted by PhD students and postdoctoral fellows, providing us with insights on their researches and their personal experience from their postgraduate programs.
- Each group was allocated with a supervisor whose role was to guide us through our research projects presented below:

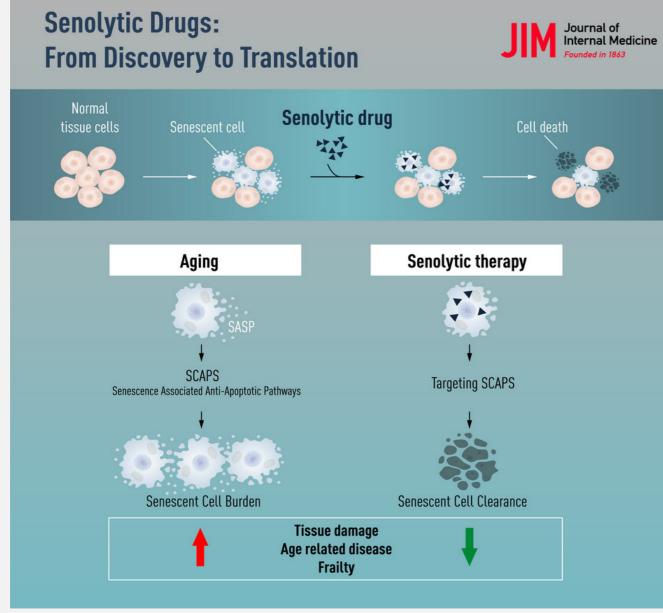
Senolytics in Cardiovascular Disease; Systematic



Angiogenesis refers to formation of new blood vessels from preexisting vessels. Sprouting angiogenesis is initiated in response to a hypoxic environment by secreting proangiogenic growth factor called Vascular Endothelial Growth Factor A (VEGF-A)

Review

Senolytics are a class of drugs that selectively clear senescent cells by inducing apoptosis that regulates senescence cells gene (BCL-2, p16, p53 and p21)



Kirkland, J. L., & Tchkonia, T. (2020). Senolytic drugs: from discovery to translation. Journal of internal medicine, 288(5), 518-536.

CONCLUSION

Navitoclax

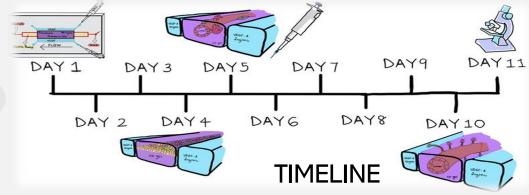
Dasatinib + Quercetin

Senolytics represent a potential new treatment for age-related diseases, but more research is needed to facilitate translation into clinics.

Blood-Retina-Barrier: Changes in Diabetic

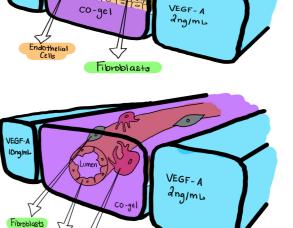
GOAL

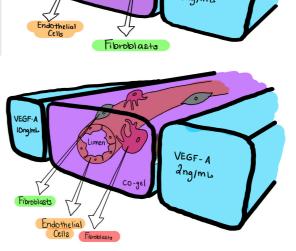
To create a model that represents in vivo conditions of angiogenesis in the presence of a parent vessel under VEGF A concentration gradient.

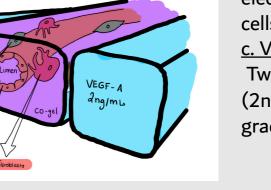




a. CHIP ASSEMBLY The Chip is made from Polyethylene glycol (PEG) b. Co-Culture Endothelial cells are cocultured with fibroblasts on an extracellular matrix to aid in lumen formation and an electric pump is attached to create a flow of endothelial cells from the inlet to the outlet. c. VEGF-A Gradient Two syringes are used to add high (10ng/ml) and low (2ng/ml) concentration of VEGF-A, hence forming a gradient.







CONCLUSION

VESSEL ON A CHIP

VEGF

VEGF

FLOW

Endothelial Cell

scaffold engineering with microfluidics and co-

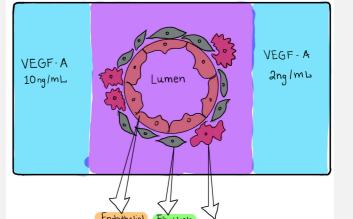
culture

The advantages (parent vessel, flow) of our proposed in vitro model- vessel on a chip will expand our current knowledge on how angiogenesis occurs in the human body. • Secondly the model can also be used to replicate various disease conditions for example cancer and gain deeper insights about its angiogenic influence.

1. Establishing a parent vessel

RESULTS

2. Sprouting angiogenesis takes place towards higher VEGF-A concentration gradient

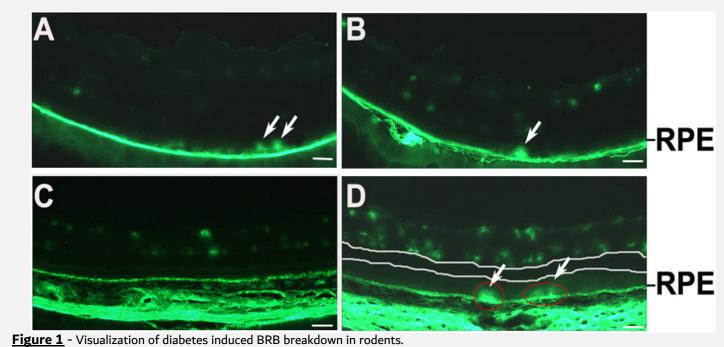


Al derived gene networks to identify markers

Retinopathy; Systematic Review

Blood-retina-Barrier (BRB) functions to maintain homeostasis in the retina to avoid entrance of blood-borne proteins, maintain metabolic + ionic gradients. BRB is divided into inner BRB (2/3^{rds} of retina) and outer BRB (1/3rd of retina) The earliest and most significant change in Diabetic Retinopathy is **BRB disruption/breakdown**.

RESULTS



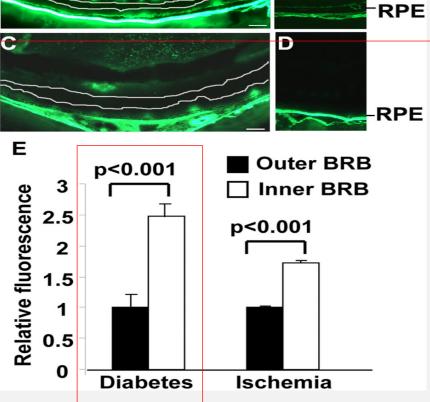


Figure 2 – Significance of BRB leakage in diabetic rodents (1 minute after intravenous injection)

GOAL

METHODS

Reviewing articles published in 2020 and

2021 about senolytics targeting vascular

and endothelial components of

cardiovascular diseases.

RESULTS

Inhibition of BCL-2 antiapoptotic

protein

Clearing cells positive for p21

(Cdkn1a) and alleviating renal

dysfunction and damage

Visualize and evaluate outer vs inner BRB breakdown as a consequence of Diabetic Retinopathy. Determined via "BRBspecific leakage in diabetic rodents by fluorescent microscopy"

METHODS

Visualize and evaluate outer vs inner BRB breakdown as a consequence of Diabetic Retinopathy. Determined by systematic review of articles on "BRB-specific leakage in diabetic rodents by fluorescent microscopy"

CONCLUSION

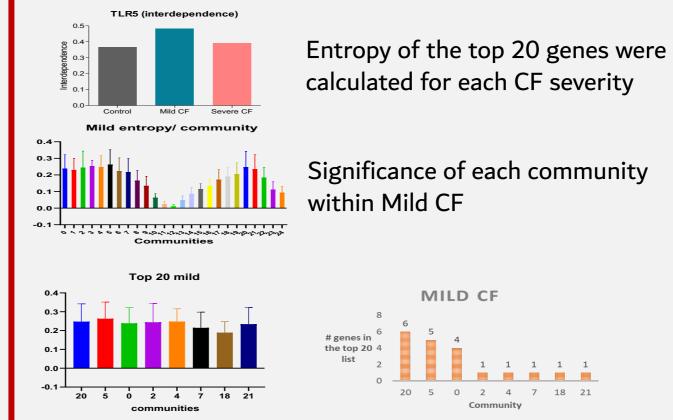
Fig. 1,2 show that FITC-dextran fluorescent dye leaked through outer BRB of DR rodents, suggesting blood-retina-barrier breakdown as a significant pathophysiological component of Diabetic Retinopathy.

REFERENCE:

Xu, Hui-Zhuo, and Yun-Zheng Le. "Significance of Outer Blood-Retina Barrier Breakdown in Diabetes and Ischemia. Investigative Ophtalmology & Visual Science, 1 Apr. 2011.

of phenotypical variability in Cystic Fibrosis (CF) inflammation

RESULTS



Even though communities 20, 5, and 0 contained the majority of the 20 most significant genes, they were not more significant than other communities as a whole

Role of TLR-5 in influencing CF severity

LPS	Control	Control Community 10				
/	- Control - Mild	GO:BP	response to organic substance cellular response to cytokine stimulus	GO:0010033 GO:0071345	3.874×10 ⁻¹³ 1.111×10 ⁻¹²	
	Severe	KEGG	Intestinal immune network for IgA production Pathways in cancer	KEGG:04672 KEGG:05200	5.426×10 ⁻⁴ 7.635×10 ⁻⁴	
		REAC	Cytokine Signaling in Immune system TNFs bind their physiological receptors	REAC:R-HSA-12 REAC:R-HSA-56		
			Mild Community 5		1	
4 24 Hours	0.4] TLR-5	GO:BP	response to organic substance cytokine-mediated signaling pathway	GO:0010033 GO:0019221	1.520×10 ⁻⁹ 9.392×10 ⁻⁹	
	0.3-	KEGG	Epstein-Barr virus infection Measles	KEGG:05169 KEGG:05162	5.215×10 ⁻⁴ 1.474×10 ⁻³	
R5 (interdependence)	on the state of th	REAC	Cytokine Signaling in Immune system TNFs bind their physiological receptors	REAC:R-HSA-12 REAC:R-HSA-56	1.272×10 ⁻⁴	
			Severe Community 16			
		GO:BP	response to organic substance cytokine-mediated signaling pathway	GO:0010033 GO:0019221	1.520×10 ⁻⁹ 9.392×10 ⁻⁹	
		KEGG	Epstein-Barr virus infection Measles	KEGG:05169 KEGG:05162	5.215×10 ⁻⁴ 1.474×10 ⁻³	
	Kruskal-Wallis statistic = 28.99	REAC	Cytokine Signaling in Immune system TNFs bind their physiological receptors	REAC:R-HSA-12 REAC:R-HSA-56	1.272×10-4 8.051×10-3	
Mild CF Severe CF						
Change of	entrony he	tween	the groups \rightarrow con	tributi	on to the	

Change of entropy between the groups \rightarrow contribution to the network and phenotypic variability

GOAL

To identify patterns among the modifier genes within communities that influence the severity of cystic fibrosis

METHODS

- 1. Gene expression data is collected from CF primary nasal epithelial cells for the Δ F508/ Δ F508 genotype with cystic fibrosis lung disease and controls
- 2. Self-supervised machine learning is applied
- 3. 10 RT² Profiler PCR Arrays (commercially available) to identify relevant inflammatory genes (total n=637 genes)
- 4. The top genes involved in the inflammatory networks in control, mild CF and severe CF are identified
- 5. Gene expression is confirmed using quantitative PCR
- 6. Bioinformatical analyses to gain further knowledge into the 'known' connection of these genes using pathways analyses g:profiler

CONCLUSION						
	Network interdependence	Contribution to phenotypic variability	Epithelial gene expression			
LR5	High in mild CF	yes	Low but variable			



- During iENGAGE we generated networks and links with students from all around the world, worked in groups on various projects and presented our results by the end of the program.
 - The program aided in enhancing our presentation skills, scientific writing, critical thinking and teamwork spirit.

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