Characterizing the Composition of the Corneal Limbal Epithelial **Stem Cell Niche**

Background

Limbal epithelial stem cells (LESCs) are essential for the maintenance of the corneal epithelium and are required to regenerate the corneal epithelium post injury. The loss of LESCs or their niche can lead to limbal stem cell deficiency (LSCD). LSCD causes corneal erosions and conjunctivalization of the cornea, which leads to ocular pain and vision loss. Hyaluronan (HA) is a constituent of the LSCN that is necessary to maintain LESCs in their stem cell state.

We sought to characterize the HA-specific LSCN by identifying HA bound proteins and proteoglycans (PGs) within the limbal region of human and porcine corneas (Fig 1). We created a protocol for PG extraction withing the limbal regions and corneas in order to analyze the extracts through Western Blot and agarose gel electrophoresis (Fig 2).

Methodology

Tissue Digestion

- 4 samples (Human Limbus, Human Cornea, Pig Limbus, and Pig Cornea) were subjected to tissue digestion in a solution of 4M guanidinium chloride, 0.05M sodium acetate, pH 6 containing 2% Triton X-100.
- Scissors used to break down tissues followed by low intensity sonication. Tissues were vortexed, centrifuged once they were homogenous, and the supernatant was collected.

Buffer Exchange

- A 10mL chromatography column was packed with Sephadex G-50 resin.
- The column was equilibrated with 7M Urea.
- Samples were added in 3mL increments and washed through with Urea (flowthrough was collected).

Fast Flow Anion Exchange

- A 50mL chromatography column was packed with 8mL of Q-Sepharose resin.
- Column was equilibrated with a low salt buffer.
- Sample was added in 15mL increments and the flow through was discarded.
- High salt buffer was used to wash out the PGs and GAGs (flow through collected).

Desalting

- Samples were put in snakeskin dialysis bags and placed into a 1L beaker filled with DI water.
- Water was changed twice a day and this process was repeated 3 days, allowing the samples to desalt.
- Samples were then concentrated through lyophilization.

PG Analysis

- Samples were quantified using a BCA kit.
- Western Blot was done using an approximated 20-50 ug of protein.
- Agarose gel electrophoresis also performed.

Arian Parsaie, Isabel Moreno, Mingxia Sun, Nadine Mutoji, and Vivien Coulson-Thomas

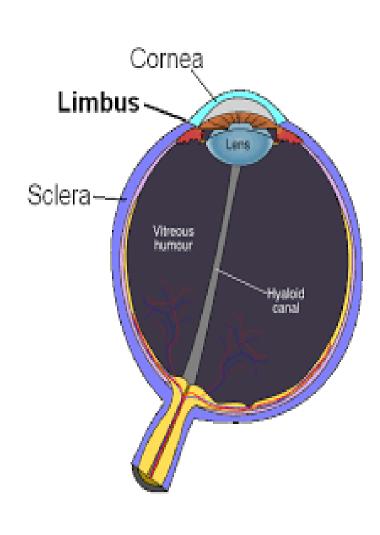


Figure 1. Anatomy of an eye distinguishing the limbus and cornea

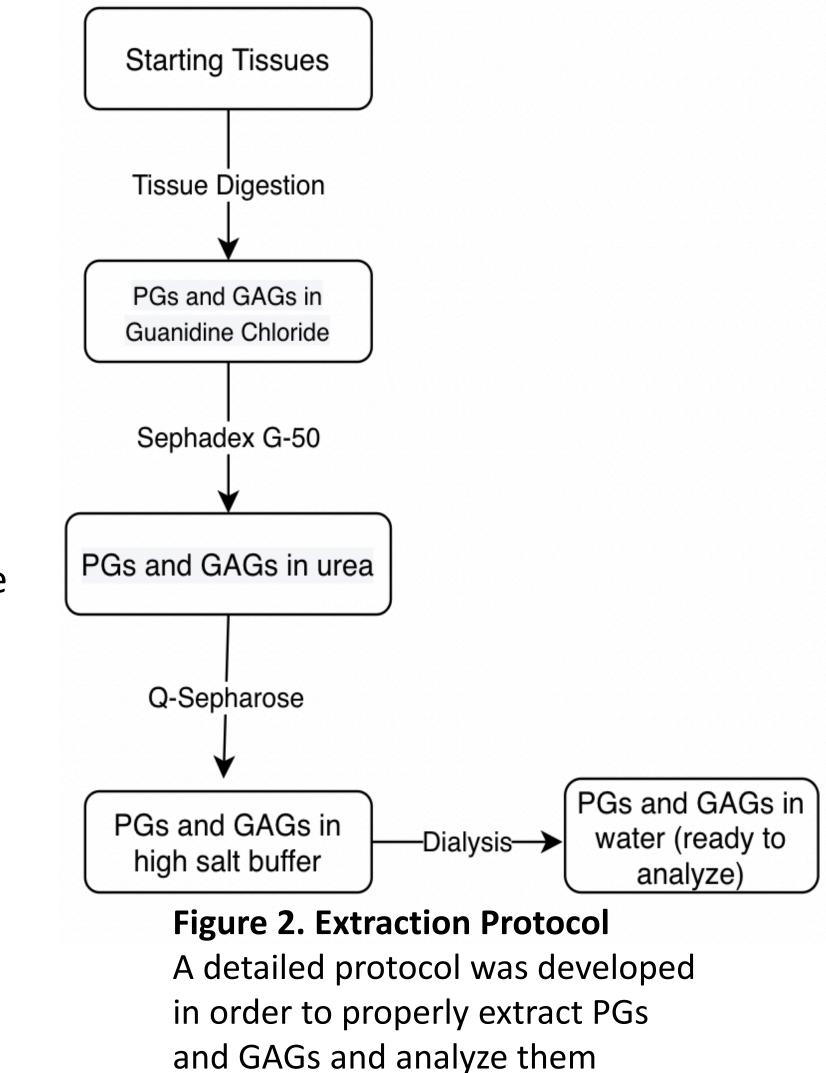
Wet Weight

Samples

Human Cornea 80 mg Human Limbus 90 mg Pig Cornea 830 mg Pig Limbus 958 mg

Table 1. Wet Weights

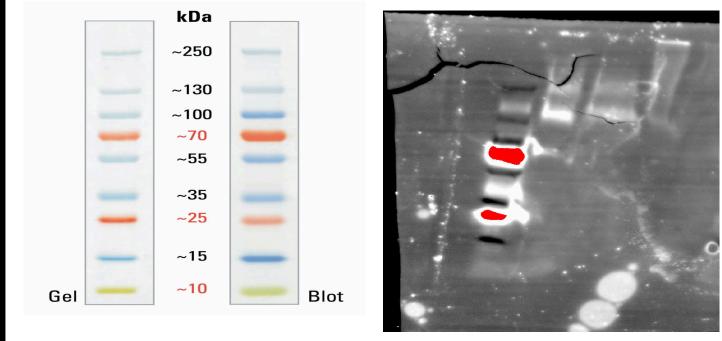
4 sample tissues were obtained, and the mass was recorded.



UNIVERSITY of HOUSTON

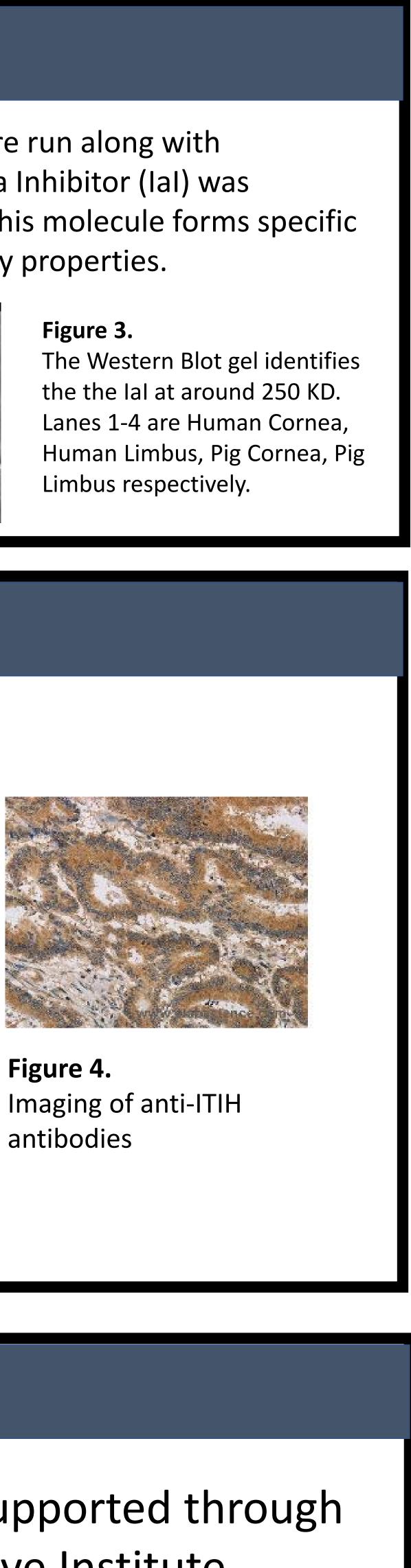
Results

After several Western Blot gels were run along with immunohistochemistry, Inter-Alpha Inhibitor (Ial) was identified in our samples (Fig. 3). This molecule forms specific HA matrices with anti-inflammatory properties.



Conclusion

With the identification of the lal (Fig. 4), future studies will work on characterizing the role of this HA/Ial and HA/TSG-6/lal matrix within the corneal limbus. Characterizing the role of the LSCN is vital for establishing novel mechanisms for treating LSCD.



Acknowledgements

Summer fellowship supported through the University Eye Institute

Thank you, Isabel Moreno, Mingxia Sun, and Nadine Mutoji for the assistance, and Vivien Coulson-Thomas for the opportunity.

