

Supplemental Figures and Table for:

Epithelial stem cell homeostasis in Meibomian gland development, dysfunction, and dry eye disease

Edem Tchegnon^{1,2,+}, Chung-Ping Liao^{1,5,+,#}, Elnaz Ghotbi¹, Tracey Shipman¹, Yong Wang¹, Renee M. McKay¹, and Lu Q. Le^{1,2,3,4,#}

¹Department of Dermatology, ²Genetics, Development and Disease Graduate Program, ³Hamon Center for Regenerative Science and Medicine, ⁴Simmons Comprehensive Cancer Center, University of Texas Southwestern Medical Center, Dallas, TX 75390, USA. ⁵Graduate Institute of Medical Sciences, College of Medicine, Taipei Medical University, Taipei 110, Taiwan

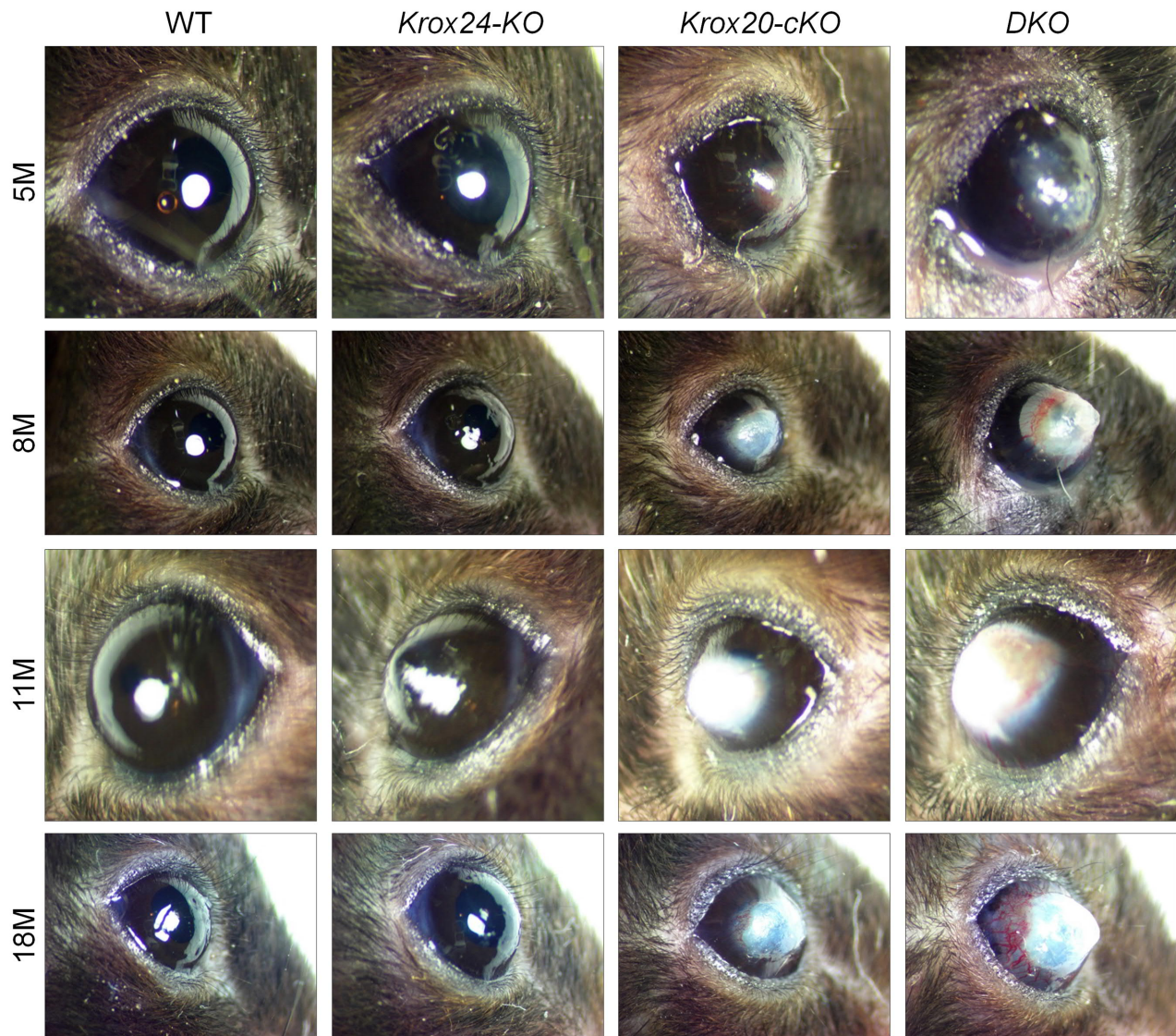
⁺ Authors contributed equally

Running title: Developmental origin of the Meibomian Gland

Keywords: Meibomian gland, Krox20, Dry Eye Disease, Meibomian gland dysfunction, Meibomian gland stem cells, Meibomian gland progenitor cells, Meibomian gland homeostasis, corneal lesions

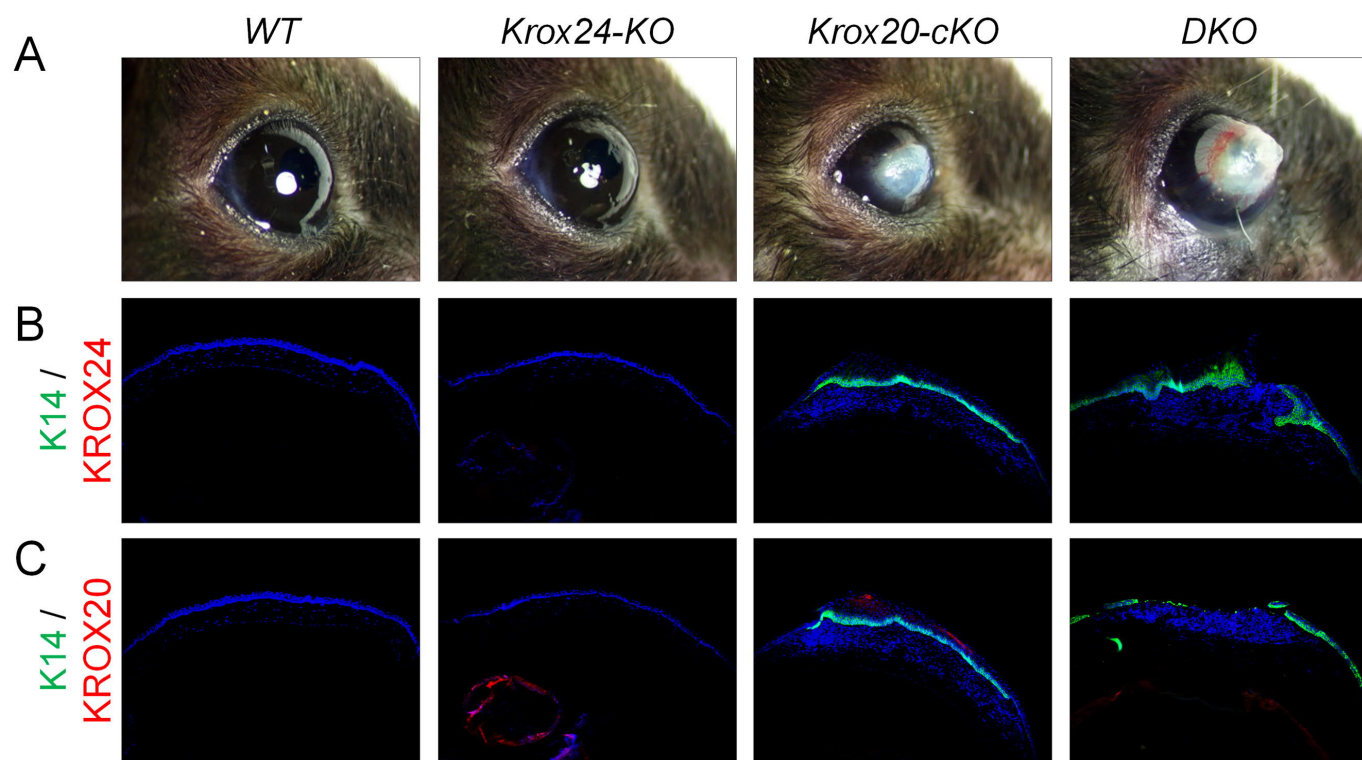
Conflict of interest: The authors have declared that no conflict of interest exists.

Supplemental Figure 1



Supplemental Figure 1. Loss of both *Krox20* and *Krox24* accelerates development and progression of corneal lesion. Gross images of the eye in wild type (WT; *Krox20*^{fl/fl}), *Krox24-KO* (*Krox24*^{-/-}), *Krox20-cKO* (*Krox20*^{fl/fl}; *K14-Cre*), and *DKO* (*Krox24*^{-/-}; *Krox20*^{fl/fl}; *K14-Cre*) mice at various ages (5, 8, 11, and 18 months), showing the progression of corneal lesion. Gross images shown for 8-month-old mice (8 M) are the same as in Figure 7 (panel **A**), Figure 8 (panel **A**), and Supplemental Figure 2 (panel **A**). n = 50 mice for each genotype. Among mice analyzed, 100% of *Krox20-cKO* and *DKO* mice developed corneal lesions. Representative images are shown. Scale bar represents 100 μ m.

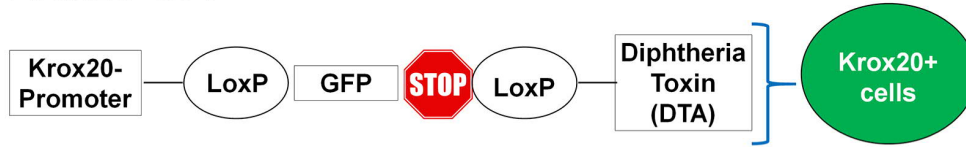
Supplemental Figure 2



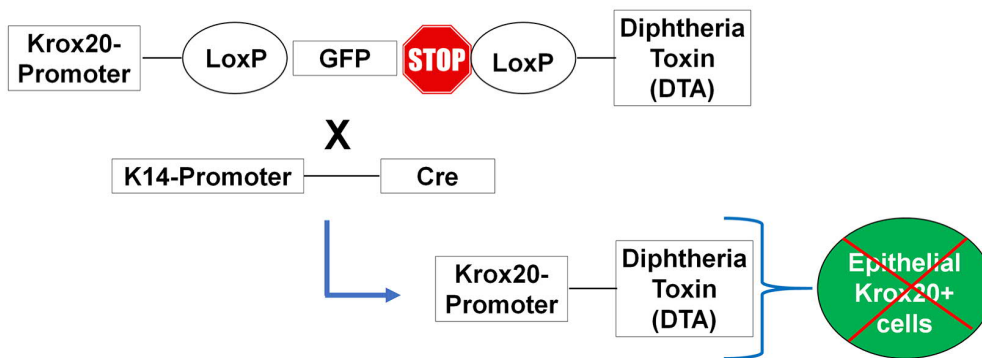
Supplemental Figure 2. Acceleration of phenotype severity is independent of KROX20 and KROX24 expression in the cornea. (A) Gross images of the eye in wild type (WT; *Krox20^{fl/fl}*), *Krox24-KO* (*Krox24^{-/-}*), *Krox20-cKO* (*Krox20^{fl/fl};K14-Cre*), and *DKO* (*Krox24^{-/-}; Krox20^{fl/fl};K14-Cre*) mice. Immunofluorescence staining for (B) K14 and KROX24, and (C) K14 and KROX20 in the indicated genotypes. n = 50 mice for each genotype. Among mice analyzed, 100% of *Krox20-cKO* and *DKO* mice developed corneal lesions. Representative images are shown. Gross images in (A) are the same as in Figure 7 (panel A), Figure 8 (panel A), and Supplemental Figure 1 (8M).

Supplemental Figure 3

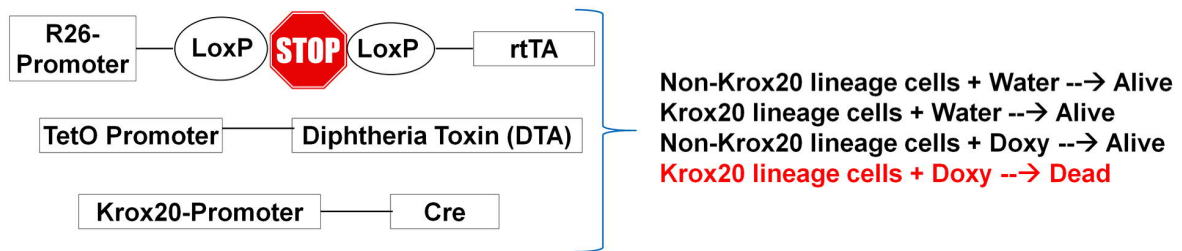
A *Krox20-GFP*



B *Krox20-DTA; K14-Cre*



C *R26-rtTA; tetO-DTA; Krox20-Cre*



Supplemental Figure 3. Schematic illustration of Krox20 cell depletion. (A) Illustration of *Krox20-GFP* (*Krox20-Flox-GFP-Flox-DTA*). In the absence of Cre expression, GFP serves as a reporter for *Krox20* expression. However, (B) when crossed with *K14-Cre* to generate *Krox20-DTA; K14-Cre*, the diphtheria toxin A (DTA) is expressed, resulting in the ablation of epithelial *Krox20*-expressing cells from K14 lineage. (C) Illustration of *R26-rtTA; tetO-DTA; Krox20-Cre⁺*. When Cre is expressed under the *Krox20* promoter, it results in the DTA-mediated death of *Krox20* lineage cells upon doxycycline (doxy) treatment.

Genotype	Primer Sequence	Size
Cre	Fwd: CAC CCT GTT ACG TAT AGC CG Rev: GAG TCA TCC TTA GCG CCG TA	Cre: 300bp
Beta-Actin	Fwd: CCT AGG CAC CAG GGT GTG AT Rev: TCA CGG TTG GCC TTA GGG TT	ActB: 239bp
GFP	Fwd: GAG CTG GAC GGC GAC GTA AAC Rev: CGT TGT GGC TGT TGT TAG TTG TAC	eGFP: 400bp
R26-TdTomato	Fwd (Tom): CTG TTC CTG TAC GGC ATG G Rev (Tom): GGC ATT AAA GCA GCG TAT CC Fwd (WT): AAG GGA GCT GCA GTG GAG TA Rev (WT): CCG AAA ATC TGT GGG AAG TC	WT: 300bp Tomato: 200bp
<u>Krox20 fl/fl</u>	Fwd (WT): GGG CTT GCA TTC TAC AGT GGT GGT C Rev (Krox20-ΔFlox): AGT TGA CAG CCC GAG TCC AGT GG Fwd (WT; Krox20-ΔFlox): GTG TCG CGC GTC AGC ATG CGT G Rev (WT): GGG AGC GAA GCT ACT CGG ATA CGG	Krox20-Flox: 195bp Krox20-ΔFlox: 210bp WT: 162bp
Egr1 +/-	Fwd (WT): AAC CGG CCC AGC AAG ACA CC Fwd (KO): CTC GTG CTT TAC GGT ATC GC Rev (Common): GGG CAC AGG GGA TGG GAA TG	WT: 414bp KO: 470bp
K14-Cre	Fwd: TTC CTC AGG AGT GTC TTC GC Rev: GTC CAT GTC CTT CCT GAA GC	K14-Cre: 494 bp
Krox20-Cre	Fwd (WT): CGC TTC CTC GTG CTT TAC GGT AT Rev (WT): TCA TCA GTC GGG TTA GAG CTG Rev (Mut): GGG CTG AGG AAG ACG ACT TTA	Krox20-Cre: 480bp WT: 312bp
TetO-DTA	Fwd: GGC GTG TAC GGT GGG AGG Rev: GGC ATT ATC CAC TTT TAG TGC	TetO-DTA: 420bp
Supplemental Table 1. Genotyping Primer Sequences		