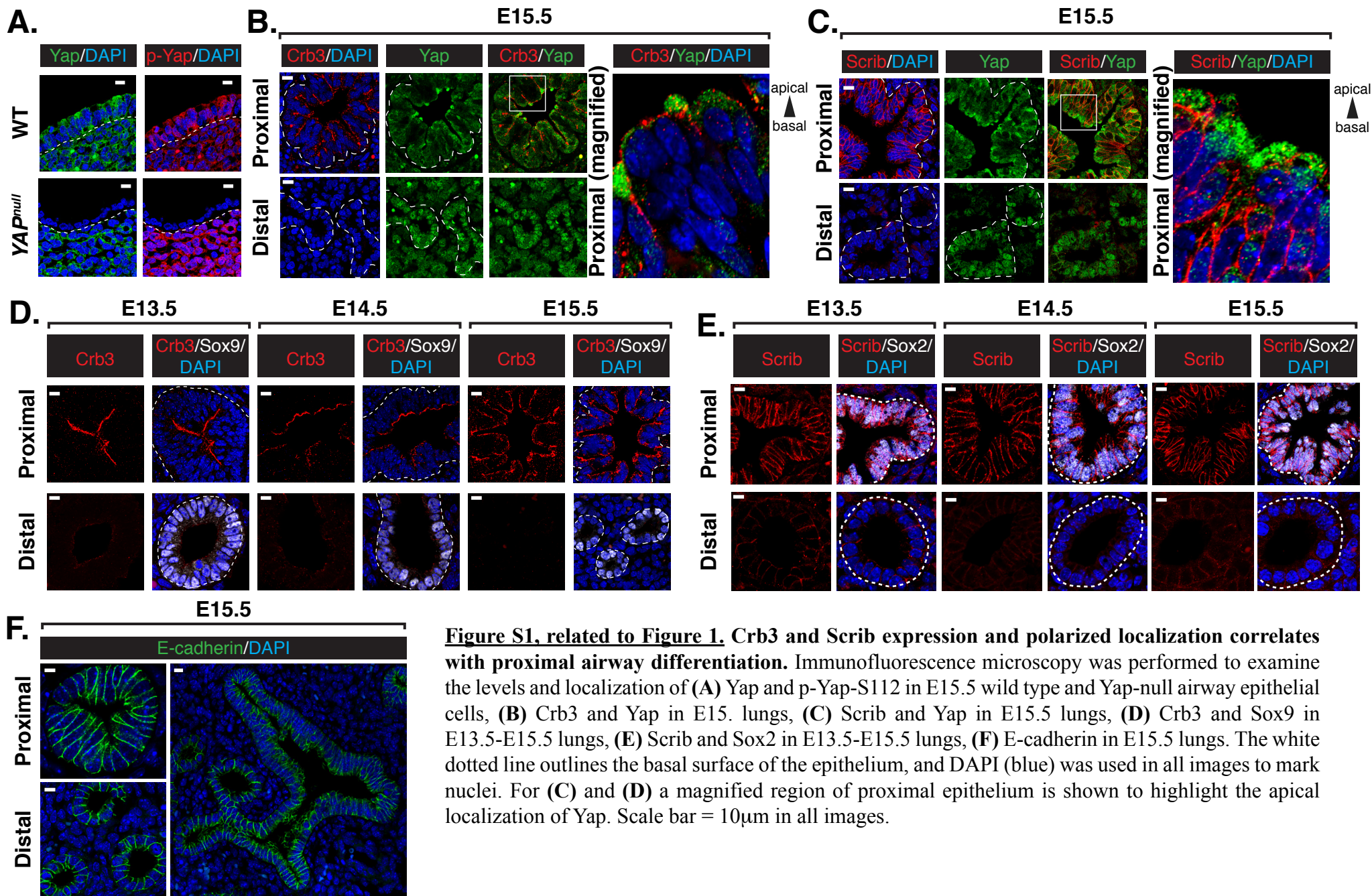


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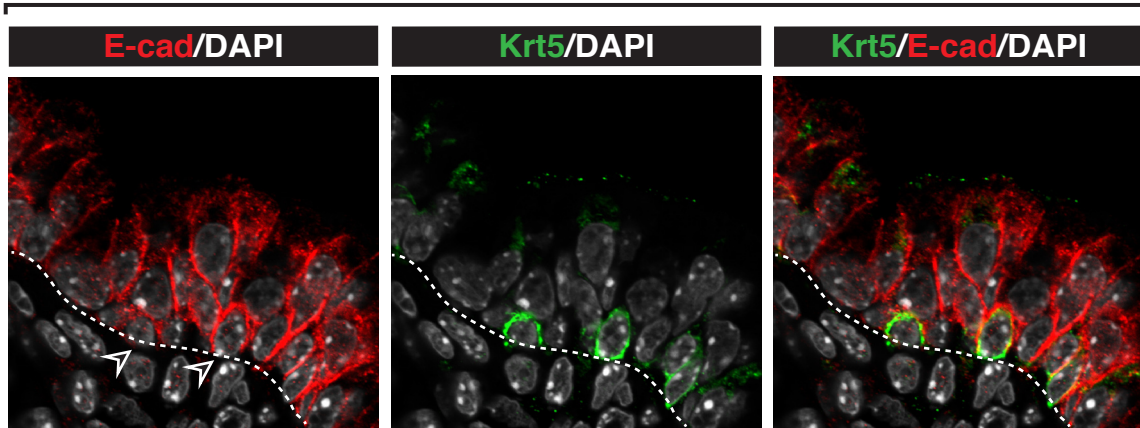
**Crumbs3-mediated polarity directs airway epithelial cell fate through the Hippo pathway  
effector Yap**

Aleksander D. Szymaniak, John E. Mahoney, Wellington V. Cardoso, Xaralabos Varelas

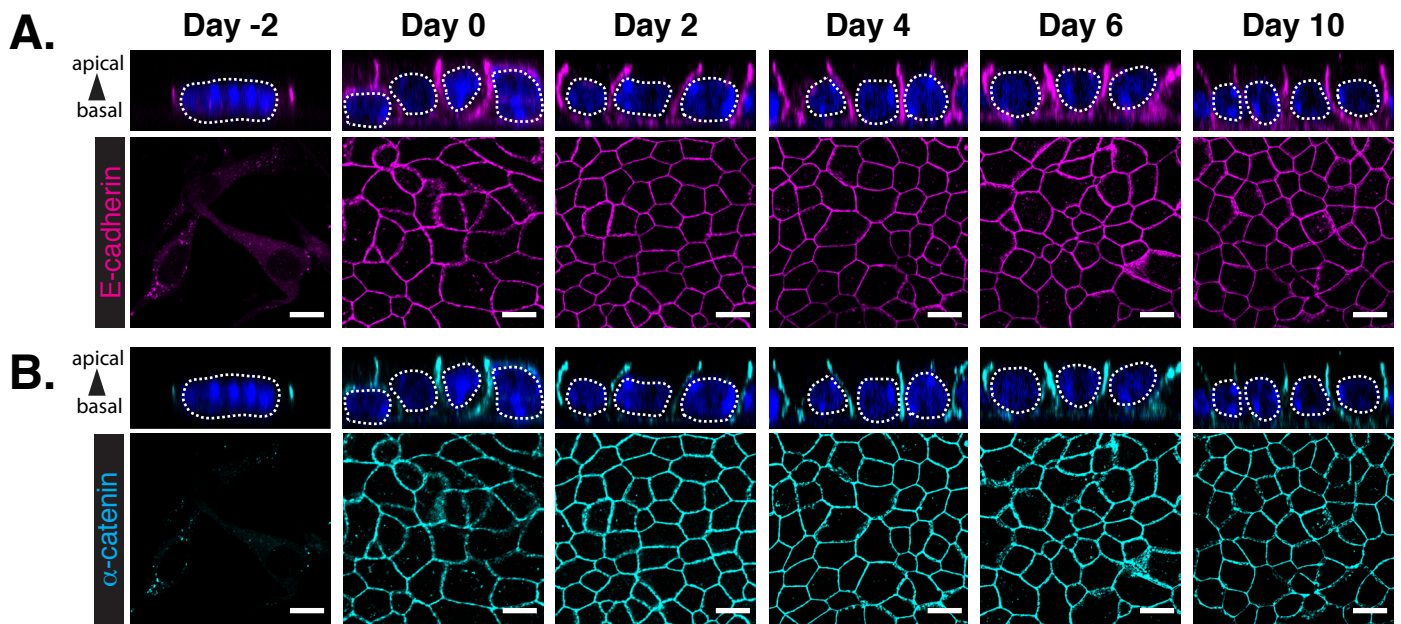


**Figure S1, related to Figure 1. Crb3 and Scrib expression and polarized localization correlates with proximal airway differentiation.** Immunofluorescence microscopy was performed to examine the levels and localization of (A) Yap and p-Yap-S112 in E15.5 wild type and Yap-null airway epithelial cells, (B) Crb3 and Yap in E15. lungs, (C) Scrib and Yap in E15.5 lungs, (D) Crb3 and Sox9 in E13.5-E15.5 lungs, (E) Scrib and Sox2 in E13.5-E15.5 lungs, (F) E-cadherin in E15.5 lungs. The white dotted line outlines the basal surface of the epithelium, and DAPI (blue) was used in all images to mark nuclei. For (C) and (D) a magnified region of proximal epithelium is shown to highlight the apical localization of Yap. Scale bar = 10µm in all images.

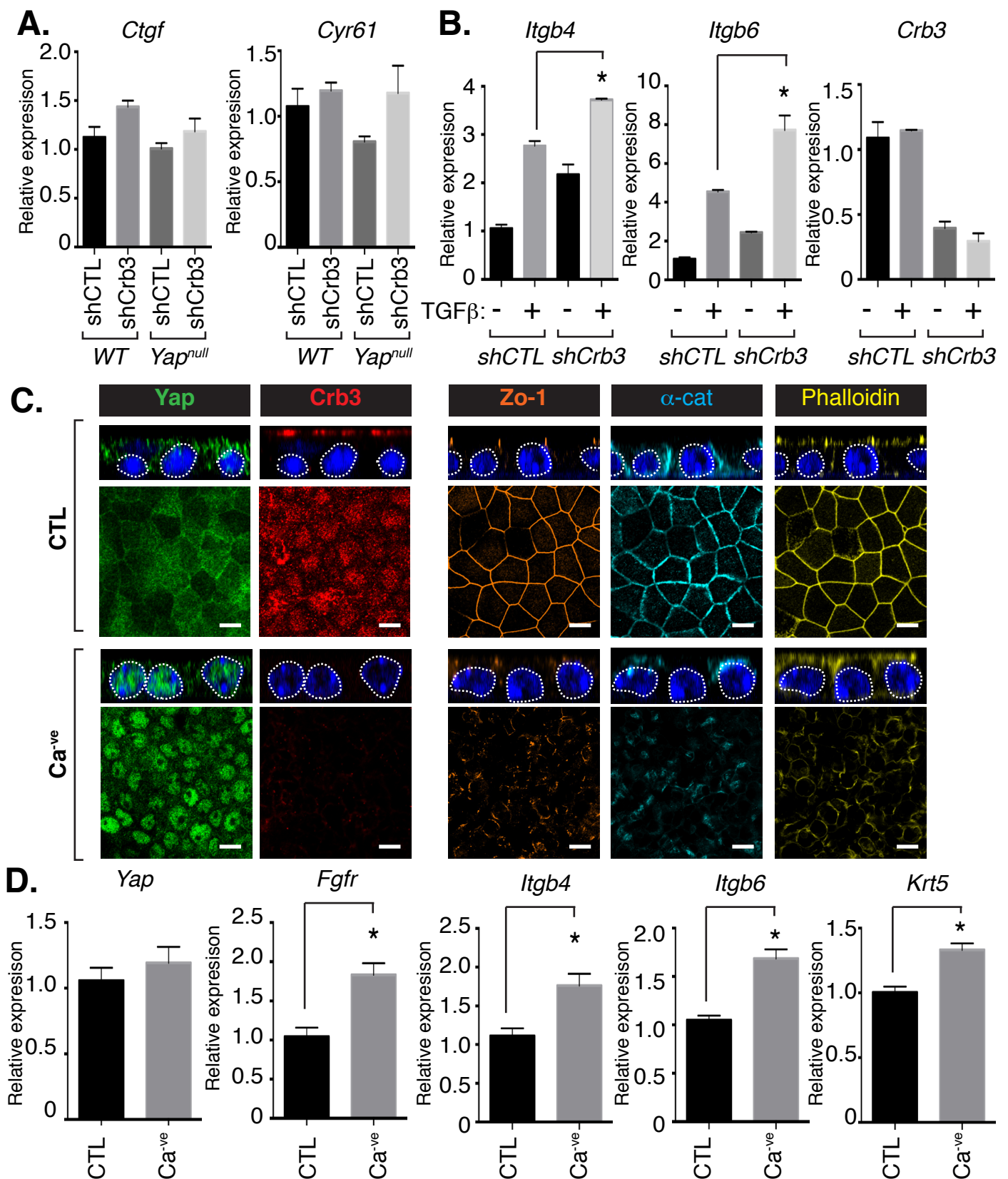
E18.5



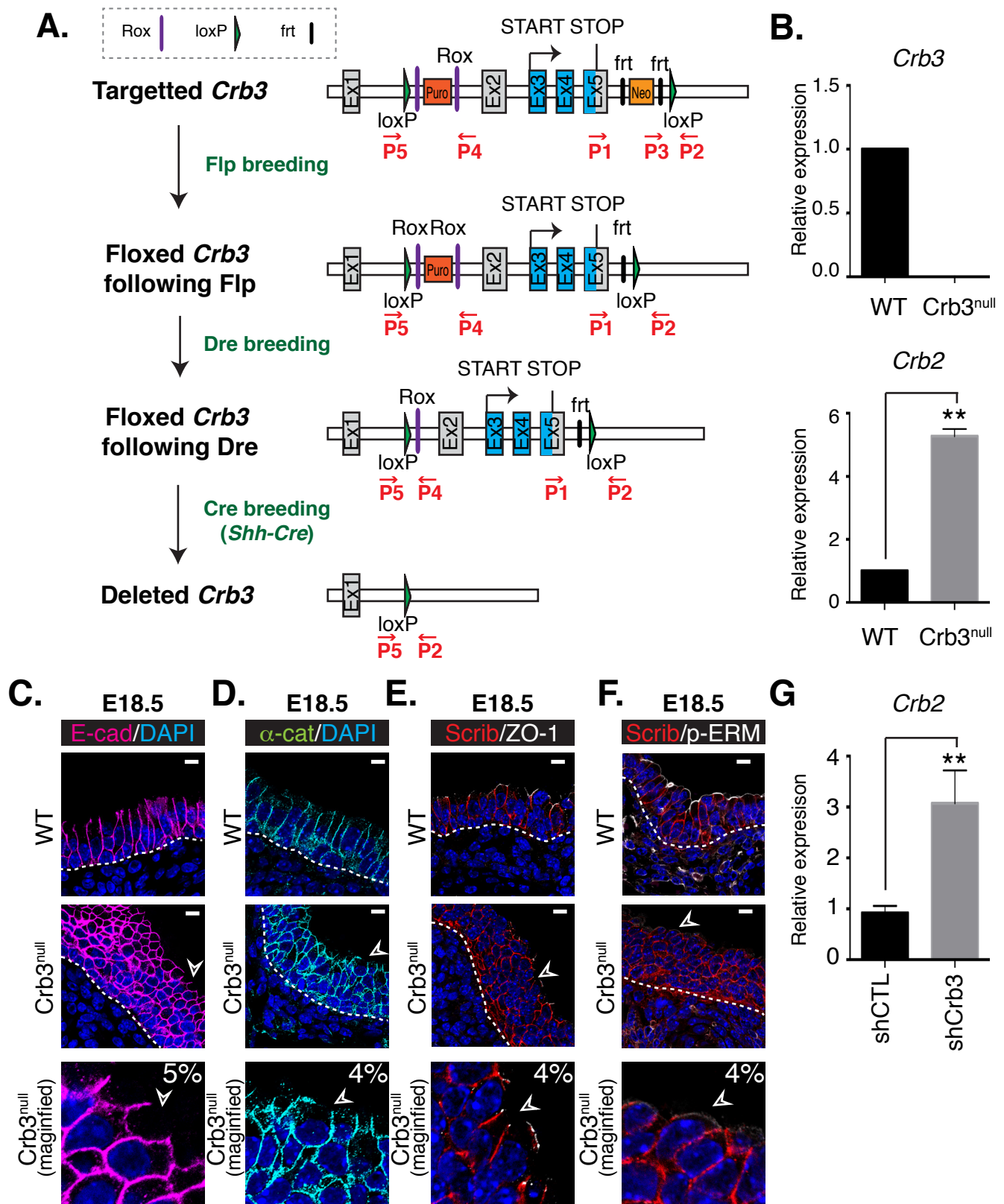
**Figure S2, related to Figure 2. Expression and localization of the adherens junction protein E-cadherin in proximal airway epithelial cells.** Similar E-cadherin (red) expression is observed between Krt5+ (green) basal progenitors and differentiated cells in developing proximal airways. Note, however, the cortical distribution of E-cadherin in the Krt5+ cells. White arrows point to the Krt5+ cells in the left image, the basal surface of the epithelium is outlined with a white dotted line, and DAPI (white) was used in all images to mark nuclei.



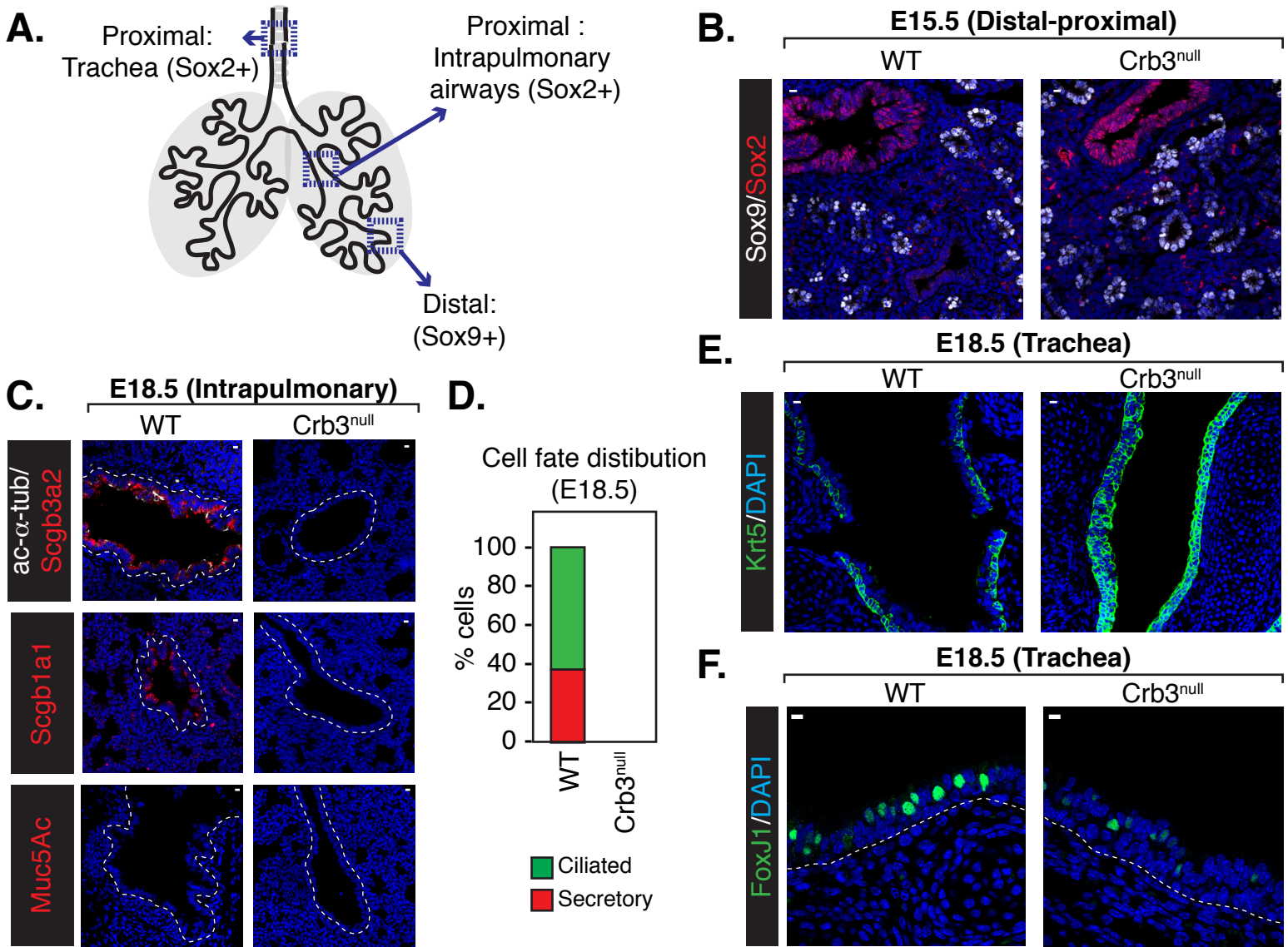
**Figure S3. Related to Figure 3. Adherens junction formation precedes airway epithelial progenitor differentiation.** Immunofluorescence confocal microscopy was performed to examine the levels and localization of (A) E-cadherin and (B)  $\alpha$ -catenin in ALI cultures. The X-Y view (bottom panels) and the Z-plane apical-basal view (top panels) for each are shown. DAPI was used to mark the nuclei (blue), which are outlined with a thin white dotted line in the Z-plane images. Scale bar = 10 $\mu$ m.



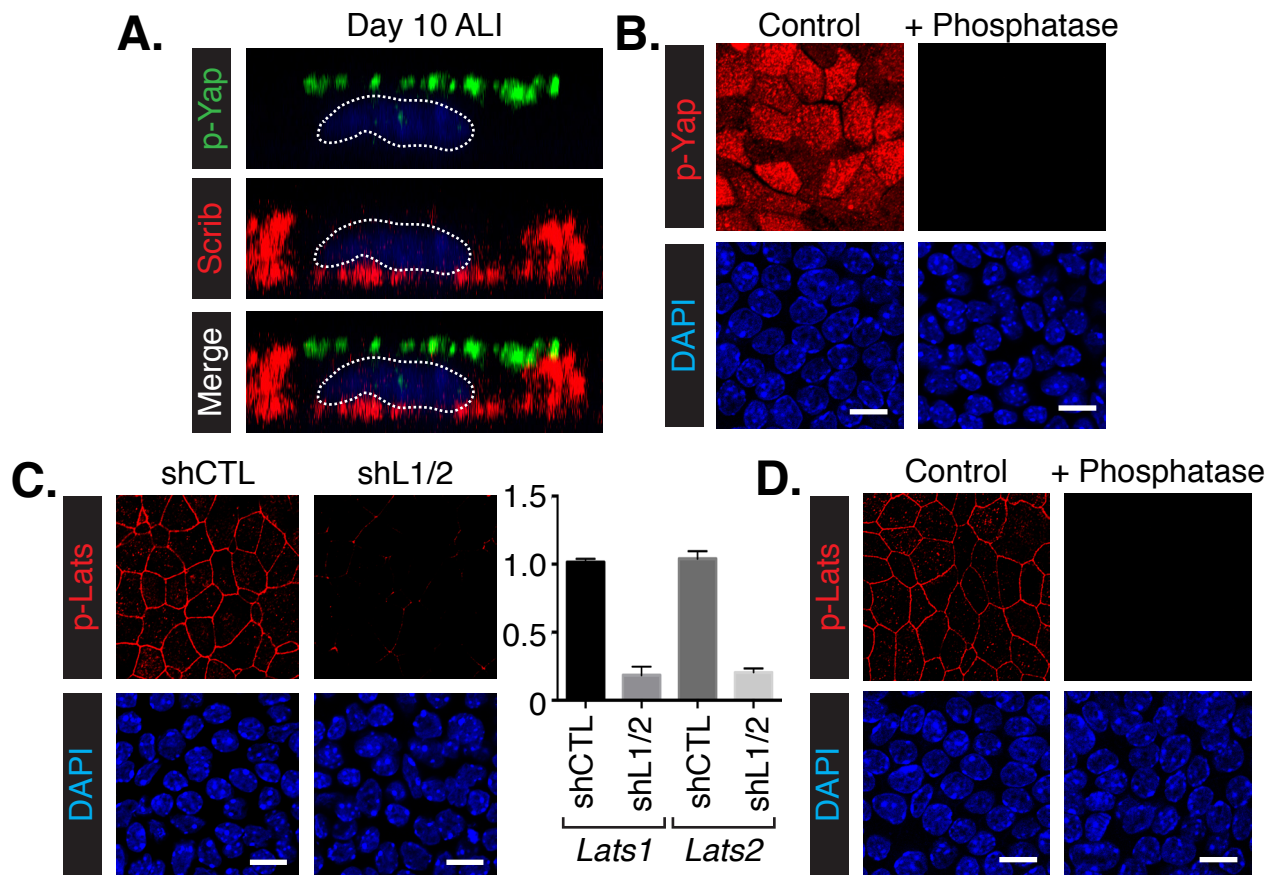
**Figure S4, Related to Figure 4.** (A) Lentiviruses were used to transduce control shRNA (shCTL) or shRNA targeting Crb3 in wild type (WT) or Yap-null airway progenitor cells, and following 6 days of ALI culture the expression of *Ctgf* and *Cyr61* was determined from these cells (same samples as in Figure 4C). (B) Airway progenitors expressing shCTL or shCrb3 were treated with or without 500pMol TGFβ for 2 hours, and the expression of the indicated Yap-p63-regulated target genes was determined (n=3, average +SEM, \* p<0.01). (C) Cadherin interactions were disrupted in differentiated airway progenitors (10 day ALI cultures) by depleting calcium from the media (*Ca*<sup>ve</sup>), and the levels and localization of the indicated proteins were determined by confocal immunofluorescence microscopy. DAPI was used to mark the nuclei (blue), which are outlined with a white dotted line in the Z-plane images. Scale bar = 10μm. (D) Cadherin interactions were disrupted in differentiated airway progenitors (10 day ALI cultures) by depleting calcium from the media (*Ca*<sup>ve</sup>) and the expression of Yap-p63-regulated target genes was determined (n=3, average +SEM, \* p<0.01).



**Figure S5, Related to Figure 5. Conditional knockout of *Crb3* in the mouse lung epithelium. (A)** Schematic of the targeting strategy used to delete the *Crb3* gene. Genotyping primer combinations are depicted as red arrows (see Supplementary Experimental Procedures for further details). **(B)** Wild type or *Crb3*<sup>null</sup> tissues from E15.5 mice were examined by quantitative real-time PCR for *Crb3* and *Crb2* expression levels (n=3, average +SEM, \*\*p<0.0001). **(C, D, E, F)** E18.5 wild type (WT) or *Crb3*-null proximal airways were analyzed by immunofluorescence microscopy for the localization of **(C)** E-cadherin, **(D)** α-catenin, **(E)** Scrib and ZO-1, and **(F)** Scrib and p-ERM. An arrow marks cells that appeared to have an apical domain, and the quantitated percentage of these cells is shown on the top right of the panels (n=400 cells over three experiments). The basal surface of the epithelium is outlined with a thick white dotted line, and DAPI (blue) was used to mark nuclei. Scale bar = 10μm. **(G)** *Crb2* levels were determined by qPCR in *Crb3* knockdown ALI cultures. (n=3, average +SEM, \*\*p<0.0001)



**Figure S6. Related to Figure 6. Knockout of Crb3 results in airway epithelium cell fate defects.** (A) Illustration depicting the different regions examined for cell fate defects in the Crb3-null lung epithelium. (B) E15.5 wild type (WT) or Crb3-null lungs were analyzed by immunofluorescence microscopy for Sox9 and Sox2 expression. (C) The intrapulmonary regions of E18.5 WT or Crb3-null proximal airways were analyzed by immunofluorescence microscopy for the presence of cells expressing Scgb1a1, Scgb3a2, Muc5A, or apical acetylated  $\alpha$ -tubulin, and (D) based on these markers the fate of the epithelial cells was quantitated (n=800 cells from 3 experiments) (E, F) The tracheas of E18.5 wild type (WT) or Crb3-null proximal airways were analyzed by immunofluorescence microscopy for the presence of (E) cells expressing the progenitor marker Krt5, and (F) the multi-ciliated cell marker FoxJ1. DAPI (blue) was used to mark nuclei in all images, and the basal epithelial surface in (C and F) is outlined with a white dotted line. Scale bar = 10 $\mu$ m.



**Figure S7, Related to Figure 7. p-YAP and p-Lats1/2 antibody specificity.** (A) p-Yap and Scrib localization was assessed by confocal immunofluorescence microscopy in a single differentiated airway epithelial cell (10 days of ALI culture). The Z-plane view is shown to highlight the apical accumulation of p-Yap and the basal-lateral localization of Scrib. DAPI marked nuclei are outlined with a white dotted line. (B) Differentiated airway epithelial cells (10 days of ALI culture) were treated with or without Antarctic phosphatase for 2 h and p-Yap levels were determined by confocal immunofluorescence microscopy. (C) Lentiviruses were used to transduce control shRNA (shCTL) or shRNA targeting Lats1 and Lats2 (shL1/2) in airway progenitor cells isolated from mouse tracheas, and following 6 days of differentiation, the levels and localization of p-Lats1/2 was determined (the X-Y view is shown), and the level of Lats1 and Lats2 knockdown was determined in parallel cultures by qPCR (n=3, average+SEM). (D) Differentiated airway epithelial cells (10 days of ALI culture) were treated with or without Antarctic phosphatase for 2 h, and p-Lats1/2 levels were assessed using immunofluorescence microscopy. The X-Y view is shown. DAPI was used to mark nuclei. Scale bar = 10 $\mu$ m in all images.



## SUPPLEMENTAL EXPERIMENTAL PROCEDURES

### Genotyping details for the *Crb3*-null mice

#### *Primers*

Primer Name	Nucleotide sequence (5'– 3')
1. Crb3-wtflpF	GAAGGGAAGAAAGTTCTTCAAAGACC
2. Crb3-R	CCTCATCCACCACTCCAAGTGACC
3. Crb3-vF	GCTGGGGCTCGACTAGAGCTTG
4. Crb3-dreR	GGAAGTGTCTGTGAGGAAGCATGG
5. Crb3-F	CAAGCTGTGCCCATACATCTGTGG

#### *Genotyping PCRs*

Primer Combinations	PCR product size (bp)	Genotype
1 & 2	236	Wild Type
2 & 3	340	Targeted Floxed <i>Crb3</i> mutant
1 & 2	601	Floxed <i>Crb3</i> mutant following Flp
4 & 5	228	Floxed <i>Crb3</i> mutant following Dre
2 & 5	351	Floxed <i>Crb3</i> mutant following Cre ( <i>Crb3</i> <sup>null</sup> )

## Antibodies used in the study

Antigen	Species	Company	Cat #	Dilution	Lot #	Use
Acetylated alpha-tubulin	Mouse	Sigma	T7415	1/1000		IF
Crb3	Rabbit	Custom	Targeted sequence: VGARAPPPNLIK LPPEERLI	1/5000	N/A	IF
CTNNA1 ( $\alpha$ -catenin)	Mouse	BD Bioscience	610194	1/100	20174	IF
Phospho-Ezrin/Radixin Moesin (41A3)	Rabbit	Cell Signaling Technologies	3149S	1/100	8	IF
FoxJ1	Mouse	eBioscience	14-9965-80	1/300	E10108-1630	IF
GAPDH	Rabbit	Sigma	G8795	1/5000		IB
Ki67	Mouse	BD Bioscience	550609	1/100	38	IF
Krt5	Rabbit	Covance	PRB-160P-100	1/300	D12KF03035	IF
Krt8	Rat	Developmental Studies Hybridoma Bank	TROMA-1c	1/500	42004	IF
p63	Mouse	Santa Cruz	8431	1/50	I1112 & D0412	IF
Phospho-LATS1 T1079	Rabbit	Cell Signaling Technologies	9159S	1/100	1	IB
Phospho-LATS1/2 (T1079/T1041)	Rabbit	Assay BioTech	A8125	1/100	118125	IF, PLA
Scgb1a1	Goat	Santa Cruz	9772	1/300	C2913	IF
Scgb3a2	Rabbit	Custom antibody	N/A	1/5000	N/A	IF
Scrib	Goat	Santa Cruz	11049	1/100	L0513	IF
Sox2	Rabbit	Seven Hills	WRAB-1236	1/500	1013A	IF
Sox9	Goat	R&D Systems	AF3075	1/100	WIL0413111	IF
Yap	Mouse	Santa Cruz	101199	1/100	B2713 & A0512	IF, PLA
Yap-p-S127 (recognizes mouse p-S112)	Rabbit	Cell Signaling Technologies	13008S	1/100	1 & 2	IF
Zo-1	Rabbit	Zymed	61-7300	1/100		IF

## Primers used for SYBR Green RT-qPCR

Target	Direction	Sequence
<i>Gapdh</i>	Forward	TGTTCCCTACCCCCAATGTGT
<i>Gapdh</i>	Reverse	GGTCCTCAGTGTAGCCCAAG
<i>Yap</i>	Forward	AATGTGGACCTTGGCACACT
<i>Yap</i>	Reverse	ACTCCACGTCCAAGATTTTCG
<i>Crb2</i>	Forward	GATCCTAACAGCTTCCGTTGC
<i>Crb2</i>	Reverse	GCACTCGTAGTGATCTGCCA
<i>Crb3</i>	Forward	CCGGACCCTTTCACAAATAG
<i>Crb3</i>	Reverse	TCGCATGAGCAGAAACAGTC
<i>Lats1</i>	Forward	GCGATGTCTAGCCCATTCTC
<i>Lats1</i>	Reverse	GGTTGTCCCACCAACATTTTC
<i>Lats2</i>	Forward	ACAGAGACGCAGCTGAAGGT
<i>Lats2</i>	Reverse	CACAGCTTCGTGATGAGGTC
<i>Ctgf</i>	Forward	GGGCCTCTTCTGCGATTTTC
<i>Ctgf</i>	Reverse	ATCCAGGCAAGTGCATTGGTA
<i>Cyr61</i>	Forward	TAAGGTCTGCGCTAAACAACCTC
<i>Cyr61</i>	Reverse	CAGATCCCTTTCAGAGCGGT
<i>Fgfr</i>	Forward	CCTCGATGTGCGTTGAACGGTC
<i>Fgfr</i>	Reverse	CAGCATCCATCTCCGTCACA
<i>Itga6</i>	Forward	GCTGTTCTTGCCGGGATTCT
<i>Itga6</i>	Reverse	AGTATGGATCTCAGCCTTGT
<i>Itgβ4</i>	Forward	ACGATTGCCCTTTAAAGTC
<i>Itgβ4</i>	Reverse	GCAACAGGAGGAAGATGAGC