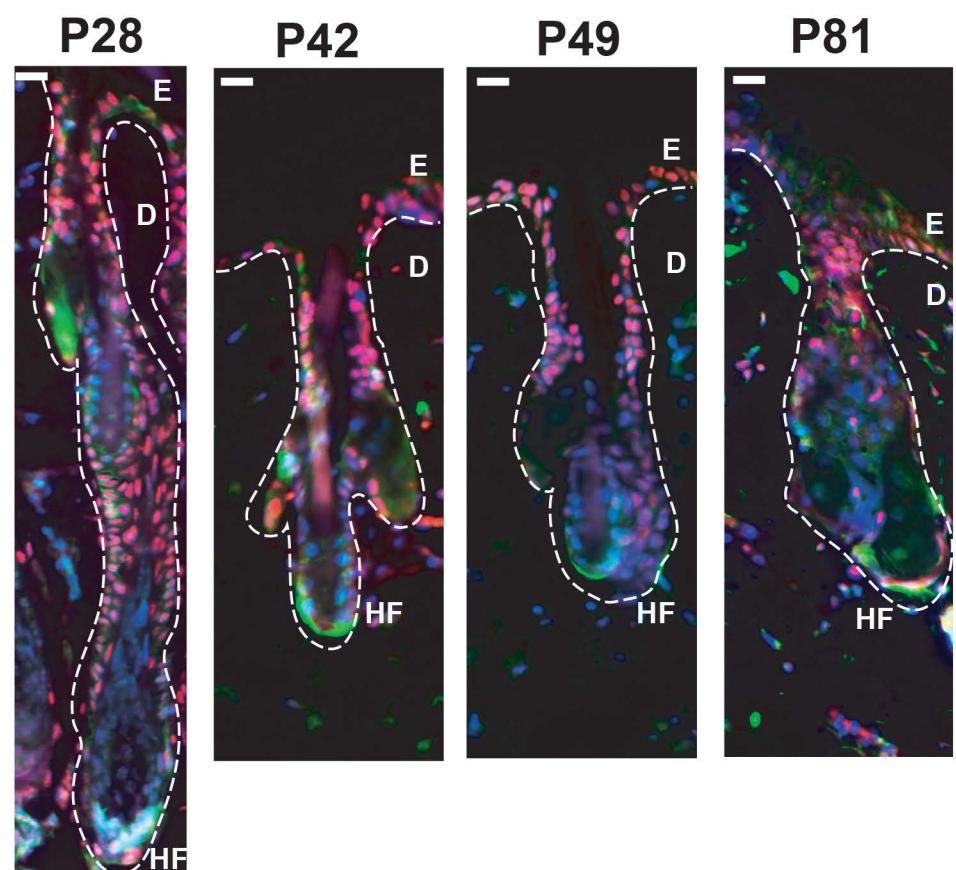
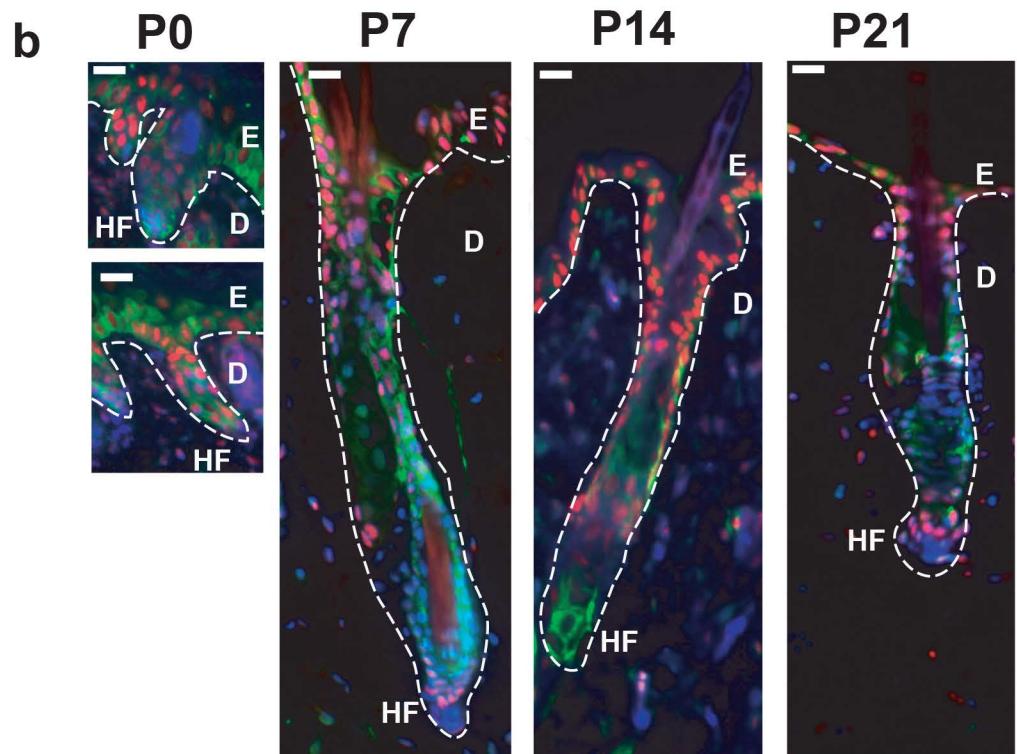
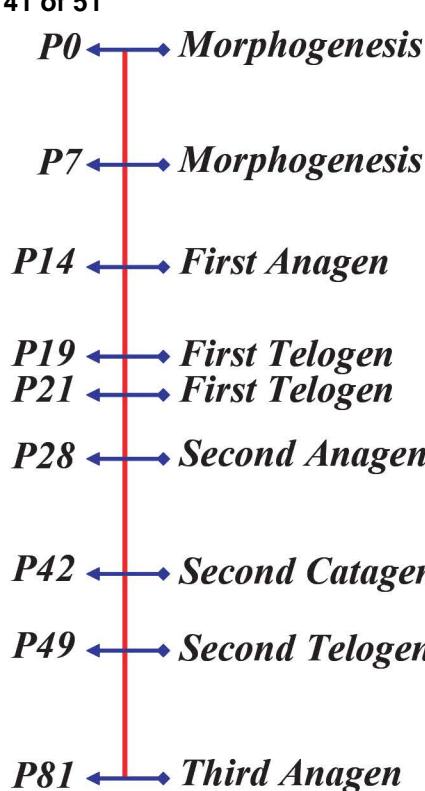


Supplementary Figure 1. Histological analysis and expression of factors involved in HF morphogenesis in *Ctip2*-null skin

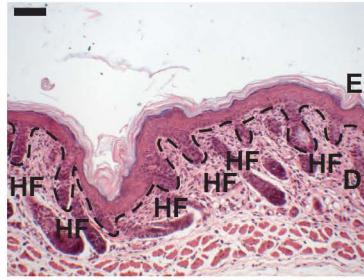
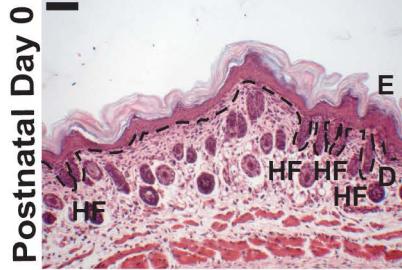
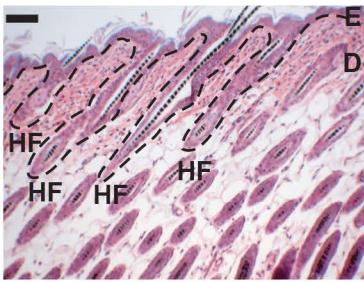
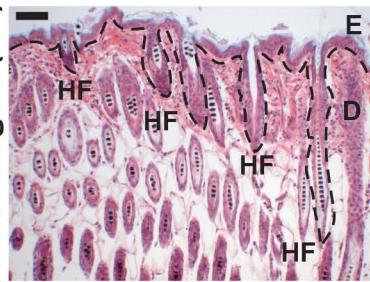
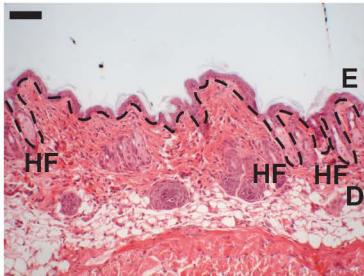
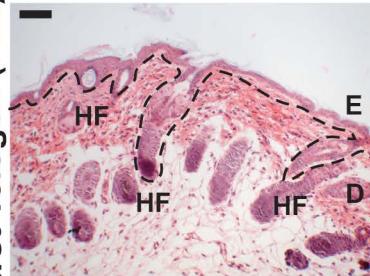


CTIP2/K15/ DAPI

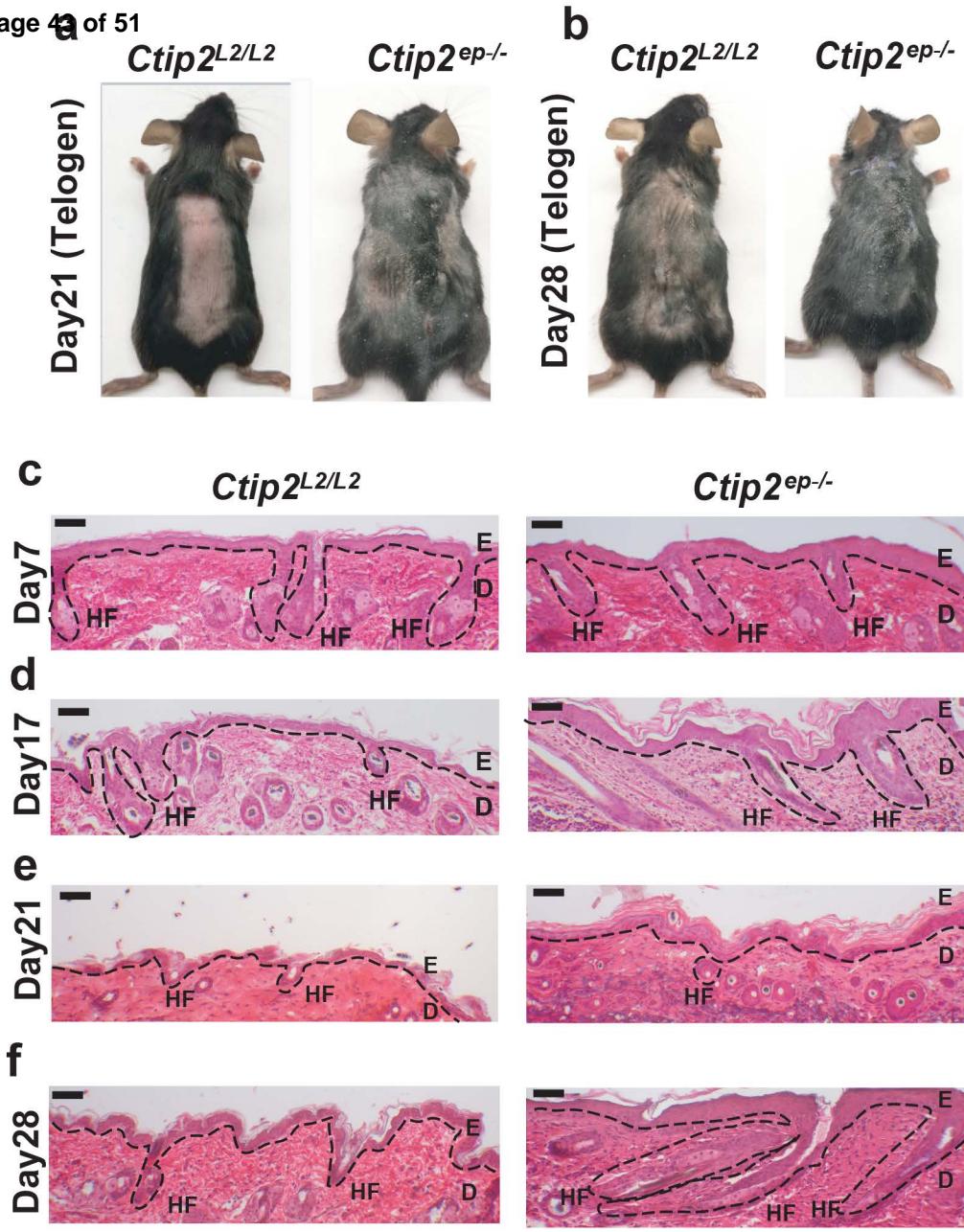
Supplementary Figure 2. CTIP2 expression during follicular morphogenesis and hair cycling

a*Ctip2*^{L2/L2}*Ctip2*^{ep-/-}

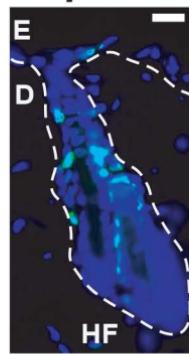
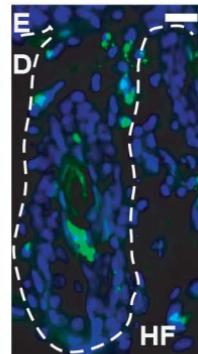
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**b****c**

Supplementary Figure 3. Hair follicle defects were not observed during follicular establishment and during first hair cycling phase in *Ctip2*^{ep-/-} mice

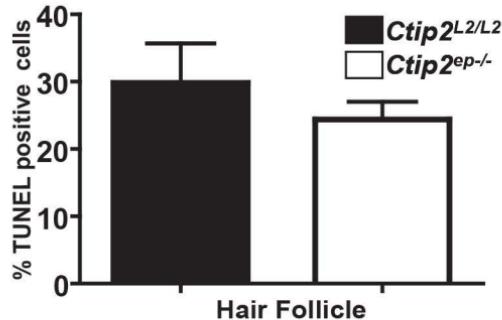


Supplementary Figure 4. *Ctip2* has a role in depilation induced hair cycling in adult mice

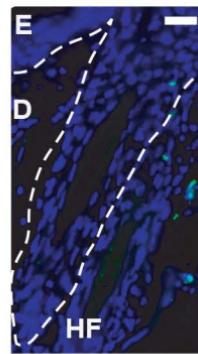
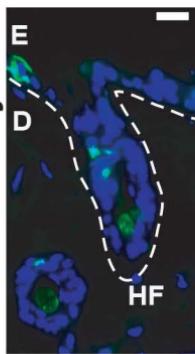
a *Ctip2^{L2/L2}**Ctip2^{ep-/-}*

TUNEL/DAPI

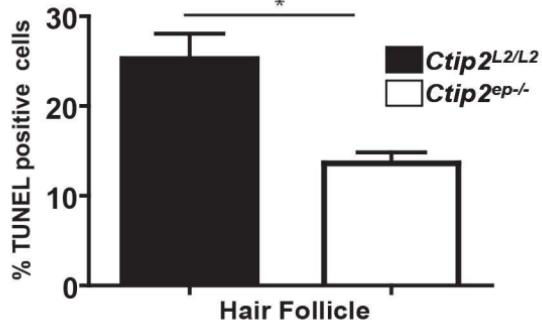
Postnatal day 42

b

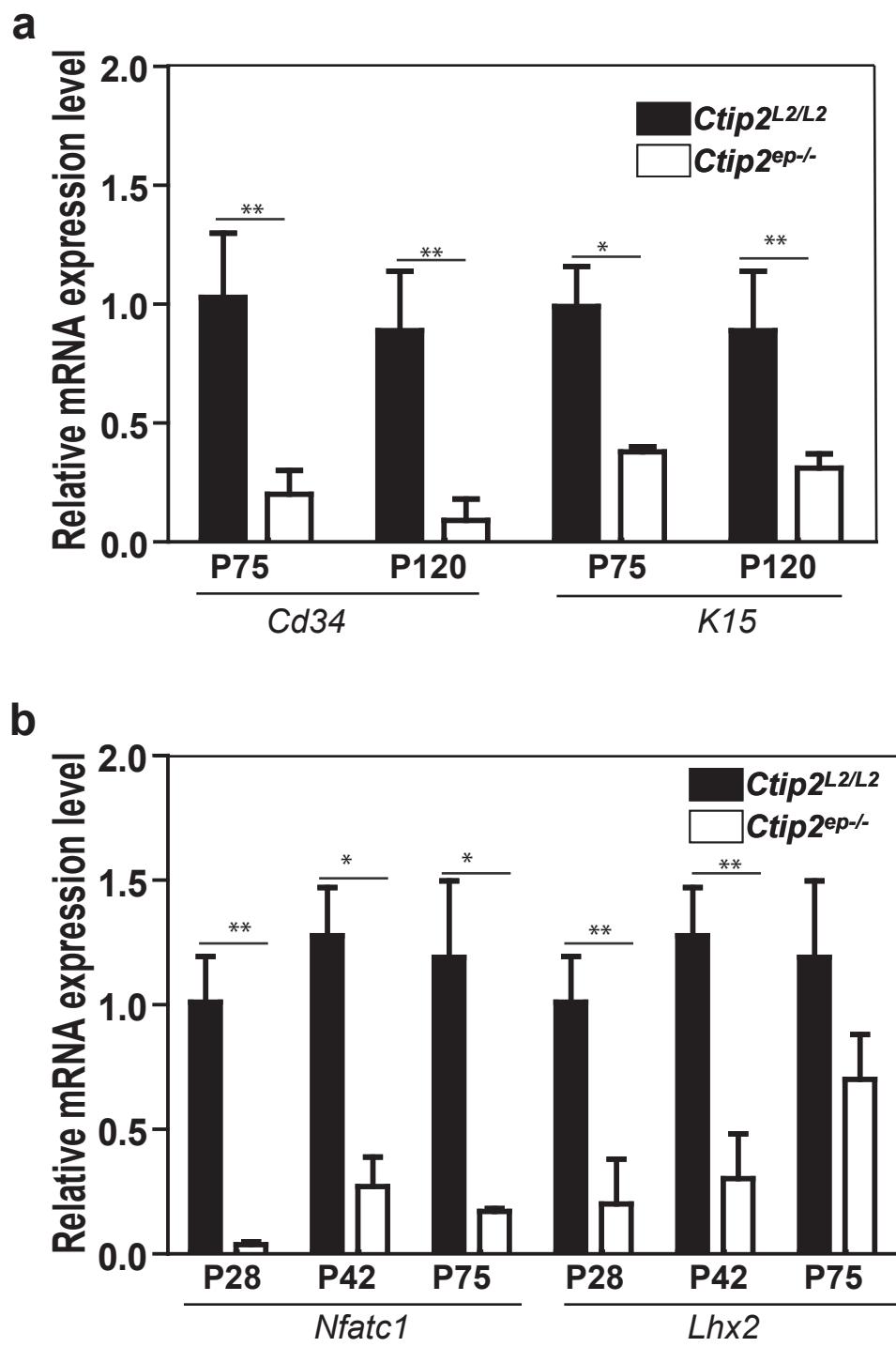
Postnatal day 75

c

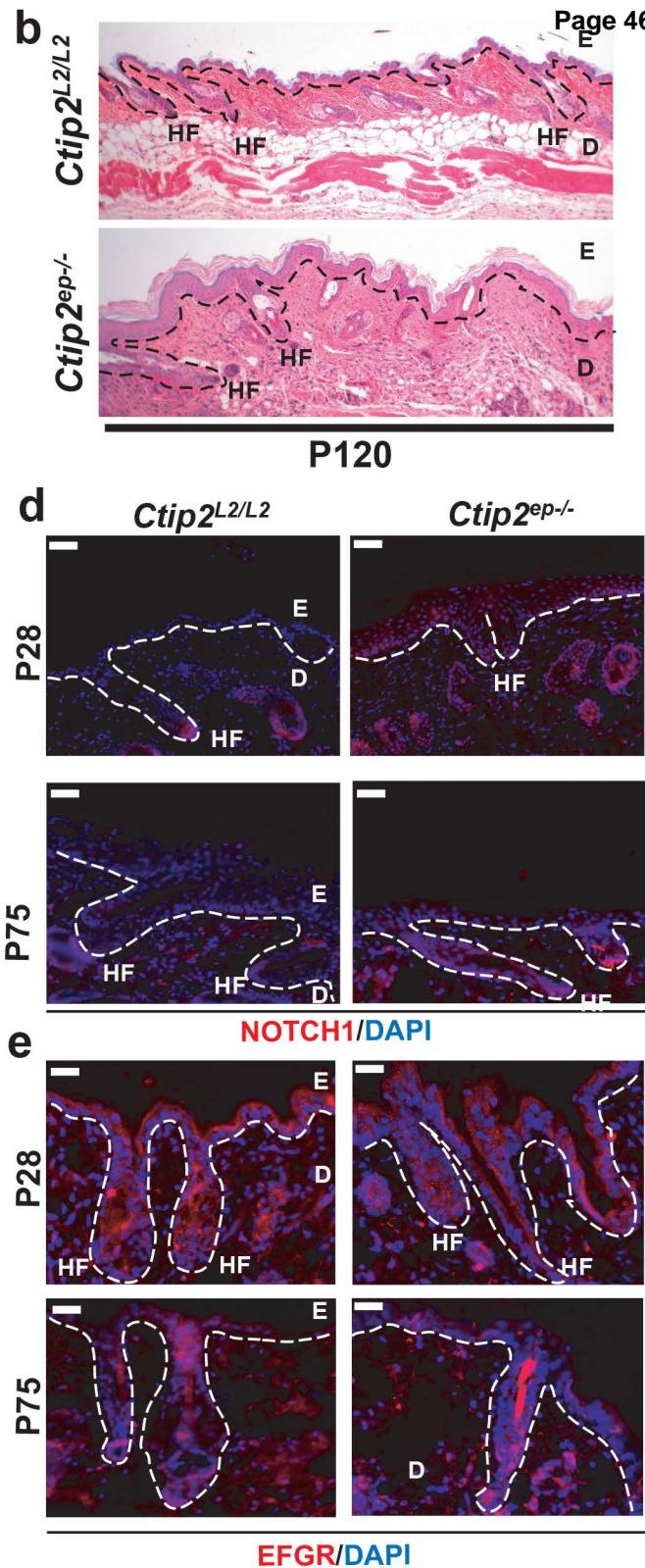
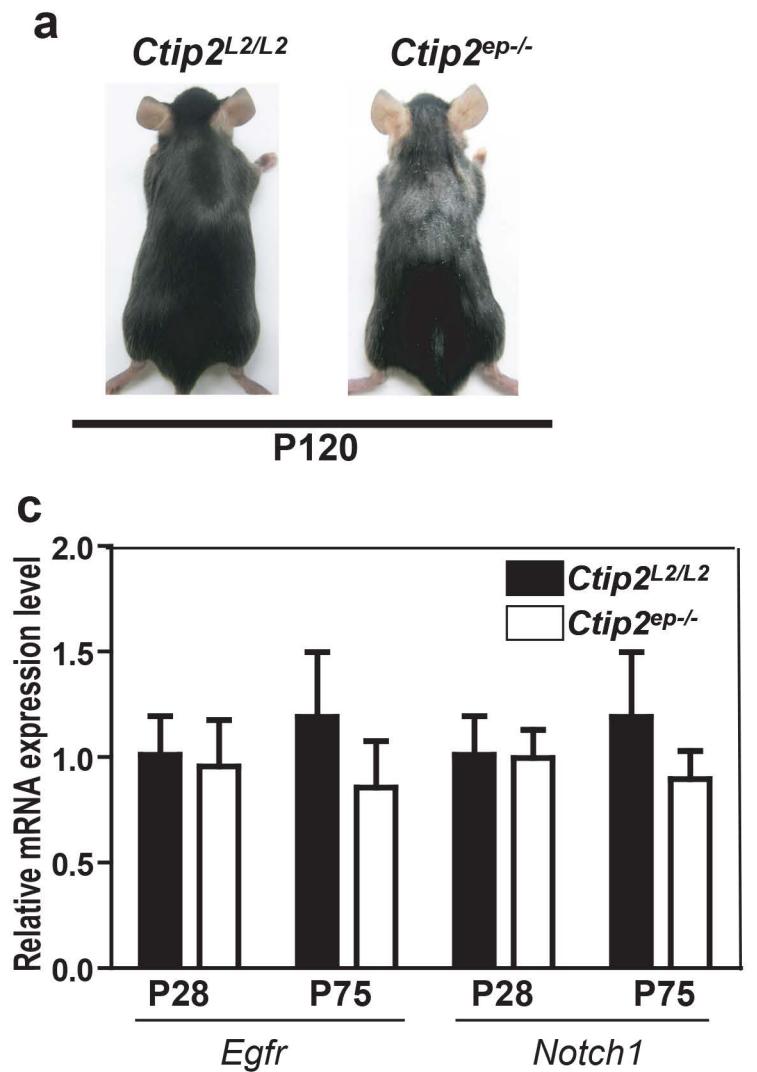
TUNEL/DAPI

d

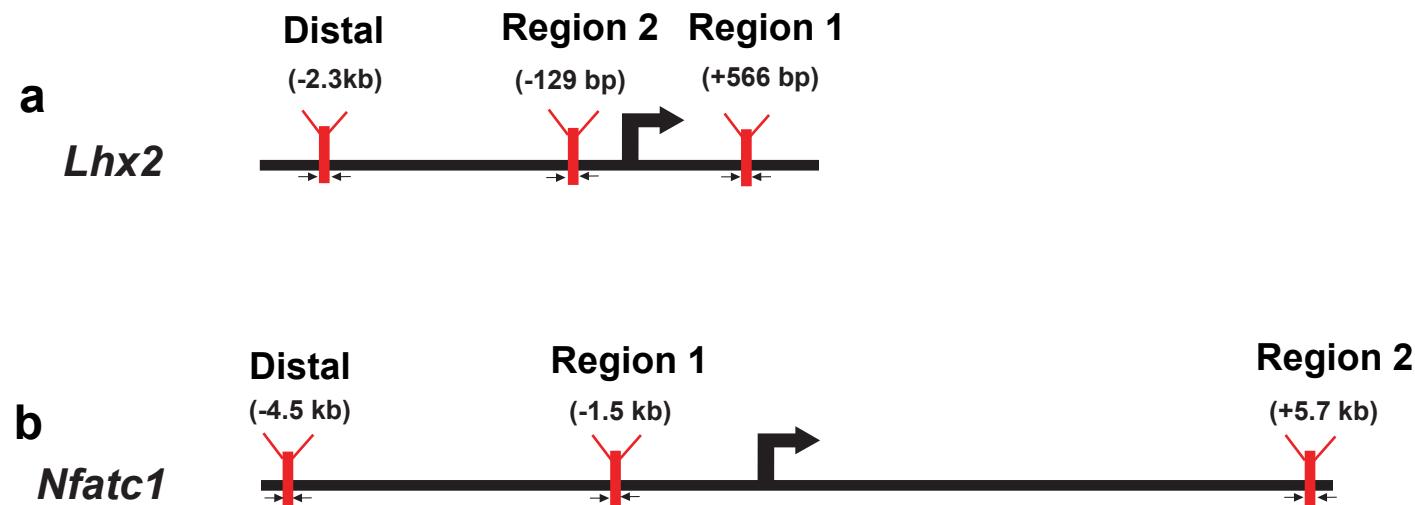
Supplementary Figure 5. Ablation of Ctip2 from epidermis and hair follicle lead to altered cell survival specifically in HF



Supplementary Figure 6. Expression of genes encoding stem cell markers and regulators of hair cycling in adult mice skin



Supplementary Figure 7. Loss of hair coat in *Ctip2^{ep-/-}* mice and expression of factors involved in hair cycling in the *Ctip2^{ep-/-}* skin



Supplementary Figure 8. Schematic diagram indicating binding region of CTIP2 on the *Lhx2* and *Nfatc1* promoters

Table S1. List of antibodies

| Antibody | Species | Source/Company | Dilution |
|-----------------|----------------|--------------------------|-----------------|
| anti-BrdU | Rat | Serotec | 1:200 |
| anti-CTIP2 | Rat | Abcam | 1:300 |
| anti-EGFR | Rabbit | Cell Signaling | 1:200 |
| anti-NOTCH1 | Rabbit | Cell Signaling | 1:400 |
| anti- LHX2 | Goat | Santa Cruz Biotechnology | 1:100 |
| anti-NFATC1 | Mouse | Santa Cruz Biotechnology | 1:200 |
| anti-CD34 | Rabbit | Santa Cruz Biotechnology | 1:200 |
| anti-K15 | Chicken | Covance | 1:500 |
| anti-K6hf | Guinea Pig | Progen | 1:200 |

Table S2. List of Primers for RT-qPCR

| Gene | Strand | Primer Sequences |
|---------------|---------------|----------------------------------|
| <i>Bmp2</i> | Forward | 5' – GGACCCGCTGTCTTCTAGTG – 3' |
| | Reverse | 5' – GTCTCTGCTTCAGGCCAAC – 3' |
| <i>Cd34</i> | Forward | 5' – AGACTCAGGGAAAGGCCAAT – 3' |
| | Reverse | 5' – TTCTGTGTCAGCCACCACAT – 3' |
| <i>Egfr</i> | Forward | 5' – GCCATCTGGGCCAAAGATAACC – 3' |
| | Reverse | 5' – GTCTTCGCATGAATAGGCCAAT – 3' |
| <i>K15</i> | Forward | 5' – AGCTTTGGTGGAGGGAGTCT – 3' |
| | Reverse | 5' – ACCAAAGCCACAACCATGC – 3' |
| <i>Lhx2</i> | Forward | 5' – CTGTTCCACAGTCTGTCGGG – 3' |
| | Reverse | 5' – CAGCAGGTAGTAGCGGTCAG – 3' |
| <i>Nfatc1</i> | Forward | 5' – GCACATTGAGTCCGTGATG – 3' |
| | Reverse | 5' – GCAGAGCAAATGACTGTGGA – 3' |
| <i>Notch1</i> | Forward | 5' – TCAATGCCGTGGATGACCTA – 3' |
| | Reverse | 5' – CCTTGTGGCTCCGTTCTTC – 3' |
| <i>Shh</i> | Forward | 5' – AAGCAGGTTTCGACTGGTC – 3' |
| | Reverse | 5' – CCGGGACGTAAGTCCTTCAC – 3' |
| <i>Sox9</i> | Forward | 5' – ATAAGTTCCCCGTGTGCATC – 3' |
| | Reverse | 5' – TACTGGTCTGCCAGCTTCCT – 3' |
| <i>Tcf3</i> | Forward | 5' – GTCACCACAGGGCTGTCTCT – 3' |
| | Reverse | 5' – CATGCTGAAGTCCAGGAGGT – 3' |
| <i>Wnt10b</i> | Forward | 5' – CCTTAAACTTCCCAGGTGAGC – 3' |
| | Reverse | 5' – TGGTGCTGACACTCGTGAAC – 3' |

Table S3. List of primers used for ChIP

| Gene and location | Region | Strand | Sequence |
|-----------------------------------|----------|---------|--------------------------------|
| <i>Lhx2</i> (-2.38 to -2.28kb) | Distal | Forward | 5'– TGGTAGGGAGGTAGGCAGTG – 3' |
| | | Reverse | 5'– AGGGGTAAAGCACCCCTCAGT – 3' |
| <i>Lhx2</i> (+517 to +616bp) | Proximal | Forward | 5' – CCTACTCCAGTTGCCCTTG – 3' |
| | | Reverse | 5' – GGTCACCCAGGAACAGCTA – 3' |
| <i>Lhx2</i> (3'UTR) | 3' UTR | Forward | 5' – GGTATTGCTGCCTACCAA – 3' |
| | | Reverse | 5' – TCAAGTGCAAGACCTGGATG – 3' |
| <i>Nfact1</i> (-4.6 to -4.5kb) | Distal | Forward | 5' – GAAAACCTTGGCAGTGAG – 3' |
| | | Reverse | 5' – GAAGCAAAAGGAAGGGAGGT – 3' |
| <i>Nfatc1</i> (-1.56 to -1.46 kb) | Proximal | Forward | 5' – CCATCCTCTCAAAGCCAGAG – 3' |
| | | Reverse | 5' – ATGTCTCAGGCGCACTACCT – 3' |
| <i>Nfact1</i> (3'UTR) | 3' UTR | Forward | 5' – TGCTGGGCAGTCTATGTGAG – 3' |
| | | Reverse | 5' – GAAACAGCCAGAGCCATGTT – 3' |

Table S4. List of primers used for Promoter Cloning for Luciferase Assay

| Gene and location | Region | Strand | Sequence |
|---|----------|---------|--|
| <i>Lhx2</i> <i>(-399 to +618bp)</i> | Proximal | Forward | 5' – AGATCTAACGCGTTGCACGTACATGGGAGAGAG – 3' |
| | | Reverse | 5'– TGAGACCTAAAGCTTGGTTCACCCAGGAACAGCTA – 3' |
| <i>Nfatec1</i> <i>(-3.2 to -1.1kb)</i> | Proximal | Forward | 5' – ACTACTAGGTACCCCCAGGTGAGGTAAACAGGAG – 3' |
| | | Reverse | 5'– CTTACATGCTAGCGGTTGGGCAGTTAGAACGTC – 3' |

Note: Black is the part of the respective gene sequence.

Green is the restriction digestion site for particular restriction enzyme.

Blue is additional nucleotide for optimal restriction digestion