



Supplementary Materials for
Foxc1 reinforces quiescence in self-renewing hair follicle stem cells

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This PDF file includes

Materials and Methods
Figs. S1 to S12
Tables S1 to S4
References

Supplementary Materials:

Materials and Methods

Mice

Mice were bred and housed according to the guidelines of IACUC at a pathogen-free facility at the University of Colorado (Boulder, CO, USA). The following mouse lines were used: *K14-Cre* (E. Fuchs, Rockefeller University), *K14-H2BGFP* (E. Fuchs, Rockefeller University), *Foxc1^{fl/fl}*(32), *Foxc1-LacZ*(9), *K15-CrePR* (Jackson lab, #005249), *Foxn1-Cre*(33) (Nancy Manley, University of Georgia) and *Rosa26-LSL-tdTomato* (Jackson lab, #021876).

K15-CrePR induction was performed by topical application of 4% RU486 (dissolved in ethanol) for 7–9 days. For anagen deletion, RU486 treatment started at P18-P20 and proceeded for 7 days or 3-4 days after the back skin turned dark color, whichever came first. For telogen deletion, RU486 treatment started after back skin turn pink/white color, which was around P37-P40 and proceeded for 7-9 days. BrdU (50 mg/g body weight) or EdU (25 mg/g body weight) was injected i.p. twice within 24h before lethal administration of CO₂. Gender matched littermates were analyzed for histology and FACS. Hair dyeing experiments were previously described(20). For counting BrdU/EdU (+) cells in Bulge+hair germ related to Fig 2G, 2H, at least 10 hair follicles with a median of 23 hair follicles were counted for each animal.

Histology and Immunofluorescence

Immunofluorescence was performed as described previously(34). The following antibodies and dilutions were used: Foxc1 (rabbit, 1:200, Cell Signaling D8A6), Keratin 5 (K5, chicken, 1:2000, Covance, SIG-3475), BrdU (1:500, rat, Abcam, ab6326), Keratin 6 (K6, Ganna Bilousova and Dennis Roop, University of Colorado Anschutz Medical Campus), CD34 (1:200, rat, eBiosciences, 13-0341), β 4-integrin (β 4, rat, 1:200, BD Biosciences, 553745) Scd1 (1:200, Santa Cruz, sc-14719), active Caspase3 (1:200, R&D Systems, AF835), P-cadherin (1:200, R&D Systems, AF761). Nuclei were stained using 4060-diamidino-2-phenylindole (DAPI).

For β -gal staining, thick sections (25 μ m) were fixed at 0.2% glutaraldehyde for 10min at 4°C and washed twice with detergent rinse, incubated in staining solution containing 1mg/ml X-gal at 37°C for 24h. Nuclear fast red was used as counter stain. Microscopy images were obtained using a Leica DM5500B microscope with either a Leica camera (bright field) or Hamamatsu C10600-10B camera (fluorescence) and processed with MetaMorph (MDS Analytical Technologies) and Fiji software.

FACS

Back skin was dissected and subcutaneous fat was removed by a scalpel. Hair cycle stage was confirmed by examining sample under dissection microscope based K14-H2BGFP-labeled hair follicle morphology, or by embedding part of the sample in OCT and examining morphology by cryosection. The remaining skin sample was minced and incubated with 0.25% collagenase (Worthington, LS004188) in 1x HBSS buffer at 37°C for 2h and then 0.05% Trypsin (GIBCO) at 37°C for 8min. Single cell suspensions were

obtained by filtration and centrifugation, and cells were incubated with the appropriate antibodies for 30-45 min on ice. DAPI was used to exclude dead cells. Hair follicle stem cells (HFSCs) of *K14Cre*-based and *Foxn1Cre*-based experiments were isolated by enriching DAPI^{neg}, K14-H2BGFP^{hi}, Sca1^{lo}, $\alpha 6^{\text{hi}}$ and CD34^{hi} cells. HFSCs of *K15CrePR*-based experiments were isolated by enriching DAPI^{neg}, K14-H2BGFP^{hi} (if available), tdtomato^{hi}, Sca1^{lo}, $\alpha 6^{\text{hi}}$ (if available) and CD34^{hi} cells. The following antibodies were used: integrin $\alpha 6$ (CD49f, 1:75; eBioscience, PE-conjugated, 12-0495; APC-conjugated, 17-0495), CD34 (1:50; eBioscience, eFluor 660-conjugated, 50-0341), Sca1 (Ly-6A/E, 1:500; eBiosciences, PerCP-Cy5.5-conjugated, 45-5981). FACS was performed on MoFlo XDP machine (Beckman Coulter). FACS data were analyzed with FlowJo.

Chromatin Immunoprecipitation

ChIP experiments are performed following previous procedures(34). Foxc1 antibody was the same antibody used in IF and rabbit IgG control was from Cell Signaling #2729.

Real-Time PCR

Total RNAs from FACS-purified cells were isolated using TRIZOL reagent (Invitrogen). 200ng of RNA (when available) was reverse-transcribed and qPCR was performed using the iQ SYBR Green Supermix system (BioRad) and BioRad CFX-384 machine. Fold-changes were computed using the $\Delta\Delta\text{Ct}$ formula normalized to Hprt values. In all qPCR figures error bars denote standard errors of the mean. All PCR primers are listed in table S4.

Statistics

The following statistical tests were performed: two-tailed Student t-test (Fig 2G, 2L, 3C, 4I, S4B, S4C, S4D); two sample Z-test (Fig 2H, S3A, S6D, S8B) or otherwise described in bioinformatics analysis. The 0.05 level of confidence was accepted for statistical significance. For all measurements, at least 3 biological replicates (gender-match, age-matched animals, usually littermates) were chosen as the minimum number necessary to achieve statistically significant differences between groups. No statistical method was used to pre-determine sample size. The experiments were not randomized. The investigators were not blinded to allocation during experiments and outcome assessment.

RNA-seq assay

Total RNAs from FACS-purified cells with RNA integrity numbers (RIN) > 8 were used to perform RNA-seq assays. Libraries were prepared using NEBNext Ultra Directional RNA Library Prep Kit and sequenced at BioFrontiers Next Generation Sequencing Facility at University of Colorado Boulder and Genomics and Microarray Core Facility at University of Colorado Denver on Illumina HiSeq 2000 and 2500 machines.

ATAC-seq assay

Assay for transposase accessible chromatin followed by sequencing (ATAC-seq) was performed as previously described (17) with following modifications: An average of 100,000 FACS-sorted HFSCs were collected into PBS containing 3% chelated fetal bovine serum (FBS) and pelleted by centrifugation for 5 min at 500g at 4°C. Cell pellets were re-suspended in 50 µl of lysis buffer per 100,000 cells processed (10 mM Tris-HCl pH 7.4,

10mM NaCl, 3mM MgCl₂, 0.1% Igepal CA-630) and nuclei were pelleted by centrifugation for 25min at 500g, 4°C using a swing rotor with low acceleration and brake settings. Supernatant was carefully discarded and nuclei were re-suspended in 50 ul reaction buffer containing 2ul of Tn5 transposase and 22.5ul of TD buffer (Nextera DNA Sample Preparation Kit, Illumina). The reaction was incubated at 37°C for 30min and terminated by adding 10ul of clean-up buffer (900mM NaCl, 300mM EDTA) and immediately purified using MinElute PCR Purification Kit (QIAGEN). Library amplification was performed following the manufacturer's protocol (Nextera DNA Sample Preparation Kit, Illumina) except using 2.5ul of each primer and 2min of extension time in PCR reaction. PCR cycle number was determined by a pilot 15ul qPCR reaction containing 1x EvaGreen (Biotium) on 10% of unamplified library. Libraries were sized-selected to enrich 250-800bp of size and sequenced as single-end or paired-end for at least 40 million reads per sample.

RNA-seq analysis

RNA-seq reads (101 bp, single-ended) were aligned to the mouse genome (NCBI37/mm10) using TopHat2 (version 2.0.13). Expression measurement of each gene was calculated from the resulting alignment bam file by HT-seq(35). Differentially expressed genes were determined using DEseq2(36) with inclusion of littermates factors and p-value cutoff of 0.05 (B-H adjustment).

Gene Ontology (GO) and GSEA analysis

GO-term enrichment analysis was performed by DAVID Bioinformatics Resources 6.7 (37) using gene symbols as input. GSEA (15) analysis was performed using ranked fold change values (cKO vs. WT) for the displayed dataset, and gene sets selected from the referenced publications.

Principle component analysis (PCA)

Transcriptome profiling of HFSCs and stem cell lineages were obtained from microarray dataset in the following publications: Greco et al., Cell Stem Cell, 2009 (12); Zhang et al., Cell Stem Cell, 2009 (16) and Lien et al., Cell Stem Cell, 2011 (8). For each dataset, fold change values were calculated using following normalization sample: Greco et al: P43_Bulge; Zhang et al: P18-P21 Bulge 0 division replicate #1; Lien et al: quiescent HFSC replicate #1. Presence/absence call was performed and genes with both counterparts of the fold change comparison classified as presence were included in downstream analysis. Fold change were calculated using the difference of log transformed intensity measurements. If multiple probes were mapped to a single gene, fold change of each individual probes were first calculated and the median of all values was selected as the representative fold change of the gene. Fold change data of P47 Foxc1 cKO were obtained from DEseq2 output. All fold change data were merged in to a table, and lines containing missing values were excluded. The remaining data were further down-sized by selecting genes with >2 fold change in at least one of the following samples: aHFSC, qHFSC and TAC from Lien et al. (8); P56 bulge, P69 bulge, P43 HG, P56 HG and P69 HG from Greco et al. (12). A total of 2684 genes were selected for PCA plotting. Data matrix was centered by subtracting the mean of each column. PCA was

performed using the function of `prcomp` in R and the first two principal components were used for plotting. Linear Discriminant Analysis (LDA) was performed using R library MASS (38).

ATAC-seq, ChIP-seq and motif analysis

ATAC-seq reads (single-end or paired-end) and ChIP-seq reads (21) were aligned to mouse genome (NCBI37/mm10) using Bowtie2 (version 2.2.3) (39). Duplicate reads were removed by Picard tools (<http://broadinstitute.github.io/picard/>). Mitochondrial reads were removed and peak calling was performed on each individual sample by MACS (version 2.0.9) (40). Peaks from different ATAC-seq samples were merged for downstream analysis. De novo motif searching was performed using the HOMER (41). For Foxc1 motif discovery, the top 500 peaks (based on MACS2 output p value) were used with regions of 500bp around peak center. For motif analysis using ATAC-peaks, all ATAC peaks were first used for motif discovery, with mm10 genome sequence as background. Then ATAC peaks associated with differentially expressed gene in P30 cKO-vs-WT RNA-seq were used as target sequence set and motif discovery was performed against sequenced of ATAC peaks that associated with non-RNAseq hits genes. In both cases, enriched motifs from the default HOMER motif were analyzed, as shown in Fig 5h. Sequencing data tracks were presented by Integrative Genomics Viewer (42).

Data Accession

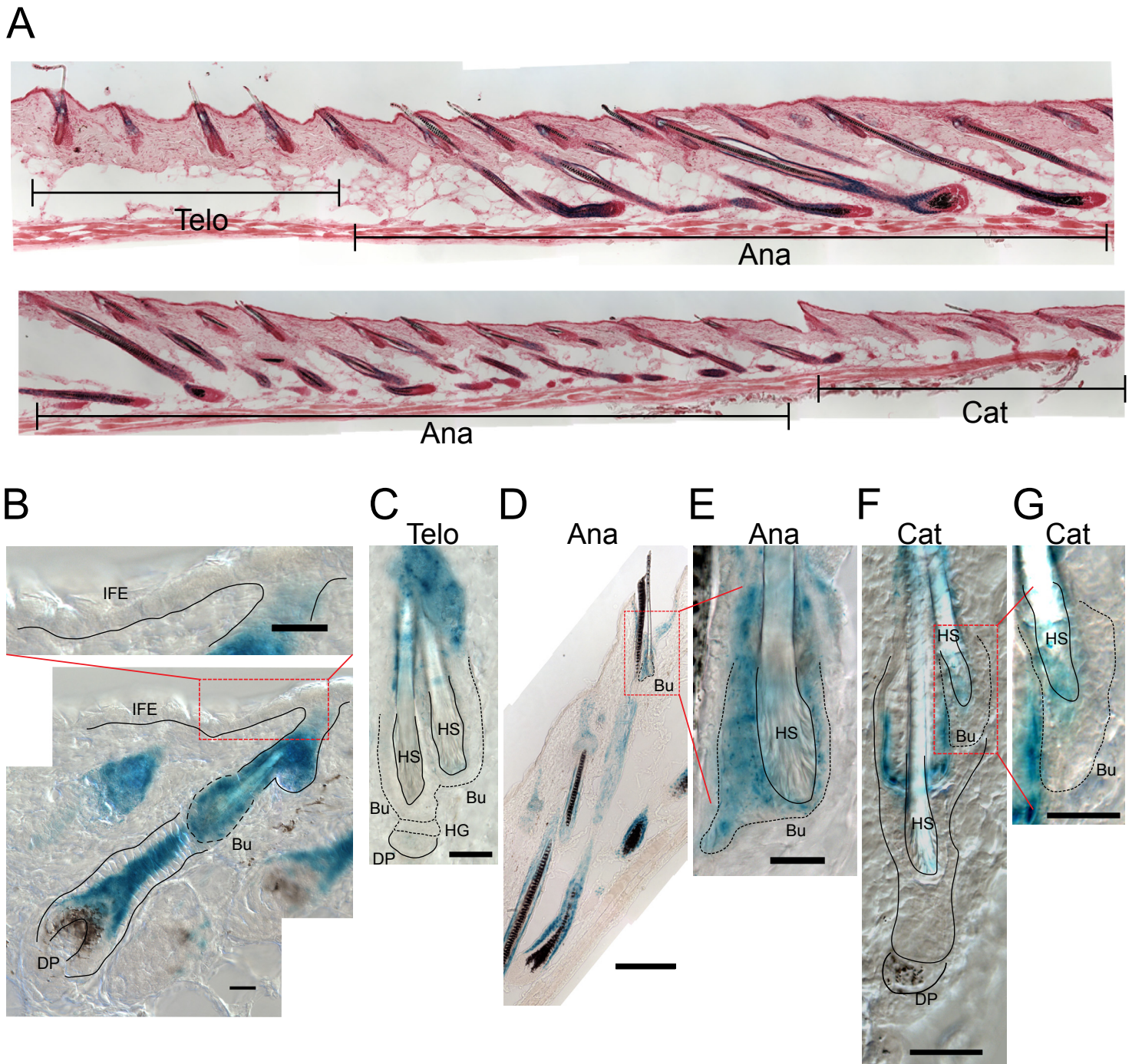
Sequencing data have been deposited to GEO with the accession number GSE67404 and GSE68288.

Figs. S1-S12

Tables S1-S4

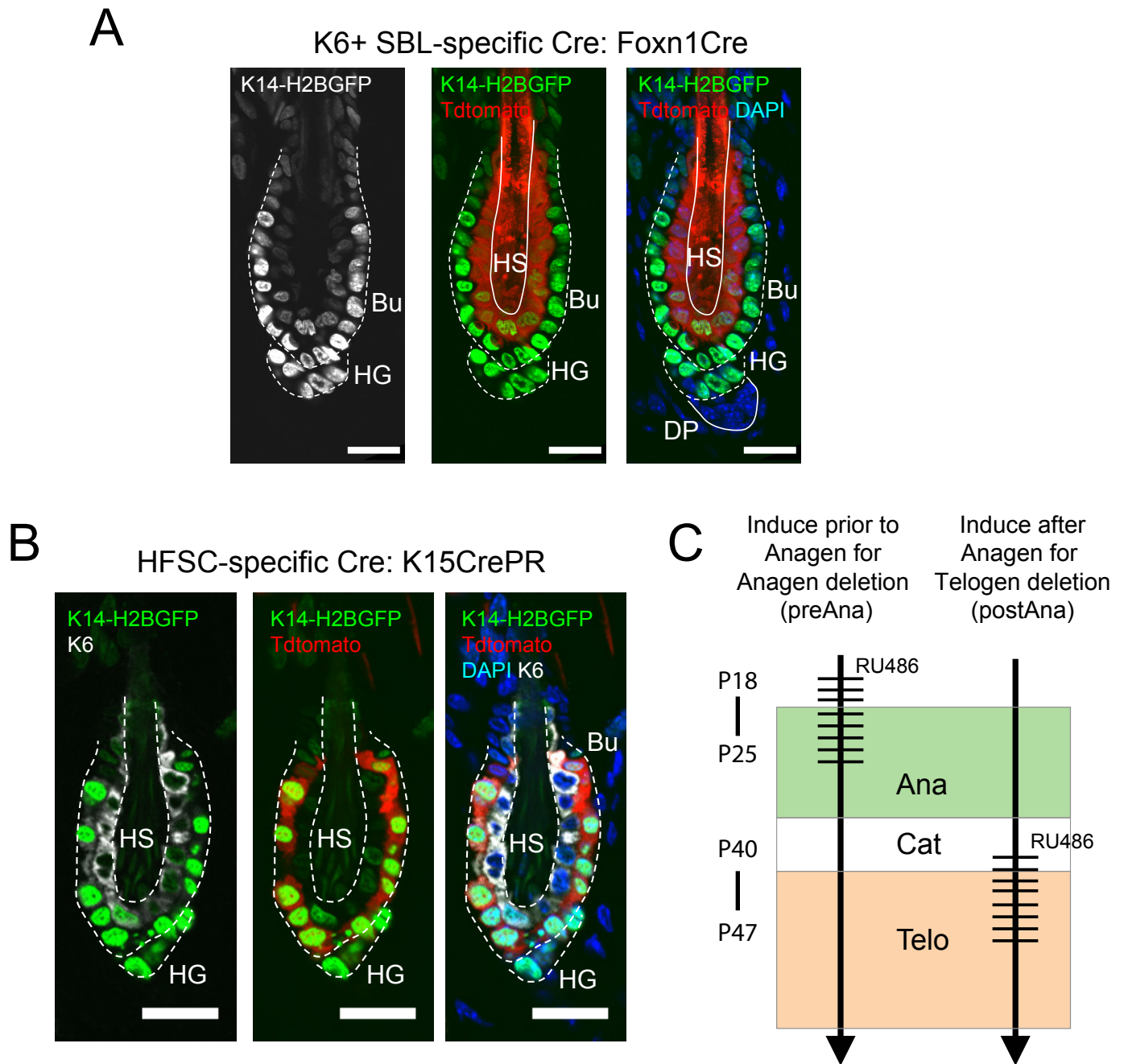
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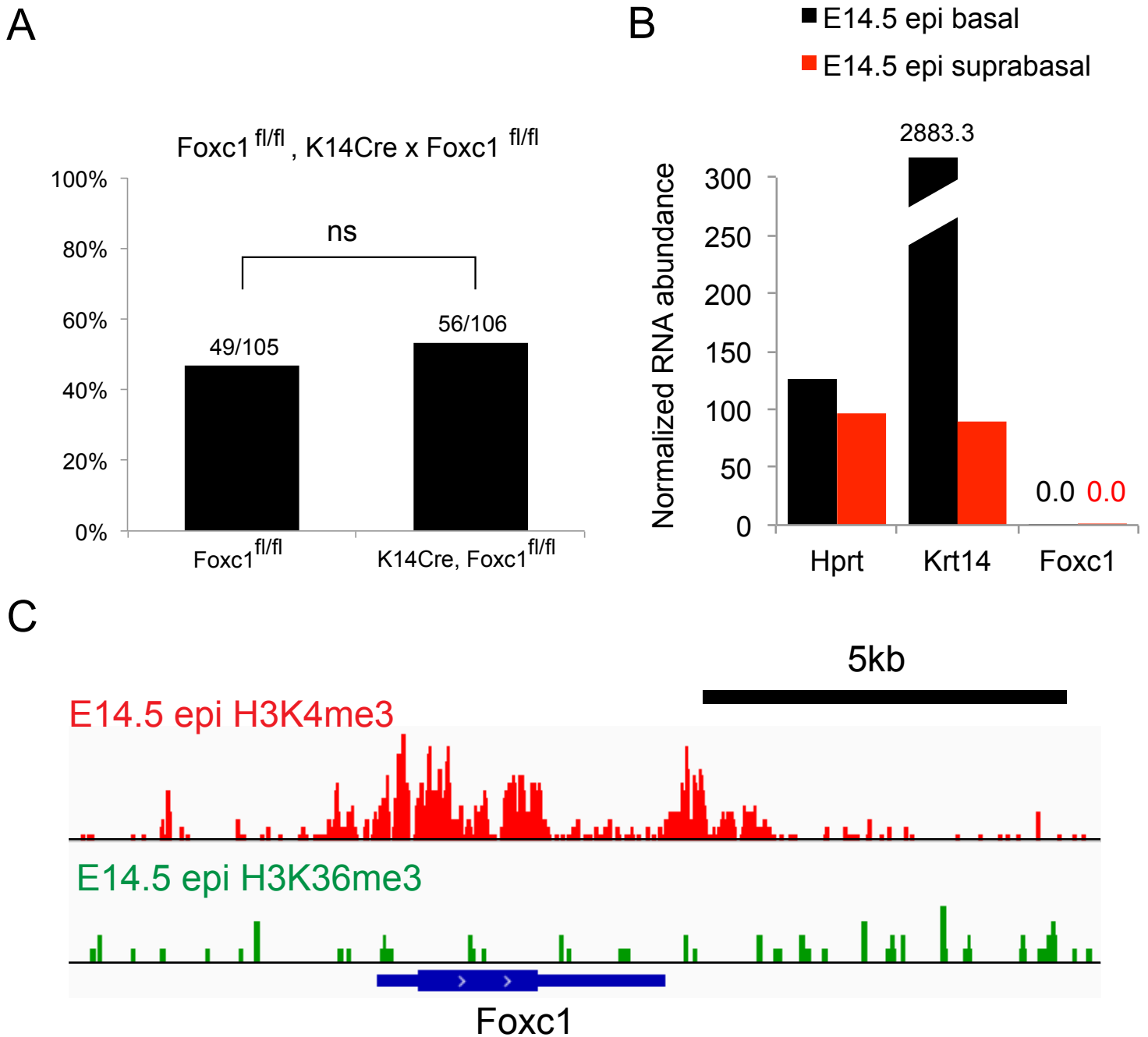
Supplementary Figure 1 Foxc1 Expression levels correlates with hair cycle progression.

(A) Hair follicle morphology of a 15-month old mouse. Section was stained with LacZ and nuclear fast red. (B-G) Foxc1 expression is detected by LacZ staining and IF staining in skin sections of Foxc1-LacZ knock-in mice. (B) Foxc1 expression is absent in the interfollicular epidermis (IFE). (C) Foxc1 expression is absent in the bulge and hair germ of telogen hair follicle. (D-E) Foxc1 is induced in the bulge of anagen hair follicle. (F-G) Bulge expression of Foxc1 wanes in catagen hair follicle. Red dashed square indicates the amplified region in the next panel. Telo, Telogen; Ana, Anagen; Cat, Catagen; Bu, bulge; HG, hair germ; DP, dermal papillae; HS, hair shaft; Scale bar: 25 μ m in C, 200 μ m in D, 20 μ m in B and E, 50 μ m in F and 10 μ m in G.



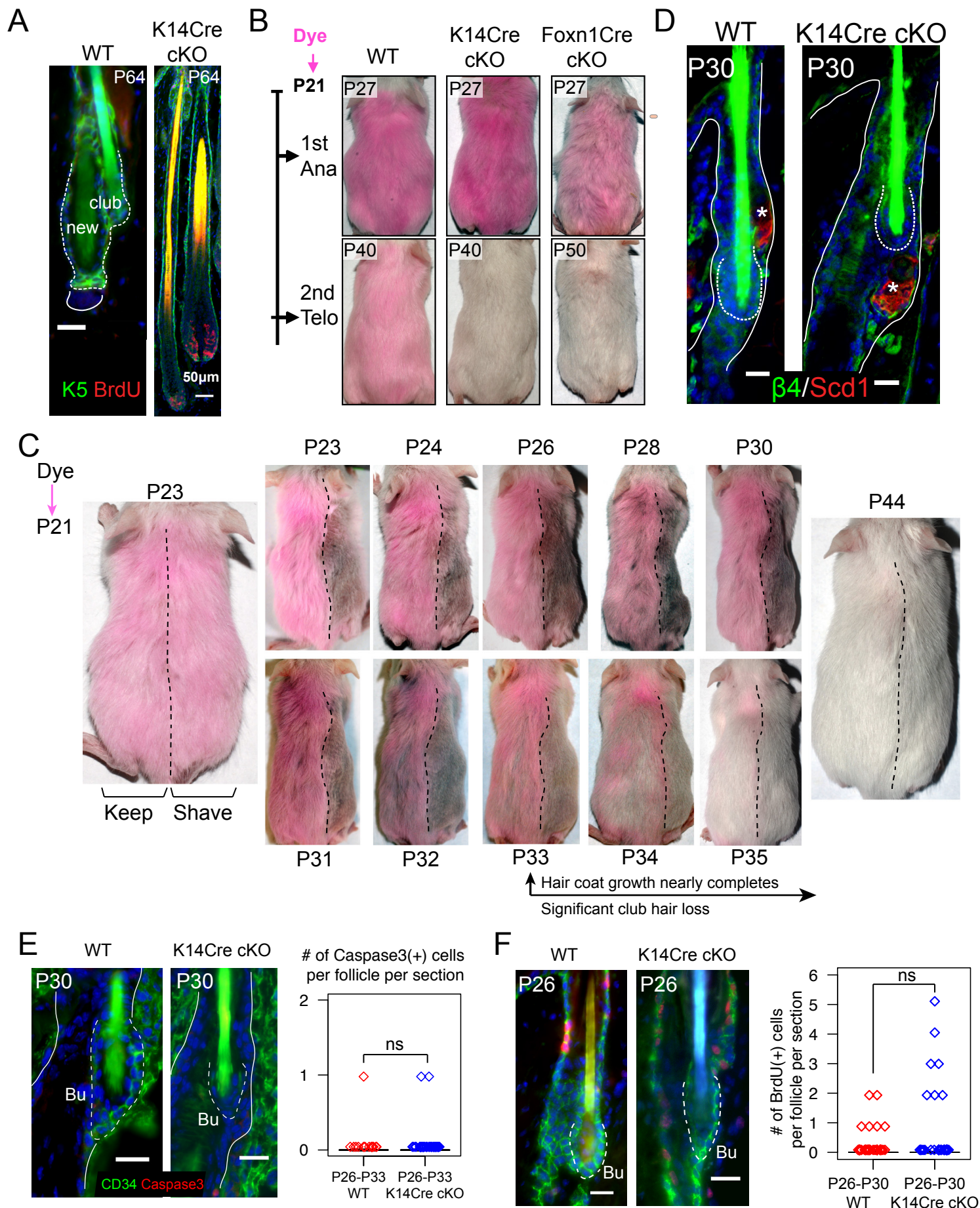
Supplementary Figure 2 Lineage-specific cre recombinase in HF bulge.

(A) Foxn1Cre specifically induces Cre expression in K6+ suprabasal bulge layer (SBL). Note that K14-H2BGFP transgene has high GFP expression in CD34+ HFSC layer and low GFP expression in SBL. (B) K15CrePR, upon RU486 induction, specifically induces Cre expression in CD34+, HFSC layer. (C) K15CrePR induction scheme. horizontal bars across the time arrow indicate single RU486 topical treatment. Bu, bulge; HG, hair germ; DP, dermal papillae; HS, hair shaft; Scale bar: 20 μ m.

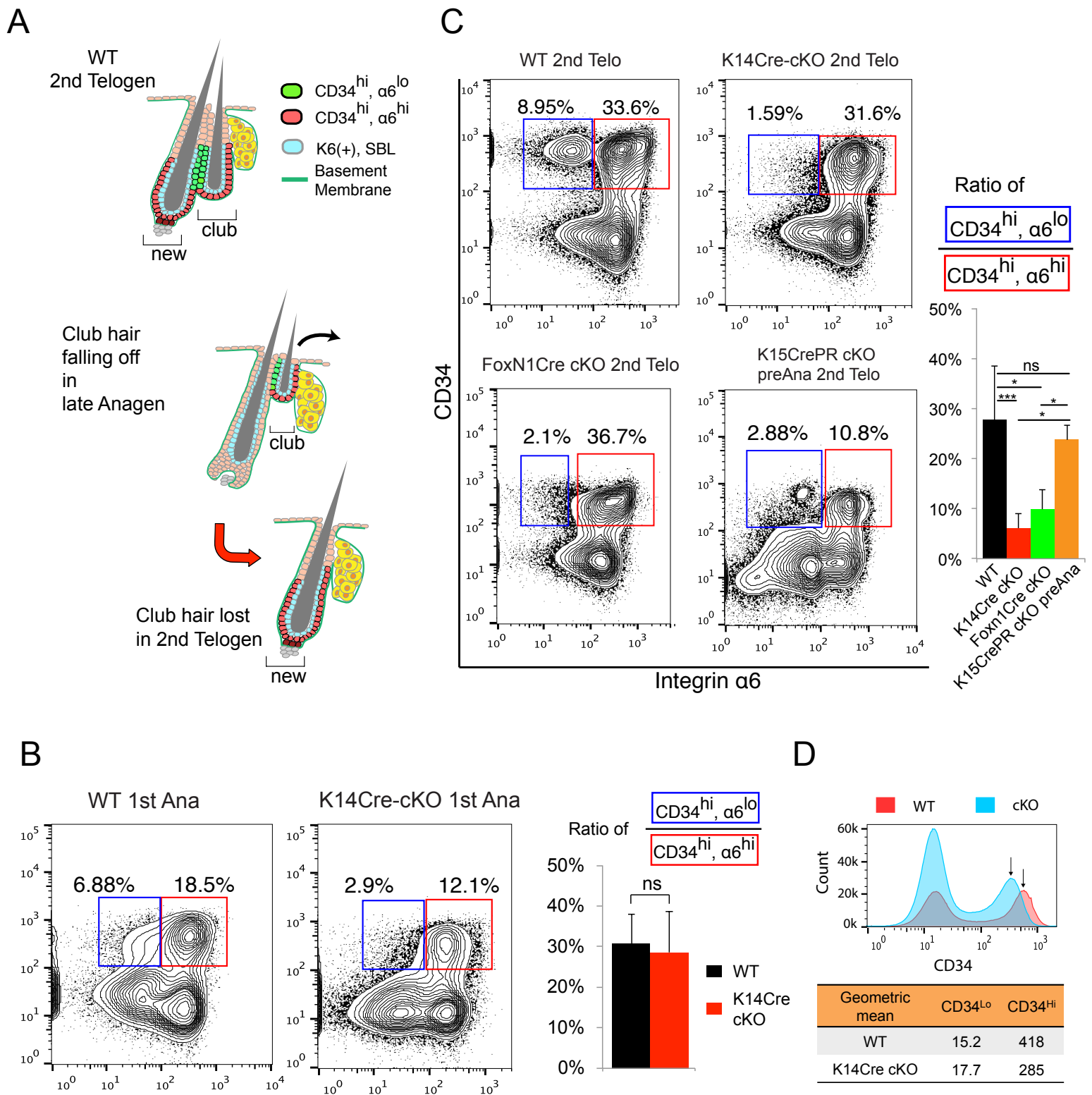


Supplementary Figure 3 Lack of *Foxc1* expression in embryonic skin progenitors

(A) Both WT and cKO animals were born at the expected Mendelian ratio. (B) Normalized RNA abundance is calculated using read per million reads mapped (RPM) in 3Seq data from E14.5 epidermis (epi) lineages. (C) Snapshot of H3K4me3 and H3K36me3 ChIP-seq data from E14.5 epidermis lineages indicates absence of active transcriptional elongation at *Foxc1* locus.



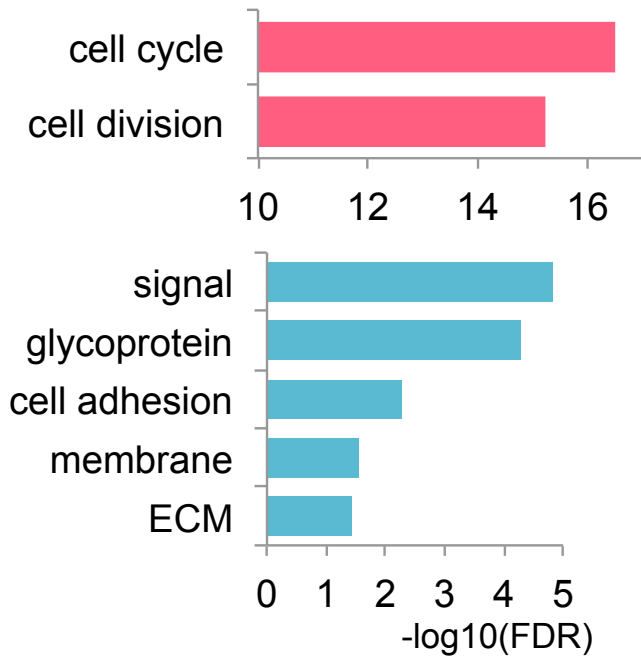
Supplementary Figure 4 Deletion of Foxc1 in the suprabasal bulge layer leads to loss of the club hair. (A) Immunofluorescence of EdU incorporation for WT and K14Cre cKO HFs at P64. The cKO HFs have entered anagen while WT HFs are still in telogen and showed no sign of proliferation. (B) Dyed hair coat shows both K14Cre cKO and Foxn1Cre cKO animals lost the club hairs during mid to late anagen. (C) Timing of the club hair loss. Hair coat from the first cycle was dyed at P21. The right half of back hair coat was shaved and monitored to the next cycle. Significant amount of dyed hair started to shed at P33-P35, when the new hair shafts nearly complete the growth phase. (D) Loss of club hair correlates with upward movement of the hair. Note the relative position of the club hair and the sebaceous gland. (E-F) No defects in apoptosis (E) or proliferation (F) were observed in middle to late anagen in K14Cre cKO Bulge compared to WT. n=4 in B and n=3 in C for each genotypes. Scale bar: 20 μ m in A, D, E, F or otherwise labeled.



Supplementary Figure 5 Using FACS to examine the club hair loss phenotype. (A) Schematic of the loss of club hair. Loss of club hair is likely due to defect of club hair anchorage in late anagen and can be examined by FACS profiling of CD34^{hi}, α6^{lo} population (green). (B) Flow cytometry profile of HFSCs during the first anagen for WT and K14Cre cKO mice. Ratio of CD34^{hi}, α6^{lo} (blue rectangle) over CD34^{hi}, α6^{hi} (red rectangle) is used for quantification. (C) Flow cytometry analysis and quantification of the loss of club hair by the second telogen. Anagen: n=3; telogen: n=8. (D) CD34 histogram from flow cytometry shows decreased level of CD34 expression. * p<0.05, *** p<0.001, ns: not significant.

A

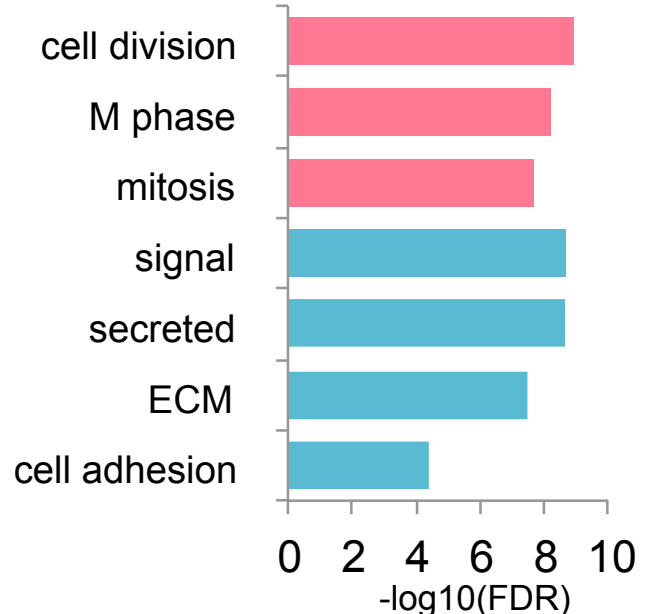
P47 HFSC RNA-seq



■ Upregulated
■ Downregulated

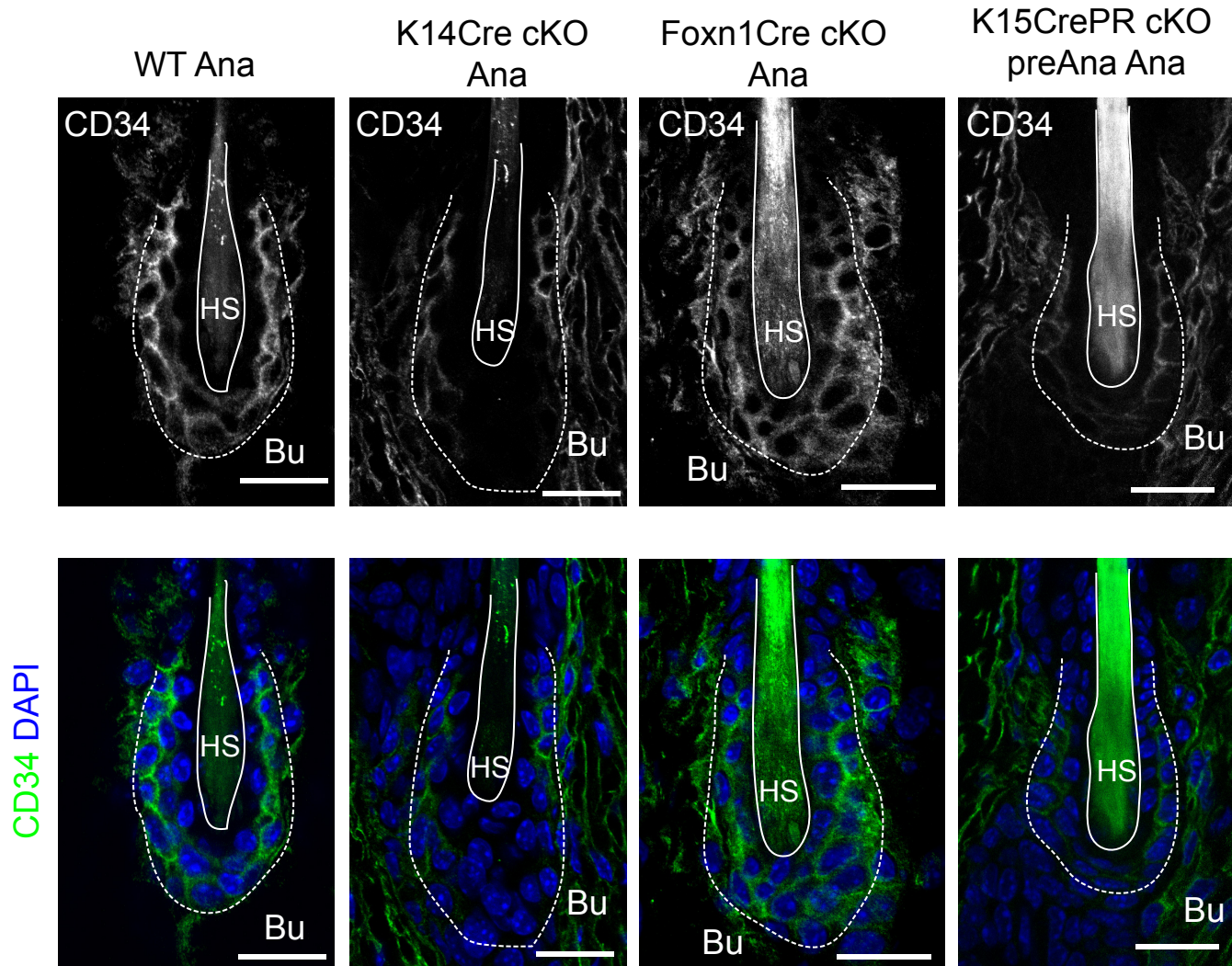
B

P30 HFSC RNA-seq

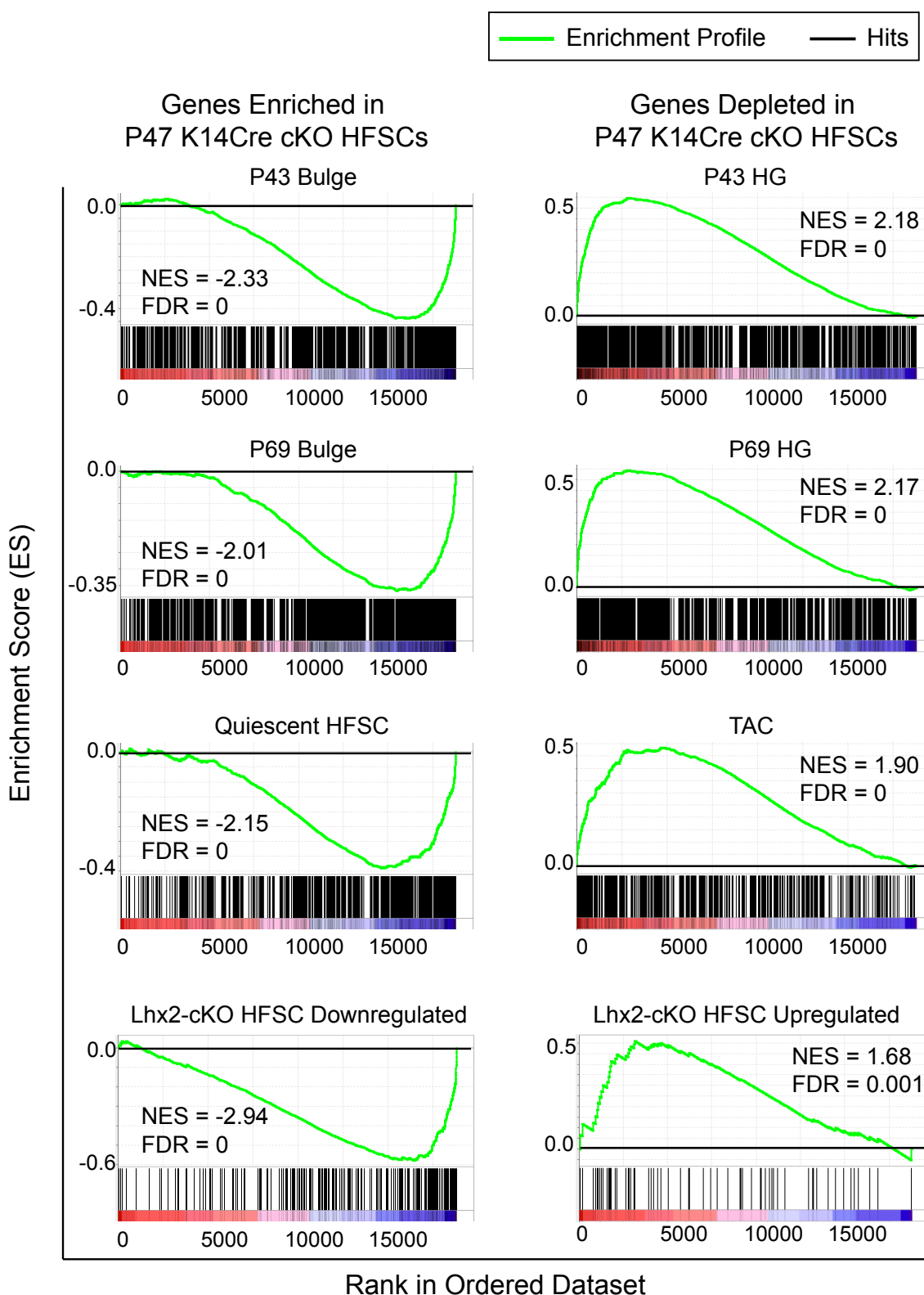


■ Upregulated
■ Downregulated

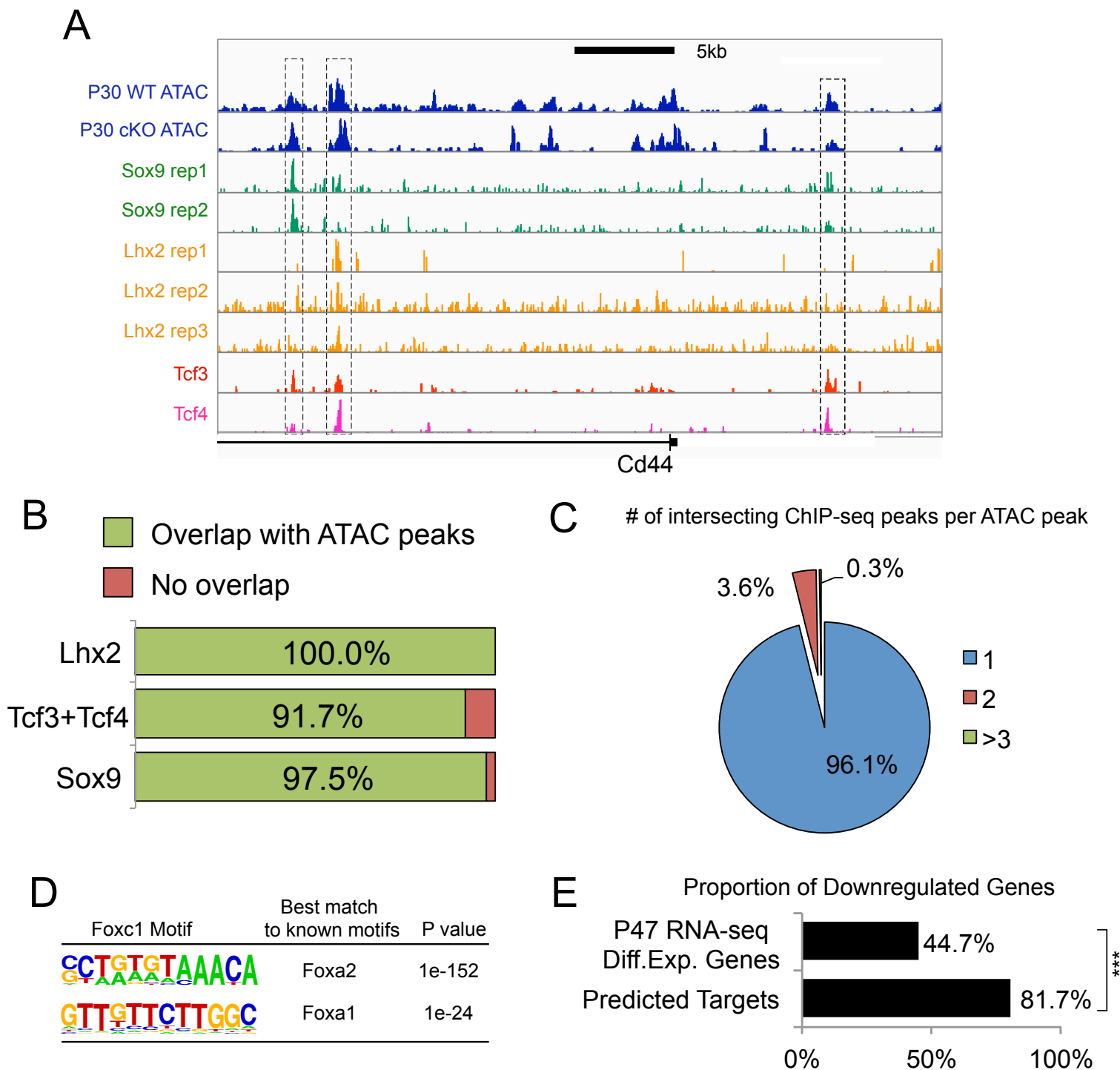
Supplementary Figure 6 GO term analysis of HFSC RNA-seq. (A) Gene Ontology of all differentially expressed genes in P47 KO HFSCs. (B) Gene Ontology of all differentially expressed genes in P30 KO HFSCs.



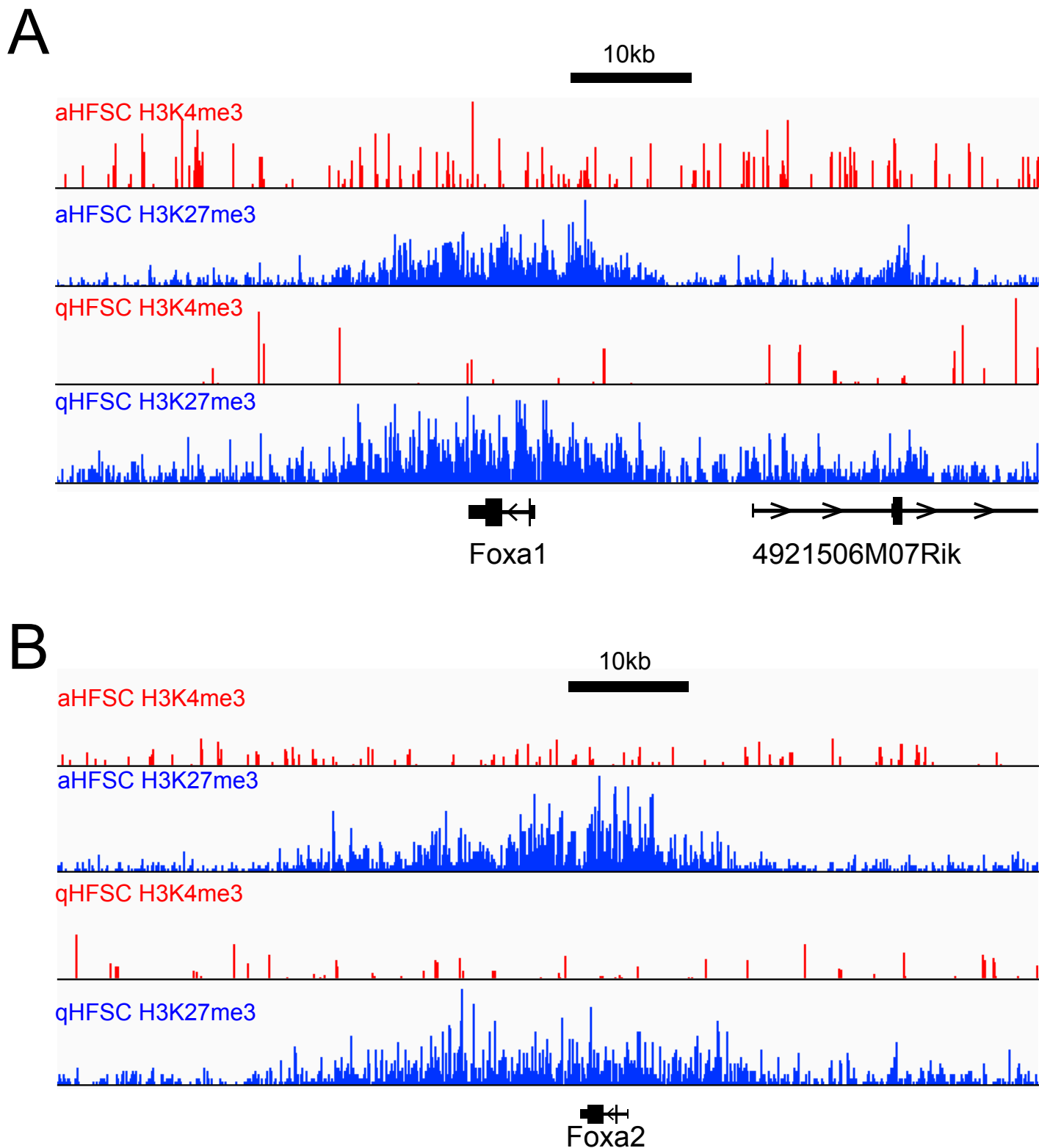
Supplementary Figure 7 Downregulation of CD34 when Foxc1 is depleted in activated HFSCs. CD34 immunofluorescence was examined in anagen bulge of WT, K14Cre cKO, Foxn1Cre cKO and K15CrePR preAna cKO mice. Bulge (Bu) is outlined with dashed line and hair shaft (HS) is outlined with solid line. Hair shaft has auto fluorescence for the green channel. CD34 staining intensity was decreased in K14Cre cKO and K15CrePR preAna cKO but not for Foxn1Cre cKO, compared to WT control. Scale bar: 20 μ m.



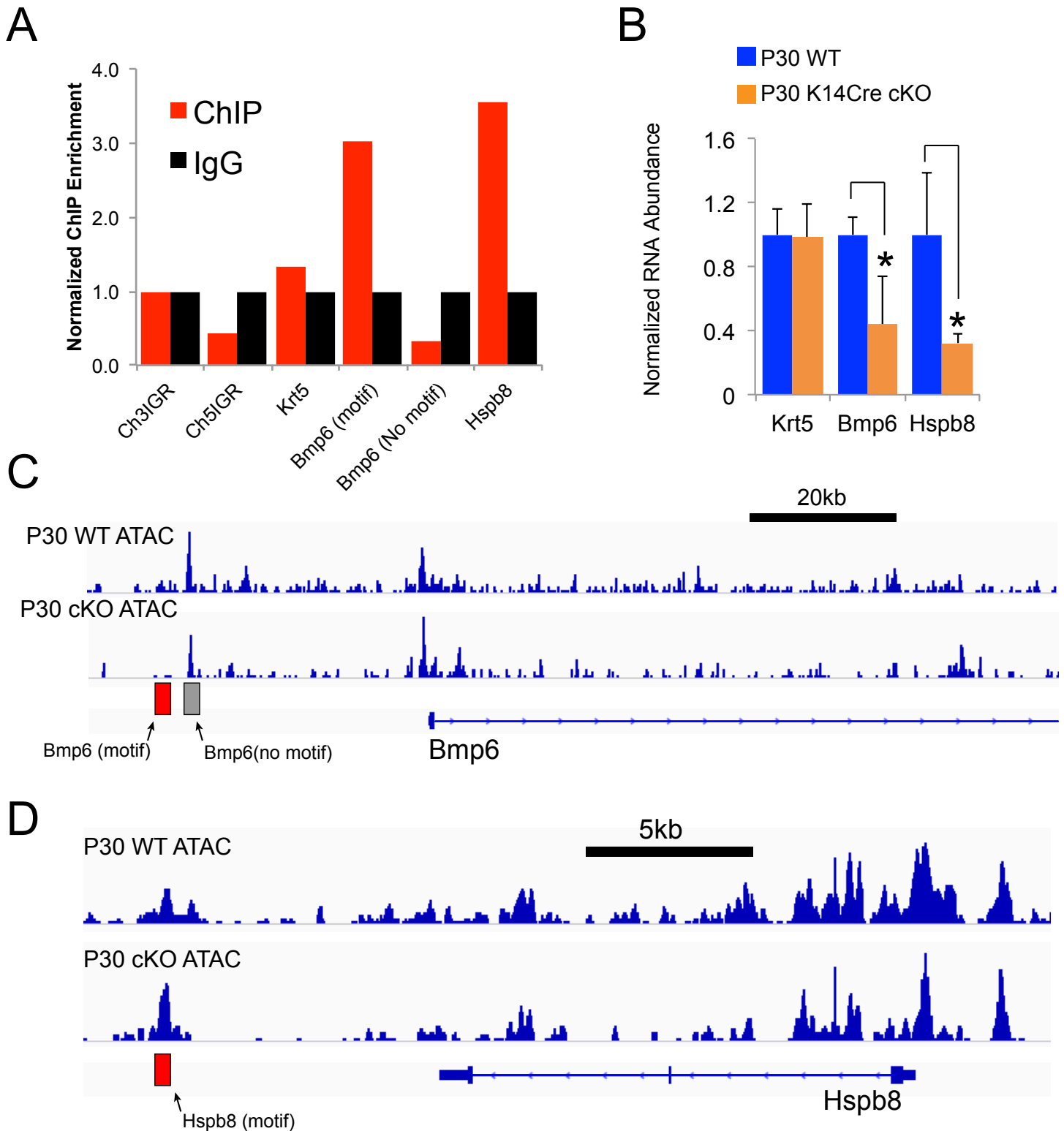
Supplementary Figure 8 GSEA of P47 HFSC RNA-seq. Ranked fold changes of gene expression of P47 HFSC RNA-seq were tested using GSEA with signature genes of quiescent (left panels) and activated (right panels) HFSCs. All quiescent signature genes were depleted (preferentially downregulated) in P47 cKO HFSCs and all activated signature genes were enriched (preferentially upregulated) in P47 cKO HFSCs.



Supplementary Figure 9 Using ATAC-Seq to predict Foxc1 targets. (A) Cd44 locus shows the overlapping peaks of ATAC-seq and ChIP-Seq signals of Sox9, Lhx2, Tcf3/4. (B) Percentage of ChIP-Seq peaks of Lhx2, Tcf3/4, and Sox9 that were detected by ATAC-seq. (C) Number of intersecting ChIP-Seq peaks per ATAC peak in All ChIP seq peaks examined in A and B. (D) De novo motif discovery of Foxc1 ChIP-seq data identifies Foxc1 recognition motifs. (E) Percentage of downregulated gene in P47 Foxc1 HFSC cKO RNA-seq and among predicted Foxc1 targets in P30, compared to WT.



Supplementary Figure 10 *Foxa1* and *Foxa2* loci are epigenetically silenced in the HFSCs. Snapshot of H3K4me3 and H3K27me3 ChIP-seq of *Foxa1* (A) and *Foxa2* (B) loci in activated HFSCs (aHFSC) and quiescence HFSCs (qHFSC)) shows depletion of H3K4me3 marks accumulation of H3K27me3 marks at both locus, indicating strong gene silencing. ChIP-seq data are from Lien et al., *Cell Stem Cell*, 2011.

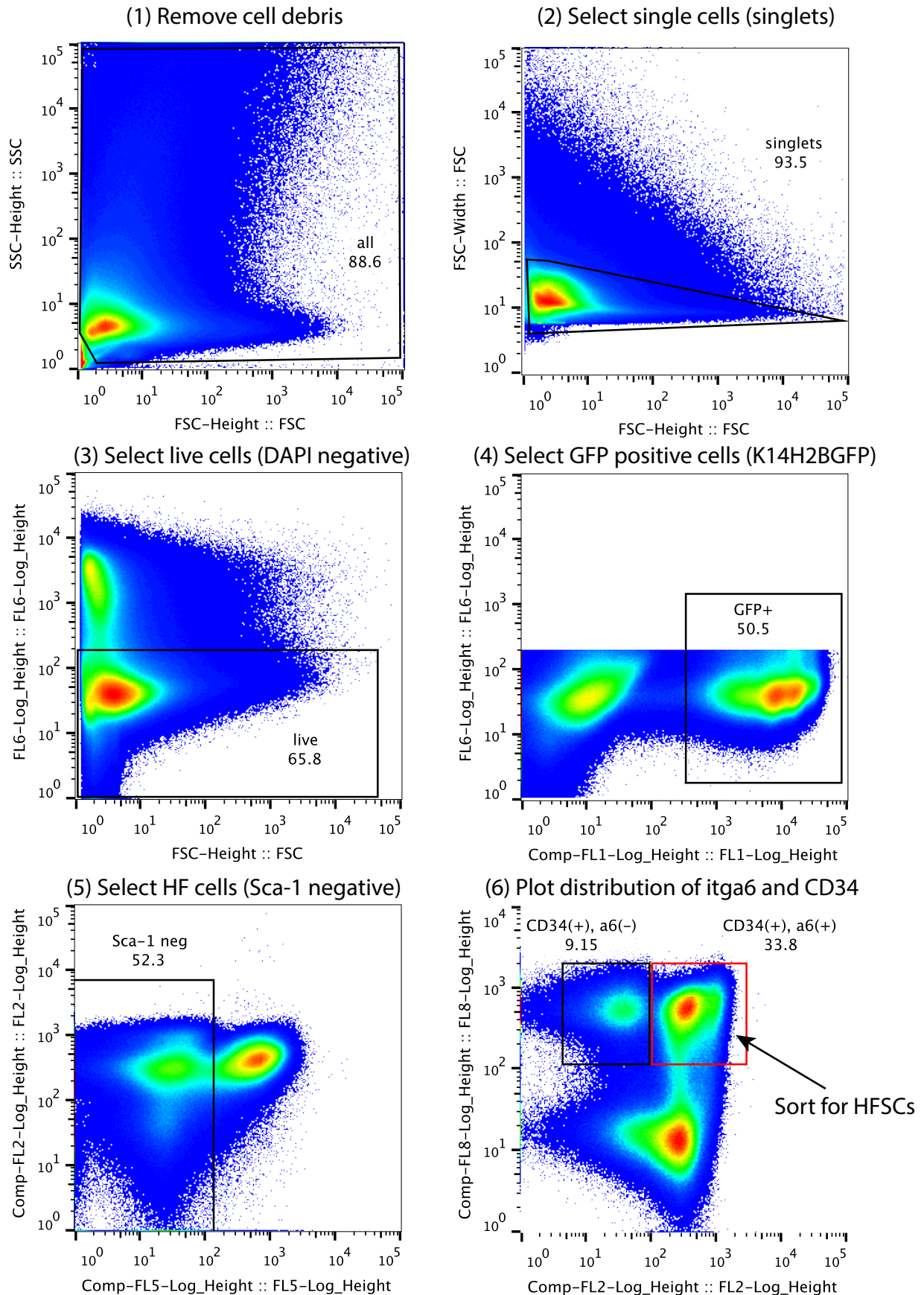


Supplementary Figure 11 Foxc1 directly binds Bmp6 and Smads target Hspb8

(A) Foxc1 ChIP-PCR of intergenic region in chr3 (Chr3IGR) and chr5 (Chr5IGR), Krt5, Bmp6 and Hspb8. ChIP quantities were normalized to Chr3IGR and fold change in ChIP over IgG control is displayed.

(B) Normalized RNA abundance of Krt5, Bmp6 and Hspb8 from P30 RNA-seq. * $p < 0.05$, two-tail t test.

(C-D) ATAC-Seq track of P30 WT and K14Cre cKO HFSCs of Bmp6 locus (upstream and 5' end, C) and Hspb8. Primers targeting ATAC-Seq peaks that contains Foxc1 motif (red rectangle) or does not contain a motif (grey rectangle) are labeled.



Supplementary Figure 12 Sorting logic for HFSC isolation

Supplementary Table S1 Predicted *Foxc1* target genes

Gene	Fold Change (P30 cKO vs WT)	p value (B-H)
Tgm5	-28.25	8.94E-65
Atp2a3	-26.49	5.36E-06
Gfra1	-11.25	5.93E-41
Ccdc88c	-10.37	1.51E-22
Trpm3	-9.30	7.94E-05
Npas2	-8.67	1.94E-09
Fam46c	-7.25	2.90E-28
Tmem108	-6.97	2.51E-05
Cacna1c	-6.64	5.07E-08
Pknox2	-6.00	0.001779
Col4a3	-5.98	0.000307
Duox1	-5.58	8.39E-15
Pvt1	-5.52	0.003601
Bdnf	-5.29	1.06E-11
Adamts17	-5.28	5.05E-06
Me3	-5.15	0.000956
Aff3	-4.79	6.96E-09
Aqp3	-4.70	7.08E-15
Ngf	-4.58	2.13E-11
Bmp2	-4.53	1.65E-14
Ogdhl	-4.27	0.001555
Ptpre	-4.24	1.80E-07
Fam25c	-4.22	1.64E-10
Hivep3	-4.15	6.43E-08
Arhgap44	-4.14	3.99E-11
Zbtb16	-4.09	2.80E-05
Ptn	-3.93	2.33E-11
Slc6a4	-3.93	0.000687
Cadm1	-3.91	4.08E-14
Lypd6	-3.87	0.000508
Cdon	-3.74	1.18E-11
Antxr1	-3.73	9.28E-10

Slc39a8	-3.69	1.91E-08
Trim16	-3.64	6.25E-11
Robo2	-3.61	9.18E-05
Hspb8	-3.38	5.74E-12
Agpat9	-3.34	0.001227
Slc38a3	-3.29	0.000433
Nrcam	-3.29	3.73E-05
Fam213a	-3.20	4.56E-06
Moxd1	-3.18	5.22E-09
Trim2	-3.17	1.76E-07
Tle4	-3.16	3.24E-05
Sdk2	-3.14	1.60E-07
S100a3	-3.11	1.30E-05
Dap	-3.07	7.94E-08
Tspan2	-3.06	0.008170
Scel	-3.03	2.16E-09
Cgnl1	-2.99	7.95E-07
Calml3	-2.91	1.68E-09
Fgf1	-2.90	0.000507
Vit	-2.84	0.003654
Ank	-2.84	5.94E-05
Gm973	-2.78	5.12E-06
Tns4	-2.75	5.11E-07
Piezo1	-2.74	0.000303
Asap2	-2.73	2.48E-05
Fst	-2.72	0.000292
Itgb6	-2.66	1.84E-07
Serpinb2	-2.62	4.68E-08
Inpp4b	-2.61	0.006124
Nfatc1	-2.59	0.000198
Bmp6	-2.59	0.008821
Crispld1	-2.52	0.004250
Gmds	-2.51	0.000286
Sh3kbp1	-2.51	0.001433
Lmo7	-2.49	0.000160

Hmcn1	-2.44	3.40E-05
Mfap3l	-2.37	9.07E-05
Sulf2	-2.36	2.89E-05
Igfbp3	-2.36	2.58E-06
Ppp2r3a	-2.34	6.01E-05
Kank1	-2.27	1.30E-05
Sema3e	-2.22	0.000272
Prlr	-2.22	0.000172
Tmem40	-2.20	0.009422
Gpc6	-2.20	0.001063
Gpr161	-2.17	0.004268
Rora	-2.15	0.000478
Enox1	-2.15	0.001287
Ankrd10	-2.14	0.000188
Efnb2	-2.09	0.000188
Cyr61	-2.06	0.000125
Cry1	-2.05	0.004932
Actb	-2.03	0.000972
Myo1e	-1.98	0.009521
Ccdc3	-1.97	0.002518
Hopx	-1.96	0.001558
Net1	-1.90	0.001539
Slc6a6	-1.90	0.001418
Cd44	-1.88	0.004205
Gpr126	-1.85	0.008605
Arid5b	-1.83	0.009456
Peli2	-1.82	0.009282
Tfap2b	-1.82	0.009285
Trabd2b	-1.81	0.008936
Insig2	1.76	0.008605
Gja1	1.88	0.002947
Tnc	2.07	0.000325
Bhlhe41	2.08	0.002151
Hist1h1c	2.09	0.001025
Smox	2.12	0.000877

Prickle1	2.15	0.000227
Car12	2.20	0.003233
Abcg2	2.22	0.001400
Parm1	2.29	0.007433
Slc22a23	2.30	0.000275
Btbd3	2.43	0.000200
Pla2g7	2.45	3.64E-05
Nrg1	2.75	1.30E-05
Ppp1r3b	3.11	4.06E-09
Serpine2	3.22	1.87E-07
Ctsc	3.23	3.35E-10
AU018091	3.34	0.001194
Pamr1	3.65	0.007829
Slc4a4	5.20	1.01E-05
Serpina3k	18.45	0.000227

**Supplementary Table S2 Foxc1/Nfatc1/Smad motif distribution in differentially expressed genes of P30
Foxc1 cKO HFSC RNA-seq**

gene	log2FoldChange	pval	Foxc1 motif	Nfatc1 motif	Smad motif
Abcg2	1.1538	0.001399551	Foxc1	Nfatc1	Smad
Adamts17	-2.4005	5.05E-06	Foxc1	Nfatc1	Smad
Aff3	-2.2600	6.96E-09	Foxc1	NA	Smad
Agpat9	-1.7415	0.001226964	Foxc1	Nfatc1	Smad
Ank	-1.5062	5.94E-05	Foxc1	Nfatc1	Smad
Ankrd10	-1.0999	0.000188118	Foxc1	NA	Smad
Ankrd35	-1.5166	0.00630418	Foxc1	NA	Smad
Antxr1	-1.8982	9.28E-10	Foxc1	Nfatc1	Smad
Arhgap44	-2.0485	3.99E-11	Foxc1	Nfatc1	Smad
Arid5b	-0.8712	0.009456243	Foxc1	Nfatc1	Smad
Atp2a3	-4.7274	5.36E-06	Foxc1	NA	NA
AU018091	1.7410	0.001193734	Foxc1	Nfatc1	Smad
Bdnf	-2.4042	1.06E-11	Foxc1	NA	Smad
Bhlhe41	1.0592	0.002150978	Foxc1	NA	NA
Bmp2	-2.1788	1.65E-14	Foxc1	Nfatc1	Smad
Bmp6	-1.3723	0.008821217	Foxc1	NA	Smad
Btbd3	1.2787	0.000200031	Foxc1	Nfatc1	Smad
Cacna1c	-2.7313	5.07E-08	Foxc1	NA	Smad
Cadm1	-1.9675	4.08E-14	Foxc1	Nfatc1	Smad
Calml3	-1.5433	1.68E-09	Foxc1	NA	Smad
Ccdc88c	-3.3743	1.51E-22	Foxc1	NA	Smad
Cd44	-0.9092	0.004204617	Foxc1	Nfatc1	Smad
Cdc42ep1	-1.8977	5.90E-10	Foxc1	NA	Smad
Cdon	-1.9048	1.18E-11	Foxc1	NA	Smad
Col4a3	-2.5801	0.000307394	Foxc1	Nfatc1	Smad
Col4a4	-2.7337	5.65E-07	Foxc1	NA	Smad
Crispld1	-1.3313	0.004250351	Foxc1	NA	Smad
Ctsc	1.6931	3.35E-10	Foxc1	Nfatc1	Smad
Cyr61	-1.0393	0.000124504	Foxc1	Nfatc1	Smad
Efnb2	-1.0643	0.000188118	Foxc1	Nfatc1	Smad
Emp1	-1.2376	1.89E-05	Foxc1	Nfatc1	Smad
Enox1	-1.1016	0.001287267	Foxc1	Nfatc1	Smad
Esyt3	-2.6480	3.98E-14	Foxc1	NA	Smad
Fabp5	1.5796	5.78E-08	Foxc1	Nfatc1	NA
Fam171b	-3.0589	3.85E-14	Foxc1	Nfatc1	Smad
Fam46c	-2.8571	2.90E-28	Foxc1	NA	Smad
Flnc	-1.0055	0.001396379	Foxc1	NA	Smad
Fmn1	-1.4249	0.002484435	Foxc1	Nfatc1	Smad
Foxp1	-0.9264	0.009790922	Foxc1	Nfatc1	Smad
Fst	-1.4416	0.000291843	Foxc1	Nfatc1	Smad
Gfra1	-3.4913	5.93E-41	Foxc1	Nfatc1	Smad
Gja1	0.9106	0.002947016	Foxc1	Nfatc1	Smad
Gjb4	-1.3712	2.08E-06	Foxc1	NA	Smad
Gmcs	-1.3293	0.000286321	Foxc1	Nfatc1	Smad
Gpc6	-1.1358	0.001062881	Foxc1	Nfatc1	Smad
Gpr126	-0.8860	0.008605465	Foxc1	Nfatc1	Smad
Hmcn1	-1.2883	3.40E-05	Foxc1	Nfatc1	Smad
Hmox1	-1.2904	8.94E-05	Foxc1	NA	Smad

Hopx	-0.9706	0.001558148	Foxc1	Nfatc1	Smad
Hspb8	-1.7579	5.74E-12	Foxc1	Nfatc1	Smad
Igfbp3	-1.2387	2.58E-06	Foxc1	Nfatc1	Smad
Il31ra	-4.3966	4.95E-30	Foxc1	Nfatc1	Smad
Inpp4b	-1.3816	0.006123833	Foxc1	Nfatc1	Smad
Itgb6	-1.4119	1.84E-07	Foxc1	NA	Smad
Kcnma1	-4.4234	2.99E-07	Foxc1	Nfatc1	Smad
Lmo7	-1.3157	0.000160327	Foxc1	Nfatc1	Smad
Lrrn3	-2.0658	0.000313448	Foxc1	NA	Smad
Lypd6	-1.9522	0.000508392	Foxc1	Nfatc1	Smad
Marveld3	-1.6722	3.77E-05	Foxc1	NA	NA
Me3	-2.3637	0.000956227	Foxc1	Nfatc1	Smad
Moxd1	-1.6703	5.22E-09	Foxc1	Nfatc1	Smad
Net1	-0.9269	0.001538596	Foxc1	NA	Smad
Nfatc1	-1.3733	0.000198344	Foxc1	Nfatc1	Smad
Ngf	-2.1949	2.13E-11	Foxc1	Nfatc1	Smad
Nrcam	-1.7182	3.73E-05	Foxc1	Nfatc1	Smad
Nrg1	1.4580	1.30E-05	Foxc1	Nfatc1	Smad
Pamr1	1.8670	0.007828742	Foxc1	NA	Smad
Parm1	1.1936	0.007433048	Foxc1	Nfatc1	Smad
Peli2	-0.8656	0.009282301	Foxc1	Nfatc1	Smad
Pla2g7	1.2955	3.64E-05	Foxc1	NA	Smad
Ppp1r3b	1.6387	4.06E-09	Foxc1	Nfatc1	Smad
Prickle1	1.1016	0.000227008	Foxc1	Nfatc1	Smad
Prlr	-1.1498	0.000172146	Foxc1	Nfatc1	Smad
Ptn	-1.9761	2.33E-11	Foxc1	NA	Smad
Pvt1	-2.4656	0.003601033	Foxc1	Nfatc1	Smad
Robo2	-1.8520	9.18E-05	Foxc1	Nfatc1	Smad
Rora	-1.1060	0.000477547	Foxc1	Nfatc1	Smad
S100a3	-1.6367	1.30E-05	Foxc1	NA	Smad
Scel	-1.6002	2.16E-09	Foxc1	Nfatc1	Smad
Sdk2	-1.6498	1.60E-07	Foxc1	NA	Smad
Sema3c	-0.8774	0.006123833	Foxc1	Nfatc1	Smad
Sema3e	-1.1526	0.000272114	Foxc1	Nfatc1	Smad
Serpib8	-1.1560	0.000565233	Foxc1	Nfatc1	Smad
Serpine2	1.6874	1.87E-07	Foxc1	Nfatc1	Smad
Sh3kbp1	-1.3274	0.001433448	Foxc1	Nfatc1	Smad
Slc22a23	1.1987	0.000275335	Foxc1	Nfatc1	Smad
Slc39a8	-1.8838	1.91E-08	Foxc1	NA	Smad
Slc4a4	2.3778	1.01E-05	Foxc1	Nfatc1	Smad
Slc6a6	-0.9257	0.001417723	Foxc1	NA	Smad
Smarca2	-1.3758	2.23E-07	Foxc1	NA	Smad
Smox	1.0845	0.000877018	Foxc1	Nfatc1	Smad
Stfa3	-2.0422	5.29E-10	Foxc1	Nfatc1	Smad
Tnc	1.0467	0.000325187	Foxc1	Nfatc1	Smad
Tns4	-1.4574	5.11E-07	Foxc1	Nfatc1	Smad
Trabd2b	-0.8557	0.008935975	Foxc1	Nfatc1	Smad
Trim16	-1.8632	6.25E-11	Foxc1	NA	Smad
Trim2	-1.6633	1.76E-07	Foxc1	Nfatc1	Smad
Trpm3	-3.2175	7.94E-05	Foxc1	Nfatc1	Smad
Tspan2	-1.6116	0.00816966	Foxc1	NA	Smad
Tuba4a	-0.9092	0.001865625	Foxc1	NA	NA

Ucp2	0.9052	0.003800326	Foxc1	NA	Smad
Unc5b	-1.0096	0.007893428	Foxc1	NA	Smad
Vit	-1.5071	0.00365376	Foxc1	NA	Smad
Zdhhc2	-2.0589	0.000558137	Foxc1	Nfatc1	Smad
1810011O10Rik	1.2752	0.003996684	NA	Nfatc1	Smad
4732456N10Rik	-1.9007	0.005718684	NA	NA	NA
Ablim1	-1.3446	2.79E-05	NA	Nfatc1	Smad
Ache	-2.9973	2.65E-06	NA	Nfatc1	NA
Ackr4	-1.2066	0.000111881	NA	NA	Smad
Acsbg1	-1.1076	0.003437838	NA	Nfatc1	Smad
Acta2	1.6405	1.34E-06	NA	Nfatc1	Smad
Actb	-1.0188	0.000971648	NA	NA	Smad
Actg1	-0.8728	0.003521374	NA	NA	Smad
Adamtsl4	-2.2282	2.34E-15	NA	Nfatc1	Smad
Adcy1	-3.9658	4.57E-41	NA	NA	Smad
Adrb2	-1.6122	2.55E-08	NA	Nfatc1	Smad
Afap111	1.3140	0.008056303	NA	NA	Smad
AI661453	-0.9177	0.004557753	NA	NA	Smad
Ak3	-0.9796	0.003315456	NA	NA	Smad
Aldh1a2	2.7177	6.56E-11	NA	NA	Smad
Aldh3a1	-2.2093	4.68E-08	NA	NA	NA
Alox12e	-1.2808	0.000188118	NA	NA	Smad
Alox8	-2.0492	0.000164332	NA	NA	NA
Aloxe3	-2.0016	3.40E-10	NA	NA	Smad
Alpl	1.6288	0.000500133	NA	NA	NA
Ano1	-2.7121	4.38E-11	NA	NA	Smad
Anpep	-1.3105	2.79E-05	NA	NA	Smad
Anxa3	1.4131	0.00010305	NA	NA	Smad
Anxa8	-1.6437	8.61E-11	NA	NA	Smad
Aqp3	-2.2338	7.08E-15	NA	NA	Smad
Arhgap23	-1.2650	9.60E-06	NA	NA	Smad
Arrdc3	1.1492	0.000209491	NA	Nfatc1	Smad
Arrdc4	0.9930	0.001046049	NA	Nfatc1	Smad
Asap2	-1.4468	2.48E-05	NA	Nfatc1	Smad
Aspn	-1.7531	0.000641614	NA	NA	Smad
Atoh8	-2.3714	1.20E-05	NA	Nfatc1	Smad
Atp6v0e2	-2.0111	0.002422271	NA	NA	Smad
Avpi1	-1.2316	9.16E-06	NA	NA	NA
Baiap2	-1.0197	0.008192995	NA	Nfatc1	Smad
BC064078	0.9992	0.00875729	NA	NA	Smad
Bok	-1.7146	8.35E-08	NA	Nfatc1	NA
Cacna2d2	-2.6330	9.47E-08	NA	Nfatc1	Smad
Cadm4	-1.9973	0.000662869	NA	NA	Smad
Camk4	-3.0750	1.80E-26	NA	NA	NA
Camkk1	-1.9972	1.30E-07	NA	Nfatc1	Smad
Car12	1.1394	0.0032333	NA	NA	Smad
Card10	-1.2558	9.07E-05	NA	Nfatc1	NA
Ccdc3	-0.9785	0.002518273	NA	NA	Smad
Ccl27a	-1.3031	0.00813563	NA	Nfatc1	Smad
Ccnb2	1.2562	0.003266714	NA	NA	NA
Cd200	0.9038	0.004505948	NA	Nfatc1	Smad
Cd207	-1.9370	0.000188362	NA	NA	NA

Cd276	1.0544	0.009731004 NA	Nfatc1	NA
Cd59a	0.9637	0.004884904 NA	NA	NA
Cdca8	1.2315	0.005315007 NA	NA	Smad
Cdh13	-1.9387	8.71E-09 NA	Nfatc1	Smad
Cdk1	1.2660	0.002938117 NA	NA	Smad
Cenpe	1.3297	0.001870907 NA	Nfatc1	Smad
Cers4	1.0109	0.003282894 NA	Nfatc1	Smad
Cers5	-0.9224	0.005099877 NA	NA	NA
Ces2g	-1.0170	0.007495317 NA	NA	Smad
Cgnl1	-1.5813	7.95E-07 NA	NA	Smad
Cgref1	-1.5555	0.008935975 NA	NA	Smad
Chat	-3.9175	2.70E-18 NA	NA	NA
Clca2	-1.3276	0.000445915 NA	NA	NA
Clcf1	-1.3269	2.59E-06 NA	Nfatc1	Smad
Cldn4	-1.2522	2.13E-05 NA	NA	Smad
Clic3	-2.1515	2.76E-09 NA	NA	Smad
Cmah	1.2364	1.02E-05 NA	NA	Smad
Cnksr3	-1.6787	4.27E-10 NA	Nfatc1	Smad
Cnn3	0.9587	0.001553566 NA	Nfatc1	NA
Cntn2	2.1407	4.98E-07 NA	NA	Smad
Col14a1	-1.4099	9.91E-05 NA	Nfatc1	Smad
Col6a1	-3.4520	2.04E-32 NA	Nfatc1	Smad
Col6a2	-1.8754	1.05E-09 NA	NA	NA
Col7a1	-0.9908	0.000458064 NA	NA	NA
Col8a2	-2.7821	1.51E-22 NA	NA	Smad
Cpa4	-2.8090	4.16E-27 NA	Nfatc1	Smad
Crif1	-2.9592	1.51E-22 NA	NA	NA
Cry1	-1.0390	0.0049324 NA	NA	Smad
Csn3	2.0179	0.001514014 NA	NA	NA
Ctgf	-1.2436	8.33E-06 NA	Nfatc1	Smad
Cxcl10	1.5832	0.001845649 NA	NA	Smad
Cyp4b1	-1.7728	2.77E-06 NA	NA	NA
Cystm1	-1.8845	5.45E-07 NA	NA	Smad
D17H6S56E-5	1.1336	0.001898451 NA	NA	NA
Dach1	0.8977	0.007495317 NA	Nfatc1	Smad
Dap	-1.6185	7.94E-08 NA	NA	Smad
Dedd2	-1.1377	0.000135529 NA	NA	Smad
Dhrs2	-3.2561	2.19E-07 NA	Nfatc1	Smad
Dhx32	-1.1036	0.004387595 NA	NA	Smad
Dmkn	-1.9037	1.60E-15 NA	Nfatc1	Smad
Duox1	-2.4797	8.39E-15 NA	NA	Smad
Dusp4	1.5049	1.35E-05 NA	Nfatc1	NA
Dusp7	-1.0112	0.000790902 NA	NA	Smad
Ecm1	-2.0504	3.34E-11 NA	NA	Smad
Eda	-1.1451	0.003208001 NA	Nfatc1	NA
Ednra	-3.5106	3.50E-18 NA	NA	Smad
Egfl6	-1.3751	5.05E-06 NA	NA	Smad
Ehd3	1.3201	0.002540937 NA	Nfatc1	Smad
Eif2s3y	-8.6144	7.95E-32 NA	NA	NA
Eml1	-2.5642	7.11E-07 NA	NA	Smad
Eps8l1	-1.2890	0.006377176 NA	NA	Smad
Ereg	-1.7765	0.00883146 NA	NA	NA

Evpl	-1.3468	8.74E-07	NA	Smad
Fads3	1.3608	0.000405697	NA	Smad
Fads6	-1.8731	0.005774605	NA	NA
Faim2	-2.9180	0.000110424	NA	NA
Fam19a5	-2.3746	0.001948144	NA	Smad
Fam213a	-1.6769	4.56E-06	Nfatc1	Smad
Fam25c	-2.0770	1.64E-10	NA	Smad
Fam84a	0.9338	0.001803975	Nfatc1	Smad
Fcgbp	-2.0408	1.54E-08	NA	Smad
Fchsd1	-1.0630	0.000411714	NA	NA
Fgf1	-1.5364	0.000506602	Nfatc1	Smad
Fgf18	-3.1202	1.16E-05	NA	Smad
Foxc1	-7.1912	6.64E-67	NA	Smad
Galnt15	-3.0004	5.52E-05	NA	NA
Gfra2	-1.6068	0.007709063	NA	Smad
Ggt1	1.5550	0.00020566	NA	NA
Gjb2	1.3988	3.79E-07	NA	Smad
Gjb6	1.2831	0.000200031	NA	Smad
Gm6969	-6.6848	2.45E-13	NA	NA
Gm7694	-2.2975	1.25E-06	Nfatc1	NA
Gm973	-1.4771	5.12E-06	NA	Smad
Gpr161	-1.1198	0.004267601	Nfatc1	Smad
Gpx2	2.7084	7.09E-09	NA	NA
Grem1	-2.7895	7.88E-20	NA	NA
Grik1	-4.1114	8.71E-05	Nfatc1	Smad
Guca2a	-2.8281	1.48E-06	NA	NA
Hhip	-2.9837	2.00E-06	NA	Smad
Hid1	-3.4540	5.78E-08	NA	NA
Hist1h1c	1.0614	0.001024664	Nfatc1	Smad
Hivep3	-2.0519	6.43E-08	Nfatc1	Smad
Hk2	-1.2723	1.80E-05	Nfatc1	Smad
Hoxa7	-1.5478	2.39E-06	NA	Smad
Hoxc13	0.8382	0.00875729	NA	Smad
Hs3st3a1	2.2853	1.25E-06	Nfatc1	Smad
Hsp90aa1	-0.8907	0.001942955	NA	Smad
Hspa1a	-1.1093	4.20E-05	NA	Smad
Hspa1b	-1.2023	8.36E-06	NA	Smad
Hspb1	-1.1102	2.79E-05	NA	Smad
Igdcc4	-1.7351	1.07E-07	NA	NA
Igfbp5	-2.3138	8.31E-18	NA	Smad
Insig2	0.8175	0.008605465	Nfatc1	Smad
Irx4	0.8895	0.003982278	Nfatc1	Smad
Itga8	2.0963	0.004412929	Nfatc1	Smad
Kank1	-1.1843	1.30E-05	NA	Smad
Kcnc4	-2.4302	4.39E-07	NA	Smad
Kcnf1	2.2435	3.10E-11	NA	Smad
Kcnh3	-2.1602	0.000143913	NA	NA
Kctd9	-0.9499	0.007330706	NA	NA
Kif26a	-1.6792	0.000303326	NA	Smad
Klf2	-1.1946	0.00342966	NA	NA
Klhl21	-0.8942	0.007330706	NA	NA
Klk10	-1.1816	0.004412929	NA	Smad

Klk7	-1.2616	4.74E-05	NA	NA
Knstrn	1.5310	0.000288715	NA	Smad
Kprp	-1.4687	0.001117229	NA	Smad
Krt15	-1.8538	1.17E-14	NA	NA
Krt16	-1.0508	0.000441076	NA	NA
Krt24	-3.8062	7.14E-52	NA	NA
Krt80	-2.0225	2.32E-09	Nfatc1	Smad
Lamb3	-1.2186	5.38E-06	Nfatc1	Smad
Lgals9	1.0764	0.000718787	NA	NA
Lgr6	-2.1222	5.53E-09	Nfatc1	Smad
Lrrfp1	-0.9523	0.003276225	Nfatc1	Smad
Ltbp1	-1.8357	9.61E-07	Nfatc1	Smad
Ltbp2	-1.2902	0.000410499	Nfatc1	Smad
Lurap1l	1.6887	2.51E-05	Nfatc1	Smad
Mafk	-1.0359	0.000218007	Nfatc1	NA
Map2k3	-0.8822	0.00635124	Nfatc1	Smad
Marveld1	-1.1969	6.09E-05	NA	Smad
Mbd1	1.0436	0.003648519	NA	NA
Mcm6	1.3417	0.000581044	NA	NA
Mertk	-1.0333	0.002775385	NA	Smad
Mettl20	1.5652	0.009171888	NA	Smad
Mfap3l	-1.2449	9.07E-05	Nfatc1	Smad
Mir27a	-2.0848	0.00348925	NA	NA
Mitf	-2.8100	1.48E-13	Nfatc1	Smad
Mki67	1.0714	0.000854876	Nfatc1	Smad
Mme	-2.9208	5.44E-06	NA	Smad
Mrgprf	-2.6605	3.05E-16	NA	NA
Myh14	-2.0567	1.47E-16	Nfatc1	Smad
Myo1e	-0.9890	0.009520936	Nfatc1	Smad
Ncald	1.3182	0.00829638	Nfatc1	Smad
Neu2	-1.9509	8.87E-06	NA	NA
Ngef	-4.3897	2.09E-17	NA	Smad
Niacr1	1.1377	7.72E-05	NA	NA
Nkd1	-2.0431	2.51E-05	NA	Smad
Nlrp10	1.0614	0.002181163	NA	NA
Nod2	-1.3086	5.48E-05	NA	Smad
Nog	-3.0027	1.72E-09	NA	Smad
Npas2	-3.1153	1.94E-09	Nfatc1	Smad
Npnt	-1.9054	6.33E-07	Nfatc1	Smad
Nppb	-1.1608	0.007513545	NA	Smad
Nptx1	-3.3346	1.16E-16	NA	NA
Nr1d1	-0.9595	0.000909645	Nfatc1	Smad
Nr3c2	-2.5421	0.006407435	NA	Smad
Nrep	-1.0811	0.002634776	Nfatc1	Smad
Nrip3	-1.4876	0.000156313	NA	Smad
Nrtn	-1.8142	1.80E-05	NA	Smad
Nt5dc2	-1.0626	0.002921347	NA	NA
Nt5e	-2.3542	4.89E-18	Nfatc1	Smad
Nxpe4	-1.9277	0.000303667	NA	Smad
Ogdhl	-2.0957	0.001555434	Nfatc1	Smad
Ovol1	-1.0054	0.00951081	Nfatc1	Smad
Pacsin3	-1.2760	0.000167485	NA	Smad

Papln	-1.7603	1.72E-08 NA	NA	Smad
Pard6b	-1.2572	3.31E-05 NA	Nfatc1	Smad
Pcbp3	-1.0543	0.008127229 NA	NA	NA
Pcdh7	-1.1347	0.000797489 NA	Nfatc1	Smad
Pcolce2	-1.7459	0.007996183 NA	NA	Smad
Pcp4	3.3145	2.98E-09 NA	NA	Smad
Pcsk6	-2.5444	9.81E-17 NA	NA	Smad
Pdlim3	-1.3478	0.00011219 NA	NA	Smad
Pdzrn4	-1.8741	2.59E-09 NA	NA	Smad
Peg3	-1.5076	1.25E-06 NA	NA	NA
Piezo1	-1.4518	0.000303326 NA	Nfatc1	Smad
Pknox2	-2.5854	0.001778778 NA	Nfatc1	Smad
Plod1	-1.0240	0.001538596 NA	NA	Smad
Pmaip1	1.1217	0.000490253 NA	NA	Smad
Ppap2a	-1.4717	0.000122945 NA	NA	NA
Ppap2b	-0.9793	0.007709063 NA	Nfatc1	Smad
Ppif	-1.4567	9.35E-09 NA	NA	Smad
Ppp2cb	0.9670	0.007330706 NA	NA	Smad
Ppp2r3a	-1.2247	6.01E-05 NA	Nfatc1	Smad
Prkcz	-2.0200	0.000913641 NA	NA	Smad
Prnp	-1.2943	1.49E-05 NA	Nfatc1	Smad
Ptges	-1.6783	1.54E-08 NA	NA	Smad
Ptpre	-2.0837	1.80E-07 NA	Nfatc1	Smad
Racgap1	1.1218	0.004150883 NA	NA	Smad
Ramp3	-1.7892	6.07E-05 NA	NA	NA
Rapgef3	-1.4508	0.002526823 NA	NA	Smad
Rarres1	-1.8002	0.003069944 NA	NA	Smad
Rbp1	1.0599	0.004687749 NA	NA	Smad
Rcan1	-1.6713	3.65E-07 NA	NA	Smad
Rhov	1.0892	0.002673519 NA	NA	Smad
Rnf208	-1.7779	5.20E-09 NA	NA	Smad
Rnf39	-0.9473	0.002794214 NA	NA	Smad
Rorc	-1.5659	0.000275335 NA	NA	NA
RP24-390G17.1	-2.0922	1.91E-08 NA	NA	NA
Rufy4	-1.2887	0.002526823 NA	NA	NA
S100a4	-2.5757	1.85E-16 NA	NA	NA
Sbsn	-1.4048	2.90E-08 NA	NA	NA
Scd2	1.3157	6.49E-05 NA	Nfatc1	Smad
Scn5a	-4.5545	4.42E-09 NA	NA	Smad
Scrg1	1.1714	0.000492458 NA	NA	Smad
Scube2	-1.2724	0.001033256 NA	NA	Smad
Sepp1	0.8660	0.005502974 NA	Nfatc1	Smad
Serpina3k	4.2055	0.000227008 NA	NA	NA
Serpinb10	-1.5309	1.88E-05 NA	NA	Smad
Serpinb11	-2.5536	6.36E-16 NA	NA	Smad
Serpinb2	-1.3907	4.68E-08 NA	Nfatc1	Smad
Serpinb3b	-2.6127	1.43E-05 NA	Nfatc1	Smad
Serpinf1	1.0805	0.003375481 NA	NA	NA
Sgms2	-1.8189	5.36E-06 NA	Nfatc1	NA
Sh3rf2	-1.8322	6.16E-07 NA	Nfatc1	Smad
Skint10	-2.0353	0.000751229 NA	NA	NA
Slc16a2	-1.3747	0.000134742 NA	Nfatc1	Smad

Slc16a6	1.9522	1.21E-07 NA	NA	NA
Slc25a45	1.3656	0.00816966 NA	NA	Smad
Slc26a9	-1.4587	0.004284346 NA	NA	Smad
Slc35e4	-0.8768	0.007495317 NA	NA	NA
Slc38a3	-1.7199	0.0004334 NA	NA	NA
Slc6a2	1.0200	0.007712176 NA	Nfatc1	Smad
Slc6a4	-1.9745	0.000686606 NA	NA	Smad
Slc6a9	-1.0553	0.003718224 NA	NA	Smad
Smad9	-3.4743	2.37E-07 NA	Nfatc1	Smad
Smim5	-1.7751	0.002674666 NA	NA	Smad
Sms	-2.0005	5.38E-07 NA	NA	Smad
Sncg	-3.2016	1.41E-05 NA	NA	NA
Sned1	-1.2716	0.003331641 NA	NA	NA
Snn	1.1657	0.000976669 NA	NA	Smad
Sostdc1	-1.3113	2.98E-05 NA	NA	Smad
Sphkap	-5.2819	1.49E-13 NA	Nfatc1	Smad
Spock2	-1.2343	0.006681372 NA	NA	NA
Spon2	-1.1411	0.002076387 NA	NA	NA
Srd5a2	4.9740	1.27E-07 NA	NA	Smad
Srf	-0.9443	0.007437061 NA	NA	NA
Srgap3	-1.2459	0.003731033 NA	Nfatc1	Smad
Sulf2	-1.2417	2.89E-05 NA	NA	Smad
Susd2	-1.6430	1.12E-05 NA	NA	NA
Tead4	-1.5703	0.003437838 NA	NA	Smad
Tekt2	-1.8341	4.44E-10 NA	Nfatc1	Smad
Tenc1	-1.0725	0.000885688 NA	NA	Smad
Tfap2b	-0.8629	0.00928501 NA	Nfatc1	Smad
Tgm5	-4.8200	8.94E-65 NA	Nfatc1	Smad
Tgm7	-1.9056	0.005000353 NA	NA	Smad
Timp3	1.1480	1.53E-05 NA	NA	NA
Tle4	-1.6602	3.24E-05 NA	Nfatc1	Smad
Tmem108	-2.8003	2.51E-05 NA	Nfatc1	Smad
Tmem40	-1.1385	0.009421823 NA	Nfatc1	Smad
Tmem45a	-1.6595	7.46E-07 NA	Nfatc1	Smad
Tmtc1	-1.5493	0.000529905 NA	Nfatc1	Smad
Tnfrsf11b	-1.9695	8.35E-07 NA	Nfatc1	Smad
Tns1	-1.8510	8.01E-13 NA	NA	Smad
Top2a	0.9077	0.007893428 NA	NA	NA
Trf	1.6195	4.35E-06 NA	NA	NA
Tslp	-1.6753	0.00238777 NA	NA	NA
Ttc39b	-1.2855	0.0004334 NA	Nfatc1	Smad
Tubb2b	1.4947	2.90E-06 NA	NA	Smad
Tubb4b	-0.8132	0.008196179 NA	NA	Smad
Ubash3b	-1.5722	0.000200031 NA	NA	Smad
Ugdh	-0.9265	0.006123833 NA	NA	Smad
Unc13b	-1.2943	0.008935975 NA	NA	Smad
Upb1	-2.2315	9.20E-06 NA	NA	NA
Usp2	-0.9588	0.002526823 NA	Nfatc1	Smad
Uty	-5.7140	1.69E-15 NA	NA	NA
Vps37b	-1.0612	0.00020566 NA	Nfatc1	Smad
Vwa2	-2.7765	1.11E-23 NA	NA	NA
Wfdc3	1.9477	4.20E-05 NA	NA	NA

Wif1	-1.2033	0.006516727 NA	NA	Smad
Wipi2	-0.8385	0.008965218 NA	NA	Smad
Wnt11	-1.2791	0.002217554 NA	Nfatc1	Smad
Wnt4	-1.1996	0.000437973 NA	NA	Smad
Wnt7b	-1.2350	0.000383246 NA	NA	Smad
Zbtb16	-2.0306	2.80E-05 NA	NA	Smad
Zfand1	1.1012	0.006233126 NA	NA	Smad
Zfp750	1.5096	0.000976669 NA	NA	Smad
Zyx	-0.8839	0.003428783 NA	Nfatc1	Smad

**Supplementary Table S3 ATAC peaks containing motifs of
Foxc1/Nfatc1/Smad in differentially expressed gene of P30 Foxc1 cKO
HFSC RNA-seq**

chromosome	start	end	Associated gene	Matched motif
chr1	17740379	17740788	Crispld1	Smad
chr1	17743770	17744179	Crispld1	Foxc1
chr1	19208694	19211434	Tfap2b	Nfatc1,Smad
chr1	19211650	19214230	Tfap2b	Nfatc1
chr1	19215753	19218865	Tfap2b	Smad
chr1	19219324	19221477	Tfap2b	Smad
chr1	19233823	19234282	Tfap2b	Smad
chr1	19269562	19270067	Tfap2b	Smad
chr1	19438290	19438490	Tfap2b	Nfatc1
chr1	19450770	19451204	Tfap2b	Smad
chr1	19567816	19568222	Tfap2b	Nfatc1
chr1	38251505	38252268	Aff3	Smad
chr1	38291951	38292472	Aff3	Foxc1
chr1	38307904	38308816	Aff3	Smad
chr1	38478289	38478935	Aff3	Foxc1
chr1	38628163	38628647	Aff3	Smad
chr1	39192494	39193385	Npas2	Nfatc1
chr1	39193969	39195478	Npas2	Nfatc1
chr1	39228678	39230100	Npas2	Nfatc1
chr1	39251343	39252168	Npas2	Smad
chr1	59549141	59549626	Gm973	Smad
chr1	72871735	72872207	Igfbp5	Smad
chr1	72887788	72888393	Igfbp5	Smad
chr1	73947002	73948411	Tns1	Smad
chr1	74027420	74028051	Tns1	Smad
chr1	75214317	75215297	Tuba4a	Foxc1
chr1	79819080	79819280	Serpine2	Smad
chr1	79858901	79859347	Serpine2	Nfatc1
chr1	79869750	79870049	Serpine2	Smad
chr1	79895151	79895358	Serpine2	Smad
chr1	79898224	79898424	Serpine2	Foxc1,Nfatc1,Smad
chr1	82476438	82477044	Col4a4	Smad
chr1	82578021	82578221	Col4a4	Foxc1
chr1	82597058	82597781	Col4a3	Foxc1
chr1	82634475	82635099	Col4a3	Foxc1,Smad
chr1	82687406	82688338	Col4a3	Nfatc1
chr1	82711849	82712156	Col4a3	Foxc1,Nfatc1

chr1	83407802	83408552	Sphkap	Smad
chr1	83611671	83614068	Sphkap	Nfatc1
chr1	87483465	87483712	Ngef	Smad
chr1	91024079	91024504	Lrrfip1	Nfatc1
chr1	91044089	91044703	Lrrfip1	Nfatc1
chr1	91077180	91078510	Lrrfip1	Smad
chr1	91078699	91079135	Lrrfip1	Nfatc1,Smad
chr1	91126759	91127362	Lrrfip1	Smad
chr1	93679650	93680815	Bok	Nfatc1
chr1	107129933	107130197	Serpinb3b	Nfatc1
chr1	107214703	107215080	Serpinb3b	Smad
chr1	107365147	107366621	Serpinb11	Smad
chr1	107495389	107496000	Serpinb2	Smad
chr1	107501967	107502422	Serpinb2	Nfatc1
chr1	107512178	107519995	Serpinb2	Nfatc1
chr1	107562218	107562476	Serpinb10	Smad
chr1	107592120	107595723	Serpinb8	Smad
chr1	107600510	107600741	Serpinb8	Foxc1
chr1	107602980	107603387	Serpinb8	Smad
chr1	107607395	107608454	Serpinb8	Nfatc1
chr1	107634585	107635166	Serpinb8	Smad
chr1	107644439	107644893	Serpinb8	Smad
chr1	121241055	121241438	Insig2	Smad
chr1	121243588	121243864	Insig2	Smad
chr1	121249791	121251315	Insig2	Smad
chr1	121261791	121262637	Insig2	Nfatc1
chr1	121275969	121278395	Insig2	Smad
chr1	121280457	121280865	Insig2	Smad
chr1	121301294	121302248	Insig2	Smad
chr1	121320474	121320908	Insig2	Smad
chr1	121329375	121331057	Insig2	Smad
chr1	131733323	131733814	Slc26a9	Smad
chr1	131751449	131752690	Slc26a9	Smad
chr1	132544414	132545042	Cntn2	Smad
chr1	135018802	135020929	Lgr6	Nfatc1
chr1	135021953	135022365	Lgr6	Smad
chr1	150536265	150536680	Hmcn1	Nfatc1
chr1	150577187	150577387	Hmcn1	Smad
chr1	150907481	150908343	Hmcn1	Nfatc1
chr1	150918021	150918782	Hmcn1	Nfatc1
chr1	150951422	150953484	Hmcn1	Smad
chr1	150958978	150960961	Hmcn1	Smad
chr1	150971366	150972070	Hmcn1	Smad

chr1	150992929	150994103	Hmcn1	Foxc1,Nfatc1
chr1	151008824	151009757	Hmcn1	Nfatc1,Smad
chr1	151019245	151019840	Hmcn1	Smad
chr1	151062109	151062749	Hmcn1	Foxc1,Nfatc1,Smad
chr1	165277323	165278646	Gpr161	Nfatc1
chr1	165301043	165303770	Gpr161	Smad
chr1	165319897	165320195	Gpr161	Smad
chr1	170292409	170292934	Gm7694	Nfatc1
chr1	193294045	193294246	Lamb3	Smad
chr1	193298095	193298613	Lamb3	Smad
chr1	193300013	193300568	Lamb3	Smad
chr1	193300711	193302458	Lamb3	Nfatc1
chr1	193326638	193327729	Lamb3	Smad
chr2	5165377	5166531	Ccdc3	Smad
chr2	5176517	5177194	Ccdc3	Smad
chr2	11963091	11963291	Itga8	Smad
chr2	12053608	12054743	Itga8	Nfatc1
chr2	12236452	12236728	Itga8	Smad
chr2	12249359	12249743	Itga8	Nfatc1,Smad
chr2	12255563	12256199	Itga8	Smad
chr2	12300838	12302430	Itga8	Smad
chr2	25223781	25225713	Tubb4b	Smad
chr2	25243384	25244014	Rnf208	Smad
chr2	25455706	25457800	Clic3	Smad
chr2	30919950	30920755	Ptges	Smad
chr2	50015457	50015889	Lypd6	Nfatc1
chr2	50049084	50049384	Lypd6	Smad
chr2	50093620	50094432	Lypd6	Foxc1
chr2	50101548	50101941	Lypd6	Smad
chr2	50104106	50104542	Lypd6	Smad
chr2	50110023	50110590	Lypd6	Smad
chr2	60600885	60601207	Itgb6	Smad
chr2	60639253	60641040	Itgb6	Smad
chr2	60667393	60669060	Itgb6	Smad
chr2	60672161	60674943	Itgb6	Smad
chr2	60688558	60689773	Itgb6	Smad
chr2	60699743	60700552	Itgb6	Smad
chr2	60720203	60720634	Itgb6	Smad
chr2	60735197	60735465	Itgb6	Foxc1,Smad
chr2	83812278	83813117	Fam171b	Smad
chr2	83813351	83814591	Fam171b	Smad
chr2	83815319	83815749	Fam171b	Smad
chr2	83883588	83885179	Fam171b	Foxc1,Nfatc1

chr2	91254917	91255510	Pacsin3	Smad
chr2	91255604	91258334	Pacsin3	Smad
chr2	102530965	102531696	Pamr1	Smad
chr2	102544144	102544533	Pamr1	Smad
chr2	102552054	102552364	Pamr1	Foxc1
chr2	102807499	102808039	Cd44	Nfatc1
chr2	102827074	102827501	Cd44	Smad
chr2	102837614	102839379	Cd44	Nfatc1
chr2	102849405	102851947	Cd44	Smad
chr2	102862310	102862963	Cd44	Nfatc1
chr2	102871664	102875784	Cd44	Smad
chr2	102893168	102895958	Cd44	Foxc1,Nfatc1
chr2	102908909	102909929	Cd44	Smad
chr2	102939661	102941496	Cd44	Smad
chr2	102947938	102949108	Cd44	Smad
chr2	102957877	102960017	Cd44	Nfatc1
chr2	109524051	109524900	Bdnf	Foxc1
chr2	109563121	109563377	Bdnf	Smad
chr2	109693264	109695325	Bdnf	Smad
chr2	109721750	109723159	Bdnf	Smad
chr2	113363605	113364908	Fmn1	Smad
chr2	113374131	113374761	Fmn1	Nfatc1
chr2	113386562	113387656	Fmn1	Foxc1,Nfatc1,Smad
chr2	113397454	113398374	Fmn1	Smad
chr2	113404815	113405522	Fmn1	Smad
chr2	113420589	113422045	Fmn1	Nfatc1,Smad
chr2	113428242	113429074	Fmn1	Smad
chr2	113449414	113450548	Fmn1	Smad
chr2	113469274	113469898	Fmn1	Nfatc1
chr2	113494621	113495124	Fmn1	Smad
chr2	113544831	113545040	Fmn1	Smad
chr2	113548607	113549213	Fmn1	Nfatc1,Smad
chr2	118813886	118814694	Knstrn	Smad
chr2	119262974	119263575	Rhov	Smad
chr2	121056000	121056436	Tgm5	Nfatc1,Smad
chr2	121068357	121070962	Tgm5	Smad
chr2	121071627	121072366	Tgm5	Nfatc1,Smad
chr2	121117933	121118152	Tgm7	Smad
chr2	122336530	122337180	Duox1	Smad
chr2	122340886	122341738	Duox1	Smad
chr2	122344416	122346658	Duox1	Smad
chr2	128709076	128711091	Mertk	Smad
chr2	131495175	131495426	Smox	Nfatc1,Smad

chr2	131509491	131510520	Smox	Smad
chr2	131522750	131522950	Smox	Foxc1
chr2	131902518	131903623	Prnp	Smad
chr2	131909328	131911015	Prnp	Smad
chr2	131934987	131936157	Prnp	Nfatc1
chr2	133263672	133263942	Bmp2	Smad
chr2	133305231	133306599	Bmp2	Smad
chr2	133326299	133327093	Bmp2	Foxc1
chr2	133336130	133336448	Bmp2	Nfatc1
chr2	133378483	133379178	Bmp2	Smad
chr2	133391560	133393711	Bmp2	Smad
chr2	133397491	133398051	Bmp2	Smad
chr2	133431586	133433044	Bmp2	Smad
chr2	133443401	133444063	Bmp2	Smad
chr2	133455575	133456106	Bmp2	Smad
chr2	133551933	133553444	Bmp2	Smad
chr2	137710184	137710973	Btbd3	Smad
chr2	137715909	137716365	Btbd3	Smad
chr2	137719802	137720320	Btbd3	Nfatc1
chr2	137757028	137757986	Btbd3	Smad
chr2	138241022	138241738	Btbd3	Smad
chr2	138247741	138248394	Btbd3	Nfatc1
chr2	138279675	138280014	Btbd3	Smad
chr2	138281498	138282216	Btbd3	Nfatc1,Smad
chr2	138283327	138283661	Btbd3	Smad
chr2	138291194	138291983	Btbd3	Nfatc1
chr2	138294494	138294702	Btbd3	Smad
chr2	138320628	138322168	Btbd3	Foxc1,Nfatc1,Smad
chr2	138437714	138438089	Btbd3	Smad
chr2	138477151	138477515	Btbd3	Smad
chr2	138477747	138477976	Btbd3	Nfatc1
chr2	138483785	138484365	Btbd3	Smad
chr2	138485775	138486056	Btbd3	Smad
chr2	138503820	138504208	Btbd3	Nfatc1
chr2	138548772	138549484	Btbd3	Smad
chr2	166116090	166116307	Sulf2	Smad
chr2	168045709	168046189	Pard6b	Nfatc1
chr2	168072802	168073488	Pard6b	Smad
chr2	168080412	168081317	Pard6b	Smad
chr3	9974263	9974681	Fabp5	Foxc1,Nfatc1
chr3	10349630	10349830	Zfand1	Smad
chr3	10350308	10351945	Zfand1	Smad
chr3	54754846	54756703	Smad9	Smad

chr3	54764145	54764718	Smad9	Nfatc1
chr3	63412987	63414338	Mme	Smad
chr3	67481392	67481688	Rarres1	Smad
chr3	84190604	84192025	Trim2	Smad
chr3	84259431	84259901	Trim2	Smad
chr3	84287162	84287362	Trim2	Smad
chr3	84314603	84315153	Trim2	Foxc1,Smad
chr3	84318543	84320503	Trim2	Smad
chr3	84323798	84325274	Trim2	Nfatc1,Smad
chr3	90597377	90598757	S100a3	Foxc1
chr3	90599980	90602594	S100a3	Smad
chr3	92822429	92822654	Kprp	Smad
chr3	92825968	92826168	Kprp	Smad
chr3	95674266	95676237	Adamtsl4	Nfatc1
chr3	95696260	95696950	Adamtsl4	Smad
chr3	95732774	95735708	Ecm1	Smad
chr3	95738365	95739875	Ecm1	Smad
chr3	96669549	96670734	Ankrd35	Smad
chr3	96675127	96675363	Ankrd35	Foxc1
chr3	100505968	100508026	Fam46c	Foxc1,Smad
chr3	102430258	102430663	Ngf	Smad
chr3	102469716	102470537	Ngf	Foxc1,Smad
chr3	102500888	102501826	Ngf	Nfatc1
chr3	102574403	102574788	Ngf	Nfatc1
chr3	102738328	102739035	Tspan2	Foxc1,Smad
chr3	107399857	107400428	Kcnc4	Smad
chr3	107445714	107446341	Kcnc4	Smad
chr3	107458291	107458915	Kcnc4	Smad
chr3	121424645	121425960	Cnn3	Nfatc1
chr3	131327475	131328308	Sgms2	Nfatc1
chr3	132880758	132881517	Npnt	Nfatc1
chr3	132893735	132894605	Npnt	Smad
chr3	132915301	132916323	Npnt	Smad
chr3	132945578	132946312	Npnt	Smad
chr3	135155059	135156338	Cenpe	Smad
chr3	135245575	135245998	Cenpe	Nfatc1,Smad
chr3	135816492	135817156	Slc39a8	Foxc1
chr3	135824354	135826924	Slc39a8	Smad
chr3	135886543	135886961	Slc39a8	Smad
chr3	145651232	145651864	Cyr61	Nfatc1
chr3	145681266	145681847	Cyr61	Smad
chr3	145684570	145685704	Cyr61	Foxc1
chr4	41082750	41083562	Aqp3	Smad

chr4	41813695	41814008	Ccl27a	Smad
chr4	41822346	41823476	Ccl27a	Nfatc1,Smad
chr4	43058407	43059566	Unc13b	Smad
chr4	43208997	43209996	Unc13b	Smad
chr4	63949691	63950648	Tnc	Smad
chr4	63958528	63959370	Tnc	Smad
chr4	63982513	63982715	Tnc	Smad
chr4	63982979	63983212	Tnc	Smad
chr4	63996128	63996493	Tnc	Nfatc1,Smad
chr4	64009739	64010294	Tnc	Smad
chr4	64046476	64047562	Tnc	Smad
chr4	64048275	64048616	Tnc	Nfatc1
chr4	64092202	64093402	Tnc	Smad
chr4	64131955	64132869	Tnc	Smad
chr4	64142234	64142985	Tnc	Smad
chr4	64156300	64156614	Tnc	Smad
chr4	64175972	64176172	Tnc	Smad
chr4	64219886	64220419	Tnc	Foxc1
chr4	64237660	64238634	Tnc	Smad
chr4	64281118	64281352	Tnc	Smad
chr4	64306891	64307202	Tnc	Nfatc1,Smad
chr4	64324324	64324815	Tnc	Nfatc1
chr4	64431369	64432797	Tnc	Smad
chr4	64451189	64452268	Tnc	Nfatc1,Smad
chr4	64497798	64498210	Tnc	Nfatc1
chr4	80886533	80887195	Lurap1l	Smad
chr4	80910606	80911605	Lurap1l	Nfatc1
chr4	83301849	83303883	Ttc39b	Nfatc1,Smad
chr4	83342500	83343263	Ttc39b	Smad
chr4	83350458	83351578	Ttc39b	Nfatc1,Smad
chr4	105156872	105157872	Ppap2b	Smad
chr4	105158154	105158579	Ppap2b	Smad
chr4	105175524	105175724	Ppap2b	Smad
chr4	105213937	105214983	Ppap2b	Nfatc1,Smad
chr4	105225609	105225928	Ppap2b	Smad
chr4	105242158	105242618	Ppap2b	Smad
chr4	105245595	105246122	Ppap2b	Smad
chr4	105328660	105329597	Ppap2b	Nfatc1
chr4	105352790	105353204	Ppap2b	Smad
chr4	105480057	105480618	Ppap2b	Smad
chr4	105608389	105608642	Ppap2b	Smad
chr4	105640822	105641781	Ppap2b	Smad
chr4	105692362	105692942	Ppap2b	Nfatc1

chr4	114348883	114349980	Trabd2b	Foxc1,Smad
chr4	114361016	114361920	Trabd2b	Smad
chr4	114370984	114371192	Trabd2b	Smad
chr4	114377400	114378924	Trabd2b	Smad
chr4	114382460	114383473	Trabd2b	Nfatc1
chr4	114384745	114388832	Trabd2b	Smad
chr4	114394539	114394905	Trabd2b	Smad
chr4	114405589	114407889	Trabd2b	Smad
chr4	114410359	114410884	Trabd2b	Foxc1,Nfatc1
chr4	114411946	114412415	Trabd2b	Smad
chr4	114509121	114510935	Trabd2b	Smad
chr4	114545501	114546123	Trabd2b	Smad
chr4	114564277	114564648	Trabd2b	Foxc1,Smad
chr4	114601309	114602245	Trabd2b	Smad
chr4	114610858	114613273	Trabd2b	Nfatc1
chr4	114613947	114614789	Trabd2b	Smad
chr4	114635829	114636285	Trabd2b	Smad
chr4	114665598	114666164	Trabd2b	Nfatc1,Smad
chr4	114688535	114688836	Trabd2b	Nfatc1
chr4	117829534	117829959	Slc6a9	Smad
chr4	119739194	119740384	Hivep3	Smad
chr4	119785027	119785319	Hivep3	Smad
chr4	119786926	119787126	Hivep3	Smad
chr4	119789435	119790125	Hivep3	Smad
chr4	119932189	119933693	Hivep3	Smad
chr4	119998137	119999164	Hivep3	Nfatc1
chr4	120031532	120033545	Hivep3	Smad
chr4	120071414	120072838	Hivep3	Smad
chr4	120131495	120131995	Hivep3	Smad
chr4	124936645	124937483	Cdca8	Smad
chr4	126309812	126310246	Col8a2	Smad
chr4	126320972	126322221	Tekt2	Smad
chr4	126324934	126325752	Tekt2	Nfatc1
chr4	126326813	126328494	Tekt2	Nfatc1,Smad
chr4	127347519	127350559	Gjb4	Smad
chr4	127352240	127353224	Gjb4	Foxc1
chr4	137271820	137272098	Wnt4	Smad
chr4	147936265	147939183	Plod1	Smad
chr4	147993200	147994426	Nppb	Smad
chr4	155275617	155276234	Prkcz	Smad
chr4	155360827	155362119	Prkcz	Smad
chr5	13915207	13915448	Sema3e	Nfatc1
chr5	13932380	13932793	Sema3e	Smad

chr5	13974630	13975059	Sema3e	Smad
chr5	13977235	13977506	Sema3e	Smad
chr5	13983740	13984631	Sema3e	Smad
chr5	13992563	13993515	Sema3e	Smad
chr5	13993932	13994430	Sema3e	Smad
chr5	14022737	14023168	Sema3e	Smad
chr5	14026626	14026969	Sema3e	Foxc1
chr5	14032168	14032493	Sema3e	Smad
chr5	14048671	14049165	Sema3e	Smad
chr5	14059709	14061390	Sema3e	Foxc1,Nfatc1,Smad
chr5	14062529	14064922	Sema3e	Smad
chr5	14072876	14073557	Sema3e	Smad
chr5	14074875	14075226	Sema3e	Nfatc1
chr5	14075807	14076115	Sema3e	Smad
chr5	14087487	14089104	Sema3e	Smad
chr5	14089534	14091535	Sema3e	Nfatc1
chr5	14177799	14178015	Sema3e	Nfatc1
chr5	14198614	14198916	Sema3e	Smad
chr5	14198922	14199224	Sema3e	Nfatc1
chr5	14230663	14231760	Sema3e	Nfatc1
chr5	14271300	14271500	Sema3e	Smad
chr5	14306421	14306999	Sema3e	Smad
chr5	14313687	14314118	Sema3e	Foxc1
chr5	14315630	14315842	Sema3e	Nfatc1
chr5	14317196	14317464	Sema3e	Smad
chr5	17539260	17539460	Sema3c	Nfatc1
chr5	17544246	17545022	Sema3c	Smad
chr5	17571212	17572240	Sema3c	Smad
chr5	17573476	17576757	Sema3c	Foxc1
chr5	17623236	17623859	Sema3c	Nfatc1
chr5	17641773	17642344	Sema3c	Smad
chr5	17711339	17711706	Sema3c	Smad
chr5	30945194	30946272	Cgref1	Smad
chr5	57713917	57720265	Pcdh7	Smad
chr5	57721991	57722848	Pcdh7	Smad
chr5	57732544	57733819	Pcdh7	Smad
chr5	57755645	57756334	Pcdh7	Nfatc1,Smad
chr5	57765325	57765626	Pcdh7	Smad
chr5	57801732	57803684	Pcdh7	Smad
chr5	57884765	57885442	Pcdh7	Nfatc1
chr5	57906963	57907177	Pcdh7	Smad
chr5	57918968	57919406	Pcdh7	Nfatc1
chr5	57940474	57940902	Pcdh7	Nfatc1,Smad

chr5	57942689	57944629	Pcdh7	Smad
chr5	57961984	57963205	Pcdh7	Smad
chr5	57976129	57976843	Pcdh7	Smad
chr5	57986686	57987919	Pcdh7	Nfatc1
chr5	57996296	57996496	Pcdh7	Smad
chr5	58034026	58034791	Pcdh7	Smad
chr5	58050906	58051298	Pcdh7	Smad
chr5	65428301	65429506	Ugdh	Smad
chr5	65434755	65437500	Ugdh	Smad
chr5	77074817	77075430	Hopx	Foxc1
chr5	77080350	77081744	Hopx	Smad
chr5	77085657	77085857	Hopx	Smad
chr5	77089256	77089680	Hopx	Nfatc1
chr5	77091997	77093170	Hopx	Smad
chr5	77094672	77096426	Hopx	Nfatc1
chr5	77114669	77115772	Hopx	Smad
chr5	77122932	77123232	Hopx	Foxc1
chr5	77136614	77137087	Hopx	Nfatc1
chr5	88853307	88854053	Slc4a4	Foxc1,Smad
chr5	88886041	88886290	Slc4a4	Smad
chr5	88949947	88950740	Slc4a4	Nfatc1
chr5	88956130	88956390	Slc4a4	Smad
chr5	88956820	88957445	Slc4a4	Smad
chr5	88994576	88994788	Slc4a4	Smad
chr5	89002353	89002553	Slc4a4	Nfatc1,Smad
chr5	89101979	89102188	Slc4a4	Nfatc1
chr5	89139914	89140114	Slc4a4	Smad
chr5	91476661	91477158	Parm1	Smad
chr5	91491486	91492151	Parm1	Smad
chr5	91517292	91518059	Parm1	Smad
chr5	91574272	91574552	Parm1	Nfatc1,Smad
chr5	91606305	91607449	Parm1	Smad
chr5	91620322	91621733	Parm1	Smad
chr5	91651226	91651625	Parm1	Foxc1
chr5	91653042	91653960	Parm1	Foxc1
chr5	92348747	92349192	Cxcl10	Smad
chr5	96790188	96790539	Anxa3	Smad
chr5	96792737	96793107	Anxa3	Smad
chr5	96804238	96804786	Anxa3	Smad
chr5	96905947	96906631	Anxa3	Smad
chr5	100865421	100866686	Agpat9	Smad
chr5	100869127	100869744	Agpat9	Foxc1
chr5	100886193	100886922	Agpat9	Smad

chr5	100917458	100917731	Agpat9	Smad
chr5	100919215	100919492	Agpat9	Nfatc1
chr5	101020822	101021077	Agpat9	Foxc1
chr5	101057045	101057373	Agpat9	Nfatc1
chr5	101091130	101091722	Agpat9	Smad
chr5	101126268	101126611	Agpat9	Smad
chr5	101163115	101163601	Agpat9	Nfatc1
chr5	101172680	101173411	Agpat9	Smad
chr5	101173839	101174744	Agpat9	Smad
chr5	101216805	101217649	Agpat9	Nfatc1
chr5	101241167	101241367	Agpat9	Smad
chr5	116392643	116393284	Hspb8	Smad
chr5	116394881	116395423	Hspb8	Smad
chr5	116399982	116400653	Hspb8	Foxc1
chr5	116410163	116412080	Hspb8	Smad
chr5	116417171	116418420	Hspb8	Nfatc1,Smad
chr5	116419124	116421289	Hspb8	Smad
chr5	116421623	116424555	Hspb8	Smad
chr5	124006568	124007926	Vps37b	Nfatc1,Smad
chr5	134946914	134947295	Cldn4	Smad
chr5	134951110	134951646	Cldn4	Smad
chr5	134965916	134966909	Cldn4	Smad
chr5	135887295	135891657	Hspb1	Smad
chr5	137293157	137295011	Ache	Nfatc1
chr5	139796011	139797621	Mafk	Nfatc1
chr5	142628981	142629939	Wipi2	Smad
chr5	142920648	142921315	Actb	Smad
chr6	29430831	29433705	Finc	Foxc1,Smad
chr6	30567930	30571307	Cpa4	Nfatc1,Smad
chr6	36691605	36691869	Ptn	Smad
chr6	36701104	36701453	Ptn	Smad
chr6	36771865	36772801	Ptn	Smad
chr6	36797439	36797785	Ptn	Foxc1,Smad
chr6	36808136	36810529	Ptn	Smad
chr6	42339750	42340965	Zyx	Nfatc1,Smad
chr6	42351453	42352677	Zyx	Smad
chr6	42352844	42353747	Zyx	Smad
chr6	48537144	48537936	Atp6v0e2	Smad
chr6	52216863	52218400	Hoxa7	Smad
chr6	58536380	58536580	Abcg2	Nfatc1
chr6	58594057	58594517	Abcg2	Smad
chr6	58608499	58608721	Abcg2	Nfatc1
chr6	58610590	58611039	Abcg2	Smad

chr6	58652455	58653044	Abcg2	Foxc1
chr6	58751264	58751707	Abcg2	Nfatc1
chr6	72220352	72220788	Atoh8	Nfatc1
chr6	72242432	72243768	Atoh8	Smad
chr6	72261174	72261641	Atoh8	Smad
chr6	82702681	82703064	Hk2	Smad
chr6	82740116	82740856	Hk2	Smad
chr6	82750623	82751483	Hk2	Nfatc1,Smad
chr6	82771842	82775390	Hk2	Smad
chr6	87160006	87160758	Antxr1	Smad
chr6	87161190	87161732	Antxr1	Smad
chr6	87186388	87187603	Antxr1	Smad
chr6	87189842	87190130	Antxr1	Foxc1,Nfatc1,Smad
chr6	87222985	87223324	Antxr1	Smad
chr6	87277483	87277863	Antxr1	Foxc1
chr6	87281902	87282660	Antxr1	Smad
chr6	91617142	91617572	Slc6a6	Smad
chr6	91647177	91648075	Slc6a6	Smad
chr6	91652072	91653069	Slc6a6	Smad
chr6	91675291	91675578	Slc6a6	Smad
chr6	91679252	91679660	Slc6a6	Foxc1,Smad
chr6	91683690	91684731	Slc6a6	Smad
chr6	97743724	97744709	Mitf	Smad
chr6	97894509	97894905	Mitf	Nfatc1
chr6	98121021	98121316	Mitf	Smad
chr6	98666523	98666744	Foxp1	Smad
chr6	98667317	98667640	Foxp1	Foxc1,Smad
chr6	98726621	98726965	Foxp1	Smad
chr6	98779104	98779373	Foxp1	Smad
chr6	98842823	98843306	Foxp1	Smad
chr6	98850711	98851440	Foxp1	Smad
chr6	98947295	98949341	Foxp1	Foxc1,Nfatc1,Smad
chr6	98949586	98950517	Foxp1	Smad
chr6	99030099	99030527	Foxp1	Smad
chr6	99084433	99085389	Foxp1	Smad
chr6	99171783	99172303	Foxp1	Foxc1
chr6	99217777	99218352	Foxp1	Smad
chr6	99266334	99266534	Foxp1	Smad
chr6	99284127	99284340	Foxp1	Smad
chr6	99350601	99350933	Foxp1	Smad
chr6	99388448	99389198	Foxp1	Smad
chr6	99392542	99393035	Foxp1	Smad
chr6	99440186	99440513	Foxp1	Foxc1,Nfatc1

chr6	99527983	99529705	Foxp1	Foxc1
chr6	112818312	112818953	Srgap3	Smad
chr6	112822996	112823580	Srgap3	Nfatc1
chr6	112842560	112842761	Srgap3	Nfatc1
chr6	112946145	112947639	Srgap3	Smad
chr6	115760853	115761729	Tmem40	Nfatc1,Smad
chr6	115762343	115762625	Tmem40	Smad
chr6	118791798	118792578	Cacna1c	Smad
chr6	119018217	119019607	Cacna1c	Smad
chr6	119081714	119082102	Cacna1c	Foxc1,Smad
chr6	119089103	119089709	Cacna1c	Smad
chr6	119104347	119105132	Cacna1c	Smad
chr6	119174864	119175920	Cacna1c	Smad
chr6	128281878	128283035	Tead4	Smad
chr6	128308320	128308673	Tead4	Smad
chr6	128992040	128993590	BC064078	Smad
chr6	135343990	135344622	Emp1	Smad
chr6	135377364	135386988	Emp1	Nfatc1
chr6	135387827	135388027	Emp1	Foxc1
chr6	135398698	135399385	Emp1	Smad
chr6	135430533	135431024	Emp1	Smad
chr6	135444717	135445039	Emp1	Smad
chr6	135504549	135504951	Emp1	Nfatc1,Smad
chr6	145881688	145882362	Bhlhe41	Foxc1
chr6	148315684	148315959	Tmtc1	Smad
chr6	148335262	148335663	Tmtc1	Nfatc1
chr6	148336622	148337043	Tmtc1	Smad
chr6	148360482	148361244	Tmtc1	Smad
chr6	148442551	148444899	Tmtc1	Smad
chr6	148454769	148455639	Tmtc1	Smad
chr6	148468994	148470640	Tmtc1	Smad
chr6	149132367	149132841	Mettl20	Smad
chr6	149138010	149138210	Mettl20	Smad
chr7	3149945	3151278	AU018091	Foxc1
chr7	3184418	3184627	AU018091	Nfatc1,Smad
chr7	4463565	4465762	Eps8l1	Smad
chr7	4475642	4476687	Eps8l1	Smad
chr7	24480154	24482051	Cadm4	Smad
chr7	24499957	24500808	Cadm4	Smad
chr7	25208795	25209959	Dedd2	Smad
chr7	25217518	25221889	Dedd2	Smad
chr7	28100599	28101012	Fcgbp	Smad
chr7	30762524	30764212	Dmkn	Nfatc1

chr7	30775613	30777804	Dmkn	Smad
chr7	43788166	43788470	Klk10	Smad
chr7	44661794	44663359	Myh14	Nfatc1,Smad
chr7	44667948	44670002	Myh14	Smad
chr7	44672263	44672672	Myh14	Smad
chr7	65872489	65874155	Pcsk6	Smad
chr7	65881215	65881614	Pcsk6	Smad
chr7	66842074	66842417	Adamts17	Foxc1
chr7	66933305	66934548	Adamts17	Nfatc1
chr7	67024850	67025099	Adamts17	Smad
chr7	67064584	67064784	Adamts17	Nfatc1,Smad
chr7	67186356	67186568	Adamts17	Smad
chr7	68681974	68683532	Arrdc4	Smad
chr7	68691271	68691829	Arrdc4	Smad
chr7	68717595	68718517	Arrdc4	Smad
chr7	68722864	68723409	Arrdc4	Smad
chr7	68736659	68736973	Arrdc4	Smad
chr7	68747624	68750204	Arrdc4	Smad
chr7	68823212	68823658	Arrdc4	Smad
chr7	68829999	68830284	Arrdc4	Smad
chr7	68833524	68833845	Arrdc4	Nfatc1,Smad
chr7	69086799	69086999	Arrdc4	Smad
chr7	79829362	79829575	Anpep	Smad
chr7	88214367	88215678	Ctsc	Nfatc1,Smad
chr7	88248816	88249201	Ctsc	Smad
chr7	88270918	88272256	Ctsc	Nfatc1
chr7	88344711	88344911	Ctsc	Smad
chr7	88346261	88346563	Ctsc	Nfatc1
chr7	88365590	88365883	Ctsc	Foxc1,Smad
chr7	89581046	89581260	Me3	Smad
chr7	89610872	89611914	Me3	Nfatc1,Smad
chr7	89687812	89688126	Me3	Smad
chr7	89698302	89698637	Me3	Smad
chr7	89707587	89708651	Me3	Smad
chr7	89712557	89713273	Me3	Smad
chr7	89769218	89769801	Me3	Foxc1
chr7	89787121	89788781	Me3	Smad
chr7	89799611	89801570	Me3	Smad
chr7	89809282	89809907	Me3	Smad
chr7	89835470	89836226	Me3	Nfatc1,Smad
chr7	89837297	89838976	Me3	Foxc1,Smad
chr7	98788449	98789159	Wnt11	Smad
chr7	98815057	98818366	Wnt11	Smad

chr7	98837660	98839201	Wnt11	Nfatc1,Smad
chr7	100493707	100494250	Ucp2	Foxc1
chr7	100495091	100496081	Ucp2	Smad
chr7	100500125	100500340	Ucp2	Smad
chr7	109781028	109782620	Nrip3	Smad
chr7	109836690	109838016	Scube2	Smad
chr7	109852508	109853695	Scube2	Smad
chr7	109872655	109873165	Scube2	Smad
chr7	133780704	133780916	Dhx32	Smad
chr7	135537275	135538537	Ptpre	Smad
chr7	135572073	135572624	Ptpre	Smad
chr7	135605114	135605752	Ptpre	Nfatc1
chr7	135609574	135611046	Ptpre	Smad
chr7	135612357	135613538	Ptpre	Smad
chr7	135631865	135632224	Ptpre	Nfatc1
chr7	135653300	135654737	Ptpre	Smad
chr7	135659291	135660029	Ptpre	Smad
chr7	135699262	135699836	Mki67	Nfatc1
chr7	135728013	135728667	Mki67	Nfatc1,Smad
chr7	135734452	135736784	Mki67	Smad
chr7	135740158	135740949	Mki67	Nfatc1,Smad
chr7	135748948	135749626	Mki67	Smad
chr7	135768439	135768946	Mki67	Nfatc1,Smad
chr7	135889142	135889510	Mki67	Nfatc1,Smad
chr7	144684577	144684827	Ano1	Smad
chr8	4433082	4433785	Cers4	Nfatc1
chr8	4436375	4436887	Cers4	Smad
chr8	4481349	4482298	Cers4	Nfatc1,Smad
chr8	4483088	4484444	Cers4	Smad
chr8	4491498	4494318	Cers4	Nfatc1
chr8	4495204	4496781	Cers4	Smad
chr8	4505111	4505812	Cers4	Smad
chr8	4507592	4508123	Cers4	Smad
chr8	4512029	4512663	Cers4	Smad
chr8	7469096	7469874	Efnb2	Nfatc1
chr8	7876408	7876608	Efnb2	Smad
chr8	7944179	7944751	Efnb2	Smad
chr8	7944889	7945412	Efnb2	Nfatc1
chr8	7946043	7946903	Efnb2	Smad
chr8	7977010	7977644	Efnb2	Foxc1,Smad
chr8	8040827	8041253	Efnb2	Foxc1,Nfatc1,Smad
chr8	8050735	8051531	Efnb2	Nfatc1,Smad
chr8	8102348	8103078	Efnb2	Smad

chr8	8107731	8107977	Efnb2	Smad
chr8	8109298	8110448	Efnb2	Smad
chr8	8122874	8123627	Efnb2	Smad
chr8	8132373	8133685	Efnb2	Smad
chr8	8142017	8143490	Efnb2	Nfatc1
chr8	8246074	8247233	Efnb2	Smad
chr8	8254505	8255363	Efnb2	Nfatc1
chr8	8260018	8261089	Efnb2	Smad
chr8	8273663	8275760	Efnb2	Smad
chr8	8303241	8303442	Efnb2	Smad
chr8	8307155	8308193	Efnb2	Smad
chr8	8315775	8315986	Efnb2	Nfatc1
chr8	8365544	8366223	Efnb2	Foxc1
chr8	8452094	8452923	Efnb2	Smad
chr8	8457641	8458048	Efnb2	Nfatc1,Smad
chr8	8461901	8462101	Efnb2	Nfatc1
chr8	8488429	8489536	Efnb2	Smad
chr8	8499926	8500269	Efnb2	Smad
chr8	8567238	8569036	Efnb2	Smad
chr8	8639356	8640658	Efnb2	Smad
chr8	8648767	8650957	Efnb2	Nfatc1,Smad
chr8	8652448	8652809	Efnb2	Nfatc1
chr8	8653072	8653341	Efnb2	Smad
chr8	11628262	11629031	Ankrd10	Smad
chr8	11632466	11634138	Ankrd10	Foxc1
chr8	11634353	11638290	Ankrd10	Smad
chr8	24317539	24318425	1810011O10Rik	Nfatc1
chr8	24364355	24365786	1810011O10Rik	Smad
chr8	24419739	24421324	1810011O10Rik	Nfatc1
chr8	24427702	24428315	1810011O10Rik	Nfatc1
chr8	24430189	24431896	1810011O10Rik	Smad
chr8	24450246	24450874	1810011O10Rik	Smad
chr8	31713940	31715065	Nrg1	Foxc1
chr8	31720464	31721130	Nrg1	Smad
chr8	31812869	31813069	Nrg1	Foxc1
chr8	31847087	31847287	Nrg1	Foxc1
chr8	31875790	31875990	Nrg1	Smad
chr8	31906815	31907103	Nrg1	Smad
chr8	31907851	31908051	Nrg1	Smad
chr8	31927654	31928515	Nrg1	Nfatc1
chr8	33601485	33601778	Ppp2cb	Smad
chr8	33606160	33606505	Ppp2cb	Smad
chr8	33607250	33607678	Ppp2cb	Smad

chr8	34804242	34809185	Dusp4	Nfatc1
chr8	35205408	35205866	Ppp1r3b	Smad
chr8	35223966	35224680	Ppp1r3b	Smad
chr8	35243048	35244309	Ppp1r3b	Foxc1,Nfatc1
chr8	35275993	35277030	Ppp1r3b	Smad
chr8	35300383	35301335	Ppp1r3b	Smad
chr8	35358393	35358628	Ppp1r3b	Smad
chr8	35401401	35402098	Ppp1r3b	Smad
chr8	35419838	35420435	Ppp1r3b	Smad
chr8	40424788	40425180	Zdhhc2	Foxc1
chr8	40464474	40465768	Zdhhc2	Smad
chr8	40485384	40486492	Zdhhc2	Nfatc1,Smad
chr8	45875101	45875858	Pdlim3	Smad
chr8	57428804	57429875	Scrg1	Smad
chr8	60597683	60597903	Mfap3l	Nfatc1,Smad
chr8	60605419	60605821	Mfap3l	Nfatc1
chr8	60627427	60628575	Mfap3l	Smad
chr8	60639497	60640818	Mfap3l	Nfatc1
chr8	60728926	60729503	Mfap3l	Smad
chr8	60773322	60773949	Mfap3l	Smad
chr8	75083211	75084769	Hmox1	Foxc1,Smad
chr8	75089248	75090304	Hmox1	Smad
chr8	75098353	75098554	Hmox1	Smad
chr8	75100491	75101753	Hmox1	Smad
chr8	76973150	76973617	Nr3c2	Smad
chr8	77132742	77133430	Nr3c2	Smad
chr8	77723489	77724781	Ednra	Smad
chr8	77774645	77775452	Ednra	Smad
chr8	77775497	77775814	Ednra	Smad
chr8	77787639	77788120	Ednra	Smad
chr8	79903865	79904176	Hhip	Smad
chr8	80057351	80058059	Hhip	Smad
chr8	81341454	81342098	Inpp4b	Smad
chr8	81342500	81343031	Inpp4b	Smad
chr8	81383965	81384556	Inpp4b	Smad
chr8	81480010	81480247	Inpp4b	Smad
chr8	81624462	81624868	Inpp4b	Smad
chr8	81625716	81625925	Inpp4b	Nfatc1
chr8	81736291	81736576	Inpp4b	Smad
chr8	81745500	81745783	Inpp4b	Nfatc1,Smad
chr8	81756572	81758183	Inpp4b	Smad
chr8	81767078	81768255	Inpp4b	Foxc1
chr8	81812210	81812628	Inpp4b	Smad

chr8	81849687	81850762	Inpp4b	Nfatc1,Smad
chr8	81854102	81855842	Inpp4b	Smad
chr8	81859828	81861993	Inpp4b	Foxc1,Smad
chr8	81947021	81947379	Inpp4b	Foxc1
chr8	81991965	81992581	Inpp4b	Smad
chr8	81996352	81997371	Inpp4b	Smad
chr8	81998864	81999122	Inpp4b	Smad
chr8	88536982	88537446	Nkd1	Smad
chr8	88683446	88684248	Nod2	Smad
chr8	92960337	92961516	Slc6a2	Nfatc1
chr8	92965649	92966520	Slc6a2	Nfatc1
chr8	92972756	92973533	Slc6a2	Smad
chr8	92984275	92985020	Slc6a2	Smad
chr8	92994458	92994658	Slc6a2	Smad
chr8	104961247	104962326	Ces2g	Smad
chr8	109947506	109948824	Marveld3	Foxc1
chr8	118285482	118287880	Cdh13	Nfatc1
chr8	118316398	118317254	Cdh13	Smad
chr8	118374702	118375336	Cdh13	Smad
chr8	118376560	118377383	Cdh13	Smad
chr8	118435223	118435667	Cdh13	Smad
chr8	118518725	118519156	Cdh13	Nfatc1
chr8	118565150	118565645	Cdh13	Smad
chr8	119064564	119064841	Cdh13	Smad
chr8	122525865	122526958	Piezo1	Smad
chr8	122536767	122537948	Piezo1	Nfatc1,Smad
chr9	35424693	35425091	Cdon	Smad
chr9	35467361	35467980	Cdon	Foxc1
chr9	35493257	35494066	Cdon	Smad
chr9	35522635	35524918	Cdon	Smad
chr9	36890496	36890949	Pknox2	Smad
chr9	36894668	36895603	Pknox2	Nfatc1
chr9	36933703	36934242	Pknox2	Nfatc1
chr9	36941584	36942176	Pknox2	Smad
chr9	37145257	37146069	Pknox2	Smad
chr9	37176830	37177504	Pknox2	Smad
chr9	41020680	41020969	Ubash3b	Smad
chr9	41021642	41022184	Ubash3b	Smad
chr9	41028526	41028726	Ubash3b	Smad
chr9	41031413	41032389	Ubash3b	Smad
chr9	44066599	44068560	Usp2	Nfatc1
chr9	44087628	44088608	Usp2	Smad
chr9	47283550	47284095	Cadm1	Foxc1

chr9	47343595	47344039	Cadm1	Nfatc1,Smad
chr9	47350159	47350708	Cadm1	Smad
chr9	47357726	47358640	Cadm1	Nfatc1,Smad
chr9	47389833	47390503	Cadm1	Smad
chr9	47529552	47530575	Cadm1	Smad
chr9	47589308	47590465	Cadm1	Smad
chr9	47625078	47626157	Cadm1	Foxc1,Nfatc1,Smad
chr9	47674766	47675120	Cadm1	Smad
chr9	47792547	47793037	Cadm1	Nfatc1
chr9	47813010	47813937	Cadm1	Smad
chr9	48407748	48408099	Nxpe4	Smad
chr9	48707486	48707956	Zbtb16	Smad
chr9	48739997	48740618	Zbtb16	Smad
chr9	54629618	54630184	Acsbg1	Smad
chr9	54635080	54636519	Acsbg1	Nfatc1,Smad
chr9	54642336	54643848	Acsbg1	Smad
chr9	54668544	54669097	Acsbg1	Smad
chr9	58555421	58556008	Cd276	Nfatc1
chr9	66713437	66714147	Car12	Smad
chr9	66722317	66722695	Car12	Smad
chr9	68786832	68787307	Rora	Smad
chr9	68944111	68944455	Rora	Smad
chr9	68972336	68972634	Rora	Smad
chr9	69010999	69011199	Rora	Foxc1
chr9	69213715	69214174	Rora	Smad
chr9	69289166	69290064	Rora	Nfatc1
chr9	69306654	69307340	Rora	Foxc1,Smad
chr9	70207004	70208976	Myo1e	Smad
chr9	70211206	70212353	Myo1e	Smad
chr9	70248707	70250592	Myo1e	Nfatc1
chr9	70290505	70291143	Myo1e	Smad
chr9	71321420	71322014	Aldh1a2	Smad
chr9	71355548	71356425	Aldh1a2	Smad
chr9	71771006	71771754	Cgnl1	Smad
chr9	88058232	88059333	Nt5e	Smad
chr9	88177634	88178283	Nt5e	Nfatc1
chr9	88259060	88261232	Nt5e	Smad
chr9	88276777	88277806	Nt5e	Smad
chr9	88325607	88327157	Nt5e	Smad
chr9	88330060	88331316	Nt5e	Smad
chr9	88332278	88333256	Nt5e	Smad
chr9	88339711	88340199	Nt5e	Nfatc1
chr9	95624111	95624445	Pcolce2	Smad

chr9	95672370	95672701	Pcolce2	Smad
chr9	98417930	98419086	Rbp1	Smad
chr9	98421431	98422353	Rbp1	Smad
chr9	98432753	98434824	Rbp1	Smad
chr9	99319515	99320080	Esyt3	Smad
chr9	99339812	99340870	Esyt3	Smad
chr9	99354723	99355430	Esyt3	Foxc1
chr9	99357537	99359523	Esyt3	Smad
chr9	101168682	101168882	Ppp2r3a	Smad
chr9	101228859	101229079	Ppp2r3a	Nfatc1,Smad
chr9	101250986	101252205	Ppp2r3a	Smad
chr9	101265410	101266545	Ppp2r3a	Nfatc1,Smad
chr9	103506876	103507737	Tmem108	Smad
chr9	103550414	103551618	Tmem108	Smad
chr9	103597869	103598106	Tmem108	Smad
chr9	103847798	103848805	Tmem108	Nfatc1,Smad
chr9	104102094	104102628	Ackr4	Smad
chr9	106367021	106373131	Dusp7	Smad
chr9	106376656	106379627	Dusp7	Smad
chr9	106392345	106392632	Dusp7	Smad
chr9	106394365	106394942	Dusp7	Smad
chr9	107396277	107396929	Cacna2d2	Nfatc1,Smad
chr9	107526543	107526972	Cacna2d2	Smad
chr9	119511058	119511351	Scn5a	Smad
chr9	119558172	119558639	Scn5a	Smad
chr9	119579083	119580039	Scn5a	Smad
chr9	119580472	119581181	Scn5a	Smad
chr10	7118085	7118665	Cnksr3	Smad
chr10	7214586	7215216	Cnksr3	Smad
chr10	7219588	7219788	Cnksr3	Smad
chr10	7257848	7259388	Cnksr3	Nfatc1
chr10	7269157	7269676	Cnksr3	Smad
chr10	7289512	7290096	Cnksr3	Smad
chr10	7295635	7297195	Cnksr3	Smad
chr10	14331799	14333283	Gpr126	Foxc1
chr10	14385706	14386120	Gpr126	Smad
chr10	14413122	14413767	Gpr126	Smad
chr10	14418273	14419751	Gpr126	Foxc1
chr10	14482375	14482886	Gpr126	Nfatc1,Smad
chr10	14491641	14492515	Gpr126	Smad
chr10	14543635	14545635	Gpr126	Nfatc1
chr10	14582889	14583863	Gpr126	Nfatc1
chr10	24210545	24211427	Moxd1	Smad

chr10	24223059	24223848	Moxd1	Smad
chr10	24245286	24245536	Moxd1	Nfatc1
chr10	24259882	24260121	Moxd1	Smad
chr10	24276434	24276847	Moxd1	Foxc1
chr10	24298474	24298984	Moxd1	Foxc1,Smad
chr10	24379415	24380129	Moxd1	Smad
chr10	24445920	24448269	Moxd1	Smad
chr10	24463101	24463862	Ctgf	Smad
chr10	24478545	24479585	Ctgf	Nfatc1
chr10	24573739	24574037	Ctgf	Smad
chr10	24595157	24596280	Ctgf	Smad
chr10	56339070	56339554	Gja1	Smad
chr10	56364084	56364361	Gja1	Nfatc1,Smad
chr10	56373022	56373525	Gja1	Foxc1,Nfatc1
chr10	56376088	56379512	Gja1	Nfatc1
chr10	56384809	56386417	Gja1	Smad
chr10	56396761	56397180	Gja1	Smad
chr10	56415832	56416485	Gja1	Smad
chr10	56424800	56425634	Gja1	Nfatc1,Smad
chr10	56431235	56432224	Gja1	Nfatc1,Smad
chr10	56455894	56456495	Gja1	Nfatc1
chr10	56474954	56475424	Gja1	Smad
chr10	56484630	56485791	Gja1	Smad
chr10	56495136	56495877	Gja1	Smad
chr10	56500789	56501267	Gja1	Smad
chr10	56535261	56535639	Gja1	Smad
chr10	56566327	56567270	Gja1	Smad
chr10	56597978	56598722	Gja1	Nfatc1
chr10	56606333	56607213	Gja1	Nfatc1,Smad
chr10	56611579	56612881	Gja1	Smad
chr10	56642590	56642994	Gja1	Nfatc1
chr10	56660182	56661465	Gja1	Nfatc1
chr10	56724602	56724944	Gja1	Nfatc1
chr10	56774978	56775547	Gja1	Foxc1,Smad
chr10	56781906	56782735	Gja1	Nfatc1
chr10	56783834	56784368	Gja1	Nfatc1
chr10	56796483	56796813	Gja1	Foxc1
chr10	56831083	56832545	Gja1	Foxc1,Smad
chr10	56910702	56911692	Gja1	Smad
chr10	56913981	56915466	Gja1	Nfatc1
chr10	60760105	60760855	Unc5b	Smad
chr10	60800450	60800822	Unc5b	Foxc1
chr10	60805172	60806280	Unc5b	Smad

chr10	68110116	68111711	Arid5b	Smad
chr10	68121521	68122647	Arid5b	Smad
chr10	68136083	68138132	Arid5b	Smad
chr10	68142110	68142931	Arid5b	Smad
chr10	68156224	68158443	Arid5b	Smad
chr10	68229480	68231335	Arid5b	Foxc1,Nfatc1,Smad
chr10	68237711	68239137	Arid5b	Smad
chr10	68248185	68249901	Arid5b	Foxc1,Smad
chr10	68268546	68269202	Arid5b	Smad
chr10	68282488	68283728	Arid5b	Smad
chr10	69321158	69322357	Cdk1	Smad
chr10	69383233	69383780	Cdk1	Smad
chr10	69398041	69399617	Cdk1	Smad
chr10	76668932	76669189	Col6a1	Smad
chr10	76729831	76730944	Col6a1	Nfatc1,Smad
chr10	76732855	76733786	Col6a1	Smad
chr10	85161482	85162348	Cry1	Smad
chr10	121160128	121162152	Wif1	Smad
chr11	6898883	6900659	Adcy1	Smad
chr11	7205664	7206147	Igfbp3	Nfatc1,Smad
chr11	7206439	7206900	Igfbp3	Foxc1,Smad
chr11	7211371	7211673	Igfbp3	Smad
chr11	7215845	7216698	Igfbp3	Smad
chr11	7226566	7226991	Igfbp3	Foxc1,Smad
chr11	7423148	7424528	Igfbp3	Nfatc1,Smad
chr11	7425299	7425702	Igfbp3	Smad
chr11	7446116	7446502	Igfbp3	Smad
chr11	7455041	7455794	Igfbp3	Nfatc1,Smad
chr11	7526469	7526669	Igfbp3	Smad
chr11	7603829	7604288	Igfbp3	Smad
chr11	7794056	7794657	Igfbp3	Smad
chr11	32998998	33000354	Fgf18	Smad
chr11	33146295	33148911	Fgf18	Smad
chr11	60941068	60941793	Map2k3	Nfatc1,Smad
chr11	60953717	60954159	Map2k3	Smad
chr11	62814071	62814285	Trim16	Smad
chr11	62819920	62821432	Trim16	Smad
chr11	62841299	62841575	Trim16	Foxc1,Smad
chr11	62841684	62841892	Trim16	Smad
chr11	64498304	64498642	Hs3st3a1	Nfatc1
chr11	64502702	64503634	Hs3st3a1	Smad
chr11	64518049	64518629	Hs3st3a1	Smad
chr11	64644206	64644594	Hs3st3a1	Smad

chr11	64718657	64719147	Hs3st3a1	Smad
chr11	65023949	65024956	Arhgap44	Smad
chr11	65039241	65040200	Arhgap44	Foxc1
chr11	65048648	65049619	Arhgap44	Smad
chr11	65067281	65067879	Arhgap44	Smad
chr11	65128128	65129467	Arhgap44	Nfatc1
chr11	65161207	65163210	Arhgap44	Foxc1
chr11	69133601	69134787	Aloxe3	Smad
chr11	70288071	70288507	Alox12e	Smad
chr11	72994320	72995196	Atp2a3	Foxc1
chr11	73018124	73020107	Camkk1	Nfatc1,Smad
chr11	73020884	73023969	Camkk1	Smad
chr11	76997351	76998294	Slc6a4	Smad
chr11	89206183	89207700	Nog	Smad
chr11	97434771	97436202	Arhgap23	Smad
chr11	97468461	97469203	Arhgap23	Smad
chr11	98768459	98769305	Nr1d1	Smad
chr11	98772526	98774717	Nr1d1	Nfatc1,Smad
chr11	99061776	99063870	Tns4	Foxc1,Smad
chr11	99080562	99084111	Tns4	Nfatc1
chr11	99092329	99092672	Tns4	Smad
chr11	99098811	99099339	Tns4	Nfatc1
chr11	113835190	113835528	Sdk2	Smad
chr11	113865571	113865927	Sdk2	Smad
chr11	113892762	113893243	Sdk2	Foxc1
chr11	113951626	113952357	Sdk2	Smad
chr11	114066770	114067434	Sdk2	Smad
chr11	114080552	114081190	Sdk2	Smad
chr11	115899830	115900204	Smim5	Smad
chr11	116237848	116238998	Evpl	Smad
chr11	119942436	119944082	Baiap2	Nfatc1
chr11	119991187	119991580	Baiap2	Smad
chr11	120346202	120351684	Actg1	Smad
chr11	121515893	121516745	Zfp750	Smad
chr11	121517079	121517844	Zfp750	Smad
chr12	13928382	13928760	Fam84a	Smad
chr12	13957134	13957383	Fam84a	Smad
chr12	14068616	14069937	Fam84a	Smad
chr12	14091264	14091464	Fam84a	Smad
chr12	14106119	14106749	Fam84a	Smad
chr12	14155155	14155838	Fam84a	Smad
chr12	14157537	14158334	Fam84a	Smad
chr12	14189219	14190155	Fam84a	Nfatc1

chr12	17112704	17113538	Kcnf1	Smad
chr12	21092443	21093915	Asap2	Nfatc1
chr12	21097191	21098583	Asap2	Smad
chr12	21135837	21136683	Asap2	Smad
chr12	21159890	21160832	Asap2	Nfatc1
chr12	21166804	21167911	Asap2	Smad
chr12	21171527	21172432	Asap2	Smad
chr12	36308265	36309376	Sostdc1	Smad
chr12	36315428	36316136	Sostdc1	Smad
chr12	36317002	36317996	Sostdc1	Smad
chr12	36343270	36344965	Sostdc1	Smad
chr12	41483475	41484999	Lrrn3	Smad
chr12	41485204	41486039	Lrrn3	Foxc1
chr12	44325648	44325849	Nrcam	Smad
chr12	44328889	44329493	Nrcam	Smad
chr12	44331973	44332493	Nrcam	Foxc1
chr12	44336055	44336284	Nrcam	Smad
chr12	44345867	44347364	Nrcam	Smad
chr12	44355136	44355756	Nrcam	Smad
chr12	44524592	44525655	Nrcam	Nfatc1
chr12	44536132	44536640	Nrcam	Foxc1
chr12	44547905	44548216	Nrcam	Smad
chr12	44572772	44573089	Nrcam	Smad
chr12	44599675	44600035	Nrcam	Smad
chr12	44601602	44602420	Nrcam	Smad
chr12	44617621	44617821	Nrcam	Foxc1
chr12	44637711	44637911	Nrcam	Smad
chr12	83767655	83769008	Papln	Smad
chr12	83770066	83771033	Papln	Smad
chr12	84805340	84806015	Ltbp2	Smad
chr12	84847421	84847933	Ltbp2	Nfatc1,Smad
chr12	84858674	84859103	Ltbp2	Smad
chr12	84860573	84862610	Ltbp2	Smad
chr12	84864890	84866199	Ltbp2	Smad
chr12	84876164	84877309	Ltbp2	Nfatc1,Smad
chr12	100925549	100926660	Ccdc88c	Smad
chr12	100952186	100953010	Ccdc88c	Foxc1
chr12	100994254	100995100	Ccdc88c	Smad
chr12	100999477	100999757	Ccdc88c	Smad
chr12	101026361	101028278	Ccdc88c	Foxc1,Smad
chr12	101028635	101029884	Ccdc88c	Smad
chr12	108370432	108371274	Eml1	Smad
chr12	108393769	108394076	Eml1	Smad

chr12	108497106	108497501	Eml1	Smad
chr12	110706381	110707253	Hsp90aa1	Smad
chr12	112182474	112182829	Kif26a	Smad
chr12	112183529	112184723	Kif26a	Smad
chr13	3742773	3743616	Calml3	Smad
chr13	3756595	3756989	Calml3	Smad
chr13	3769423	3771205	Calml3	Foxc1,Smad
chr13	3782277	3782970	Calml3	Smad
chr13	3803751	3805140	Calml3	Foxc1
chr13	3888759	3889357	Net1	Foxc1,Smad
chr13	3891381	3892594	Net1	Foxc1,Smad
chr13	23736197	23736730	Hist1h1c	Nfatc1,Smad
chr13	23738241	23740536	Hist1h1c	Smad
chr13	24397146	24397637	Cmah	Smad
chr13	24401870	24402752	Cmah	Smad
chr13	24414686	24414915	Cmah	Smad
chr13	24429945	24430246	Cmah	Smad
chr13	24521604	24522615	Cmah	Smad
chr13	31750921	31751282	Foxc1	Smad
chr13	31876857	31877613	Gmds	Smad
chr13	32025433	32025992	Gmds	Foxc1
chr13	32118246	32118717	Gmds	Smad
chr13	32169296	32170180	Gmds	Smad
chr13	32242137	32242872	Gmds	Nfatc1
chr13	32298462	32299086	Gmds	Nfatc1,Smad
chr13	32312457	32313068	Gmds	Nfatc1,Smad
chr13	32337784	32339432	Gmds	Smad
chr13	32370372	32371373	Gmds	Smad
chr13	32478798	32479598	Gmds	Smad
chr13	34129415	34132042	Tubb2b	Smad
chr13	34189985	34190892	Slc22a23	Smad
chr13	34191887	34193108	Slc22a23	Smad
chr13	34211338	34211880	Slc22a23	Nfatc1
chr13	34214796	34215609	Slc22a23	Nfatc1,Smad
chr13	34281471	34281707	Slc22a23	Nfatc1
chr13	34281872	34282095	Slc22a23	Smad
chr13	34344437	34346182	Slc22a23	Smad
chr13	34370012	34370891	Slc22a23	Smad
chr13	34371862	34372582	Slc22a23	Smad
chr13	34392505	34392998	Slc22a23	Smad
chr13	34397886	34398511	Slc22a23	Foxc1,Smad
chr13	34446354	34446891	Slc22a23	Smad
chr13	38344685	38345351	Bmp6	Foxc1

chr13	38349634	38350513	Bmp6	Smad
chr13	38410492	38410936	Bmp6	Smad
chr13	49549674	49550009	Aspn	Smad
chr13	73254478	73255287	Irx4	Smad
chr13	73257821	73258918	Irx4	Nfatc1,Smad
chr13	73273258	73273592	Irx4	Smad
chr13	79885151	79885477	Arrdc3	Smad
chr13	79908703	79908903	Arrdc3	Smad
chr13	80256079	80256834	Arrdc3	Smad
chr13	80267528	80268001	Arrdc3	Smad
chr13	80289204	80289433	Arrdc3	Smad
chr13	80387653	80388359	Arrdc3	Smad
chr13	80388481	80389537	Arrdc3	Smad
chr13	80393678	80394106	Arrdc3	Smad
chr13	80394161	80394363	Arrdc3	Nfatc1
chr13	80519913	80520138	Arrdc3	Smad
chr13	80525478	80526191	Arrdc3	Smad
chr13	80542998	80543198	Arrdc3	Smad
chr13	80557109	80557515	Arrdc3	Smad
chr13	80862128	80862640	Arrdc3	Nfatc1
chr13	80881651	80886662	Arrdc3	Smad
chr13	112556568	112557439	Ii31ra	Smad
chr13	112578105	112578420	Ii31ra	Smad
chr13	112584629	112585461	Ii31ra	Foxc1,Nfatc1
chr13	114490570	114491406	Fst	Nfatc1
chr13	114500947	114501465	Fst	Nfatc1,Smad
chr13	114523034	114523234	Fst	Foxc1
chr14	23305544	23306460	Kcnma1	Smad
chr14	23307780	23309555	Kcnma1	Smad
chr14	23389791	23389991	Kcnma1	Nfatc1,Smad
chr14	23494206	23495026	Kcnma1	Nfatc1,Smad
chr14	23624134	23625205	Kcnma1	Nfatc1,Smad
chr14	23626674	23628502	Kcnma1	Smad
chr14	23633131	23633332	Kcnma1	Foxc1
chr14	23707399	23707772	Kcnma1	Nfatc1
chr14	23903182	23903417	Kcnma1	Smad
chr14	23911144	23911630	Kcnma1	Foxc1
chr14	25700138	25701702	Ppif	Smad
chr14	32313328	32313716	Ogdhl	Nfatc1,Smad
chr14	32321876	32322136	Ogdhl	Smad
chr14	34085217	34090836	Anxa8	Smad
chr14	34355035	34356024	Fam25c	Smad
chr14	40995297	40997233	Fam213a	Smad

chr14	40999730	41000562	Fam213a	Nfatc1
chr14	41004135	41006469	Fam213a	Smad
chr14	41030679	41031851	Fam213a	Nfatc1,Smad
chr14	48038251	48040103	Peli2	Nfatc1,Smad
chr14	48092568	48092859	Peli2	Smad
chr14	48098969	48099836	Peli2	Smad
chr14	48118992	48120411	Peli2	Smad
chr14	48137550	48137750	Peli2	Smad
chr14	48155128	48155581	Peli2	Smad
chr14	48172784	48175077	Peli2	Smad
chr14	48203710	48204088	Peli2	Foxc1
chr14	48212764	48213968	Peli2	Smad
chr14	48224796	48225379	Peli2	Smad
chr14	48241250	48241856	Peli2	Smad
chr14	48318336	48318987	Peli2	Nfatc1
chr14	55210819	55211027	Dhrs2	Smad
chr14	55318154	55318402	Dhrs2	Nfatc1
chr14	57083932	57084433	Gjb2	Smad
chr14	57105848	57106424	Gjb2	Smad
chr14	57158544	57158914	Gjb6	Smad
chr14	57190438	57190738	Gjb6	Smad
chr14	70966073	70966432	Gfra2	Smad
chr14	70988703	70989012	Gfra2	Smad
chr14	77163763	77165510	Enox1	Smad
chr14	77174130	77174683	Enox1	Smad
chr14	77203961	77204607	Enox1	Smad
chr14	77209553	77211639	Enox1	Nfatc1
chr14	77218065	77218935	Enox1	Smad
chr14	77225171	77225782	Enox1	Smad
chr14	77353238	77353890	Enox1	Smad
chr14	77406183	77407149	Enox1	Smad
chr14	77608397	77608871	Enox1	Foxc1
chr14	97544255	97544850	Dach1	Smad
chr14	97731263	97731523	Dach1	Smad
chr14	97792641	97792970	Dach1	Smad
chr14	97982776	97983299	Dach1	Smad
chr14	98160672	98161796	Dach1	Nfatc1
chr14	98170552	98171287	Dach1	Nfatc1
chr14	98202015	98202495	Dach1	Smad
chr14	98380340	98380774	Dach1	Nfatc1
chr14	98442096	98443133	Dach1	Nfatc1,Smad
chr14	98451857	98452606	Dach1	Smad
chr14	98471718	98472836	Dach1	Smad

chr14	98482683	98483092	Dach1	Smad
chr14	98483181	98483941	Dach1	Smad
chr14	98505330	98505964	Dach1	Nfatc1,Smad
chr14	98589729	98590299	Dach1	Smad
chr14	101738375	101738954	Lmo7	Smad
chr14	101787540	101789398	Lmo7	Smad
chr14	101797749	101799189	Lmo7	Smad
chr14	101804947	101805750	Lmo7	Foxc1
chr14	101822188	101822999	Lmo7	Smad
chr14	101831910	101832767	Lmo7	Nfatc1,Smad
chr14	101839557	101840998	Lmo7	Smad
chr14	101842244	101843104	Lmo7	Nfatc1,Smad
chr14	101908144	101909162	Lmo7	Nfatc1,Smad
chr14	101917666	101918943	Lmo7	Smad
chr14	101930451	101931362	Lmo7	Smad
chr14	101936909	101937418	Lmo7	Nfatc1
chr14	101993734	101994221	Lmo7	Smad
chr14	102013300	102013642	Lmo7	Smad
chr14	102014566	102015598	Lmo7	Smad
chr14	102093049	102093493	Lmo7	Nfatc1,Smad
chr14	102117920	102118936	Lmo7	Smad
chr14	102135098	102136020	Lmo7	Smad
chr14	102142983	102143242	Lmo7	Nfatc1
chr14	102149129	102149475	Lmo7	Smad
chr14	103451904	103452208	Scel	Smad
chr14	103468980	103470620	Scel	Smad
chr14	103499691	103499891	Scel	Smad
chr14	103499993	103501335	Scel	Smad
chr14	103502209	103503138	Scel	Smad
chr14	103503273	103504051	Scel	Smad
chr14	103526631	103527453	Scel	Smad
chr14	103529317	103530532	Scel	Foxc1
chr14	103531687	103534079	Scel	Smad
chr14	103545862	103546611	Scel	Foxc1
chr14	103567994	103569293	Scel	Nfatc1
chr14	116736161	116736361	Gpc6	Smad
chr14	116765774	116766689	Gpc6	Smad
chr14	116914067	116914600	Gpc6	Nfatc1
chr14	116916676	116917853	Gpc6	Smad
chr14	116928006	116929599	Gpc6	Smad
chr14	116978297	116979760	Gpc6	Foxc1
chr14	116980875	116982199	Gpc6	Smad
chr14	116987311	116987511	Gpc6	Smad

chr14	117027846	117028663	Gpc6	Nfatc1
chr14	117097698	117098544	Gpc6	Foxc1
chr14	117243544	117244069	Gpc6	Smad
chr14	117396167	117396599	Gpc6	Foxc1,Smad
chr14	117413408	117414224	Gpc6	Nfatc1,Smad
chr14	117706450	117706995	Gpc6	Smad
chr14	117710330	117710875	Gpc6	Smad
chr14	117781673	117782342	Gpc6	Nfatc1,Smad
chr14	117783774	117784246	Gpc6	Smad
chr14	117831739	117832158	Gpc6	Smad
chr14	117929879	117931475	Gpc6	Smad
chr14	117951492	117952435	Gpc6	Smad
chr15	3204082	3204356	Sepp1	Smad
chr15	3214524	3215593	Sepp1	Nfatc1,Smad
chr15	3227234	3229352	Sepp1	Smad
chr15	3232837	3233820	Sepp1	Smad
chr15	10057835	10058608	Prlr	Nfatc1,Smad
chr15	10087926	10088454	Prlr	Foxc1
chr15	10088490	10089573	Prlr	Nfatc1,Smad
chr15	10116667	10117447	Prlr	Nfatc1
chr15	10125881	10126488	Prlr	Smad
chr15	10177332	10179347	Prlr	Smad
chr15	10189194	10189567	Prlr	Smad
chr15	10213608	10213954	Prlr	Nfatc1,Smad
chr15	10238258	10238977	Prlr	Smad
chr15	10295325	10296051	Prlr	Foxc1,Smad
chr15	10313719	10314034	Prlr	Nfatc1,Smad
chr15	27189786	27190139	Ank	Nfatc1,Smad
chr15	27325793	27327012	Ank	Smad
chr15	27381413	27381986	Ank	Nfatc1,Smad
chr15	27405330	27406610	Ank	Smad
chr15	27418313	27419251	Ank	Nfatc1
chr15	27488177	27488620	Ank	Smad
chr15	27506282	27507377	Ank	Smad
chr15	27525309	27526058	Ank	Smad
chr15	27536745	27538068	Ank	Smad
chr15	27563566	27566698	Ank	Nfatc1
chr15	27569127	27570936	Ank	Foxc1,Smad
chr15	31150769	31151093	Dap	Smad
chr15	31183168	31183368	Dap	Smad
chr15	31197218	31197418	Dap	Smad
chr15	31245997	31246377	Dap	Smad
chr15	31255817	31256141	Dap	Smad

chr15	31278124	31278444	Dap	Smad
chr15	37432184	37433038	Ncald	Nfatc1,Smad
chr15	37515846	37516120	Ncald	Nfatc1,Smad
chr15	37607577	37607833	Ncald	Smad
chr15	54177210	54177433	Tnfrsf11b	Smad
chr15	54208354	54208627	Tnfrsf11b	Nfatc1
chr15	54278108	54279495	Tnfrsf11b	Smad
chr15	54293846	54295173	Tnfrsf11b	Nfatc1
chr15	55396273	55396570	Col14a1	Smad
chr15	55464522	55466117	Col14a1	Smad
chr15	55493180	55493702	Col14a1	Smad
chr15	55513504	55513861	Col14a1	Nfatc1
chr15	62043492	62044238	Pvt1	Nfatc1,Smad
chr15	62092493	62093148	Pvt1	Smad
chr15	62106563	62106774	Pvt1	Foxc1
chr15	62165833	62166129	Pvt1	Nfatc1
chr15	62421139	62422007	Pvt1	Smad
chr15	62465419	62465844	Pvt1	Smad
chr15	62466668	62467301	Pvt1	Smad
chr15	62768129	62768533	Pvt1	Nfatc1
chr15	78815217	78816149	Card10	Nfatc1
chr15	78826486	78827006	Cdc42ep1	Foxc1,Smad
chr15	78844252	78846402	Cdc42ep1	Smad
chr15	85523355	85523889	Wnt7b	Smad
chr15	85581595	85582835	Wnt7b	Smad
chr15	87179131	87179843	Fam19a5	Smad
chr15	87623825	87624526	Fam19a5	Smad
chr15	92386123	92386757	Pdzrn4	Smad
chr15	92596696	92597818	Pdzrn4	Smad
chr15	92639857	92641111	Pdzrn4	Smad
chr15	92867310	92867647	Pdzrn4	Smad
chr15	93513890	93514353	Prickle1	Nfatc1,Smad
chr15	93551959	93552587	Prickle1	Smad
chr15	93575810	93576307	Prickle1	Smad
chr15	93681707	93681959	Prickle1	Smad
chr15	93705759	93706037	Prickle1	Smad
chr15	93708107	93709318	Prickle1	Foxc1
chr15	93710380	93711070	Prickle1	Smad
chr15	93733812	93734639	Prickle1	Smad
chr15	97774887	97775925	Rapgef3	Smad
chr15	99651535	99651860	Racgap1	Smad
chr15	101336318	101336960	Krt80	Nfatc1,Smad
chr15	101368387	101370828	Krt80	Smad

chr15	102098995	102101308	Tenc1	Smad
chr15	102108510	102108710	Tenc1	Smad
chr15	102108821	102109393	Tenc1	Smad
chr15	102839564	102840376	Hoxc13	Smad
chr15	102920423	102922329	Hoxc13	Smad
chr16	11049994	11050821	Snn	Smad
chr16	11050975	11051678	Snn	Smad
chr16	36461300	36461639	Stfa3	Smad
chr16	36463407	36464863	Stfa3	Nfatc1
chr16	36469004	36469896	Stfa3	Foxc1
chr16	45335979	45336309	Cd200	Smad
chr16	45346262	45347468	Cd200	Smad
chr16	45352914	45353189	Cd200	Smad
chr16	45360491	45361139	Cd200	Nfatc1,Smad
chr16	45408414	45409377	Cd200	Smad
chr16	56807771	56808251	Tmem45a	Smad
chr16	56808680	56809319	Tmem45a	Nfatc1
chr16	56842954	56843542	Tmem45a	Smad
chr16	56869753	56872240	Tmem45a	Smad
chr16	56872681	56874276	Tmem45a	Nfatc1
chr16	56879643	56881921	Tmem45a	Smad
chr16	56884442	56885361	Tmem45a	Nfatc1
chr16	56891637	56891944	Tmem45a	Nfatc1
chr16	56897441	56898056	Tmem45a	Smad
chr16	56919912	56920678	Tmem45a	Smad
chr16	56942107	56942917	Tmem45a	Smad
chr16	73483428	73483985	Robo2	Smad
chr16	73542389	73544441	Robo2	Nfatc1,Smad
chr16	73583735	73584001	Robo2	Nfatc1
chr16	73762201	73764124	Robo2	Foxc1,Nfatc1
chr16	73764654	73764854	Robo2	Smad
chr16	73766129	73766340	Robo2	Nfatc1
chr16	73775351	73775713	Robo2	Smad
chr16	73786722	73787240	Robo2	Smad
chr16	73855561	73856474	Robo2	Nfatc1
chr16	73861337	73861653	Robo2	Smad
chr16	73924150	73924605	Robo2	Smad
chr16	74130684	74130953	Robo2	Nfatc1,Smad
chr16	74235174	74235403	Robo2	Nfatc1,Smad
chr16	74307105	74308031	Robo2	Smad
chr16	74388831	74389744	Robo2	Smad
chr16	74400667	74401571	Robo2	Foxc1
chr16	74411066	74413484	Robo2	Smad

chr16	74426741	74427444	Robo2	Smad
chr16	74553841	74554041	Robo2	Nfatc1
chr16	74628352	74628926	Robo2	Smad
chr16	74664457	74664667	Robo2	Foxc1
chr16	74801265	74801800	Robo2	Smad
chr16	74821401	74821788	Robo2	Foxc1
chr16	87915186	87915547	Grik1	Smad
chr16	88005171	88005371	Grik1	Smad
chr16	88064083	88064389	Grik1	Nfatc1
chr16	88215373	88215765	Grik1	Smad
chr16	88243835	88246313	Grik1	Nfatc1
chr16	88252721	88253030	Grik1	Smad
chr16	88358844	88360292	Grik1	Smad
chr16	92427979	92429082	Rcan1	Smad
chr16	96467546	96467835	Pcp4	Smad
chr17	34953621	34961762	Hspa1b	Smad
chr17	34967616	34972947	Hspa1a	Smad
chr17	36942052	36944647	Rnf39	Smad
chr17	43557633	43559223	Pla2g7	Foxc1
chr17	43567507	43567967	Pla2g7	Smad
chr17	43569729	43570439	Pla2g7	Smad
chr17	47436065	47437354	Al661453	Smad
chr17	56747879	56748338	Nrtn	Smad
chr17	73822495	73822695	Ehd3	Smad
chr17	73838715	73839511	Ehd3	Nfatc1,Smad
chr17	73989843	73990113	Srd5a2	Smad
chr17	74043837	74044037	Srd5a2	Smad
chr17	75006370	75006746	Ltbp1	Smad
chr17	75046772	75047215	Ltbp1	Smad
chr17	75094615	75094915	Ltbp1	Smad
chr17	75105791	75105991	Ltbp1	Nfatc1
chr17	75163353	75163607	Ltbp1	Nfatc1
chr17	75378842	75379084	Ltbp1	Smad
chr17	78554595	78555939	Vit	Smad
chr17	78576849	78577049	Vit	Smad
chr17	78603847	78604326	Vit	Foxc1
chr18	33350435	33350653	Nrep	Smad
chr18	33377893	33378724	Nrep	Smad
chr18	33463129	33463341	Nrep	Nfatc1
chr18	33463613	33464437	Nrep	Smad
chr18	36335391	36335904	Cystm1	Smad
chr18	38846487	38847878	Fgf1	Nfatc1
chr18	38882183	38882802	Fgf1	Nfatc1,Smad

chr18	38917819	38920014	Fgf1	Smad
chr18	42033762	42034225	Sh3rf2	Smad
chr18	42042674	42042874	Sh3rf2	Smad
chr18	42042936	42043551	Sh3rf2	Smad
chr18	42056544	42057938	Sh3rf2	Smad
chr18	42116701	42117139	Sh3rf2	Smad
chr18	42123862	42124724	Sh3rf2	Smad
chr18	42128283	42128773	Sh3rf2	Nfatc1,Smad
chr18	61744342	61745083	Afap111	Smad
chr18	61745972	61747212	Afap111	Smad
chr18	62131365	62131775	Adrb2	Smad
chr18	62146179	62147050	Adrb2	Nfatc1
chr18	66453261	66453491	Pmaip1	Smad
chr18	66457953	66459307	Pmaip1	Smad
chr18	66468909	66469131	Pmaip1	Smad
chr18	80653878	80657414	Nfatc1	Nfatc1
chr18	80682708	80683263	Nfatc1	Smad
chr18	80718092	80718920	Nfatc1	Foxc1,Smad
chr19	4212273	4212547	Clcf1	Nfatc1,Smad
chr19	5544008	5544490	Ovol1	Nfatc1,Smad
chr19	5560026	5561573	Ovol1	Nfatc1
chr19	5877767	5878624	Slc25a45	Smad
chr19	10039548	10039882	Fads3	Smad
chr19	10043570	10044015	Fads3	Smad
chr19	14469507	14470213	Tle4	Nfatc1,Smad
chr19	14511651	14512979	Tle4	Smad
chr19	14588252	14588887	Tle4	Smad
chr19	14622634	14623945	Tle4	Smad
chr19	14657119	14657515	Tle4	Smad
chr19	14692031	14692630	Tle4	Nfatc1
chr19	14777131	14778036	Tle4	Nfatc1
chr19	14791821	14792207	Tle4	Smad
chr19	14884224	14884424	Tle4	Smad
chr19	15003648	15004482	Tle4	Smad
chr19	15035601	15036007	Tle4	Smad
chr19	22249020	22249432	Trpm3	Nfatc1,Smad
chr19	22576475	22576769	Trpm3	Smad
chr19	22915775	22917452	Trpm3	Foxc1,Smad
chr19	22940516	22940998	Trpm3	Smad
chr19	22977076	22977726	Trpm3	Nfatc1
chr19	23000034	23000778	Trpm3	Foxc1,Smad
chr19	23006627	23007343	Trpm3	Smad
chr19	23054434	23054925	Trpm3	Smad

chr19	25341612	25342691	Kank1	Smad
chr19	25383351	25383956	Kank1	Smad
chr19	25397886	25398090	Kank1	Smad
chr19	25406338	25408006	Kank1	Smad
chr19	26338968	26340770	Smarca2	Smad
chr19	26389464	26390339	Smarca2	Smad
chr19	26472668	26472870	Smarca2	Smad
chr19	26501061	26501904	Smarca2	Smad
chr19	26502750	26503271	Smarca2	Smad
chr19	26604369	26605644	Smarca2	Smad
chr19	26606243	26608366	Smarca2	Foxc1,Smad
chr19	26733252	26734199	Smarca2	Smad
chr19	29041001	29041930	Ak3	Smad
chr19	29047478	29048775	Ak3	Smad
chr19	34243648	34244519	Acta2	Smad
chr19	34254331	34255220	Acta2	Smad
chr19	34255545	34255907	Acta2	Smad
chr19	34267894	34268194	Acta2	Smad
chr19	34270237	34270437	Acta2	Nfatc1,Smad
chr19	42146833	42149051	Marveld1	Smad
chr19	44280718	44281529	Scd2	Smad
chr19	44283874	44284838	Scd2	Smad
chr19	44295331	44296104	Scd2	Smad
chr19	44303611	44303959	Scd2	Smad
chr19	44314325	44315425	Scd2	Nfatc1
chr19	57152434	57152699	Ablim1	Smad
chr19	57154961	57156632	Ablim1	Nfatc1
chr19	57238235	57239822	Ablim1	Smad
chr19	58189068	58189269	Gfra1	Foxc1,Smad
chr19	58214975	58215708	Gfra1	Foxc1
chr19	58226393	58227388	Gfra1	Smad
chr19	58271959	58272191	Gfra1	Smad
chr19	58453702	58454012	Gfra1	Nfatc1,Smad
chr19	58470963	58471338	Gfra1	Smad
chrX	100085271	100085892	Eda	Nfatc1
chrX	103793748	103793956	Slc16a2	Nfatc1,Smad
chrX	157465357	157465639	Sms	Smad
chrX	157490865	157492679	Sms	Smad
chrX	159731315	159731922	Sh3kbp1	Nfatc1
chrX	159839191	159840704	Sh3kbp1	Smad
chrX	159849090	159849614	Sh3kbp1	Smad
chrX	159896276	159897097	Sh3kbp1	Smad
chrX	159963190	159963445	Sh3kbp1	Foxc1

chrX	166564396	166565916	Egfl6	Smad
chrX	166584945	166585891	Egfl6	Smad

Table S4 Primers

Gene	Forwad primer	Reverse primer
Bmp6	CTGCGCACCAACCAAAGTAA	TTGGGGGAGGCGAACATTAGGTA
Ccna2	CTGTCTCTTTACCCGGAGCA	CGTTCACTGGCTTGTCTTCT
Cd34	AAGACCACACCAGCCATCTC	GGGGAAGTCTGTGGTTGTGA
Cdc20	CAGGAGGAGGAACCAGTGAC	GTGGGGGAGACCAGAGGATG
Fgf18	ATCAAGGGCAAGGAGACAGA	CAGGGCCGTGTAGTTGTTTT
Foxc1-a	CACTCGGTGCGGGAAATGT	GTGCGGTACAGAGACTGACTG
Foxc1-b	CACTCGGTGCGGGAAATGT	GGTACAGAGACTGACTGGCA
Foxm1	GAGCACTTGAATCACAGCA	AGCAGAGGCTTCATCTTTTCG
Hprt	TCAGTCAACGGGGGACATAAA	GGGGCTGTACTGCTTAACCAG
Kif23	AGCGTGTTTCAGCATTAACTTG	GCTCCCAGCAAGATCTACCA
Top2a	TGGGAGTGAAGAAGACAGCA	ACGAGCACTCAAGGCTGAAT
Nfatc1 ChIP motif	AAACCTCATGGTGCATTGTG	AAACGCAACCCAGGTGTAAG
Nfatc1 ChIP no motif	CTGCATCTACCCAGCACAGA	CCTTCCAGAAAGCCTTACCC
Hspb8 ChIP	CGATTCACCTTGGGGTTTTA	GGCCAAGGTTTCCTGTAACAA
Bmp6 ChIP motif	AACCTGGAGTCTCAGCTTCG	TGTTTATTTCTGTGTGGCTTGG
Nfatc1 ChIP no motif	GCAGCCCACAATGATCTCTT	AGAAGGCTAGCAGCACCTGA
Krt5 ChIP	CTGGAATATACCCAAGAAAAGAGG	AGGTCACCCACTTGACAATGA
Chr5IGR ChIP	CCCTCATCACAGACCCACTTCT	GTGGGAGTGGATGTATCTCTGACTT
Chr3IGR ChIP	TGTCTGGAATGTGGTGGTTTGA	GCCCACTGCTATAATTAGGAAGGA

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