Supplemental Information

Materials and reagents.

Antibodies for MLL4 (ABE167) and MLL3 (ABE1851) were purchased from Millipore; for KLF4 (AF3158) from R&D system; for SHP (ab186874) from Abcam; for F4/80 (NB600-404SS) from Novus Biologicals; for LAMIN A (sc-20680) and β -TUBULIN (sc-9104) from Santa Cruz Biotechnology, and for β -ACTIN (CST-4970) from Cell Signaling Technology. Small interfering RNAs (ON-TARGETplus SMART pool) for *Klf4* (M-066785-00-0005), *Ahr* (M-044066-01-0005) and *Lrh1* (M-047044-00-0005) were purchased from Dharmacon. The target vector for lentivirus construction for the precursor of miR-210 (MmiR3303-MR03) was purchased from GeneCopoeia and for anti-miR-210 (mm30343) from Applied Biological Materials Inc. The lentiviral packaging vectors, psPAX2 (12260) and pMD2.G (12259) were obtained from Addgene. The miR-210 miRNA precursor (mmu-miR-210-3p, AM17100) and anti-miR-210 (mmu-miR-210-3p, AM17000) were purchased from Thermo Fisher. Lentiviruses expressing shRNA for *Mll4* and expression vectors for KLF4 were purchased from VectorBuilder.

Supplemental Figures



Supplemental Figure S1. Increased hepatic BA levels in SHP-LKD mice compared to control mice. SHP was downregulated as described in the Fig.1 legend and hepatic BA levels were measured. Statistical significance was determined by the Student's t-test, SD (n=5), ** p < 0.01.



Supplemental Figure S2. MiR-210 inhibits MLL4 expression in human HEK293 cells.

HEK293 cells were transfected with the indicated luciferase plasmids and with the indicated amounts of pre-miR-210 (A), anti-miR-210 (B), or scrambled RNA (miR-Scr) as a control, and 48 h later, luciferase activity was determined and normalized to β -galactosidase activity. Statistical significance was determined by one-way ANOVA, SD (n=3), ** p < 0.01, NS, not significant.



Supplemental Figure S3. Activation of FXR in cholestatic mice can reverse the increase in miR-210 levels. To examine the effects of OCA on ANIT-induced cholestasis, mice were treated with vehicle or 10 mg/kg of OCA, i.p., and after five days, treated with 35 mg/kg ANIT by gavage daily for 2 days. Hepatic mRNA levels of indicated genes were determined by RT-qPCR analysis. Statistical significance was determined by one-way ANOVA, SD (n=5), ** p < 0.01.



Supplemental Figure S4. SHP occupancy is increased at the hepatic miR-210 promoter in response to FGF19 treatment. (A) SHP binding peak at the hepatic miR-210 promoter from published ChIP-seq data (11). (B) Mice were fasted overnight and treated with vehicle or FGF19 for 2 h and standard liver ChIP assays were done to confirm SHP occupancy at the *miR-210* gene promoter. Statistical significance was determined by two-way ANOVA, SD (n=5), ** p < 0.01.



Supplemental Figure S5. Potential transcription factor binding sites within the SHP binding peak region (312 bp) at the miR-210 promoter. Potential binding sites for transcription factors detected by ChIP-seq were determined using the JASPAR online program (2018; http://jaspar.genereg.net/). The JASPAR motif scores for the individual potential binding motifs for the indicated factors are plotted.



Supplemental Figure S6. Functional interaction between SHP and KLF4 is increased in the late fed-state. C57BL/6J mice were refed for 4 h after fasting overnight. (A) CoIP: SHP protein levels in anti-KLF4 or IgG immunoprecipitates and SHP and KLF4 protein levels in the input samples. (B) Re-ChIP: Chromatin was immunoprecipitated with KLF4 antibody and then, eluted and re-precipitated with SHP antibody. Enrichment of the *miR-210* promoter sequence determined by qPCR. Statistical significance was determined by two-way ANOVA, SD (n=5), ** p < 0.01.

Supplemental Table S1

Mouse RT-qPCR primers

	Forward (5'-3')	Reverse (5'-3')
Cxcl2	TCCAGAGCTTGAGTGTGACG	TTCAGGGTCAAGGCAAACTT
Tnfa	AGCCCCCAGTCTGTATCCTT	GGTCACTGTCCCAGCATCTT
Shp	CAAGAAGATTCTGCTGGAGG	GGATGTCAACATCTCCAATG
Klf4	CTGAACAGCAGGGACTGTCA	GTGTGGGTGGCTGTTCTTTT
Cyp7a1	AACGGGTTGATTCCATACCTGG	GTGGACATATTTCCCCATCAGTT
Cyp8b1	GAATCTAACCAGGCCATGCT	AGGAGCTGGCACCTAGACT
Fgf15	GTTTCACCGCTCCTTCTTTG	CATCCTCCACCATCCTGAAC
36b4	CGACATCACAGAGCAGGC	CACCGAGGCAACAGTTGG
Bsep	CAATGTTCAGTTCCTCCGTTCA	TTTGGTGTTGTCCCCSTSCTTG
Mrp2	TATCCCCGGGAAATCTGTTC	TAACCAACATTCTCCGCGC
Oatp1	GTCTTACGAGTGTGCTCCAGAT	GGAATACTGCCTCTGAAGTGGATT
Ntcp	TACCTCCTCCTGATGCCTTTC	TGCGTCTGCAGCTTGGATTTA
MII3	GCAACCTCTTACCGGTTGAA	GTTCTCTCGGGAACCCTTGT
MII4	GCACCGAGTGGAGAGACAAT	TAAATACCCGCGGTTCTGCTC
Ahr	TCCACAACTGGCTTTGTTTG	CCAGAATAAGCTGCCCTTTG
Lrh-1	TCATGCTGCCCAAAGTGGAGA	TGGTTTTGGACAGTTCGCTT
Cck	AAGTGACCGGGACTACATGG	CCCACTACGATGGGTATTCG
Mdr1	CTGGTGTGCTCATAGTTG	CCTAATCTTGTGTATCTGTCTT

Mouse ChIP primers

	Forward (5'-3')	Reverse (5'-3')
Shp	CAGTGAGAACCCTGGTCTT	CTGGCCAAACAACCTTGAC
Bsep	CGACCTTTCCTCTCATGTCA	CATTGAACAGAAATCAGGCTTTT

Human RT-qPCR primers

	Forward (5'-3')	Reverse (5'-3')
KLF4	ACCCTGGGTCTTGAGGAAGT	ACGATCGTCTTCCCCTCTTT
36B4	TGCTGAACATGCTCAAC	GTCGAACACCTGCTGGATGAC
SHP	CAGAGATCAGGTGGGCAGAG	TGTGGCTGAGTGAAGAGCTG
MLL4	GCACAATGCTGTCAGGAGAA	GTGCAGCAGAAGATGGTGAA