

Title: miR-34c attenuates epithelial-mesenchymal transition and kidney fibrosis with ureteral obstruction

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Supplementary Table S1. miRNAs up-regulated by UUU

systematic_name	fold change	systematic_name	fold change
mmu-miR-142-5p	75.0024	mmu-miR-877*	3.9418
mmu-miR-466a-3p	46.2183	mmu-miR-351	3.5247
mmu-miR-212	37.5707	mmu-miR-135a*	3.4993
mmu-miR-467a	28.9733	mmu-miR-28*	3.4721
mmu-miR-467e	26.6278	mmu-miR-714	3.3824
mmu-miR-205	24.4364	mmu-miR-425*	3.3360
mmu-miR-184	21.0981	mmu-miR-672	3.3236
mmu-miR-31*	20.7141	mmu-miR-466h	3.2492
mmu-miR-17	19.6553	mmu-miR-132	3.2323
mmu-miR-181a-1*	19.5967	mmu-miR-300*	3.1264
mmu-miR-127	18.3310	mmu-miR-546	3.0927
mmu-miR-744	16.6562	mmu-miR-770-3p	3.0723
mmu-miR-467c	15.5661	mmu-miR-183*	3.0623
mmu-miR-342-5p	15.1527	mmu-miR-872*	3.0125
mmu-miR-18a*	13.1978	mmu-miR-125a-3p	2.9349
mmu-miR-505	11.9566	mmu-miR-744*	2.9254
mmu-miR-379	11.7161	mmu-miR-99b*	2.9195
mmu-miR-141*	11.5208	mmu-miR-685_v14.0	2.9088
mmu-miR-214*	11.0526	mmu-miR-669h-5p	2.8887
mmu-miR-296-5p	10.5931	mmu-miR-701	2.8797
mmu-miR-15b*	10.3725	mmu-miR-31	2.8371
mmu-miR-677	9.3028	mmu-miR-26b*	2.8251
mmu-miR-20a*	8.8458	mmu-miR-224	2.7367
mmu-miR-450a-3p	8.8068	mmu-miR-883b-5p	2.7131
mcmv-miR-m88-1	8.7979	mmu-miR-342-3p	2.6943
mmu-miR-669l	8.2181	mmu-miR-223	2.6738
mmu-miR-297a*	7.6802	mmu-miR-878-3p	2.6724
mcmv-miR-M23-1-5p	6.5483	mmu-miR-214	2.6599
mmu-miR-542-3p	6.3537	mmu-miR-678	2.5721
mmu-miR-129-3p	6.0070	mmu-miR-155	2.5038
mmu-miR-1901	5.9849	mmu-miR-1188	2.4823
mmu-miR-341	5.4161	mmu-miR-375	2.4663
mmu-miR-669b	5.2025	mmu-miR-142-3p	2.4361
mmu-miR-468	5.1922	mmu-miR-190b	2.4192

mcmv-miR-M23-1-3p	5.1037	mghv-miR-M1-2	2.4110
mmu-miR-501-3p	4.9230	mmu-miR-467b	2.3321
mmu-miR-34c*	4.8129	mmu-miR-188-3p	2.2439
mmu-miR-300	4.7881	mmu-miR-421	2.2120
mmu-miR-877	4.6055	mmu-miR-680	2.1965
mmu-miR-146b	4.5654	mmu-miR-199a-5p	2.1906
mmu-miR-802	4.3538	mmu-miR-1186	2.1368
mmu-miR-21*	4.3344	mmu-miR-34c	2.1339
mmu-miR-883a-5p	4.2978	mmu-miR-669a	2.1301
mmu-miR-290-5p	4.1858	mmu-miR-210	2.1154
mmu-miR-130b	4.1544	mmu-miR-292-5p	2.0997
mmu-miR-106b*	4.0159	mmu-miR-186*	2.0665
mmu-miR-329	3.9942	mmu-miR-489	2.0274
mmu-miR-449a	3.9734	mmu-miR-199a-3p	2.0233

Supplementary Table S2. miRNAs up-regulated by TGF- β in MCT

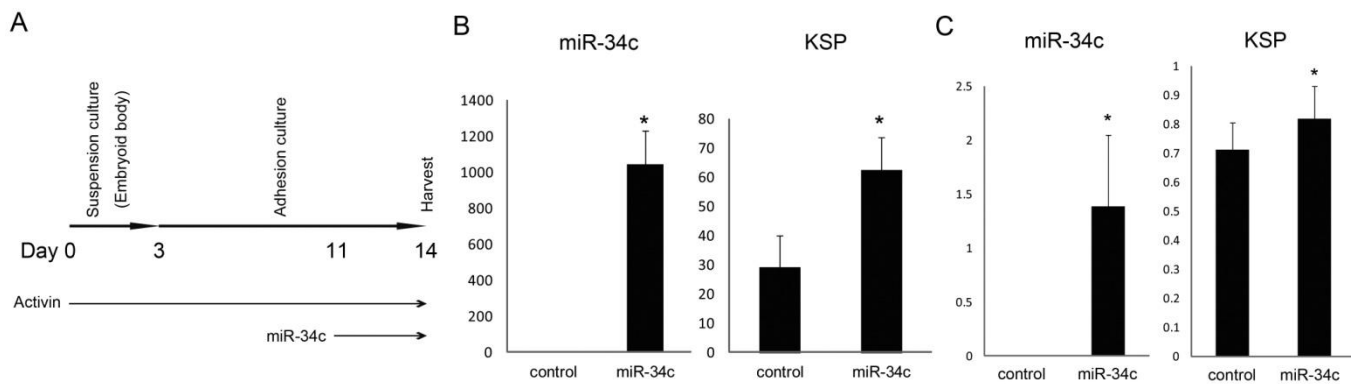
systematic_name	fold change	systematic_name	fold change
mmu-miR-494	273.1822	mmu-miR-467f	6.4509
mmu-miR-714	99.6733	mmu-miR-671-5p	6.0041
mmu-miR-188-5p	79.7492	mmu-miR-483	5.3729
mmu-miR-574-5p	78.9182	mmu-miR-450a-3p	5.3078
mmu-miR-1906	49.7573	mmu-miR-185	5.1866
mmu-miR-689	45.1235	mmu-miR-425	4.9647
mmu-miR-135a*	44.8211	mmu-miR-1892	4.9405
mmu-miR-449a	34.8543	mmu-miR-181a	4.8859
mmu-miR-140*	30.2912	mmu-miR-365	4.8526
mmu-miR-193	29.0345	mmu-miR-128	4.8222
mmu-miR-298	28.6193	mcmv-miR-M87-1	4.7572
mmu-miR-721	25.1908	mmu-miR-705	4.1271
mmu-miR-872	24.3895	mmu-miR-340-3p	4.0287
mmu-miR-29b*	14.9323	mmu-miR-30c-1*	3.9218
mmu-miR-1224	14.2491	mmu-miR-681	3.7968
mmu-miR-1895	11.9413	mmu-miR-125a-3p	3.7460
mmu-miR-1187	10.4649	mmu-let-7b*	3.7248
mcmv-miR-M23-1-5p	10.3352	mmu-miR-30c	3.4835

mmu-miR-712	9.3542	mmu-miR-181b	2.5848
mmu-miR-715	7.8953	mmu-miR-34c	2.5734
mmu-miR-1897-5p	7.7120	mmu-miR-125b-5p	2.5555
mmu-miR-370	7.6363	mmu-miR-34a	2.3150
mmu-miR-1894-3p	7.3622	mmu-miR-34b-5p	2.2151
mmu-miR-680	6.8924	mmu-miR-30d	2.1574

Supplementary Table S3. miRNAs down-regulated by Activin in mouse ES cells

systematic_name	fold change	systematic_name	fold change
mmu-miR-467b	-27.73095	mmu-miR-181b	-5.343746
mmu-miR-19a	-27.163473	mmu-miR-34b-5p	-5.2363887
mmu-miR-1904	-26.608042	mmu-miR-181c	-5.16735
mmu-miR-685_v14.0	-25.322464	mmu-miR-30c	-5.1665454
mmu-miR-223	-25.279774	mmu-miR-26b	-5.130635
mmu-miR-291a-5p	-18.82812	mmu-miR-669a	-5.101503
mmu-miR-290-5p	-18.221317	mmu-miR-483*	-5.075339
mmu-miR-705	-18.081573	mmu-miR-181d	-5.057848
mmu-miR-671-5p	-16.059662	mmu-miR-96	-5.0199513
mghev-miR-M1-2	-8.658391	mmu-miR-141	-4.9155693
mmu-miR-125a-3p	-8.328011	mmu-miR-218	-4.851454
mmu-miR-10a	-8.32481	mmu-miR-34c	-4.6313596
mmu-miR-100	-8.0890045	mmu-miR-494	-4.620617
mmu-miR-1897-5p	-7.596741	mmu-miR-367	-4.581095
mmu-miR-202-3p	-7.5897546	mmu-miR-423-5p	-4.521743
mmu-miR-1895	-7.5245156	mmu-miR-434-3p	-4.2810717
mmu-miR-10b	-6.700873	mmu-miR-677	-4.2440267
mmu-miR-301a	-6.578678	mmu-miR-702	-4.1883388
mmu-miR-139-3p	-6.2485433	mmu-miR-483	-4.1605797
mmu-miR-148a	-6.011732	mmu-miR-877	-4.013603
mmu-miR-135a*	-5.9646735	mmu-miR-712	-3.9998624
mmu-miR-124	-5.778936	mmu-miR-29a	-3.8379686
mmu-miR-429	-5.696437	mmu-miR-290-3p	-3.331493
mmu-miR-30d	-5.379591	mmu-miR-721	-3.1268754

Supplementary Figure S1. miR-34c up-regulated the expression of KSP in mouse ES cells and mouse embryonic kidneys. (A) The protocol of differentiation of mouse ES cells with Activin and miR-34c over-expression. miR-34c was transfected at day 11 of the differentiation. (B) Real-time PCR of miR-34c and KSP at day 14 of the differentiation. (C) Real-time PCR of miR-34c and KSP. E14.5 mouse embryonic kidneys were transfected with precursors of miR-34c, and samples were harvested after 3 days organ culture. Asterisks indicate statistically significant against control.



Supplementary Figure S2. The original whole blot pictures of figure 1A, B.

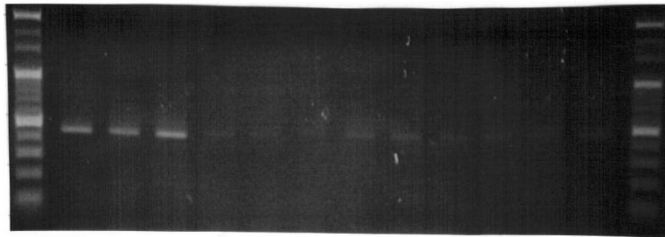
UUO: unilateral ureter obstruction. n=3. Control: contralateral kidneys in UUO mice. n=3.

Sham Left: left kidneys in sham-operated mice. Sham Right: right kidneys in sham-operated mice.

PCR

Western blot

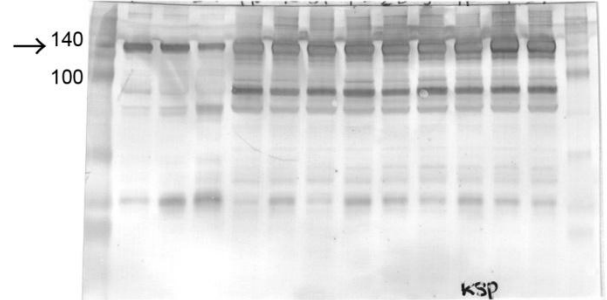
Vimentin



UUO Control Sham Left Sham Right

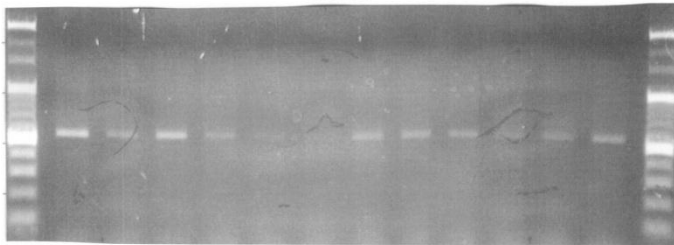
(kDa)

KSP



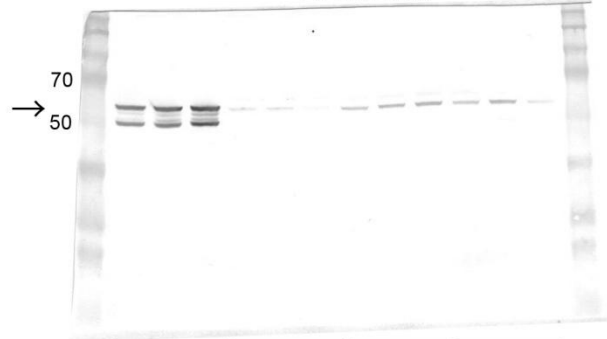
UUO Control Sham Left Sham Right

Snail1



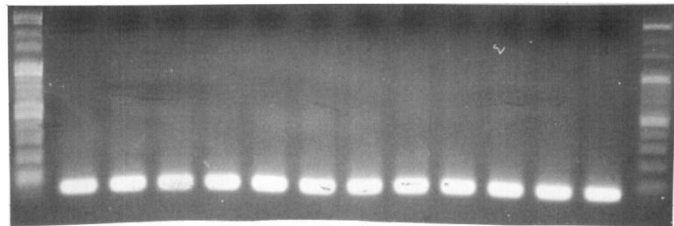
UUO Control Sham Left Sham Right

Vimentin



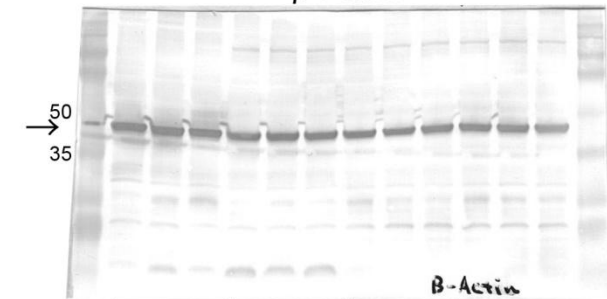
UUO Control Sham Left Sham Right

GAPDH



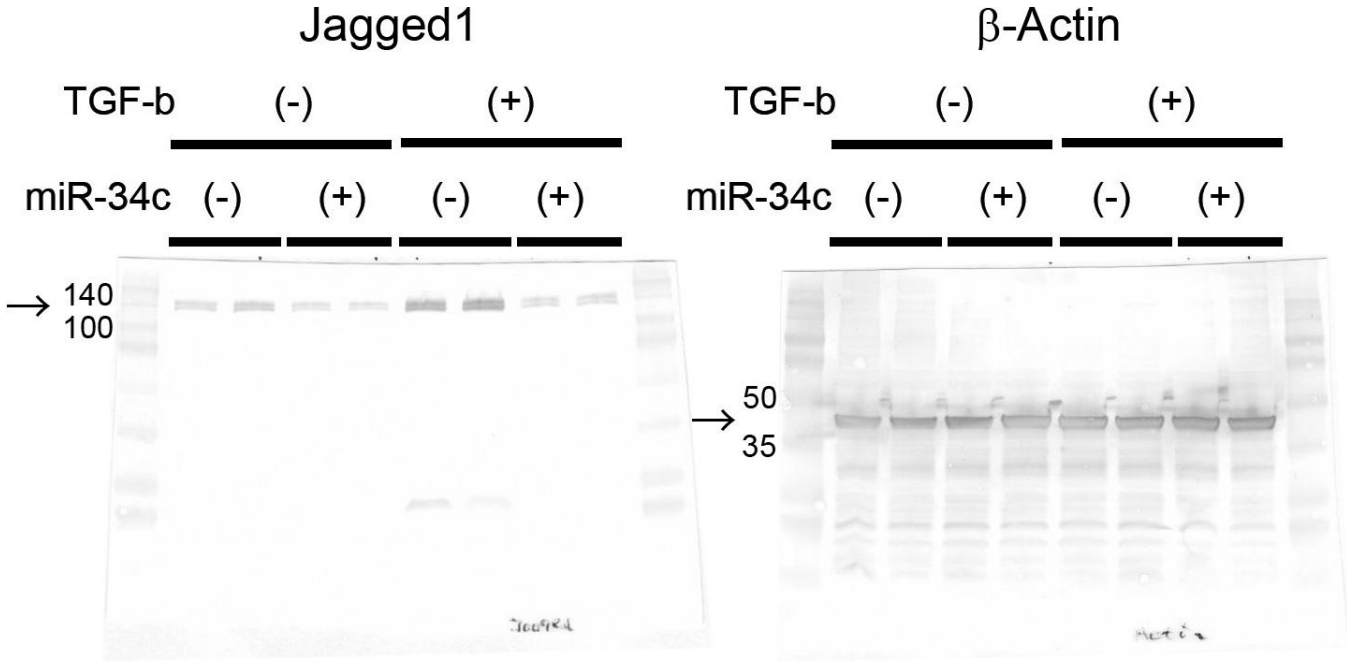
UUO Control Sham Left Sham Right

β -Actin



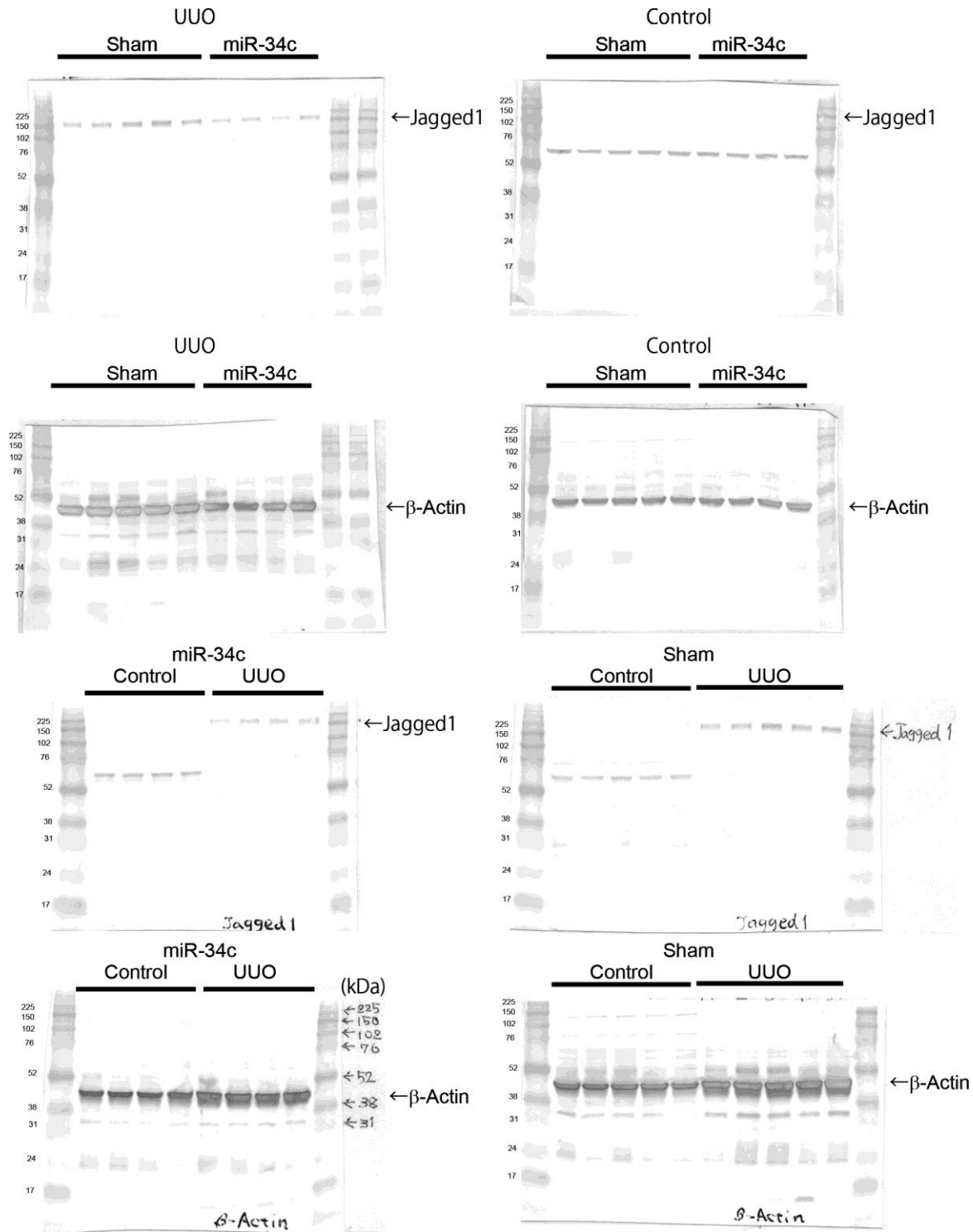
UUO Control Sham Left Sham Right

Supplementary Figure S3. The original blot pictures of figure 2F.



Supplementary Figure S4. The original blot pictures of figure 3B.

To compare miR-34c samples with sham samples, those samples were electrophoresed into a same blotting membrane. Moreover, to confirm the difference of Jagged1 between UUO and control kidneys, those samples were electrophoresed into a same blotting membrane. The former results was used for quantification. Sham-injected mice, n=5. miR-34c injected mice, n=4.



Supplementary Figure S5. Representative pictures of Masson trichrome staining.

Quantification of fibrosis area was done by ImageJ using 10 randomly taken pictures at x100. Fibrosis area was calculated by "Color Threshold" using following parameters; Hue 130-180, Saturation 0-255, Brightness 0-255.

