	Antisense sequence	miR-155 loop	Sense sequence	
5'-	(1-21 mature mi/siRNA)	GTTTTGGCCACTGACTGAC	(Nucleotides 1-8 and 11-21)	-3'

## A2

ADAM17 coding sequence	matching miA17-94
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rat	(NM_020306)	CTTGAGAAGCTTGATTCTTTG				
mouse	(NM_009615)	CTTGAGAAGCTTGATTCTTTG				
human	(NM_003183)	CT <b>C</b> GAGAAGCTTGATTCTTTG				
ADAM17 coding sequence matching miA17-450						
rat	(NM_020306)	GCCACTTTGGAGGTTTGTTAA				
mouse	(NM_009615)	GCCACTTTGGAGGTTTGT <u>C</u> AA				
human	(NM_003183)	GCCACTTTGGAGATTTGT <u>T</u> AA				
ADAM17 coding sequence matching miA17-724						
rat	(NM_020306)	GGAGAAGAGAGCACTACTACA				
mouse	(NM_009615)	GGAGAAGAGAGCAC <u>C</u> ACTACA				
human	(NM_003183)	GG <u>G</u> GAAGAGAG <u>T</u> AC <u>A</u> ACTACA				

## A3

Closest rat homologies to miA17-94

ADAM17	(NM_020306)	CTTGAGAAGCTTGATTCTTTG		
S1pr3 3'UTR	(NM_001271143)	<u>G</u> T <u>A</u> GAGAAGCTTGATTC <u>G</u> T <u>GT</u>		
TL0ADA33YG04	(FQ221658)	CTTGAGAAGCTTGA <u>GCA</u> T <u>GCT</u>		
Gcfc2	(NM_001134554)	<u>A</u> TTGAGAAGCTTGA <u>GAG</u> T <u>GGA</u>		
ADAM15	(NM_020308)	<u>TGA</u> GAGAAGCT <u>ATACA</u> CT <u>GGA</u>		
Closest rat homologies to miA17-450				
ADAM17	(NM_020306)	GCCACTTTGGAGGTTTGTTAA		
TL0ACA47YG15	(FQ215635)	<u>C</u> C <u>AGG</u> T <u>A</u> TGGAGGTTTGTTA <u>C</u>		
Dnmt1 3'UTR	(NM_053354)	<u>AAGCTG</u> TTGGAGGTTTGTT <u>CT</u>		
ADAM4	(NM_020305)	<u>C</u> C <u>TG</u> C <u>AC</u> TGGAGGT <u>C</u> TG <u>AA</u> A <u>G</u>		
Closest rat homologies to miA17-724				
ADAM17	(NM_020306)	GGAGAAGAGAGCACTACTACA		
Usp29 3'UTR	(NM_001108465)	<u>TC</u> AGAAGAGAGCACTA <u>GATCT</u>		
ADAM4	(NM_020305)	G <u>CTTTTC</u> AGAGCACTA <u>A</u> T <u>GCT</u>		

**Supplemental Fig A. 1**. Adenovirally encoded synthetic Pre-miRNA oligo structure. The ninth and tenth nucleotides of the sense target sequence are removed to create an internal loop. **2**. ADAM17 sequence homologies to rat ADAM17 mi/siRNAs. **3**. Blast results of closest rat homologies to rat ADAM17 mi/siRNAs.

**A1** 



Supplementary Figure B: 1. Enhancement of adenovirus infection by FuGene 6. VSMC were infected with adenovirus encoding GFP and miA17-450 at 50 MOI for 3 days in the presence or absence of 3% FuGene 6. GFP expression was analyzed by fluorescent microscopy. Representative results are shown from triplicated experiments. 2. VSMC were infected with adenovirus expressing miCon or miA17-724 at 100 MOI for 3 days. Cell lysates were analyzed by immunoblotting with antibodies as indicted. Representative blots are shown from triplicated experiments. 3. Densitometry analysis of the data from triplicated experiments (mean ± SD, \* p <0.05 compared with miRNA control). 4. VSMC were infected with 50 MOI adenovirus for the indicated time. GFP expression was analyzed by fluorescent microscopy. Representative results are shown from triplicated experiments. 5. Rat VSMC were infected with adenovirus encoding miA17-450 or miCon (100 MOI) together with wild type mouse ADAM17 or control GFP (10 MOI) for 3 days and immunoblotting was performed with or without 100 nM Ang II stimulation. 6. VSMC infected with adenovirus encoding HA-tagged wild-type ADAM17 were stimulated with 100 nM Ang II for indicated time. Cell lysates were immunoprecipitated with anti-HA antibody and immunoblotted with antibodies as indicated. Representative blots are shown from triplicated experiments.