

**Table S1 Sequence of primers used in the study. Red fonts indicate mutated nucleotides**

<b>cloning</b>		<b>Forward 5'-3'</b>
miR-200ba429 Cluster	ATAAGAATGCGGCCGCGCCCCACTCCGACCTAGTCCT	
MMP-2 promoter	CCGCTCGAGCACACCCACCAGACAAGCCT	
MMP-9 promoter	CCGCTCGAGACATTTGCCCGAGGTCCTGAAG	
<b>Reporter</b>		<b>Forward 5'-3'</b>
fibronectin1 3'UTR	CCGCTCGAGCAAAATAAATCCGGGTGTGC	
miR-200b target	TCGAGTAGCGGCCGCTAGTTCATCATTACCAGGCAGTATTA	
miR-429 target	TCGAGTAGCGGCCGCTAGTACGGTTTTACCAGACAGTATTA	
<b>Mutagenesis</b>		<b>Forward 5'-3'</b>
fibronectin1_MUT1	GTCTATTTTAAATGAAGTGATTCTAAGATTTGG	
fibronectin1_MUT2	GTCTATTTTATACGGAAAAAATTGTATTG	
miR-200b_MUT	TCGAGTAGCGGCCGCTAGTTCATCATTACCAGG <b>GTC</b> TATTA	
miR-429_MUT	TCGAGTAGCGGCCGCTAGTACGGTTTTACCAGA <b>GTC</b> TATTA	
<b>qRT-PCR</b>		<b>Forward 5'-3'</b>
fibronectin1	CCAGTCCTACAACCAGTATTCT	
zeb1	GGCAGAGAATGAGGGAGAAG	
<b>siRNA</b>		<b>targetsequence 5'-3'</b>
siFN1#1	CCGGTTGTTATGACAATGGAA	
siFN1#2	CCCGTTGTTATGACAATGGAA	

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**Reverse 5'-3'**

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CGGGAATTCGGCTCCGGGTATCTGTGACT  
GGAAGATCTAAGCCCCAGATGCGCAGCCT  
GGAAGATCTAGGGGCTGCCAGAGGCTCATGGT

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**Reverse 5'-3'**

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TGCTCTAGATCCCTTGCAGGCAATCTCTT  
CTAGATAAATACTGCCTGGTAATGATGAACTAGCGGCCGCTAC  
CTAGATAAATACTGTCTGGTAAAACCGTACTAGCGGCCGCTAC

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**Reverse 5'-3'**

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AGCGGTATTGAATACTTCGAAGCAGAACAGGCA  
GGAAAAATTGATAAATAACAGGTAAGAGAAAAGAT  
CTAGATAATA**GAC**CCTGGTAATGATGAACTAGCGGCCGCTAC  
CTAGATAATA**GAC**TCTGGTAAAACCGTACTAGCGGCCGCTAC

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**Reverse 5'-3'**

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CTTCTCTGTCAGCCTGTACATC  
CTTCAGACACTTGCTCACTACTC

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