

Table S1: Primary antibody specifications

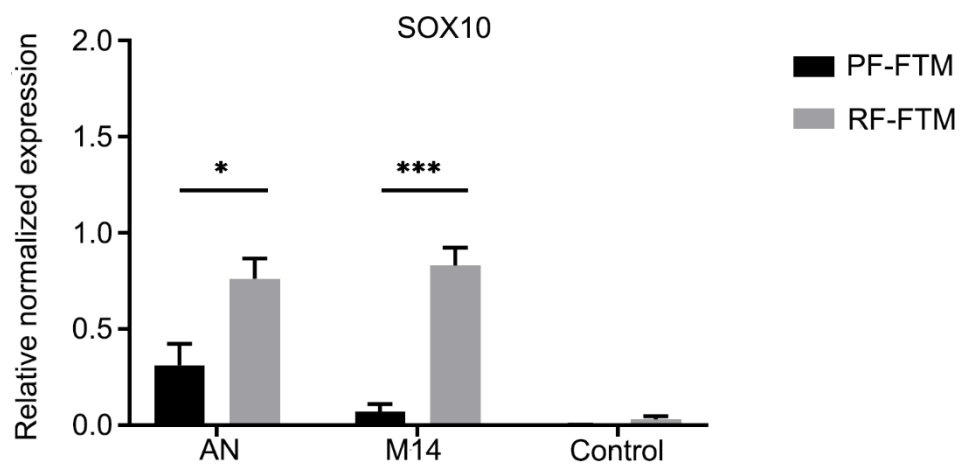
Antibody	Cat.#	Host	Dilution	Source
Collagen type IV	PHM12	Mouse	1:150	Chemicon, Temecula, USA
Ki67	MIB1	Mouse	1:75	Dako, Glostrup, Germany
Keratin 17	CK-E3	Mouse	1:25	Novus Biologicals, Colorado, USA
Keratin 10	DE-K10	Mouse	1:100	Labvision/neomarkers, California, USA
Vimentin	EPR3776	Rabbit	1:1000	Abcam
$\alpha$ -SMA	1A4	Mouse	1:400	Abcam
ZEB2	HPA003456	Rabbit	1:250	Atlas antibodies
ZEB1	HPA027524	Rabbit	1:666	Atlas antibodies
$\beta$ -catenin	610154	Mouse	1:250	BD Bioscience
SPARC	15274-1-AP	Rabbit	1:400	Proteintech
NKI/beteb	ab34165	Mouse	1:100	Abcam
Pro-COL11A1	CI0011	Mouse	1:125	Oncomatrix

Table S2: qPCR primer specifications

Gene	Forward primer 5'> 3'	Reverse primer 5'> 3'
<b>Genes of interest</b>		
SERPINB4	CAAAGGGCAGTGGGAGAATA	CCTCCAGCAAGGCAAAATTA
MMP9	CCTGGAGACCTGAGAACCAA	ATTTCGACTCTCCACGCATC
SNAI2	ACAAGCAGCTGCACTGTGAT	ACACAAGGCAATGTGTGGGT
N-cadherin	TCATTGCCATCCTGCTCTGCAT	AGTTGTTTGGCCTGGCGTTCTT
CLEC3B	CTCAAGAGCCGCTCTGGACACC	AAGCATTTTCATGTGCACCTTGG
COL7A1	GTACCAGGACCCTGAAGCTC	GTACCAGCGCAGGGTGTAG
MMP12	ATGCAGCTGTTTTTAACCCACG	TGGGATAACCAGGGTCCATCA
MMP3	AGGCTTTCCCAAGCAAATAGC	GGGTCAAACCTCAAAGTGTGA
COL11A1	AAAGCACTTCGCTTCCTGGG	GACAGTCTTTTCATAGCCTTTTCTG
HAS1	TACTTTTGGGGATGACCGGC	GACGAGGGCGTCTCTGAGTA
ITG $\beta$ 2	GGGATGGACCGCTACCTCAT	GAATGCCGATCAGCACGATG
MCAM	TCATTGATGGAGTGCCTGTG	CTGTTTCTCCACCTCCAGTAGA
POSTN	TCATTGATGGAGTGCCTGTG	CTGTTTCTCCACCTCCAGTAGA
SPP1	AGAAGTTTCGCAGACCTGACA	GGGATGGCCTTGTATGCACC
VCAM	CAGGCTGGAAGAAGCAGAAAG	TGTCTCCTTCTTTGACACTCTCAG
SOX10	TGAAGGCAGGAAGGAGTTGGC	TGGAGGTTGTAGTGGAGGAGGAC
MMP9	CGACGTCTTCCAGTACCGAG	CTGGTTCAAACCTACTCCGGG
SPARC	CTGGACCAGCACCCATTGA	AGGTCTCGAAAAAGCGGGTG
TGF $\beta$ 3	CGGGCTTTGGACACCAATTAC	CCCAGATCCTGTCTGGAAGTC
ITG $\alpha$ 5	CCTGCTCATCCAGAATGGGG	CGATGTGAATCGGCGAGAGT
TMEM132A	AAATCCTGGTGTCTGAGCGG	CTCCAGTCAGTGGTGCTGTA
TMEFF1	TGCCCTGAAAACCTCAATGGT	TTTCACAGTGCTGTCCAGTGT
ZEB2	AAGTACCAGCGGAAACAAGGA	ACAGACAGGAGTCGGAGTCT
SERPINE1	TCCACAAATCAGACGGCAGC	TCGTAGTAATGGCCATCGGG

FZD7	GCTCATGAACAAGTTCGGCTT	GAGCCGTCCGACGTGTTCT
$\alpha$ -SMA	CTCGGGCTGAAGAAGGAACCCAA	ATTGGCACACGGCAGGTGGT
CCRL1	TGAGGGTCCTACAGAGCCAACCA	CTCCCCCTTCCCCCAACCCA
TGM2	GGTGTCCCTGCAGAACCCGC	CGGGGTCTGGGATCTCCACCG
MGP	GCCATCCTGGCCGCCTTAGC	TTGGTCCCTCGGCGCTTCCT
<b>Reference genes</b>		
RPS29	TATGTGCCGCCAGTGTTTCC	TGCCCCGGATAATCCTCTGA
ZNF410	GCTGTGGTAAGCAGTTTACTACAG	CTTGGGCTTCACAAAGGAAAGG

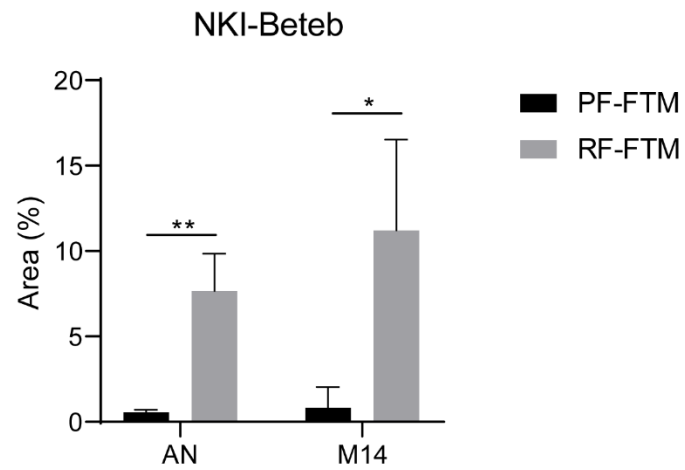
Figure S1: The mRNA expression of SOX10 in melanoma FTMs



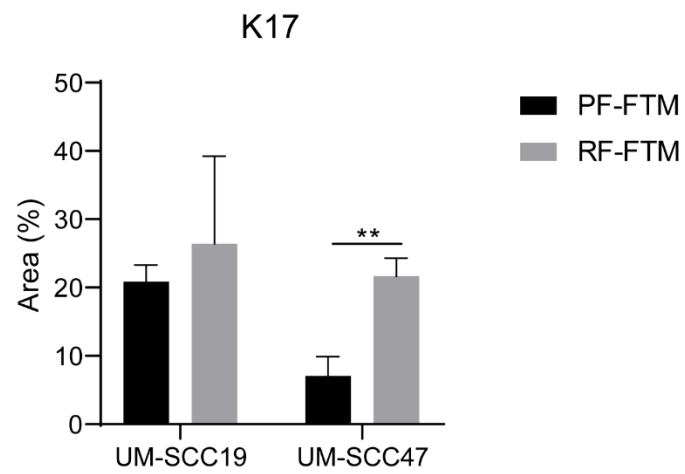
Data was collected from three independent experiments and presented as mean  $\pm$  standard deviation (SD); \*  $p < 0.05$ , \*\*\*  $p < 0.001$ .

Figure S2: The quantification of NKI-Beteb, K17 and K10 in melanoma and HNSCC FTMs

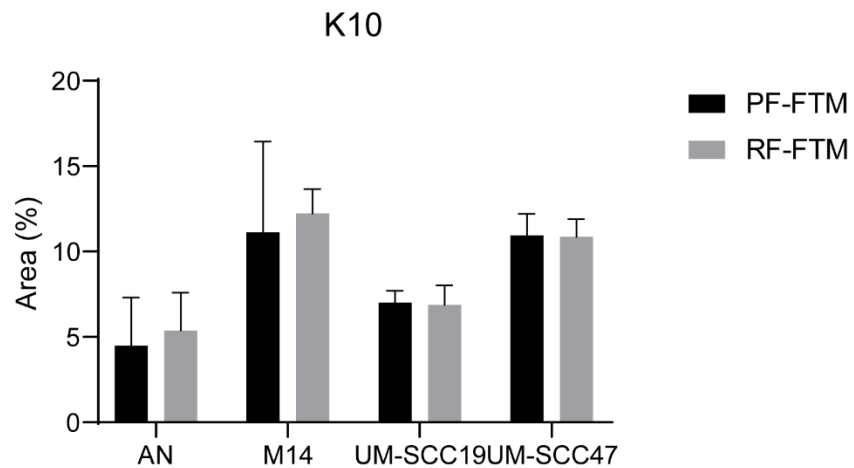
a



b



c



The percentage of positive staining area of (a) NKI-Beteb, (b) K17 and (c) K10 in each FTM was calculated. Data was obtained from three independent experiments and presented as mean  $\pm$  standard deviation (SD); \*  $p < 0.05$ ; \*\*  $p < 0.01$ .