

Supporting Information

Table S1. Gene abbreviations, NCBI reference sequences of mRNAs (used as template for primer design), amplicon size and primer sequences used for Real-Time PCR Analysis.

Gene	mRNA ID (NCBI Reference Sequence)	Amplicon Size (bp)	Forward Primer Sequence (3'→ 5')	Reverse Primer Sequence (5'→ 3')
<i>18S rRNA</i>	NR_033238.1	176	GGAAGTGGAGCCAT GATTAAG	CGGAAGTACGACGGT ATCTG
<i>COL1A1</i>	XM_008271783.1	271	CTGGTGAATCTGGAC GTGAG	TGTCTCACCCCTGTCA CCAC
<i>COL1A2</i>	NM_001195668.1	170	CCAATCACGCCTCTC AGAAC	GCAGCCATCGACAAG AACAG
<i>COL3A1</i>	XM_002712333.2	259	GCATTGCTTACATGG ATCAGG	CCAACGTCCGCACCA AATTC
<i>FN</i>	AF135404.1	270	GCAACCCACAGTGG AATACG	AGTCCTGACACAACA ACAGAC
<i>αSMA</i>	NM_001101682.2	224	CGTGACTACTGCTGA ACGTG	GGATGCCAGCAGATT CCATC
<i>TNC</i>	FJ480400.1	219	GTCATCATCACAGC TCTGG	CTGAGTGTGTATTCCG TGGC
<i>TNMD</i>	NM_001109818.1	239	GCAGTTTCCGAGTTA CAAGAC	CGACGGCAGTAAATA CAACAG
<i>DCN</i>	NM_001082330.1	255	GTGGACAATGGTTCT CTGGC	AAGGTGGATGGCTGG ATCTC
<i>ACAN</i>	L38480.1	113	GCTACGGAGACAAG GATGAG	GTAAAAGACCTCACC CTCCAT
<i>BGN</i>	NM_001195691.1	199	GGCCTGAAGCTCAAC TACCT	GGCTCCCGTTCTCAAT CATC
<i>MKI67</i>	XM_008251084.2	283	CACATCCAGCAGTGA AACGG	GTGTTAGCAGTACCT GAAGTC
<i>MMP2</i>	D63579.1	220	GAAGATCGACGCTGT GTACG	GTATCTCCAGAACTT GTCTCC
<i>MMP9</i>	D26514.1	160	GATACAGCCTGTTCC TCGTG	GGACCATATAGATGC TGGATG
<i>TIMP1</i>	NM_001082232.2	236	CTACCTTGTAACCAGC GTTATG	GAAGCTCAGACTGTT CCAGG
<i>TIMP2</i>	XM_008252510.2	224	CTGTCTGGAGGAAGC ACTTG	CTGCTCAGCGTGGAT ACAAG

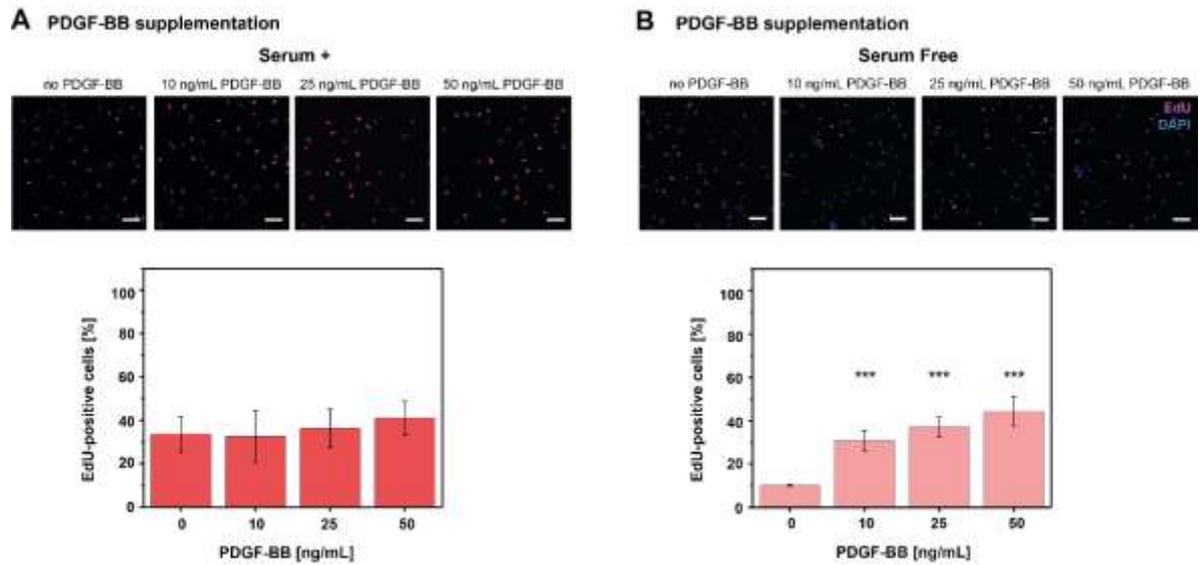


Figure S1. Effect of PDGF-BB supplementation on tenocyte proliferation. Representative CLSM images from the EdU proliferation assay and EdU-positive cells [%] for each PDGF-BB concentration (ng/mL) tested in A) serum+ conditions and B) serum-free conditions. Data was adapted from [47]. The total EdU-positive cells [%] was calculated as EdU positive cells (pink) in relation to the total cell number (DAPI stained, blue). Scale bars: 100 μ m. (Data shown as mean \pm standard deviation, ** $p < 0.01$, *** $p < 0.001$, obtained by one-way ANOVA).

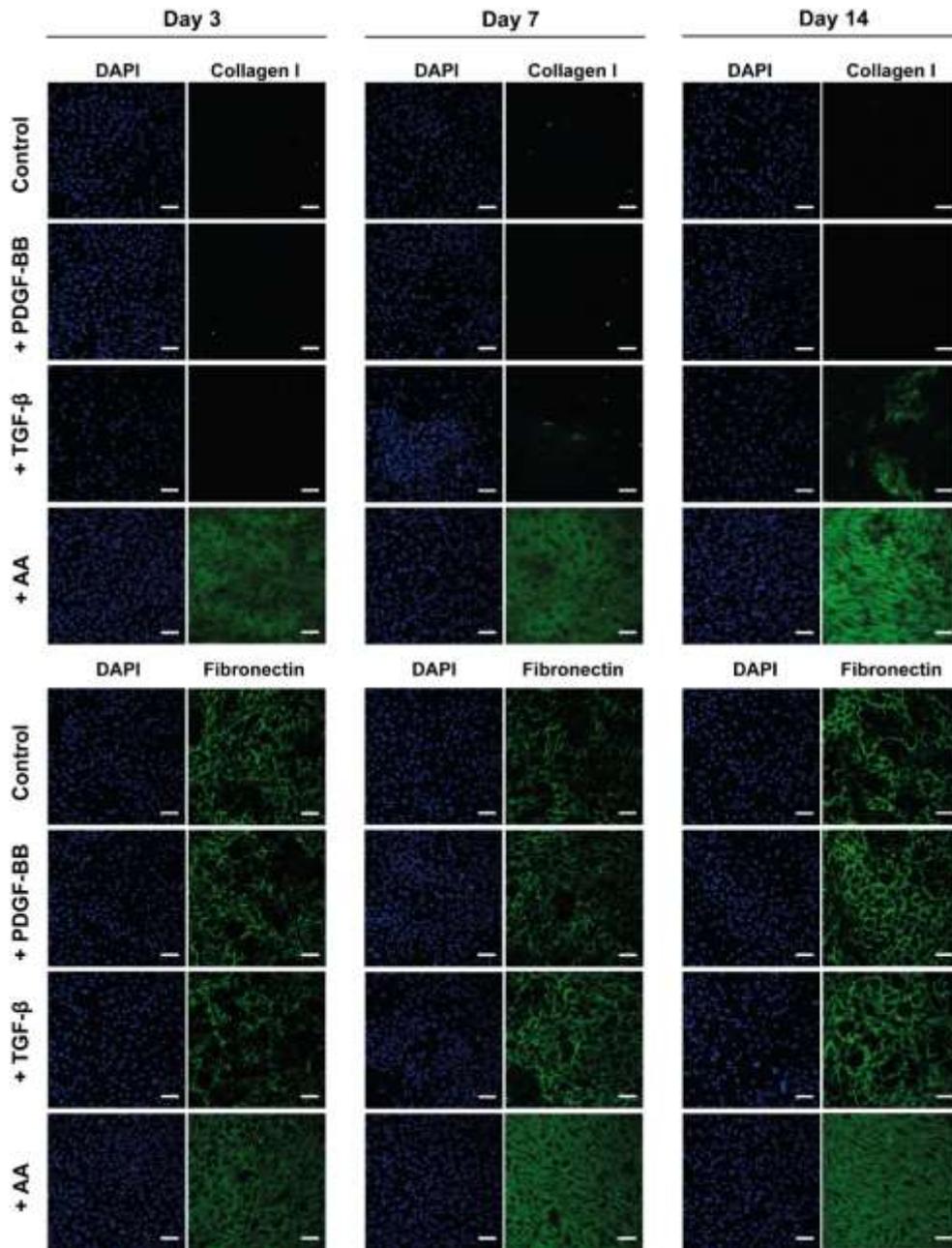


Figure S2. Collagen I and fibronectin staining (green) and cell nuclei DAPI staining (blue) of rabbit tenocytes (Animal 2) cultured in vitro in 2D, treated with PDGF-BB, TGF-β, AA or untreated (control) at day 3, 7 and 14. Scale bars: 100 μm. .

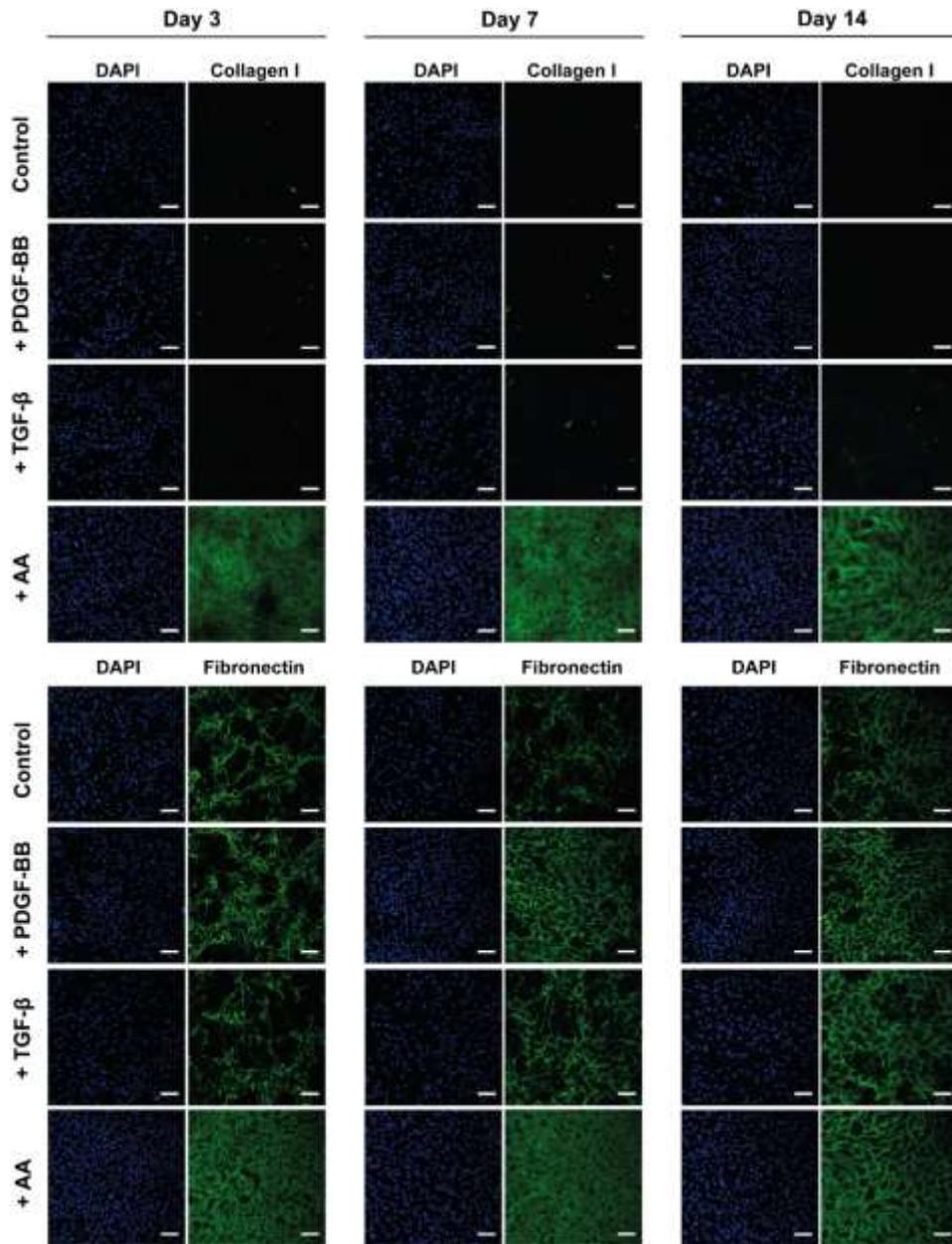


Figure S3. Collagen I and fibronectin staining (green) and cell nuclei DAPI staining (blue) of rabbit tenocytes (Animal 3) cultured in vitro in 2D, treated with PDGF-BB, TGF- β , AA or untreated control at day 3, 7 and 14. Scale bars: 100 μ m.

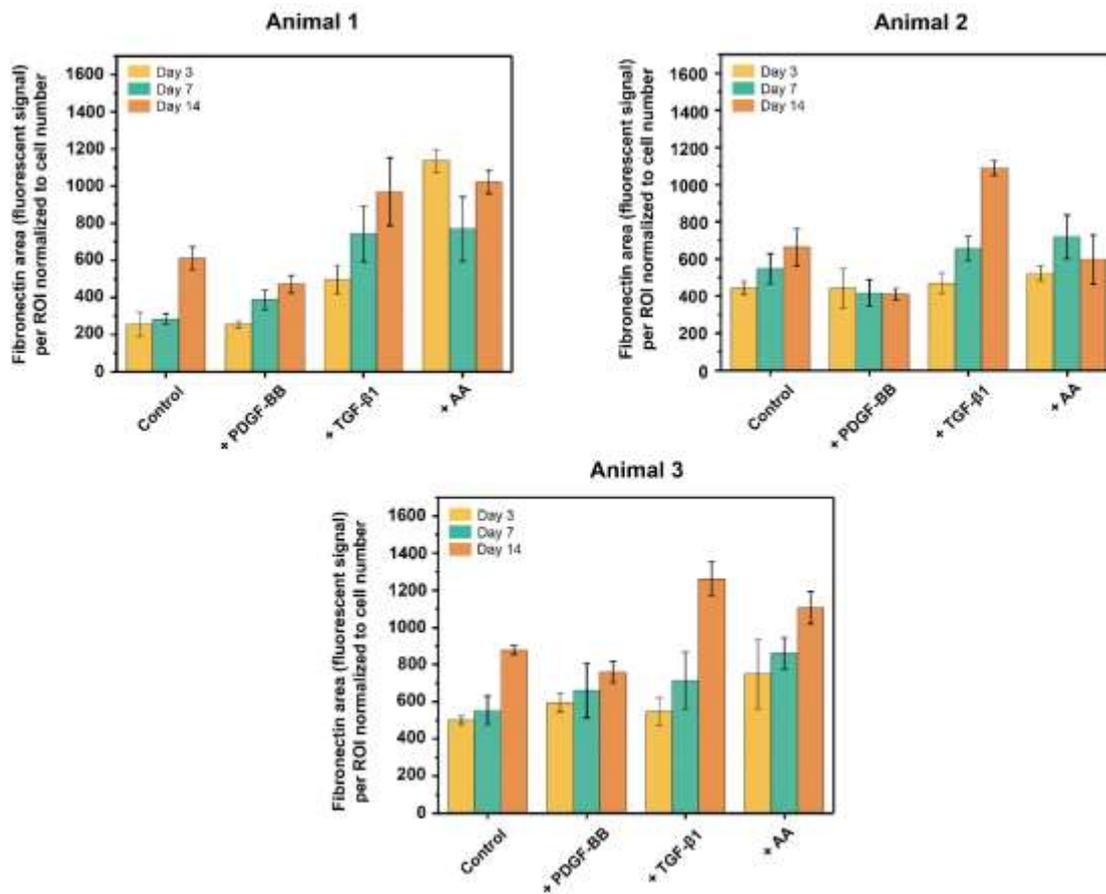


Figure S4. Quantification of fibronectin deposition by tenocytes in the different treatment conditions. The signal of fibronectin staining was assessed from images of maximum projections of the z-stacks obtained with confocal microscopy. Briefly, thresholded images were segmented and the total area of the signal in each image was assessed. The area measured was normalized to the cell number relative values for the ratio obtained are shown on the y-axis) in each image and thus the fibronectin deposition was quantified independent of cell density, as the treatments used do increase cell proliferation.