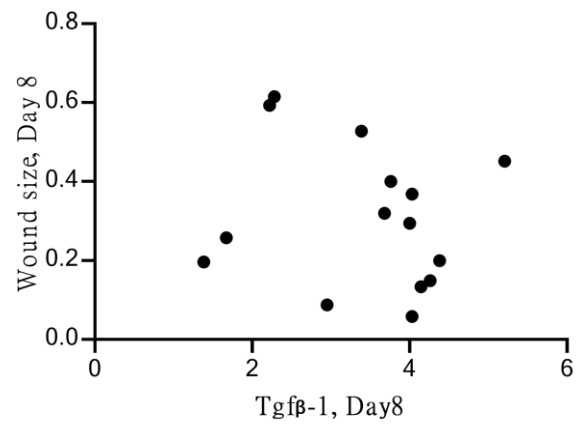




Supplementary Figure 1. Correlation between *Tgfb-1* expression and wound size at Day 8.

Each dot represents an individual sample (n = 15). Spearman correlation analysis was performed.

Table 1. Real-time qPCR primer sequences.

Gene	Forward (3' -> 5')	Reverse (5' -> 3')	Ref
<i>Col I</i>	TGACTGGAAGAGCGGAGAGT	GAATCCATCGGTCATGCTCT	[14]
<i>Col III</i>	TCCTGAAGATGTCCTTGATGTACAG	TTCAGAGACTTCTTTACATTGCCATT	[15]
<i>Mmp9</i>	CCTCTGCATGAAGACGACATAA	GGTCAGGTTTAGAGCCACGA	[16]
<i>bFgf</i>	AGCAGAAGAGAGAGGCGTTG	ACTGCCCAGTTCGTTTCAGT	[17]
<i>Ctgf</i>	AGACCTGTGCCTGCCATTA	TGTCTCCGTACATCTTCCTG	[18]
<i>Tgf-β1</i>	ACCGACCCTTCCTGCTCCTCAT	GATCCACTTCCACCCAGGTCCT	[19]
<i>Tgf-β2</i>	GAGGATAATTGCTGCCTTC	GGCTGAGGACTTTGGTGTGT	[19]
<i>Tgf-β3</i>	TGCGCCCCCTCTACATTG	GGTTCGTGGACCCATTTC	[20]
<i>Wisp1</i>	CCGACCACACATCAAGGCAGG	GGTCGGTAGGTGCGTGTGCTG	[19]
<i>Wisp2</i>	GCCCGAGGTACGCAATAGG	GCAGTTGGGTTGGAAGGACT	[19]
<i>Timp1</i>	ATCGCGGGCCGTTTAAGGA	CAAGGGATGGCTGAACAGGGA	[19]
<i>Gapdh</i>	GTTACCAGGGCTGCCTTCTC	GGGTTTCCCGTTGATGACC	[21]

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