

## Supplementary Tables

**Supplementary Table 1.** Primer sequences for reverse transcription-polymerase chain reaction

Gene	Primer sequence
Human GAPDH	Forward 5'-GAAGGTGAAGGTCGGAGTC-3'
	Reverse 5'-GAAGATGGTATGGGATTTC-3'
Human Smad7	Forward 5'-CCTTAGCCGACTCTGCGAACTA-3'
	Reverse 5'-CCAGATAATTGTTCCCCCTGT-3'
Human TSP-1	Forward 5'-AACAAACCCACACCCAGTTG-3'
	Reverse 5'-TTGAAGCAGGCATCAGGCAC-3'
Human COL1A1	Forward 5'-AGGACAAGAGGCATGTCTGGTT-3'
	Reverse 5'-TTGCAGTGGTAGGTATGTTCTG-3'
Mouse Col1A1	Forward 5'-TTCACCTACAGCACCGCGCTTGTG-3'
	Reverse 5'-GATGACTGTCTGCCCAAGTT-3'

**Supplementary Table 2.** List of genes from the Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways extracted by DAVID program\*

Probe ID	Signal intensity		dnT $\beta$ RII /GFP ratio	Gene symbol	KEGG pathway		
	dnT $\beta$ RII	GFP			<i>a</i>	<i>b</i>	<i>c</i>
202310_s_at	43.9	230.4	0.19	<i>COL1A1</i>	x	x	x
201110_s_at	91.3	454	0.20	<i>THBS1</i>	x	x	x
235086_at	42.3	179.3	0.24	<i>THBS1</i>	x	x	x
202267_at	174.1	813.2	0.21	<i>LAMC2</i>	x	x	x
209800_at	76.2	386.6	0.20	<i>KRT16</i> <sup>†</sup>	x		
203879_at	9.7	69.9	0.14	<i>PIK3CD</i> <sup>†</sup>		x	

\* *a* = cell communication; *b*, focal adhesion; *c*, extracellular matrix–receptor interaction; x, included; KEGG = Kyoto Encyclopedia of Genes and Genomes; ID = identification number.

<sup>†</sup>*KRT16* and *PIK3CD* genes encode keratin 16 and the catalytic, delta polypeptide of phosphatidylinositol-3-kinase, respectively.

**Supplementary Table 3.** Relationship between phosphorylation of Smad2 and expression of thrombospondin-1 in human gastric cancer specimen\*

TSP-1 expression	No. of specimens Phosphorylated Smad2		<i>P</i> value†
	Positive	Negative	
Positive	44	14	
Negative	20	24	.002

\* TSP-1 = thrombospondin-1.

† Obtained from a two-sided  $\chi^2$  test.