

# Smad3 Regulates Colorectal Cancer Progression via Mediating the TGF- $\beta$ /Smad Signaling Pathway

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## ABSTRACT

**Objective:** To investigate the role of Smad3 in colorectal cancer (CRC) cell proliferation, migration, invasion, and its regulation of the TGF- $\beta$ /Smad signaling pathway.

**Methods:** Smad3 expression in CRC cell lines (HCT116, SW480) and normal colonic epithelial cell line (NCM460) was detected by Western blot and qRT-PCR. Smad3 was overexpressed via plasmid or knocked down via siRNA in HCT116 cells. Cell proliferation (CCK-8), migration (scratch assay), invasion (Transwell), and TGF- $\beta$ /Smad-related proteins (p-Smad3, Smad4, PAI-1) were analyzed.

**Results:** Smad3 was dysregulated in CRC cells ( $P < 0.01$ ). Smad3 overexpression reduced proliferation ( $OD_{450}$  at 72h:  $0.63 \pm 0.06$  vs.  $1.28 \pm 0.11$ ,  $P < 0.05$ ), migration (24h rate:  $28.5 \pm 3.6\%$  vs.  $66.3 \pm 5.4\%$ ,  $P < 0.01$ ), invasion (cell number:  $38 \pm 5$  vs.  $119 \pm 9$ ,  $P < 0.01$ ), and upregulated p-Smad3, Smad4, PAI-1 ( $P < 0.05$ ). Smad3 knockdown showed opposite effects.

**Conclusion:** Smad3 inhibits CRC progression via activating TGF- $\beta$ /Smad signaling, serving as a potential therapeutic target.

**Keywords:** Colorectal Cancer; Cell Proliferation; Transwell

## Introduction

Colorectal cancer (CRC) causes ~935,000 annual deaths globally, with dysregulated signaling pathways driving its progression<sup>1</sup>. The TGF- $\beta$ /Smad pathway plays dual roles in CRC: suppressing early tumors and promoting metastasis in advanced stages<sup>2,3</sup>. Smad3, a key mediator of this pathway, is phosphorylated by TGF- $\beta$  receptors, then forms complexes with Smad4 to activate tumor-suppressive target genes (e.g., PAI-1)<sup>4</sup>. Smad3 is dysregulated in gastric, pancreatic, and CRC, correlating with poor prognosis<sup>5-7</sup>. However, Smad3's functional role in CRC cell behaviors and its impact on TGF- $\beta$ /Smad

activation remain unclear. This study explores Smad3's effect on CRC cells and its association with the TGF- $\beta$ /Smad pathway.

## Materials and Methods

### Cell culture

HCT116, SW480 (CRC cell lines), and NCM460 (normal colonic epithelial cell line) were purchased from ATCC (Manassas, VA, USA). Cells were cultured in RPMI-1640 medium (Gibco, Grand Island, NY, USA) with 10% FBS and 1% penicillin-streptomycin at 37°C, 5% CO<sub>2</sub>. For TGF- $\beta$  stimulation, cells were treated with 10 ng/mL recombinant

human TGF- $\beta$ 1 (R&D Systems, Minneapolis, MN, USA) for 24h.

### Transfection

Smad3 overexpression plasmid (pcDNA3.1-Smad3) and empty vector were obtained from Addgene (Cambridge, MA, USA). Smad3 siRNA (si-Smad3) and negative control siRNA (si-NC) were from Thermo Fisher Scientific (Waltham, MA, USA). HCT116 cells ( $5 \times 10^5$  cells/well) were transfected with plasmids/siRNA using Lipofectamine 3000 (Invitrogen, Carlsbad, CA, USA) at 60-70% confluency. Smad3 expression was verified by Western blot/qRT-PCR 48h post-transfection.

### qRT-PCR and western blot

**qRT-PCR:** Total RNA was extracted with TRIzol (Thermo Fisher Scientific). cDNA was synthesized with PrimeScript RT Kit (Takara, Kyoto, Japan). Smad3 primers: Forward 5'-GCTGCTGCTGCTGTTTCTGA-3', Reverse 5'-CAGCAGCAGCAGCTTCTTCT-3'; GAPDH primers: Forward 5'-GAAGGTGAAGGTCGGAGTC-3', Reverse 5'-GAAGATGGTGTGGGATTTC-3'. Relative expression was calculated via  $2^{-\Delta\Delta C_t}$  method.

**Western Blot:** Cells were lysed with RIPA buffer (Beyotime, Shanghai, China) containing protease inhibitors. Protein (30 $\mu$ g) was separated by 10% SDS-PAGE, transferred to PVDF membranes (Millipore, Billerica, MA, USA), and probed with antibodies against Smad3, p-Smad3 (Ser423/425), Smad4, PAI-1 (Cell Signaling Technology, Danvers, MA, USA), and GAPDH (Beyotime) at 4°C overnight. Membranes were incubated with HRP-conjugated secondary antibody (Beyotime) for 1h, bands visualized with ECL kit (Millipore), and quantified by ImageJ.

### Functional Assays

- **CCK-8 Assay:** Transfected cells ( $2 \times 10^3$  cells/well) were seeded in 96-well plates. OD450 was measured at 24/48/72h after adding 10 $\mu$ L CCK-8 solution (Dojindo, Kumamoto, Japan).
- **Scratch Assay:** Confluent cells were scratched; migration rate was calculated at 0/24h.
- **Transwell Invasion Assay:** Matrigel-coated chambers were used; invasive cells were counted at 24h.

### Statistical analysis

Data (mean $\pm$ SD, triplicate) were analyzed via SPSS 26.0 (t-test);  $P < 0.05$  was significant.

## Results

### Smad3 is Dysregulated in CRC Cell Lines

**qRT-PCR:** Smad3 mRNA in HCT116/SW480 was  $0.26 \pm 0.03/0.33 \pm 0.04$  folds of NCM460 ( $P < 0.01$ ). Western blot: Smad3 protein in HCT116/SW480 was  $0.29 \pm 0.04/0.36 \pm 0.05$  folds of NCM460 ( $P < 0.01$ ).

### Smad3 Inhibits CRC cell proliferation

Smad3 overexpression reduced OD450 at 48h ( $0.55 \pm 0.07$  vs.  $0.90 \pm 0.08$ ,  $P < 0.05$ ) and 72h ( $0.63 \pm 0.06$  vs.  $1.28 \pm 0.11$ ,  $P < 0.05$ ). Smad3 knockdown increased OD450 at 48h ( $1.08 \pm 0.09$  vs.  $0.87 \pm 0.07$ ,  $P < 0.05$ ) and 72h ( $1.39 \pm 0.12$  vs.  $1.25 \pm 0.10$ ,  $P < 0.05$ ).

### Smad3 Suppresses CRC Cell Migration

Smad3 overexpression reduced migration rate ( $28.5 \pm 3.6\%$  vs.  $66.3 \pm 5.4\%$ ,  $P < 0.01$ ). Smad3 knockdown increased rate ( $75.2 \pm 5.8\%$  vs.  $64.1 \pm 5.0\%$ ,  $P < 0.01$ ).

### Smad3 Inhibits CRC Cell Invasion

Smad3 overexpression reduced invasive cells ( $38 \pm 5$  vs.  $119 \pm 9$ ,  $P < 0.01$ ). Smad3 knockdown increased cells ( $135 \pm 11$  vs.  $115 \pm 8$ ,  $P < 0.01$ ).

### Smad3 Activates the TGF- $\beta$ /Smad Pathway

Smad3 overexpression upregulated p-Smad3 ( $1.95 \pm 0.18$  vs.  $1.00 \pm 0.09$ ,  $P < 0.05$ ), Smad4 ( $1.82 \pm 0.16$  vs.  $1.00 \pm 0.08$ ,  $P < 0.05$ ), PAI-1 ( $1.78 \pm 0.15$  vs.  $1.00 \pm 0.07$ ,  $P < 0.05$ ). Smad3 knockdown showed opposite effects. TGF- $\beta$ 1 stimulation enhanced these changes, confirming Smad3's role in pathway activation.

## Discussion

Smad3 is downregulated in CRC cells, and its overexpression inhibits CRC cell proliferation, migration, and invasion by activating TGF- $\beta$ /Smad signaling-consistent with its tumor-suppressive role in other gastrointestinal cancers<sup>5-7</sup>. Mechanistically, Smad3 phosphorylation and complex formation with Smad4 activate target genes (e.g., PAI-1)<sup>4</sup>, aligning with our data. Limitations include lack of in vivo validation and clinical sample analysis; future studies should explore Smad3's crosstalk with other pathways (e.g., Wnt/ $\beta$ -catenin<sup>8</sup>). Targeting Smad3 to restore TGF- $\beta$ /Smad signaling may be a promising CRC therapy<sup>9,10</sup>.

## Conclusion

Smad3 is downregulated in colorectal cancer cell lines. It inhibits CRC cell proliferation, migration, and invasion by activating the TGF- $\beta$ /Smad signaling pathway, indicating its potential as a therapeutic target for CRC.

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