The Niche Component Periostin Is Produced by Cancer-Associated Fibroblasts, Supporting Growth of Gastric Cancer through ERK Activation

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Overexpression of periostin (POSTN), an extracellular matrix protein, has been observed in several cancers. We investigated the importance of POSTN in gastric cancer. Genome-wide gene expression analysis using publicly available microarray data sets revealed significantly high POSTN expression in cancer tissues from stage II–IV gastric cancer, compared with background normal tissues. The POSTN/vimentin mRNA expression ratio was highly associated with gene groups that regulate the cell cycle and cell proliferation. IHC showed that periglandular POSTN deposition, comprising linear deposition abutting the glandular epithelial cells in normal mucosa, disappeared during intestinal gastric cancer progression. Stromal POSTN deposition was also detected at the invasive front of intestinal-type and diffuse-type cancers. In situ hybridization confirmed POSTN mRNA in cancer-associated fibroblasts, but not in tumor cells themselves. POSTN enhanced the in vitro growth of OCUM-2MLN and OCUM-12 diffuse-type gastric cancer cell lines, accompanied by the activation of ERK. Furthermore, coinoculation of gastric cancer cells with POSTN-expressing NIH3T3 mouse fibroblast cells facilitated tumor formation. The OCUM-2MLN orthotopic inoculation model demonstrated that tumors of the gastric wall in Postn−/− mice were significantly smaller than those in wild-type mice. Ki-67 and p-ERK positive rates were both lower in Postn−/− mice. These findings suggest that POSTN produced by cancer-associated fibroblasts constitutes a growth-supportive microenvironment for gastric cancer.

stress, such as the endocardium, fascia, periosteam, articular surface, and periodontal ligament. \(^4\) POSTN has also been recognized as playing a key role in tissue repair. Our research group has previously shown that POSTN is expressed by myofibroblasts and that it regulates tissue repair processes in myocardial infarction\(^5\) and wound healing. \(^6\) POSTN is regarded as an interventional molecular target for myocardial infarction therapy. \(^7\) In neoplastic lesions, overexpression of POSTN has been identified in various cancers, including those of the lung (non—small-cell subtypes), \(^8\) ovary, \(^9\) breast, \(^10, 11\) colon, \(^12-14, 17\) pancreas, \(^13\) bile duct, \(^15\) and head and neck. \(^19\) However, expression and function of POSTN in gastric cancer, the second most common cause of cancer death worldwide, \(^20\) have not been studied sufficiently, \(^21, 22\) especially with respect to the two distinctive histological groups, the intestinal and diffuse types. \(^23\) The morphology, molecular and genetic background, and clinical features of the two subtypes differ completely. \(^24\)

In the present study, we investigated the expression and function of POSTN in gastric cancer. First, we evaluated POSTN expression in normal and neoplastic stomach using both bioinformatics and an extensive tissue sample examination to clarify the cell types that predominantly produce POSTN. Next, we examined the function of POSTN using POSTN-producing fibroblasts and an orthotopic inoculation model of diffuse-type gastric cancer in Postn\(^−/−\) mice. Our results demonstrated that POSTN is overexpressed by cancer-associated fibroblasts (CAFs) and suggested that POSTN constitutes the primary tumor niche by supporting cancer cell proliferation through ERK signaling pathway in gastric cancer.

Materials and Methods

Genome-Wide Gene Expression Analysis Using Publicly Available Microarray Data Sets

We compared POSTN mRNA levels using the following public domain microarray data sets from the Gene Expression Omnibus (GEO, \(\text{http://www.ncbi.nlm.nih.gov/geo}\)): GSE15460, 25 gastric cancer cell lines and 161 cases of gastric cancer (8 cases in T1, 44 cases in T2, 108 cases in T3, and 1 case in T4); GSE7307, 12 cases of normal stomach; and GSE19826, 12 cases of gastric cancer and matched normal tissues in which the clinical TNM stage was available, 3 cases for each stage. Raw data were retrieved and processed using the MAS5 algorithm with target intensity of 500 (Affymetrix, Santa Clara, CA). \(^25\)

Gene Set Enrichment Analysis (GSEA) software \(^26\) implemented in a Java GSEA desktop application version 2.0 (\(\text{http://www.broadinstitute.org/gsea/index.jsp}\)) was used to determine whether POSTN has a function correlated with the cell cycle and cell proliferation. Gene sets for use with GSEA were taken from REACTOME (\(\text{http://www.reactome.org}\)) \(^27\) and Gene Ontology (GO) (\(\text{http://www.geneontology.org}\)). \(^28\) Pearson’s correlation coefficient of POSTN per stromal cell [as the ratio of POSTN to vimentin (POSTN/VIM)] and each gene expression level across all samples >100 were used to rank genes. Expression levels of probe sets 210809_s_at and 204126_s_at were used as POSTN and VIM mRNA expression, respectively.

Human Tissue Samples

Human tissue materials (32 non-neoplastic gastric mucosae, 87 intestinal-type gastric cancers, and 50 diffuse-type gastric cancers) were obtained during surgery and retrieved from the archives of the University of Tokyo Hospital during 2005 to 2006. This study was approved by the Ethics Committee of the Graduate School of Medicine of the University of Tokyo (no. 2381).

Cells and Tissue Culture

Human diffuse-type gastric cancer cell lines OCUM-2MLN and OCUM-12\(^25\) and the mouse fibroblast cell line NIH3T3 expressing POSTN (NIH3T3\(_{\text{POSTN}}\)) and its EGFP-expressing control (NIH3T3\(_{\text{EGFP}}\)) were established as described previously. \(^13\) OCUM-2MLN, OCUM-12, and the NIH3T3 transfectants were grown in Dulbecco’s modified Eagle’s medium (Sigma-Aldrich) at 37°C in a humid atmosphere saturated with 5% CO\(_2\).

Animals

C57BL/6 Postn\(^−/−\) mice (previously established in our laboratory) \(^14\) were intercrossed with C57BL/6 Rag2\(^−/−\) mice (Taconic Farms, Germantown, NY) to obtain POSTN and RAG-2 double-knockout mice. The method using Rag2\(^−/−\) mice had been adopted in a study to survey the function of every type of host-derived matrix metalloproteinase in the cancer microenvironment. \(^31\) Genotyping procedures for Postn and Rag2 were performed as described previously \(^30\) and according to the supplier’s protocol. Xenograft studies were performed using female BALB/c nude mice, obtained from Charles River Laboratories International (Kanagawa, Japan; Wilmington, MA). Mice were handled under specific-pathogen-free conditions in accordance with the policies of the Animal Ethics Committee of the University of Tokyo (no. 1822T-020).

IHC and Immunofluorescence

Tissue samples were fixed in 4% neutral formalin and were embedded in paraffin routinely. The primary antibodies used for the present study included a previously established rabbit polyclonal anti-POSTN, \(^3\) anti—α-smooth muscle actin (anti—α-SMA) (1:50; 1A4; Dako—Agilent Technologies, Glostrup, Denmark), anti—human Ki-67 (1:200; MIB-1; Dako—Agilent Technologies), and anti—p-p44/42 MAPK
(1:200; no. 9101; Cell Signaling Technology, Danvers, MA). IHC and immunofluorescence microscopy were performed (sections 5 μm thick) as described previously. For semiquantifying POSTN expression, we scored four groups according to prevalence: score 0, <10%; score 1, 10% to 49%; score 2, 50% to 80%; and score 3, >80%. We regarded scores 2 and 3 as significant and positive.

**In Situ Hybridization for POSTN mRNA**

Antisense and sense complementary (c)RNA probes were prepared by *in vitro* transcriptions of EcoRI–XbaI fragment of human and mouse POSTN cDNA using a DIG labeling mix (Roche Applied Science, Indianapolis, IN) as described previously. Nonradioactive mRNA *in situ* hybridization was performed manually on formalin-fixed, paraffin-embedded sections (5 μm thick) as described previously.

**Three-Dimensional Coculture with Cancer Cells and Fibroblasts**

OCUM-2MLN and either of the NIH3T3 transfectants (NIH3T3POSTN or NIH3T3EGFP), each at a final concentration of 1 × 10^6/mL, were suspended in 1 mL of the mixture of type I collagen gel. Three-dimensional coculturing was then performed for 2 weeks, as described previously. For semiquantifying POSTN expression, we scored four groups according to prevalence: score 0, <10%; score 1, 10% to 49%; score 2, 50% to 80%; and score 3, >80%. We regarded scores 2 and 3 as significant and positive.

**Proliferation Assay**

Proliferation assay was performed using a cell-counting kit (CCK-8; Dojindo Molecular Technologies, Kumamoto, Japan) as described in the technical manual. The final concentration was supplemented with 0.1% bovine serum albumin and either recombinant human POSTN (250 ng/mL, BioVendor, Heidelberg, Germany) dissolved in PBS or vehicle alone.

**Phosphorylated MAP Kinase Protein Array**

Phosphorylated MAP kinase protein array analysis was performed using a human p-MAPK array kit (Proteome Profiler antibody array; R&D Systems, Minneapolis, MN) according to the manufacturer’s instructions. Samples of cell lysates were obtained after 24 hours incubation in a no-serum condition, supplemented with 1 μg/mL of recombinant human POSTN (BioVendor) or equal amounts of PBS.

**Western Blotting**

For detection of ERK phosphorylation by POSTN, cells were cultured overnight in a no-serum condition after 1 hour of incubation supplemented with 250 ng/mL of recombinant human POSTN (BioVendor) or equal amounts of PBS. Cell lysate samples were prepared in radioimmunoprecipitation assay buffer separated on 8% polyacrylamide gels. They were electrophoretically transferred onto a polyvinylidene difluoride membrane. After blocking, the membranes were incubated for 1 hour at room temperature with an anti-ERK (1:1000; sc-94; Santa Cruz Biotechnology) or anti—p-p44/42 MAPK (1:1000; no. 9101; Cell Signaling Technology). The membranes were washed again and incubated for 1 hour with secondary antibodies. The antigen was then detected using enhanced chemiluminescence Western blot detection reagents (GE Healthcare, Little Chalfont, UK) according to the manufacturer’s instructions.

**Coinoculation of OCUM-2MLN or OCUM-12 Diffuse-Type Gastric Cancer Cells with NIH3T3POSTN Cells in Vivo**

OCUM-2MLN or OCUM-12 (5 × 10^6 cells) and NIH3T3 POSTN or NIH3T3EGFP (1 × 10^6 cells) were suspended in 50 μL of PBS and were coinjected subcutaneously in the flanks (left flank with NIH3T3POSTN; right flank with NIH3T3EGFP) of nude mice. Tumor volumes were calculated using the following formula: volume = (W^2 × L)/2, where W is the short diameter and L is the long diameter. Histological features of the xenografts were examined with H&E staining. H&E sections were scanned using a Nanozoomer 2.0HT system (Hamamatsu Photonics, Hamamatsu, Japan) and the tumor area was analyzed using NDP.view viewer software version 1.1.27 (Hamamatsu Photonics).

**Orthotopic Inoculation Model of Diffuse-Type Gastric Cancer Cell Line**

A total of 5 × 10^6 cells suspended in 50 μL of PBS were inoculated subserosally into the gastric walls of mice, as described previously. For the present study, we used Postn<sup>−/−</sup> Rag2<sup>−/−</sup> mice (n = 6) and Postn<sup>+/−</sup>Rag2<sup>−/−</sup> mice (n = 7), aged 4 to 5 weeks. At 5 weeks after inoculation, the mice were sacrificed for subsequent evaluation procedures.

**Statistical Analysis**

For analysis of publicly available microarray data sets, the data were evaluated using Wilcoxon signed-rank test for statistical comparisons. The trend test was performed with a Jonckheere–Terpstra test. For the distribution of POSTN expression in IHC, a Cochrane–Armitage trend test was applied. These statistical tests were performed using R software version 2.1.10 (http://www.r-project.org). In *vitro* and *in vivo* growth data were evaluated with two-way analysis of variance using GraphPad Prism software.
version 4 (GraphPad Software, San Diego, CA). All other numerical results are expressed as means ± SEM. Data were evaluated using *U*-tests for statistical comparisons using StatView software version 4 (SAS Institute, Cary, NC). *P* < 0.05 was considered significant for all statistical tests.

**Results**

**Microarray Data Set Analysis of POSTN mRNA Expression in Gastric Cancer**

To study whether POSTN expression is up-regulated in gastric cancer, we first analyzed POSTN mRNA levels in normal gastric tissue, gastric cancer tissue, and gastric cancer cell lines from publicly available microarray data sets (GSE15460 and GSE7307). POSTN expression was found in both normal and neoplastic tissues, without any marked differences, although all gastric cancer cell lines (excepting only YCC11) expressed negligible POSTN (Figure 1A). In a closer comparison of POSTN mRNA expression levels between cancer and matched normal stomach samples, we analyzed another data set including an annotation for the clinical stage (GSE19826). It is particularly interesting that no difference between cancer and normal tissue samples was found for stage I gastric cancer (*P* = 0.98, Wilcoxon signed-rank test; *n* = 3), although cancer tissue samples showed significantly higher POSTN expression than each corresponding background normal tissue for stage II, III, and IV gastric cancer (*P* = 0.049, Wilcoxon signed-rank test; *n* = 9) (Figure 1B).

We further analyzed the correlation between POSTN mRNA level and tumor progression using the GSE15460 data set. The trend test results showed that the POSTN mRNA expression in tumor samples was proportionally greater at the higher T stages (*P* = 0.0033, Jonckheere–Terpstra test) (Figure 1C).

To correlate the stromal amount and VIM expression, we analyzed the data set GSE8218, comprising 136 cases of prostate cancer for which the stromal percentage was available for each tumor sample (Supplemental Figure S1). Raw data were retrieved and processed using the MAS5 algorithm (Affymetrix). Pearson correlation coefficients indicated correlation between VIM mRNA expression and the amount of stroma in prostate cancer (*R* = 0.525,

![Figure 1](image-url)

**Figure 1** POSTN mRNA expression in normal stomach and gastric cancer from publicly available microarray data sets. **A:** Expression of POSTN mRNA in normal gastric tissue (*n* = 12), gastric cancer tissue (*n* = 161), and gastric cancer cell lines (*n* = 25). **B:** Comparison of POSTN mRNA expression levels between cancer and matched normal stomach samples. No difference was found between cancer (gray bars) and normal tissue (white bars) samples in stage I gastric cancer (*P* = 0.98), but for stage II, III, and IV gastric cancer tissue samples showed significantly higher POSTN expression than the corresponding background normal tissues, *P* = 0.049. **C:** Correlation between POSTN expression and tumor progression. POSTN mRNA expression in tumor samples was greater at the higher T stages (*P* = 0.00339). The POSTN/VIM mRNA expression ratio was also associated significantly with T stage (*P* = 0.0077). Boxplots indicate the median and first and third quartiles; whiskers indicate the 10th and 90th percentiles. Outlier values (>1.5 box-lengths from the 3rd quartile) are indicated by symbols (circles). *P* < 0.05.
POSTN deposition in the ECM of reactive gastric mucosa, in lamina propria mucosae, but not in epithelial cells. Immunostaining of normal gastric mucosa, showing POSTN expression in the lamina propria mucosae and surrounding the proper gastric glands (Figure 2, A–C). Stromal staining, however, was indicative of intense POSTN deposition in the ECM of reactive gastric mucosa, in which there is an inflammatory cell infiltrate and an increase in vascularity, in addition to more pronounced periglandular immunoreactive POSTN in the intestinal metaplasia area (Figure 2D). The stromal staining pattern was barely visible in the normal lamina propria mucosae. Immunofluorescent double staining for POSTN and α-SMA showed a focal close approximation of α-SMA+ cells to POSTN immunoreactivity, which encircled normal gastric glands (Figure 2E). No specific staining of epithelial cells or other mesenchymal cell types was noted, with the exception of occasional POSTN deposition in the vascular wall. In the submucosal and muscular layer, POSTN expression was focally detected in perivascular cells of capillaries (data not shown).

Next, we examined POSTN expression in gastric adenocarcinomas according to the two histological subtypes, the intestinal type (n = 87) and the diffuse type (n = 50).23 Periglandular POSTN staining was found in all of the non-neoplastic tissues (n = 14), although it was dramatically lower in carcinoma of the intestinal subtype at Tis, in 19% (10/52) of cases, which did not involve the basement membrane (Figure 3, A–C, and Supplemental Table S1). Furthermore, periglandular POSTN staining was observed in only 8% (1/13) of cases of the invasive intestinal-type adenocarcinomas at T1, which involved the lamina muscularis mucosae and which extended into the submucosa (Figure 3C). Periglandular POSTN expression tended to decrease with progressing stage (P = 7.62 × 10⁻⁸). Invasive intestinal-type adenocarcinomas in stages T2 to T4 showed prominent stromal POSTN deposition, rather than periglandular staining (Figure 3D), in contrast to noninvasive intestinal-type cancers. The immunoreactive POSTN was fibrillar and was confined at the invasive front of invasive tumor nests. However, all diffuse-type adenocarcinomas, especially the limitis plastica type, showed prominent stromal-type POSTN staining in the lamina propria mucosae and stroma of the deeper layers at all T stages (Figure 3, E and F). Cochran–Armitage trend testing revealed that stromal POSTN expression increased with progressing stage in both intestinal-type (P = 8.12 × 10⁻⁴) and diffuse-type (P = 7.54 × 10⁻⁴) gastric cancer. No staining of carcinoma cells for POSTN was noted at any point during the IHC investigation.

Stromal Myofibroblasts Express POSTN in Gastric Cancer

In gastric cancer tissue, fibroblastic stromal cells, which lie in a dense collagenous matrix, show strong stromal immunoreactivity for POSTN (Figure 3, G and H). Immunofluorescent double-staining for α-SMA and POSTN in advanced invasive cancer showed that α-SMA+ fibroblasts were embedded in a cancer stroma, which contained abundant immunoreactive POSTN. Focal colocalization of POSTN and α-SMA signals occurred (Figure 3G). Positive signal for POSTN mRNA was noted in fibroblastic stromal cells, but not in carcinoma cells (Figure 3I). These findings indicate that POSTN is expressed and secreted by stromal myofibroblasts in gastric cancer.

**Figure 2** POSTN expression in non-neoplastic gastric mucosa. A: Immunostaining of normal gastric mucosa, showing POSTN expression in the lamina propria mucosae, but not in epithelial cells. B–D: High-power magnification of the foveolar epithelial region (B), the proper gastric gland region (C), and the intestinal metaplasia area (D) reveals circumscribed POSTN immunostaining (arrowheads) and a fine network of immunoreactive POSTN (arrows). Note prominent periglandular POSTN immunoreactivity in the intestinal metaplasia area. E: Double immunofluorescence staining for α-SMA and POSTN. The merged image shows focal colocalization immunoreactivity of α-SMA and POSTN surrounding the gland. Original magnification: ×200 (A–D); ×400 (E).
Proliferative Function of POSTN for Gastric Cancer Cell via ERK Phosphorylation

We examined a possible correlation between POSTN transcripts and other genes using public domain microarray data sets from the GEO (http://www.ncbi.nlm.nih.gov/geo; accession no. GSE15460) with GSEA software. Among the gene groups associated with the POSTN/VIM-high mRNA expression ratio, the gene groups that regulate cell cycle (REACTOME and GO gene sets) and cell proliferation (GO gene sets) were highly ranked ($P < 0.0002$) (Figure 4A and Supplemental Tables S2 and S3). These results suggest that high POSTN expression per stromal cell is associated with the cell cycle and cell proliferation in gastric cancer.

The growth effect of POSTN for tumor cells was evaluated using OCUM-2MLN and OCUM-12 diffuse-type gastric cancer cell lines. Both OCUM-2MLN and OCUM-12 cells showed a significant increase in cell number with supplementation of a recombinant human POSTN (Figure 4B). For the assessment of cancer—stromal interaction in vitro, we performed three-dimensional coculture in type I collagen gel together with OCUM-2MLN cells and previously established POSTN-expressing NIH3T3 transfectants (NIH3T3POSTN). The cancer cells were significantly more numerous in NIH3T3POSTN coculture than in the coculture with control (Figure 4C).

To investigate the underlying mechanism of the capacity of POSTN to promote cell proliferation, we screened phosphorylated MAP kinase proteins expressed in OCUM-2MLN cells. Incubation of recombinant POSTN caused phosphorylation of ERK1 and ERK2, at levels approximately twofold and threefold greater, respectively, than...
those of controls (Figure 4D). No phosphorylation of the other MAP kinase proteins was detected. Induction of p-ERK by POSTN in OCUM-2MLN and OCUM-12 cells was confirmed by Western blotting (Figure 4E).

Coinoculation of OCUM-2MLN or OCUM-12 Diffuse-Type Gastric Cancer Cells Line with NIH3T3POSTN Cells in Vivo

To examine the role of stromal POSTN in gastric tumor growth in vivo, OCUM-2MLN or OCUM-12 gastric cancer cells were coinoculated with POSTN-expressing mouse fibroblast NIH3T3 cells. OCUM-2MLN cells coinoculated with NIH3T3EGFP or NIH3T3POSTN (OCUM-2MLN–NIH3T3EGFP and OCUM-2MLN–NIH3T3POSTN) caused tumor formation in 100% of mice (6/6) in both groups. Tumors in mice injected with OCUM-12–NIH3T3POSTN cells grew faster and were larger at the end of the experiment than tumors injected with OCUM-12–NIH3T3EGFP cells (Figure 5, A and B). However, one of the six mice injected with OCUM-12–NIH3T3EGFP did not form a tumor (Figure 5B), whereas all six of the mice injected with OCUM-12–NIH3T3POSTN formed tumors. OCUM-2MLN cells coinoculated with NIH3T3POSTN also showed a tendency to form larger tumors than the control, but the difference was not significant (Figure 5A). Intriguingly, OCUM-2MLN cells formed well-vascularized tumors, whereas OCUM-12 cells showed only moderate change (Figure 5B). H&E staining revealed that coinoculation with NIH3T3POSTN led to larger tumors, with central necrosis, in the OCUM-2MLN group, compared with the OCUM-12 group (Figure 5C). Analysis of the invasive tumor lesion area, with the necrotic area excluded from the total tumor area, revealed that coinoculation with NIH3T3POSTN led to larger tumor lesion area, compared with coinoculation with NIH3T3EGFP (Figure 5D). OCUM-12–NIH3T3POSTN cells led to macroscopically larger tumors (Figure 5, A and B), and also had the same tendency to form a larger invasive lesion area as OCUM-2MLN–NIH3T3POSTN cells (P = 0.05) (Figure 5, C and D).
Orthotopic Inoculation Model of Diffuse-Type Gastric Cancer in Postn\(^{-/-}\) and Postn\(^{+/+}\) Rag2 Knockout Mice

To further examine the function of host-derived POSTN in cancer stroma, we inoculated OCUM-2MLN cells into the gastric wall of Postn\(^{-/-}\)Rag2\(^{-/-}\) double-knockout mice. The OCUM-2MLN cells formed invasive tumors in both Postn\(^{-/-}\) and wild-type (Postn\(^{+/+}\)) Rag2 knockout mice (Figure 6A). We analyzed the tumor area inside the gastric wall, excluding the necrotic area from the total tumor area (Figure 6A). The tumor area in Postn\(^{-/-}\) mice was significantly smaller than that in Postn\(^{+/+}\) mice (Figure 6B). Stromal immunostaining for POSTN was observed only in Postn\(^{+/+}\) mice (Figure 6C). In situ hybridization showed positive signals for POSTN mRNA in stromal fibroblasts in Postn\(^{+/+}\) mice, but not in Postn\(^{-/-}\) mice (Figure 6C). No expression of POSTN protein or mRNA was noted in the inoculated tumor cells. Moreover, a significantly lower fraction of Ki-67\(^+\) cancer cells was found in Postn\(^{-/-}\) mice than in Postn\(^{+/+}\) mice (Figure 6, C and D), and immunostaining for p-ERK revealed a significantly lower number of p-ERK\(^+\) cancer cells in Postn\(^{-/-}\) mice than in Postn\(^{+/+}\) mice (Figure 6, C and E). These data suggest that host-derived POSTN was produced by CAFs and promoted cancer cell proliferation via activation of ERK signaling cascades.

Discussion

Histological features distinguish the intestinal and diffuse types of gastric carcinoma.\(^2\,^3\) The diffuse type comprises poorly cohesive or signet-ring cell carcinoma lying in an abundant desmoplastic stroma. In the present study, stromal cells in gastric cancer produced POSTN. Whether POSTN in the cancer stroma is secreted by the cancer cells or CAFs has been equivocal in certain cancer types.\(^3\,^3\,^6\,^3\,^3\,^3\,^3\) Analysis of publicly available microarray data sets showed that no gastric cancer cell lines expressed POSTN mRNA except YCC11; these cells harbor a unique, nonbenign
single-nucleotide variant in RPS6KA6, which might be related to POSTN expression via activation of CREB, a known inducer of POSTN. It is particularly interesting that POSTN mRNA expression was correlated with that of VIM but not with that of keratin 18 (data not shown), suggesting that POSTN is produced by nonepithelial cells rather than by neoplastic cells in gastric cancer. Significant correlation was found between the POSTN mRNA expression level and clinical and biological behavior. When the POSTN-expressing level was standardized per stromal cell using the VIM expression level (POSTN/VIM), gene groups regulating the cell cycle and cell proliferation were ranked highly by an association of the POSTN/VIM mRNA expression ratio and other genes using GSEA software. The strategy in bioinformatics, standardizing data for each cell type, is therefore crucial in mining data sets in cancer studies, especially in the context of cancer stromal interaction, in which mixed cell populations are involved. Our results suggest that POSTN plays a role in cell proliferation in the gastric cancer microenvironment.

Interaction between epithelial cells and ECM in the microenvironment is necessary for the maintenance of normal glandular structures. This association is recognized more specifically as a stem cell and its niche; POSTN is an ECM molecule comprising the niche. IHC investigation showed that POSTN immunoreactivity frequently circumscribed the foveolar epithelium and the gastric glands in normal gastric mucosa. In addition, stromal and more pronounced peril glandular POSTN staining was noted in gastric mucosa, indicative of intestinal metaplasia under a chronic inflammatory condition. These findings resemble those of POSTN expression in other inflammatory diseases, such as asthma and atopic dermatitis. In these allergic conditions, recruited Th2 cells (in asthma) and epithelial cells (in atopic dermatitis) respectively release IL-4 and IL-13 and growth factors such as TGF-β. These factors induce POSTN overexpression by activated myofibroblasts, which in the airway disease increases goblet cells. Our IHC findings thus strongly suggest that the alteration of POSTN expression reflects the disruption of homeostasis in the lamina propria mucosae niche.

The mucosal microenvironment exhibits minimal morphological change in carcinoma in situ. In stomach, the histological definition of carcinoma in situ is still equivocated by histopathologists, but it seems readily apparent that atypical tumor glands retain the basement membrane, and that the lamina muscularis mucosae is intact in Tis and T1 intestinal-type gastric adenocarcinomas. In these early-form or noninvasive cancers, peril glandular POSTN staining was lost during progression, and the stromal overexpression of

Figure 6 Orthotopic inoculation model of diffuse-type gastric cancer in immunodeficient Postn−/− or Postn+/+ mice. A: Histological appearance of the maximum cut surface of inoculated tumor in Postn−/− mice (POSTN KO) and wild-type Postn+/+ mice (POSTN wild). Invasive lesion area, with the necrotic area excluded from the total tumor area, is outlined in black. B: Areas of respective invasive lesions inside the gastric wall were measured using NBP.view software. C: IHC for POSTN, in situ hybridization (ISH) for POSTN mRNA, and IHC for Ki-67 and p-ERK. D and E: Quantitative analysis of Ki-67+ (D) and p-ERK+ (E) tumor cells. **P < 0.01, ***P < 0.001. Scale bars: 3 mm (A); 50 μm (C).
POSTN that is found in advanced cancer was not observed. The neoplastic glands were irregular, and some periglandular POSTN stainings were lost, but the basement membrane remained apparently intact in a reactive stroma, in which there was mildly increased fibroblast cellularity. Our findings resemble those reported for Cdx2 transgenic mouse gastric carcinomas, as well as those for human colorectal carcinomas in which immunoreactive pericytals POSTN disappeared along the colorectal adenoma–carcinoma sequence. When epithelial—periglandular fibroblast interaction is disrupted in the local mucosal niche, it is possible that periglandular POSTN staining is eventually lost during progression of the early form of intestinal-type gastric cancer.

In fact, CAFs (fibroblasts in the tumor stroma) acquire a modified phenotype, which is seen in myofibroblasts in the wound healing process. The CAFs, which interact with cancer cells to expedite their progression, are identifiable by the expression of the myofibroblast marker α-SMA. In contrast to Tis and T1 intestinal-type cancers, no T2 to T4 intestinal-type or any diffuse-type gastric cancer exhibited periglandular POSTN staining, but all exhibited stromal POSTN staining. In these tumors, increased fibroblast cellularity and α-SMA+ myofibroblasts were noted in a desmoplastic cancer stroma. Although the intestinal-type carcinoma involves and extends through the lamina propria mucosae, the tumor cells are exposed to the underlying stroma, in which there is increased stromal cellularity, an inflammatory cell infiltrate, and new vascularization. Little or no basement membrane exists in the diffuse-type gastric cancer cells. The tumor cells thus induce cancer—stromal interaction directly in the lamina propria mucosae and in the deeper layers of the gastric wall. Particularly, the linitis plastica type of diffuse gastric cancer, which contains scattered tumor cells and accumulated CAFs in desmoplastic fibrous stroma, showed intense stromal POSTN staining. These staining differences might explain the clinical behavior of the two gastric cancer types, with the diffuse type exhibiting more aggressive behavior than the intestinal type.

POSTN is involved in the epithelial—mesenchymal transition (EMT), which is responsible for dissemination of primary tumor epithelial cells to the sites of metastasis. POTN has been shown to be not only an EMT marker, but also an EMT inducer. Our histological expression analyses strongly suggested that CAFs are the primary source of POSTN, which facilitates tumor cell invasion by inducing EMT and by establishing a neoplastic niche in gastric cancers. POSTN exerts its protumorigenic effect through its binding to the integrins, αvβ3, αvβ5, and α6β4, promoting the recruitment of the epidermal growth factor receptor (EGFR) and the activation of the Akt/PKB and FAK-mediated signaling pathways. We demonstrated the cancer-supportive effects of POSTN in a diffuse-type gastric cancer cell line, OCUM-2MLN, with a supplement of the recombinant POSTN or by coculture of POSTN-overexpressing fibroblasts in vitro and in vivo. Furthermore, recombinant POSTN activated ERK signaling in OCUM-2MLN and OCUM-12 cells. Using a diffuse-type gastric cancer orthotopic inoculation model, we demonstrated that host stromal cell—derived POSTN supported tumor cell growth. The tumors in Postn+/− mice were significantly smaller than those in Postn+/+ mice. More Ki-67+ and p-ERK+ tumor cells were observed in the tumors inoculated into Postn+/+ mice than in those inoculated into Postn−/− mice. POSTN increased the ability to contract a collagen matrix, which is a characteristic of myofibroblasts and CAFs. Such mechanical strains are known to trigger the release of the active form of TGF-β1 from the stiff ECM in which the inactive form of TGF-β1 is embedded. In addition, POSTN promotes incorporation of tenascin-C into the ECM to reinforce them. We therefore speculate that POSTN promotes stiffening of ECM in the cancer niche, thereby releasing the active form of TGF-β1. This release can be expected to result in an accumulation of α-SMA+ CAFs secreting POSTN and inducing EMT and cancer cell proliferation. POSTN-expressing CAFs constitute a cancer-promoting microenvironment via regulation of cancer cell proliferation, in part, by triggering ERK signaling pathway, but also via indirect modification of ECM composition and growth factor activity in the cancer niche.

In conclusion, results of detailed bioinformatic and histological investigation showed that α-SMA+ CAFs produced POSTN in a desmoplastic stroma and formed the neoplastic niche in gastric cancer. POSTN promoted cancer cell growth via a signaling pathway involving ERK. Moreover, POSTN might facilitate EMT. With further understanding of cancer microenvironments, in which complex molecular networks exist among tumor cells and other components, POSTN is potentially an attractive therapeutic intervention target for cancer therapy.

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Supplemental Data

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