Aberrant methylation of *PSD* disturbs Rac1-mediated immune responses governing neutrophil chemotaxis and apoptosis in ulcerative colitis-associated carcinogenesis

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Abstract. We previously reported that the *Pleckstrin and Sec7* domain-containing (PSD) gene is preferentially methylated in patients with ulcerative colitis (UC) who developed colorectal cancer (CRC), and is implicated in UC-associated carcinogenesis through its inhibition of apoptosis. This study aimed to determine the potential effect of PSD methylation on its downstream molecule, Ras-related C3 botulinum toxin substrate 1 (Rac1), which governs neutrophil chemotaxis and apoptosis signaling. PSD was knocked down in a normal human fibroblast cell line (HNDF) and a neutrophil-like cell line (HL-60). Both NHDF and HL-60 cells exhibited numerous filamentous-actin (F-actin) rich membrane extensions, resulting in the activation of Rac1; this activation was hampered by PSD silencing. Lipopolysaccharide, a reactive oxygen species (ROS) inducer, stimulated NHDF cells to release ROS and activated caspase-3/7 in the presence of neutrophils, which was inhibited by PSD knockdown. Migration assays demonstrated that chemotaxis of HL-60 cells was affected by PSD silencing in NHDF cells. Tissue sections from 6 UC patients with CRC and 15 UC patients without CRC were examined. To verify Rac1-mediated chemotaxis in tissue sections, we evaluated the grade of neutrophil infiltration by histological assessment and assessed F-actin and PSD expression by immunohistochemistry. Neutrophil infiltration, F-actin and PSD expression were significantly decreased in specimens from UC patients with PSD methylation compared with those without. Decreased levels of F-actin expression were observed in colorectal mucosa, as well as in infiltrating cells with PSD methylation. PSD expression was preferentially inhibited in

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colorectal mucosa by *PSD* methylation, whereas PSD expression was rarely observed in infiltrating cells, regardless of *PSD* methylation status. These data indicate that aberrant methylation of *PSD* occurs in UC-associated colorectal mucosa, enabling circumvention of Rac1-mediated immune responses governing neutrophil chemotaxis and apoptosis, and thus plays a pivotal role in the mechanisms underlying UC-associated carcinogenesis.

Introduction

Ulcerative colitis (UC) is a chronic inflammatory bowel disease (IBD), the etiology of which is not completely understood. Patients with UC face an increased risk of developing colorectal cancer (CRC) (1,2). Recent advances in IBD research have provided genetic insights into its pathogenesis. The innate and adaptive immune system (3-5), autophagy pathway (6-8), and epithelial barrier (9,10) participate in fighting pathogens involved in IBD. Interaction between host and pathogens leads to persistent or severe inflammation, whereas insufficient interaction may result in failure to prevent cancer development.

A genome-wide association study previously identified genetic variants in Disks large homolog 5 (DLG5) gene associated with IBD (10). DLG5 regulates cell shape, polarity (11) and cell-cell contact (12), disruption of which interferes with the epithelial barrier function in the colon. In our previous study, genome-wide analysis of methylation alterations using a methylation-sensitive representational difference analysis identified the Pleckstrin and Sec7 domain-containing (PSD) gene (13), which has similar roles to DLG5, such as coordination of cell shape and polarity PSD was more frequently methylated in both UC-associated colorectal cancer tissues (71.4%) and matched normal epithelia (57.1%) than in non-neoplastic UC epithelia (27.3%) and sporadic colorectal cancer tissue (18.8%). In addition, silencing of PSD inhibited apoptosis in a fibroblast cell line and in tissue specimens from UC patients harboring PSD methylation. These findings led us to address the potential roles of PSD methylation in the mechanisms underlying UC-associated carcinogenesis.

PSD regulates Ras-related C3 botulinum toxin substrate 1 (Rac1), a Rho GTPase. Rac1 is implicated in the regulation of neutrophil functions in response to inflammatory signals, including actin remodeling, chemotaxis and production of NADPH oxidase. Rac1 is reported to induce apoptosis in response to UV light (14) and other damaging agents such as Fas (15) and TNF- α (16). These findings indirectly support our data that previously reported PSD silencing and methylation inhibited apoptosis *in vitro* and in tissue specimens, respectively. In this study, we elucidated the effect of PSD methylation on Rac1, which governs neutrophil chemotaxis and apoptosis signaling, in a normal human fibroblast cell line (HNDF) and a human promyelocytic leukemia cell line (HL-60), which look and behave like neutrophils (17-19), and in tissue sections from UC patients with and without CRC.

Materials and methods

Patients and tissues. Six samples of UC-associated colorectal cancer tissue (UCT) with matched normal epithelial tissue (UCN), and 15 samples of non-neoplastic UC epithelial tissue (UCI) were obtained from patients who had undergone surgery at Jichi Medical University Saitama Medical Center and Jichi Medical University Hospital between November 2000 and September 2006. Matched normal epithelia were taken from lesions harboring colitis adjacent to tumors. This study was approved by the Ethics Committee of Jichi Medical University, and informed consent was obtained from each participant.

Cell lines. A human skin fibroblast cell line, NHDF, was obtained from Kurabo (Osaka, Japan) and was maintained in medium 106S supplemented with low-serum growth supplement. A human promyelocytic leukemia cell line, HL-60, which orient their polarity in response to a gradient and migrate towards the stimulus (17-19), were obtained from the Japanese Collection of Research Bioresources (JCRB, Osaka, Japan). HL-60 cells were maintained in RPMI-1640 medium supplemented with 10% heat-inactivated fetal bovine serum.

Quantitative reverse transcription-PCR. Tissue specimens were immediately added to RNA later (Ambion, Austin, TX, USA) and stored at -80°C until DNA or RNA extraction. Total RNA was immediately treated with DNase I (Invitrogen, Carisbad, CA, USA) and reverse-transcribed using a Superscript II reverse transcriptase kit (Invitrogen) to prepare first-strand cDNA. The primer sequences for PSD were 5'-CCATAGACGAGGAGGAGCTG-3' (forward) and 5'-TCTTCCTGCAGTCAGGGTCT-3' (reverse). Thermal cycling conditions were 42°C for 60 min (cDNA synthesis), 95°C for 10 sec (hot start), and then 40 cycles of 95°C for 5 sec, 58°C for 10 sec, and 72°C for 30 sec. The expression level of PSD was determined using the fluorescence intensity measurements from the ABI 7900HT Real-Time PCR System Data Analysis Software. A GAPDH fragment was amplified as an internal control.

Knockdown of PSD in HNDF and HL-60 cells. PSD-specific siRNA (siPSD) was purchased from Invitrogen. RNA oligonucleotides were resuspended in 10 μ M Tris-HCl, pH 8.0, 20 mM NaCl, and 1 mM EDTA to make a 20 μ M siRNA solution. The final siRNA concentration was 30 nM in Opti-MEM I

without serum. HNDF and HL-60 cells were cultured in dishes at 30-50% confluency without antibiotics, and transfection was performed with Lipofectamine 2000 (Invitrogen) according to the manufacturer's instructions. The Block-iTTM Fluorescent Oligo (Invitrogen), a fluorescently labeled double-stranded RNA duplex with the same length, charge, and configuration, was used for the assessment of transfection efficiency, and Scrambled StealthTM RNA molecule was used as the control siRNA (siControl). Cells were incubated for 48 h after transfection at 37°C in a CO₂ incubator, and used for subsequent experiments.

Actin cytoskeleton analysis. siPSD-treated and siControl-treated cells were stained with fluorescent rhodamine-labeled phalloidin using a F-actin Visualization BiochemKit™ (Cytoskelton, Denver, USA) according to the manufacturer's instructions. Cells were stimulated with EGF (10 ng/ml for 1, 2 and 30 min) or calpeptin (0.1 mg/ml for 30 min) after treatment with siPSD or siControl for 48 h and subsequently stained with rhodamine-labeled phalloidin. Hoechst 33342 was used to observe nuclear morphology. Signals were observed by a fluorescence microscope (Fluoview FV500; Olympus) with excitation at 535 nm and emission at 585 nm for the detection of rhodamine, and with excitation at 365 nm and emission at 480 nm for the detection of Hoechst 33342. EGF was purchased from Wako (Tokyo, Japan) and Hoechst 33342 was included in the Total ROS/Superoxide Detection kit™ (Enzo, PA, USA).

Measurement of Rac1 activity. The activity of Rac1 was measured by G-LISATM Rac Activation Assay Biochem kitTM (Cytoskelton) according to the manufacturer's instructions. Cells were stimulated with EGF (10 ng/ml for 1, 2 and 30 min) or calpeptin (0.1 mg/ml for 30 min) after treatment with siPSD or siControl for 48 h and then collected in 100 μ l of ice-cold lysis buffer provided by G-LISA. Lysates were centrifuged to remove cellular debris. From each supernatant, 10 μ l was removed to measure protein content using the Precision RedTM Advanced Protein Assay Reagent included in the G-LISA Rac Activation Assay Biochem kitTM, and 50 μ l was used for the G-LISA Rac activation assay, followed by dilution with 50 μ l of cold binding buffer. Lysate (50 µl) was added into respective wells in the Raclbinding plate for duplicate assays, and then the plate was put on a cold orbital microplate shaker at 4°C for exactly 30 min. Next, the wells were washed twice with wash buffer. Antigen (200 μ l) presenting buffer was immediately added into each well and the plate was incubated at room temperature for 2 min. Wells were then washed three times with wash buffer. Next, 50 µl of diluted anti-Rac1 primary antibody was added to each well and the plate was placed on an orbital microplate shaker at room temperature for 45 min. Next, 50 μ l of diluted anti-Rac1 secondary antibody was added to each well and the plate was incubated for 45 min, followed by incubation with 50 µl of HRP detection reagent for 20 min. Immediately after addition of 50 μ l of HRP Stop Buffer, the absorbance was read using a microplate spectrophotometer.

Detection of reactive oxygen species and active caspase-3/7. Reactive oxygen species (ROS) and active caspase-3/7 were detected by Total ROS/Superoxide Detection kit (Enzo) and CaspaTagTM Caspase-3/7 Assay In Situ Assay Kit (Chemicon), respectively, according to the manufacturer's instructions.

Table I. Clinicopathological characteristics of tumor specimens from UC patients with and without colorectal cancer.

	UCT	UCN	UCI		
Number	N=6	N=6	N=15		
Average age (years)	54.8±17.1	54.8±17.1	40.8±14.0		
Gender					
Male	4	4	9		
Female	2	2	6		
Disease duration (years)	14.8 ± 7.0	14.8 ± 7.0	8.0 ± 5.0^{a}		
PSD methylation (%)	71.4%	57.1%	27.3%		

^aSignificantly short duration in UCN (= UCT) samples than UCI samples (P<0.05). UCT, UC-associated colorectal cancer tissues; UCN, matched normal epithelia; UCI, non-neoplastic, UC epithelia.

After treatment with si*PSD* or siControl for 48 h, NHDF cells were exposed to lipopolysaccharide (20 ng/ml; Wako) and were co-cultured with HL-60 cells for an additional 48-h period. Then, HL-60 cells were removed and NHDF cells were subjected to the ROS detection assay or the caspase-3/7 activity assay. Fluorescence signals were subsequently detected by a fluorescence microscope (Fluoview FV500; Olympus), with excitation at 490 nm and emission at 525 nm for the detection of ROS, and with excitation at 550 nm and emission at 580 nm for the detection of caspase-3/7, respectively. Average number of cells inducing ROS or expressing caspase-3/7 in three random fields was calculated.

Migration assay. Neutrophil chemotaxis in response to inflammation in vitro was assessed by migration assays using a BD Falcon companion plate and cell culture insert (Becton-Dickinson) with a 3-μm pore size. After NHDF cells were treated with siPSD or siControl for 48 h in the bottom wells (24-well companion plate, Becton-Dickinson), they were exposed to LPS and cultured with HL-60 cells seeded in culture inserts for an additional 48 h. Then, non-migrating cells were removed from the insert membranes by cotton swabs. The membrane was mounted onto a slide glass and

the nuclei of migrated cells were then stained with Hoechst 33342. The number of migrated cells was counted in three random fields using an inverted microscope.

Histological grades of neutrophil infiltration in tissue sections. Histological grades of neutrophil infiltration were determined using a scoring system as previously described (20): 0, normal (no inflammatory cells); 1, mild active; 2, moderate active (with cryptitis). The average grading of three regions of the colorectum (rectum, descending colon and ascending colon) were calculated where samples were available.

Detection of filamentous actin and PSD in tissue specimens. Detection of filamentous actin (F-actin) and PSD in paraffin-embedded tissue sections was performed by immunohistochemical analysis using the I-View DAB Universal kit (Roche, Rotkreuz, Switzerland). Sections were de-waxed in xylene and rehydrated with distilled water before analysis, then treated with a heat-induced epitope retrieval technique using an EDTA buffer at pH 9.0, and blocked for endogenous peroxidase activity before addition of the primary antibody. NH3 and ab5962 (Abcam, Tokyo, Japan) were used as primary antibodies for F-actin and PSD, respectively. Incubation with primary antibody was performed overnight at 37°C. Cells displaying slight staining of the cytoplasm were determined to be positive. A grading system was applied to the assessment of accumulation of F-actin and PSD expression in tissue sections. The F-actin and PSD index were calculated based on percentage of staining cell, with 0, +1, and +2 when <-5% of cells, 5-20% cells, and >20% cells demonstrated cytoplasm reactivity, respectively. The average grading of three regions of the colorectum (rectum, descending colon and ascending colon) was calculated when samples were available.

Statistical analysis. Values are shown as mean ± SE. Statistical differences between variables were determined by use of an unpaired t-test or an analysis of variance, as appropriate. Simple regression coefficient analysis was used to examine associations between two categorical variables. Values of P<0.05 were considered significant.

Table II. Clinicopathological characteristics of tumor specimens from patients with UC-associated colorectal cancer.

Group	PSD	Age	Gender	Duration	Onset	Loc	Type	Dukes	INF	Ope
UCT1	M	77	M	13	64	R	Well	A	1	Total
UCT2	M	40	M	8	32	A	Muc	A	0	Total
UCT3	M	64	F	9	55	D	Well	A	1	Total
UCT4	U	35	M	15	20	D	Poor	D	0	Partial
UCT5	M	68	F	24	44	R	Well	В	0	Total
UCT6	U	45	M	25	20	D	Well	В	0	Total

UCT, UC-associated colorectal cancer tissues; UCN, matched normal epithelia; UCI, non-neoplastic UC epithelia; U in PSD, unmethylated; M in PSD, methylated; duration, disease duration (years); M in gender, male; F in gender, female; onset, age of onset (years); loc, location of carcinoma; A, D and R in loc, ascending colon, descending colon, and rectum, respectively; type, histological findings of carcinoma; well, poor, and muc in loc, well differentiated adenocarcinoma, poorly differentiated adenocarcinoma, and mucinous adenocarcinoma, respectively; Dukes, Dukes' classification; INF, infiltration of neutrophils; ope, operation; total, partial, and right in ope, total colectomy, partial resection of the colon, and right hemi-colectomy, respectively.

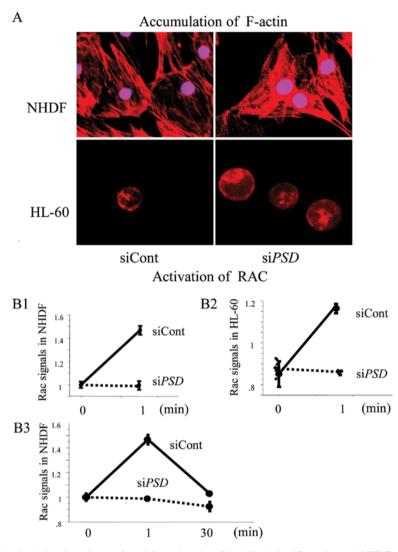
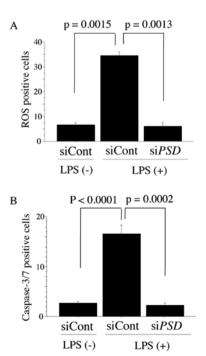


Figure 1. (A) Rhodamine-labeled actin (red) polymerizes preferentially at the edge of lamellipods in siControl-treated NHDF and HL-60 cells (left) after stimulation with EGF, whereas rhodamine-labeled actin remained inside the plasma membrane in siPSD-treated NHDF and HL-60 cells (right). Nuclei were stained with Hoechst 33342 (pink). (B) Rac1 signaling before and after stimulation with EGF in siControl-treated and siPSD-treated NHDF cells (B1) and in siControl-treated and siPSD-treated HL-60 cells (B2). (B3) Rac1 signaling declined to basal levels by 30 min after stimulation.

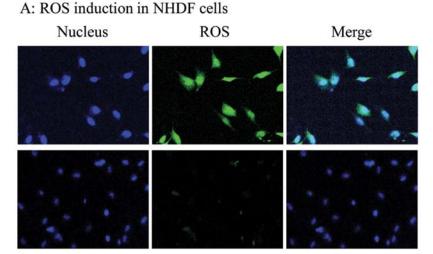


Results

Clinicopathological features. The clinicopathological features of patients recruited for this study are shown in Tables I and II. There was no significant difference in the average age. The disease duration of UCI patients $(8.0\pm5.0 \text{ years})$ was significantly shorter than that of the UCT patients $(14.8\pm7.0 \text{ years})$, P<0.05). Aberrant methylation of *PSD* was observed in 4 of 6 UCT patients (71.4%), 4 of 6 UCN patients (57.1%) and 5 of 15 UCI patients (27.3%), respectively, as previously reported.

Knockdown of PSD inhibited membrane ruffling and reduced Rac1 activity in NHDF and HL-60 cells. To determine the

Figure 2. Average number of NHDF cells inducing ROS (A) and expressing active caspase-3/7 (B). After treatment with siPSD or siControl for 48 h, NHDF cells were exposed to lipopolysaccharide (LPS; 20 ng/ml) for 48 h in the presence of HL-60 cells. Then, HL-60 cells were removed and NHDF cells were subjected to ROS detection assays or caspase-3/7 activity assays.



B: Caspase-3/7 activity in NHDF cells

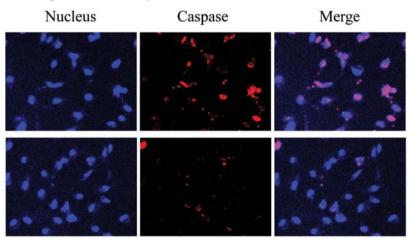


Figure 3. (A) Cells inducing reactive oxygen species (ROS) in siControl-treated (top panel) and siPSD-treated NHDF cells (bottom panel). Nuclear morphology of cells stained with Hoechst 33342 (blue, left), cells inducing ROS stained with Total ROS/Superoxide Detection kit Reagent (green, middle), and merged staining with both reagents (white green, bottom). (B) Cells expressing active caspase-3/7 in siControl-treated (top panel) and siPSD-treated NHDF cells (bottom panel). Nuclear morphology of cells stained with Hoechst 33342 (blue, left), cells expressing active caspase-3/7 stained with CaspaTag Reagent (red, middle), and merged staining with both reagents (pink, right).

inhibitory effect of PSD silencing on the activation of Rac1, NHDF and HL-60 cells were treated with PSD-specific siRNA (siPSD) or siControl. Transfection efficiency was 89.1% in NHDF cells and 73.0% in HL-60 cells, which reduced the mRNA levels of PSD by 90.1% in siPSD-transfected NHDF cells and 61.3% in siPSD-transfected HL-60 cells, respectively. PSD promotes numerous F-actin-rich membrane extensions (21), which leads to the activation of Rac1. The accumulation of F-actin by stimulation with EGF was visualized by staining with fluorescent rhodamine-labeled phalloidin. EGF-stimulated membrane ruffling was observed in both siControl-treated NHDF and HL-60 cells, whereas no morphological changes of the membrane were found in either siPSD treated-NHDF or HL-60 cells (Fig. 1A). Likewise, Rac1 activity was increased in both siControl-treated NHDF and HL-60 cells after stimulation with EGF (Fig. 1B1 and B2), whereas activation was hindered in either siPSD-treated NHDF or HL-60 cells. Stimulation with calpeptin for 30 min also resulted in increased levels of Rac1 in both siControl-treated NHDF and HL-60 cells (data not shown). Rac1 activity declined to basal levels within half an hour after stimulation with EGF (Fig. 1B3), which was consistent with the results reported by Kurokawa *et al* (22).

Knockdown of PSD in NHDF cells prevented induction of ROS and ROS-induced caspase-3/7 activity in the presence of HL-60 cells. Our previous study showed that NHDF cells were stimulated to release reactive oxygen species (ROS) by pyocyanin-harboring redox reactions, while lipopolysaccharide (LPS), which mediates the activation of NADPH oxidation in neutrophils, did not stimulate NHDF cells (13). In the present study, we attempted to elucidate whether LPS stimulates NHDF cells to release ROS when co-cultured with HL-60 cells, and if ROS induction is inhibited by PSD silencing in NHDF cells because they co-exist and interact in the body. LPS stimulated siControl-treated NHDF cells to release ROS in the presence of HL-60 cells, whereas LPS did not stimulate siPSD-treated

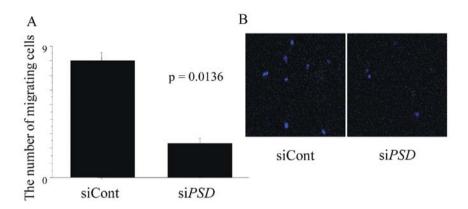


Figure 4. (A) Average number of NHDF cells migrating through membrane as detected by migration assay. After NHDF cells were treated with si*PSD* or siControl for 48 h in the bottom wells (24-well companion plate), cells were exposed to LPS and cultured with HL-60 cells in culture inserts for additional 48 h. The number of migrated HL-60 cells was counted in three random fields using an inverted microscope. (B) Microscopic images of migrated NHDF cells.

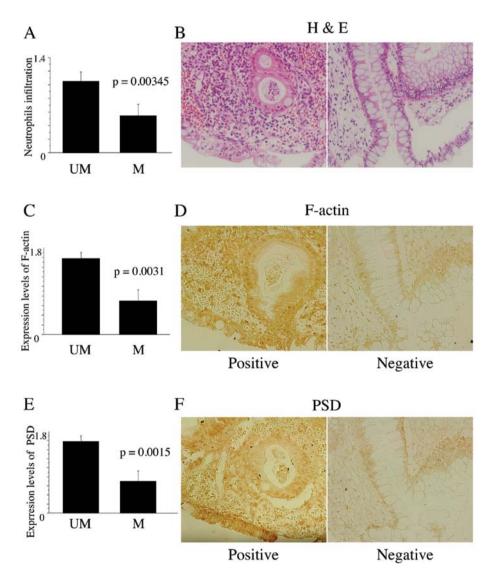


Figure 5. (A) Histological grades of neutrophil infiltration in tissue specimens from UC patients with (M) and without *PSD* methylation (UM). For evaluation, histological grades of inflammation were determined using a scoring system based on epithelial neutrophils as previously described (19): 0, normal (no inflammatory cells); 1, mild active; 2, moderate active (with cryptitis). The average grading from three regions of the colorectum was calculated. (B) Representative neutrophil infiltration in tissue specimens from UC patients without (left) and with *PSD* methylation (right). (C) The F-actin index in tissue specimens from UC patients with (M) and without *PSD* methylation (UM). For evaluation, three grades were determined as 0, +1, and +2 when <5% of cells, 5-20% cells, and >20% cells demonstrated cytoplasm reactivity, respectively. The average grading from three regions of the colorectum was calculated. (D) Representative positive (right) and negative cells for F-actin (left) in tissue specimens from UC patients without and with *PSD* methylation, respectively. (E) The *PSD* index in tissue specimens from UC patients with (M) and without *PSD* methylation, three grades were determined as 0, +1, and +2 when <5% of cells, 5-20% cells, and >20% cells demonstrated cytoplasm reactivity, respectively. The average grading from three regions of the colorectum was calculated. (F) Representative positive (right) and negative cells for *PSD* in tissue specimens from UC patients without and with *PSD* methylation, respectively.

NHDF cells to release ROS (Fig. 2A; 16.67±3.06 in siControl vs. 2.67±10.58 in siPSD, P=0.0013). Representative data for siControl-treated and siPSD-treated NHDF cells are shown in Fig. 3A. Next, we accessed ROS-mediated caspase-3/7 activation. Activation of caspase-3/7 was observed in siControl-treated NHDF cells, but was not noted in siPSD-treated NHDF cells (Fig. 2B; 34.67±2.51 in siControl vs. 6.00±2.65 in siPSD, P=0.0002). Representative caspase-3/7 positive cells detected by the CaspaTag Caspase-3/7 Assay In Situ Assay Kit (Chemicon) and cell nucleus stained with Hoechst 33342 in NHDF cells treated with siPSD or siControl are shown in Fig. 3B.

Chemotaxis of HL-60 cells was disturbed by PSD silencing in NHDF cells. NHDF cells exhibited induction of LPS-mediated ROS and activation of caspase-3/7 when cultured with HL-60 cells, which was not observed in NHDF cells cultured alone or siPSD-treated NHDF cells co-cultured with HL-60 cells, suggesting that the interaction between activated NHDF cells and HL-60 cells was involved in this process. To investigate whether LPS-mediated PSD activation in NHDF cells affects the chemotaxis of HL-60 cells, migration assays were performed. Fig. 4 shows microscopy images of cells in a migration assay for each experimental setting. The number of migrated HL-60 cells co-cultured with siControl-treated NHDF cells was 4.6 times greater than HL-60 cells co-cultured with siPSD-treated HL-60 cells (Fig. 4A; 2.667±0.882 in siControl vs. 12.333±1.453 in siPSD, P=0.0047). Microscopic images of migrated NHDF cells are shown in Fig. 4B.

The level of neutrophil infiltration was significantly decreased in specimens from patients with PSD methylation. To verify if PSD methylation affected neutrophil chemotaxis in tissue specimens, the infiltration of neutrophils was evaluated by histological assessment. Neutrophil infiltration was significantly decreased in specimens from patients with PSD methylation than in those without (Fig. 5A; 0.51±0.55 with vs. 1.01±0.55 without, P=0.0166). Regardless of cancer status, the level of neutrophil infiltration was significantly decreased in both UCT and UCN when compared with UCI (0.29±0.49 in UCT vs. 1.1±0.55 in UCI, P=0.0018; 0.39±0.32 in UCN vs. 1.1±0.55 in UCI, P=0.0034). Representative data from tissue specimens from UC patients without and with PSD methylation are shown in Fig. 5B.

Accumulation of F-actin was decreased in tissue specimens from UC patients with PSD methylation. To determine the distribution of PSD promoting accumulation of F-actin in tissue specimens, immunohistochemistry was performed. F-actin levels were significantly decreased in specimens from UC patients with PSD methylation compared to those without (Fig. 5C; 0.69±0.86 with vs. 1.57±0.51 without, P=0.0031), suggesting that accumulation of F-actin was inhibited by PSD methylation. This change was seen not only in colorectal mucosa but also in infiltrating cells. Representative F-actin-positive and -negative cells in tissue specimens are shown in Fig. 5D.

PSD expression was decreased in tissue specimens from UC patients with PSD methylation, and was preferentially suppressed in epithelial cells rather than neutrophils. To clarify the distribution of PSD expression in tissue specimens, immunohistochemical analysis was performed. The expression

level of *PSD* was significantly decreased in specimens from UC patients with *PSD* methylation compared with those without (Fig. 5E; 0.727±0.141 with vs. 1.462±0.144 without, P=0.0015), and was significantly correlated with the methylation status of *PSD*. In addition, immunohistochemical analysis revealed that PSD expression was inhibited in colorectal mucosa with methylated *PSD*, whereas PSD was rarely expressed in infiltrating cells regardless of *PSD* methylation status. These results indicated that aberrant methylation of *PSD* in UC-associated colorectal mucosa circumvented neutrophil chemotaxis, which resulted in the disturbance of neutrophil-regulated apoptosis. Representative *PSD*-positive and -negative cells are shown in Fig. 5F.

Discussion

The present study demonstrated the crucial role of *PSD* silencing in NHDF cells through its inhibitory effect on Rac1 activation, which disturbed membrane ruffling and chemotaxis of HL-60 cells and consequently hampered the apoptotic machinery. In tissue specimens from UC patients, aberrant methylation of *PSD* interfered with actin rearrangement, neutrophil infiltration and apoptosis. Taken together, these data indicate that aberrant methylation of *PSD* in UC-associated colorectal mucosa circumvents Rac1-mediated immune responses, neutrophil chemotaxis and the apoptotic machinery, and thus likely plays a pivotal role in the mechanisms underlying UC-associated carcinogenesis.

The small GTPases have been implicated in diverse biological functions such as cytoskeleton rearrangement, cell growth, transformation, cell motility, migration, metastasis, and response to stress. One of these GTPases, Rac1, is reported to play a crucial role in inducing apoptosis in response to several stimuli such as UV light (14), and other damaging agents such as Fas (15) and TNF- α (16). These findings strongly support our data showing that Rac1-mediated apoptosis was inhibited by PSD silencing in siPSD-treated cells and was decreased in tissue specimens from UC patients harboring PSD methylation. Some reports, however, have presented conflicting results (23-25), raising the possibility that Rac1 has a complex role, and is involved in both inhibition and stimulation of apoptosis. This dual role in cell proliferation and apoptosis has been observed for other Rho proteins such as oncogenic vav (16) and R-ras (26,27). Esteve et al, explain that a dual role of Rho proteins in the regulation of cell apoptosis comes from the evidence that Rho proteins induce activation of the pathway leading to the JNK/SPAK cascade in several cell systems as well as nuclear factor kB (16). The fate of cells as determined by these molecules still remains to be explored.

Rho GTPases coordinate many cellular responses, often by regulating formation of different actin assemblies. *PSD* promotes numerous F-actin-rich membrane extensions (21), which leads to the activation of Rac1. Our results revealed that the accumulation of F-actin was significantly decreased in si*PSD*-treated NHDF and HL-60 cells, and in tissue specimens from UC patients harboring methylated *PSD*, which indicated that *PSD* methylation abolished F-actin-induced membrane extensions and consequently inactivated Rac1. Our findings concur with the report by Srinivasan *et al*, that a dominant-negative Rac1 mutant inhibits chemoattachment-stimulated accumulation of F-actin and polarization (28). A significant correlation between a decreased level of apop-

tosis and the accumulation of F-actin in specimens from UC patients with *PSD* methylation indicated that *PSD* methylation inactivated Rac1 and consequently abrogated Rac1-mediated apoptosis.

In considering alternative participants in the immune system underlying inflammatory bowel disease, neutrophils should be taken into consideration. Neutrophils are recruited from the circulation to take part in the defense against infectious agents but they may also cause tissue destruction in the host by secretion of toxic granule proteins and ROS (29). In this machinery, Rac1 also plays a crucial role in neutrophil migration and oxygen radical generation through NADPH oxidase. Glogauer et al demonstrated that infiltration of neutrophils was decreased in Rac1-null mice compared to wild-type mice (30). Delayed accumulation of neutrophils was also observed in Rac1-null mice. In addition, Srinivasan et al (28) demonstrated that a dominantnegative Rac1 mutant inhibited chemoattachment-stimulation. In accordance with these observations, our results revealed that the accumulation of neutrophils was significantly decreased in specimens from patients with PSD methylation. The inactivation of Rac1 resulting from PSD methylation may lead to delayed recruitment of neutrophils or disturbance of neutrophil chemotaxis. In the present study, we demonstrated that PSD-promoted accumulation of F-actin was decreased in colorectal mucosa, as well as infiltrating cells from ulcerative colitis patients with PSD methylation. PSD expression was preferentially inhibited in colorectal mucosa by PSD methylation, whereas PSD expression was rarely observed in infiltrating cells regardless of PSD methylation status. Considering the short half-life of neutrophils, it is unlikely that they would be methylated, indicating that colorectal mucosa could be targeted for aberrant methylation under circumstances of chronic inflammation such as ulcerative colitis, leading to neutrophil dysfunction.

Oxidative stress occurs in connection with inflammatory bowel disease. Neutrophils participate in this mechanism to release ROS, leading to protein damage, lipid peroxidation, and DNA damage. This results in genetic and epigenetic alterations, which pave the way for increasing grades of dysplasia and carcinoma. Furthermore, ROS have important functions in intracellular signaling, partly with anti-carcinogenic effects such as triggering apoptosis (31). Inadequate interaction between neutrophils and colorectal mucosa by *PSD* methylation could result in the disruption of the immune system in inflammatory bowel disease, which would be implicated in the mechanisms underlying UC-associated carcinogenesis. However, apoptotic pathways mediated by *PSD* signaling still remain to be considered as direct effectors on UC-associated colorectal mucosa to induce apoptosis.

Although the number of samples included in this study was limited and further investigations are required to draw definitive conclusions, the present study demonstrates that the interaction between colorectal mucosa and neutrophils that governs neutrophil chemotaxis and apoptosis is disturbed by aberrant methylation of *PSD*, which may suppress the host immune system and result in UC-associated carcinogenesis.

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