

Research Paper

Increased Expression of Pattern Recognition Receptors and Nitric Oxide Synthase in Patients with Endometriosis

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Abstract

Objective: Endometriosis is characterized by repeated inflammatory changes and serious adhesions, inducing innate and adaptive immune responses within the abdominal cavity. To assess these immune responses, we evaluated the levels of expression of Toll-like receptors (TLR)-1, -2, -4, -5, and -9; nucleotide-binding oligomerization domains (NOD)-1 and -2; interleukins-1 β , -6, -8, -10, and -12; interferon- γ ; tumor necrosis factor- α ; inducible nitric oxide synthase (iNOS) and endothelial NOS (eNOS); and immunoglobulins (Igs) in patients with endometriosis.

Methods: The levels of TLRs, NODs, cytokines, and NOS mRNAs in peritoneal effusions were assessed by real time reverse transcription-polymerase chain reaction; and IgG, IgA and IgM concentrations were measured by enzyme-linked immunosorbent assays (ELISA) in 40 patients with and 40 without endometriosis. Findings from the two groups were compared.

Results: We observed expression of all pattern recognition receptors (PRRs), cytokines, and NOS mRNAs and Igs in the effusion fluid of patients with and without endometriosis. The levels of TLR-2 and -9; NOD-1 and -2; iNOS and eNOS mRNAs and CA 125 were significantly higher in the endometriosis than in the non-endometriosis group ($p < 0.05$ each). Moreover, PRR, cytokine, and NOS expression showed significant correlations ($p < 0.05$).

Conclusions: PRRs, cytokines, and NOS, which act cooperatively in the innate immune response, are closely associated with endometriosis. Increased expression of TLR-2, TLR -9, NOD-1, NOD-2, and NOS mRNA in peritoneal fluid may be associated with endometriosis.

Key words: Endometriosis - Pattern Recognition Receptors, Toll-like receptors - Nitric oxide synthase, cytokine, Immunoglobulin

Introduction

Endometriosis is a disease related to chronic pelvic inflammation and associated pelvic pain that may be accompanied by, for example, dysmenorrhea, dyspareunia, infertility and menstrual irregularities. Although the pathogenesis of endometriosis has not

been clearly defined, abnormal levels of immune system cells, including macrophages, dendritic cells and natural killer cells, have been observed within the abdominal cavities of patients with endometriosis. These cells, however, are unable to detect and elimi-

nate ectopic endometrial cells. Moreover, immune system cells in the abdominal cavity were found to be dysfunctional (1). Complicated reactions may occur within the abdominal cavity, due to endometriosis-induced secretion or reactions of cytokines, chemokines, nitric oxide, immunoglobulins, and immune cells. Triggered immune reactions signify the host recognition of infectious agents, but, if pathogens are not swiftly recognized, immune reactions necessary to fight infections do not occur. Thus, recognition of infectious agents is regarded as one of the crucial processes in the host immune system.

Pattern recognition receptors (PRRs) recognize unique molecular characteristics of pathogens and induce appropriate immune responses. PRRs respond to distinct molecular motifs of pathogens, their sites of expression in microorganisms and signaling (2). Among the various types of PRRs in humans are Toll-like receptors (TLRs) and cytoplasmic nucleotide-binding oligomerization domain (NOD)-like receptors (NLRs). The principal TLRs involved in endometriosis are TLR-4, present on cell surfaces, and TLR-3, present on lysosome/endosome membranes (3).

Increased concentrations of lipopolysaccharide (LPS) in the peritoneal cavity or endometriotic fluid can trigger pelvic inflammation and TLR-mediated endometriosis (4). TLRs are also activated by endogenous ligands, including heat shock protein, S100, fibronectin, fatty acid, oxidized LDL, neutrophil elastase and hyaluronan. TLRs stimulated by LPS or endogenous ligands and oxidative stress activate NF- κ B, upregulating cytokine secretion as pro-inflammatory cascades (3, 5). As this process proceeds, the adaptive immune system becomes involved, along with the innate immune system. Although several studies have assessed the expression of PRRs, cytokines, NOS, and immunoglobulins separately in patients with endometriosis, no study to date has analyzed the relationships of these molecules by measuring all of them at the same time. We therefore analyzed the expression of PRRs, which are involved in inflammatory and immune responses; NOS, which are involved in the female reproductive process; and Igs, which are involved in the adaptive immune response, in patients with and without endometriosis. We also analyzed the relationships among these molecules in the peritoneal cavities of patients with and without endometriosis according to patient age, parity and serum CA125 concentration.

Subjects & Methods

Subjects

Intraperitoneal fluid samples were obtained

from 80 patients who visited the Department of Obstetrics and Gynecology at our hospital between June 2011 and July 2012. Of these, 40 were positive for endometriosis on laparoscopy, a finding confirmed during histological examination after surgery. All patients enrolled in this study were in the proliferative stage. Of the patients with endometriosis, 27 had stage 1, 7 had stage 2, and 6 had stage 3 endometriosis; none had stage 4. The remaining 40 patients had benign tumors, with no evidence of endometriosis, including 26 patients with myomas, 5 with dermoid cysts, 2 with hydrosalpinx, 3 with paratubal cysts, 1 with a serous borderline ovarian tumor, and 3 with non-pathologic specificities. During laparoscopy, peritoneal fluid was collected aseptically from the Douglas pouch, taking care to avoid bleeding. Patients were excluded if they had inflammatory diseases or hormone producing conditions, including pregnancy; if peritoneal fluid was contaminated with blood; or if no peritoneal fluid could be obtained. The samples were centrifuged at 1800 x g for 10 min; the supernatants were stored at -80°C in 1.5 ml aliquots; and the cell pellets were stored at -80°C in 1.5 ml aliquots after adding RNase inhibitor. The study protocol was approved by the institutional review boards (IRBs) of Vincent's Hospital, The Catholic University of Korea and Kyung Hee University Hospital, and informed consent was obtained from each patient (VC11TISI0091, KMC IRB 1236-02).

Real-time reverse transcription-polymerase chain reactions

Total RNA was extracted from peritoneal fluid using RNA-Bee solution kits (Tel-Test, Friendswood, TX, USA), as described by the manufacturer. First-strand cDNA was synthesized by reverse transcription in 20 μ l of a reaction mixture containing 1 μ g of RNA, 1x reaction buffer, 1 mM dNTP, 5 μ M random primers, 20 units RNase inhibitor, and 20 units AMV reverse transcriptase (Promega, Madison, WI, USA). The reaction mixture was incubated at 42°C for 1 h, and the reaction was terminated by heating at 95°C for 5 min. Primers specific for Toll-like receptors (TLR)-1, -2, -4, -5, 6, and -9; nucleotide-binding oligomerization domains (NOD)-1 and -2; interleukins (IL)-1 β , -6, -8, -10, and -12; interferon- γ ; tumor necrosis factor- α ; inducible nitric oxide synthase (NOS); and endothelial NOS are shown in Table 1. Real-time polymerase chain reactions (PCR) were performed using a Chromo4 Detector real-time system (Bio-Rad, Hercules, CA, USA) and the SsoFast EvaGreen supermix (Bio-Rad). Each 20- μ l PCR reaction contained 2 μ l of cDNA, 10 μ l SsoFast EvaGreen supermix, 2 μ l of each primer and 6 μ l PCR grade water. The amplification protocols consisted of an initial denaturation

at 95°C for 30 sec, followed by 45 cycles of denaturation at 95°C for 5 sec and annealing and extension at 55°C to 64°C for 12 sec. The point at which expression of each cDNA crossed that of β -actin cDNA was applied to the formula, $2^{-(\text{target gene} - \beta \text{ actin})}$, and the relative amounts of each cDNA were quantitated.

Enzyme-Linked Immunosorbent Assay (ELISA)

IgG, IgA, and IgM concentrations in the supernatants of peritoneal fluid were measured by ELISA. Briefly, 50 μ l 1:100 goat anti-human IgG, IgA, and IgM in coating buffer (1.59 g Na₂CO₃+2.93 g NaHCO₃+5% NaN₃, pH 9.6) were placed into each well of a 96 well plate and incubated overnight at 4°C. The wells were washed 6 times and incubated with blocking antibody. To each well was added 50 μ l of sample, and the plates were incubated at room temperature for 3 hours. The wells were washed 6 times, followed by incubation at room temperature with purified goat

anti-human IgG, IgA, or IgM conjugated to horseradish peroxidase in PBS/Tween/BSA solution. The plates were again washed 6 times and incubated with substrate solution (2, 2'-AZINO-Bis), and the optical absorbance of each well was measured at 450 nm (Bethyl Laboratories, Montgomery, TX).

Statistical analysis

The Kolmogorov-Smirnov test was used to assess normality and Levene's test was used to assess the equality of variances between groups. Between group differences in expression were determined using independent t-tests, with correlations assessed using Pearson correlation tests. Results according to endometriosis stage were analyzed using the Kruskal-Wallis test.

All statistical analyses were performed using SPSS version 13, with a p-value less than 0.05 considered statistically significant.

Table 1. Primers for real-time RT-PCR

Name	Sequences	Annealing temperature	Product size (bp)
TLR1	F:5'-CTATACACCAAGTTGTCAGC-3' R:5'-GTCTCCAACCTCAGTAAGGTG-3'	60	220
TLR2	F:5'-GCCAAAGTCTTGATTGATIGG-3' R:5'-TTGAAGTTCTCCAGCTCCTG-3'	64	347
TLR4	F:5'-TGGATACGTTTCCTTATAAG-3' R:5'-GAAATGGAGGCACCCCTTC-3'	56	507
TLR5	F:5'-CTAGCTCCTAATCCTGATG-3' R:5'-CCATGTGAAGTCTTTGCTGC-3'	56	438
TLR6	F:5'-CCTCCCAGGATCAAGGTAAGTIG-3' R:5'-ATCAGGCCAGCCCTCTAACAC-3'	60	327
TLR9	F:5'-CCCTCAACTTCACCTTGATCT-3' R:5'-CCACATATGGCCAGTGCA-3'	64	408
IL-1 β	F:5'-TGATGGCTTATTACAGTGGCAATG-3' R:5'-GTAGTGGTGGTCGGAGATTTCG-3'	140	60
IL-6	F:5'-GTGTTGCTGCTGCCCTTC-3' R:5'-AGTGCCCTTTGCTGCTTTC-3'	60	194
IL-8	F:5'-GACATACTCCAAACCTTCCAC-3' R:5'-CTTCTCCACAACCCTCTGC-3'	60	160
IL-10	F:5'-GAACCAAGACCCAGACATC-3' R:5'-CATCTTCACCTGCTCCAC-3'	60	137
IL-12p40	F:5'-TCGGCAGGTGGAGGTCAGC-3' R:5'-CGCAGAATGTCAGGGAGAAGTAGG-3'	60	77
IFN- γ	F:5'-TGTGGAGACCATCAAGGAAGAC-3' R:5'-TGCTTTGCGTTGGACATTCAAG-3'	60	121
TNF- α	F:5'-ATCTTCTCGAACCCCGAGTG-3' R:5'-GGGTTTGCTACAACATGGGC-3'	60	51
iNOS	F:5'-TGGATGCAACCCATTGTC-3' R:5'-CCCGCTGCCCCAGTTT-3'	60	59
eNOS	F:5'-CGGCATCACCAGGAAGAAGA-3' R:5'-CATGAGCGAGGCGGAGAT-3'	67	60
β -actin	F:5'-GCGAGAAGATGACCCAGATC-3' R:5'-GGATAGCACAGCTGGATAG-3'	60	77

RT-PCR: real time-polymerase chain reaction; TLR: Toll-like receptor; NOD: nucleotide-binding oligomerization domain.

Results

Characteristics of Patients in the Endometriosis and Non-Endometriosis Groups

The 40 endometriosis patients included 18 nulliparas and 22 multiparas, of mean age 36.5 ± 8.8 years. The 40 non-endometriosis patients included 13 nulliparas and 27 multiparas, of mean age 40.6 ± 10.6 years. There were no between group differences in age, body mass index (BMI), fertility, or history of prior surgery ($p > 0.05$ each). However, mean serum CA125 concentration was significantly higher in the endometriosis than in the non-endometriosis group (51.0 ± 54.9 IU/ml vs 26.7 ± 24.8 IU/ml, $p < 0.05$).

Expression of TLR, NOD, IL, IFN- γ , TNF- α , iNOS and eNOS mRNAs in peritoneal fluid (Table 2)

We found that mRNAs encoding all PRRs, cytokines and NOS were present in the effusion fluid of patients in the endometriosis and non-endometriosis groups. The level of expression of each was higher in the endometriosis group, with the levels of expression of mRNAs encoding TLR-2 and -9; NOD-1 and -2; iNOS and eNOS being significantly higher in the endometriosis than in the non-endometriosis group ($p < 0.05$).

Concentrations of Igs in effusion fluid

The concentrations of IgG (1759 ± 379 μ g/ml vs 1670 ± 590 μ g/ml) and IgA (875 ± 449 μ g/ml vs 817 ± 359 μ g/ml) were higher, whereas the concentration of IgM was lower (242 ± 128 μ g/ml and 291 ± 151 μ g/ml), in the endometriosis than in the non-endometriosis group. However, none of these differences was statistically significant ($p > 0.05$ each).

Correlations of clinical manifestations with expression of PRRs, cytokines, NOS, and Igs (Table 3)

No significant correlation was observed between any clinical or demographic characteristic, including age, parity, or CA125 concentration, and the level of cytokine or NOS mRNA expression or Ig concentration, in the two patient groups ($p > 0.05$ each).

Correlation of PRR, cytokine, and NOS mRNAs with Ig concentrations (Table 4)

Significant correlations were observed among the expression of mRNAs encoding PRRs, including TLRs-1, -2, -4, -5, 6, and -9; NODs-1 and -2; ILs-1 β , -6, -8, -10, and -12; IFN- γ ; TNF- α , and iNOS and eNOS in the two groups ($p < 0.05$ each) (Table 4). In the endometriosis group, however, none of these mRNAs was correlated with the concentrations of IgG, IgM, and IgA ($p > 0.05$ each).

Expression of TLR, NOD, NOS, and cytokine mRNAs according to stage of endometriosis

In analyzing the expression of TLR, NOD, NOS, and cytokine mRNAs according to the stage of endometriosis, we found that the levels of expression of TLR2, TLR4, and TLR5 increased significantly with increasing stage of endometriosis ($p < 0.05$ each) (Table 5). Significant differences were also observed when these patients were divided into two groups, those with Stage 1 and those with Stage 2 and higher ($p < 0.05$ each) (Table 6).

Table 2. TLR, cytokine and NOS mRNA expression in the peritoneal fluid of patients with and without endometriosis.

	Endometriosis group		Non-endometriosis group		P value
	Mean	SD	Mean	SD	
TLR1	0.678	1.824	0.107	0.389	0.074
TLR2	0.828	2.441	0.081	0.293	0.007
TLR4	0.004	0.016	0.000	0.001	0.175
TLR5	0.013	0.046	0.000	0.001	0.153
TLR6	6.046	19.816	0.739	3.115	0.133
TLR9	8.669	19.843	0.618	1.694	0.022
NOD1	1.132	3.851	0.027	0.077	0.014
NOD2	0.008	0.021	0.004	0.014	0.048
iNOS	0.131	0.330	0.011	0.016	0.036
eNOS	5.838	13.943	0.419	1.146	0.026
IL-1 β	0.093	0.292	0.032	0.110	0.240
IL-6	0.090	0.271	0.126	0.683	0.773
IL-8	2.802	10.911	0.453	2.017	0.200
IL-10	0.056	0.103	0.078	0.249	0.632
IL-12	0.622	2.594	0.550	3.180	0.915
IFN- γ	0.004	0.007	0.019	0.084	0.322
TNF- α	0.085	0.408	0.086	0.465	0.991

SD; standard deviation

TLR: Toll-like receptor; NOD: nucleotide-binding oligomerization domain; iNOS: inducible nitric oxide synthase; eNOS: endothelial NOS; IL: interleukin; IFN- γ : interferon- γ ; TNF- α : Tumor necrosis factor- α

Table 3. Correlations among age, parity, CA125 concentration, the expression of PRR, cytokines, and NOS mRNAs, and Ig concentrations

		Endometriosis group		Control group	
		Pearson's Coefficients	P value	Pearson's coefficients	P value
Age	TLR1	-.099	.567	-.248	.171
	TLR2	-.068	.710	-.263	.176
	TLR4	-.095	.619	-.235	.247
	TLR5	-.056	.766	-.192	.338
	TLR6	-.048	.789	-.260	.165
	TLR9	.089	.611	-.153	.410
	NOD1	-.044	.797	-.146	.410
	NOD2	-.045	.796	-.280	.142
	IL-1 β	-.104	.547	-.145	.412
	IL-6	.070	.687	-.087	.625
	IL-8	-.167	.325	-.248	.152
	IL-10	.098	.558	.063	.721
	IL-12	-.075	.655	-.210	.226
	IFN- γ	.191	.259	-.014	.935
	TNF- α	.172	.303	.056	.748
	iNOS	.273	.102	-.196	.267
	eNOS	.202	.223	.062	.719
	Ig G	.209	.195	-.021	.903
	Ig A	.339	.032	.034	.844
	Ig M	.268	.094	.316	.060
Parity	TLR1	-.109	.539	-.182	.319
	TLR2	-.089	.639	-.197	.315
	TLR4	-.122	.537	-.151	.462
	TLR5	-.076	.697	-.116	.565
	TLR6	-.062	.738	-.201	.288
	TLR9	.036	.843	-.033	.861
	NOD1	-.063	.725	.028	.876
	NOD2	-.062	.733	-.211	.271
	IL-1 β	-.125	.481	-.108	.541
	IL-6	.009	.961	.077	.666
	IL-8	.168	.333	-.198	.255
	IL-10	.197	.248	-.032	.853
	IL-12	.038	.827	-.191	.272
	IFN- γ	.202	.244	-.081	.642
	TNF- α	.159	.356	-.035	.840
	iNOS	.211	.224	-.182	.302
	eNOS	.186	.278	-.040	.819
	Ig G	.290	.078	.149	.385
	Ig A	.415	.010	-.119	.491
	Ig M	.198	.232	.136	.430
CA125	TLR1	.089	.660	.345	.148
	TLR2	.165	.440	.311	.224
	TLR4	.144	.511	.288	.279
	TLR5	.127	.553	.290	.258
	TLR6	.146	.476	.318	.185
	TLR9	.372	.056	.298	.216
	NOD1	.102	.612	.237	.300
	NOD2	.116	.574	.308	.214
	IL-1 β	.138	.492	.536	.012
	IL-6	.368	.059	.198	.389
	IL-8	.104	.598	.302	.172
	IL-10	.113	.561	-.114	.615
	IL-12	.149	.440	.287	.195
	IFN- γ	.126	.515	-.030	.894
	TNF- α	-.012	.953	-.127	.565
	iNOS	.169	.390	.310	.171
	eNOS	-.011	.957	-.119	.588
	Ig G	.067	.719	-.003	.988
	Ig A	.423	.018	.242	.266
	Ig M	.213	.250	.047	.831

TLR: Toll-like receptor; NOD: nucleotide-binding oligomerization domain; iNOS: inducible nitric oxide synthase; eNOS: endothelial NOS; IL: interleukin; IFN- γ : interferon- γ ; TNF- α : Tumor necrosis factor- α

Table 4. Correlations between IgG, IgA, and IgM concentrations and mRNAs encoding TLRs-1, -2, -4, -5, -6, and -9; NODs-1 and -2; ILs-1 β , -6, -8, -10, and -12; IFN- γ ; and TNF- α in endometriosis group

	TLR1	TLR2	TLR4	TLR5	TLR6	TLR9	NOD1	NOD2	IL-1 β	IL-6	IL-8	IL-10	IL-12	IFN- γ	TNF- α	iNOS	eNOS	IgG	IgA	IgM
TLR1	1	.979**	.942**	.947**	.979**	.822**	.962**	.922**	.860**	.855**	.043	.419**	.012	.537**	.602**	.166	.878**	-.065	.098	.015
TLR2	.979**	1	.979**	.987**	.997**	.781**	.994**	.905**	.770**	.832**	.019	.419**	.012	.537**	.602**	.166	.878**	-.065	.098	.015
TLR4	.942**	.979**	1	.997**	.985**	.680**	.981**	.862**	.731**	.752**	-.007	.405**	-.009	.377**	.429**	.013	.691**	-.099	.022	.022
TLR5	.947**	.987**	.997**	1	.989**	.704**	.991**	.864**	.733**	.770**	-.011	.408**	-.010	.375**	.422**	-.001	.695**	-.121	.040	.044
TLR6	.979**	.997**	.985**	.989**	1	.755**	.989**	.921**	.769**	.809**	.026	.420**	.007	.489**	.467**	.119	.768**	-.081	.076	.042
TLR9	.822**	.781**	.680**	.704**	.755**	1	.778**	.671**	.812**	.988**	.013	.343**	-.020	.353**	.538**	.024	.830**	-.069	.250*	.124
NOD1	.962**	.994**	.981**	.991**	.989**	.778**	1	.863**	.764**	.829**	-.012	.417**	-.009	.382**	.446**	-.011	.735**	-.098	.090	.058
NOD2	.922**	.905**	.862**	.864**	.921**	.671**	.863**	1	.672**	.706**	.137	.400**	.054	.774**	.417**	.476**	.776**	-.050	.079	.048
IL-1 β	.860**	.770**	.731**	.733**	.769**	.812**	.764**	.672**	1	.824**	-.004	.337**	-.016	.344**	.901**	.006	.949**	.008	.179	-.032
IL-6	.855**	.832**	.752**	.770**	.809**	.988**	.829**	.706**	.824**	1	.014	.364**	-.017	.352**	.529**	.007	.823**	-.079	.218	.149
IL-8	.043	.019	-.007	-.011	.026	.013	-.012	.137	-.004	.014	1	.133	.482**	.044	-.034	.336**	-.030	-.063	-.098	-.052
IL-10	.419**	.417**	.405**	.408**	.420**	.343**	.417**	.400**	.337**	.364**	.133	1	.085	.949**	.919**	.027	.910**	-.067	-.062	-.008
IL-12	.012	.000	-.009	-.010	.007	-.020	-.009	.054	-.016	-.017	.482**	.085	1	.038	.013	.159	.016	.099	-.016	-.137
IFN γ	.537**	.457**	.377**	.375**	.489**	.353**	.382**	.774**	.344**	.352**	.044	.949**	.038	1	.906**	.275*	.897**	-.031	-.018	.007
TNF α	.602**	.456**	.429**	.422**	.467**	.538**	.446**	.417**	.901**	.529**	-.034	.919**	.013	.906**	1	.012	.993**	-.041	-.060	.036
iNOS	.166	.079	.013	-.001	.119	.024	-.011	.476**	.006	.007	.336**	.027	.159	.275*	.012	1	.024	.065	-.041	-.001
eNOS	.878**	.768**	.691**	.695**	.768**	.830**	.735**	.776**	.949**	.823**	-.030	.910**	.016	.897**	.993**	.024	1	-.054	-.063	.059
IgG	-.065	-.087	-.099	-.121	-.081	-.069	-.098	-.050	.008	-.079	-.063	-.067	.099	-.031	-.041	.065	-.054	1	.384**	-.122
IgA	.098	.112	.022	.040	.076	.250*	.090	.079	.179	.218	-.098	-.062	-.016	-.018	-.060	-.041	-.063	.384**	1	-.073
IgM	.015	.042	.022	.044	.042	.124	.058	.048	-.032	.149	-.052	-.008	-.137	.007	.036	-.001	.059	-.122	-.073	1

*p<0.05, ** p<0.01.

TLR: Toll-like receptor; NOD: nucleotide-binding oligomerization domain; iNOS: inducible nitric oxide synthase ; eNOS: endothelial NOS; IL: interleukin; IFN- γ : interferon- γ ; TNF- α : Tumor necrosis factor- α

Table 5. Expression of TLR, NOD, NOS, and cytokines according to stage of endometriosis

	Number	Stage	Mean	SD	P value
TLR1	25	1	0.539	1.988	0.189
	6	2	1.037	1.686	
	5	3	0.941	1.204	
TLR2	23	1	0.679	2.686	.016
	5	2	1.518	2.193	
	4	3	0.818	1.136	
TLR4	21	1	0.004	0.019	.003
	5	2	0.007	0.012	
	4	3	0.002	0.001	
TLR5	22	1	0.012	0.052	.032
	5	2	0.019	0.036	
	4	3	0.008	0.013	
TLR6	23	1	5.438	22.677	.115
	6	2	9.196	16.527	
	5	3	5.061	7.192	
TLR9	24	1	4.747	14.425	.191
	6	2	19.807	34.838	
	5	3	14.129	17.826	
NOD1	25	1	0.982	4.286	.096
	6	2	1.791	3.441	
	5	3	1.091	1.987	
NOD2	24	1	0.007	0.024	.191
	6	2	0.012	0.016	
	5	3	0.009	0.011	
IL-1 β	25	1	0.072	0.261	.561
	6	2	0.203	0.347	

	5	3	0.341	0.557	
IL-6	25	1	3.135	10.364	.191
	6	2	14.708	25.143	
	5	3	8.706	9.613	
IL-8	26	1	0.100	0.340	.510
	6	2	0.132	0.165	
	5	3	0.013	0.015	
IL-10	27	1	0.113	0.318	.075
	6	2	0.041	0.065	
	5	3	0.025	0.032	
IL-12	27	1	3.739	12.885	.195
	6	2	0.785	0.955	
	5	3	0.160	0.165	
IFN- γ	26	1	0.059	0.123	.196
	6	2	0.041	0.025	
	5	3	0.060	0.035	
TNF- α	27	1	0.718	3.041	.087
	6	2	0.161	0.254	
	5	3	0.658	1.293	
iNOS	26	1	0.003	0.008	.061
	6	2	0.006	0.007	
	5	3	0.005	0.006	
eNOS	27	1	0.109	0.484	.121
	6	2	0.021	0.027	
	5	3	0.030	0.042	
	27	1	1739429.259	431206.009	.447
IgG	7	2	1695337.143	148262.886	
	6	3	1922398.333	297959.256	
IgA	27	1	816538.519	436065.112	.423
	7	2	979875.714	382088.450	
	6	3	1017103.333	596140.403	
IgM	27	1	222407.778	107831.864	.220
	7	2	323938.571	157043.179	
	6	3	237848.333	166191.940	

TLR: Toll-like receptor; NOD: nucleotide-binding oligomerization domain; iNOS: inducible nitric oxide synthase; eNOS: endothelial NOS; IL: interleukin; IFN- γ : interferon- γ ; TNF- α : Tumor necrosis factor- α

Table 6. Expression of TLR, NOD, NOS, and cytokines in patients with endometriosis stage 1 and with stage 2 or higher

	Stage 1		Stage 2 or more		P value
	Mean	SD	Mean	SD	
TLR1	0.539	1.988	0.994	1.416	.080
TLR2	0.679	2.686	1.207	1.739	.004
TLR4	0.004	0.019	0.014	0.009	.001
TLR5	0.012	0.052	0.014	0.028	.008
TLR6	5.438	22.677	7.317	12.725	.038
TLR9	4.747	14.425	17.226	27.253	.078
NOD1	0.982	4.286	1.473	2.763	.035
NOD2	0.007	0.024	0.011	0.013	.078
iNOS	0.072	0.261	0.266	0.435	.018
eNOS	3.135	10.364	11.980	19.049	.041
IL-1 β	0.100	0.340	0.078	0.132	.523
IL-6	0.113	0.318	0.034	0.051	.095
IL-8	3.739	12.885	0.501	0.757	.339
IL-10	0.059	0.123	0.050	0.030	.028
IL-12	0.718	3.041	0.386	0.877	.076
IFN- γ	0.003	0.008	0.006	0.006	.075
TNF- α	0.109	0.484	0.025	0.033	.027

TLR: Toll-like receptor; NOD: nucleotide-binding oligomerization domain; iNOS: inducible nitric oxide synthase ; eNOS: endothelial NOS; IL: interleukin; IFN- γ : interferon- γ ; TNF- α : Tumor necrosis factor- α

Discussion

Endometriosis is a complex inflammatory disease of the pelvis, characterized by disparate morphological, histological and biochemical properties. We evaluated endometriosis immunologically by analyzing the expression of PRRs, cytokines, nitric oxide, and immunoglobulins in the peritoneal cavity and determining whether expression of any of these molecules was correlated with the clinical characteristics of endometriosis.

The inability of the immune system to promptly recognize invading pathogens can lead to various diseases since the host immune response against these pathogens may not occur or may be inappropriate. Thus, recognizing pathogens during the initial stage of disease is crucial. This process is initiated by PRRs, a group of molecules distributed in the extracellular, membrane, and cytoplasmic compartments. Humans have various PRRs, including TLRs and NOD-like receptors (NLRs), which differ by location. For example, TLRs-1, -2, -4, -5, -6 and -10 are present on the cell surface; TLRs-3, -7, -8, and -9 are located on lysosome/endosome membranes; and NODs-1 and -2 are present in the cytoplasm (2,6-7). Pathogens that escape extracellular or membrane detection systems may be recognized by PRRs in the cytoplasm or on lysosome/endosome membranes (8).

We found that patients with endometriosis expressed mRNAs encoding TLRs-1, -2, -4, -5, 6, and -9 and NODs-1 and -2, all PRRs associated with immunity against infection. Moreover, the levels of expression of mRNAs encoding TLRs-2 and -9 and NODs-1 and -2 were significantly higher in patients with than without endometriosis. Up-regulation of PRR expression and accelerated endometrial proliferation can result in tumor formation (9). Human heat-shock protein 70 has been reported to induce pelvic inflammation, involving the TLR-3 and TLR-4-mediated growth of endometrial cells (5, 10), with TLR-4 having a significant role in innate immune reactions to bacterial endotoxin in patients with endometriosis (4). Moreover, a TLR-4 polymorphism associated with hypo-responsiveness of the receptor may result in peritoneal inflammation. Thus, endometrial cells tend to adhere to the peritoneum, a condition that may induce the initiation of endometriosis (11). We found that PRRs other than TLR-3 and TLR-4 are involved in immune reactions in patients with endometriosis of the peritoneal cavity.

NO reacts to homeostatic and pathologic stimuli and is secreted by neurons, endothelial cells, platelets and neutrophils. NO also plays a crucial role in female reproductive processes, including ovulation, menstruation, implantation, pregnancy maintenance, and

labor and delivery (12). A higher level of activated peritoneal macrophages has been observed in women with endometriosis compared to those without the condition, with iNOS expression and NO production increased due to the activation of peritoneal macrophages (13). The participation of peritoneal macrophages in antimicrobial and antitumor activities increases as NO concentration increases. Furthermore, increased NO concentrations can change pro-inflammatory and peritoneal immune defense reactions and may be involved in the pathogenesis of endometriosis (14). Of the three NO-synthase isoenzymes, eNOS and nNOS are constitutively expressed, whereas iNOS, while not normally present in macrophages, can be upregulated depending on the stimulus. The latter enzyme is also involved in vasodilation and the destruction of pathogens such as bacteria (15). Although the level of NOS is dependent on endometrial phase, association with infertility, position in the menstrual cycle, and types of cells and tissues, we found that the levels of expression of iNOS and eNOS were higher in the endometriosis than the non-endometriosis group (16, 17).

Cytokines, a group of glycoproteins that participate in modulating inflammatory and immune reactions in many diseases, were recently found to be involved in endometriosis in humans and experimental animals. Although most studies have reported that the secretion of cytokines increases during and after the process of endometriosis, other studies have reported that cytokine secretion decreases or does not change significantly.

We observed no differences in the levels of expression of ILs-1 β , -6, -8, -10, and -12, INF- γ , and TNF- α between our endometriosis and non-endometriosis groups. In contrast, other studies have reported higher expression of ILs-1 β , -6, -8, and -15, TNF- α , monocyte chemotactic protein-1 (MCP-1), eotaxin, regulated upon activation normal T-cell expressed and secreted (RANTES), and intercellular adhesion molecule-1 (ICAM-1), in patients with than without endometriosis (18-21). Several other studies, however, have found that the levels of expression of ILs-1 β , -10, -12, and -18, TNF- α , RANTES, VEGF, PDGF, sFas and sFasL, and vascular cell adhesion molecule-1 (VCAM-1) were similar, or in some cases lower, in patients with than without endometriosis (18, 22-24). Differences among study results may be due to the severity of endometriosis; polymorphisms in the genes encoding these molecules; types of samples, including endometriosis tissue, peritoneal fluid, or serum; the nature of the control group; or the use of medications by patients with endometriosis. Although none of our patients showed evidence of peritonitis or other inflammation before surgery, the ab-

dominal cavities of all 80 patients were positive for IgG, IgA, and IgM secretion. The concentrations of IgG and IgA were higher, and the concentration of IgM was lower, in our endometriosis group, but none of these differences was statistically significant. Although B cells in the peritoneal cavity cannot produce antibodies in a sterile environment, without infection or external stimuli, immunoglobulins are spontaneously produced by B-1, not B-2, cells in the peritoneal cavity even in the absence of exogenous infection (25, 26). B-1 cells express the pan T-cell surface glycoprotein, CD5, and are localized in the peritoneal and pleural cavities, with few, if any, found in the spleen (27, 28). Although we did not separate B-1 from B-2 cells, the antibodies in the peritoneal cavity likely resulted from the immune responses of B-1 cells.

When B cells differentiate into plasma cells, the initial immunoglobulins produced are IgM and IgD, with IgG, IgA, and/or IgE produced following a process called "class-switching" (27). Thus, various antibodies can be present in the peritoneal cavity in the absence of any exogenous infection. We focused on three classes of antibody: IgG, which is primarily involved in chronic inflammation and auto-immune reactions; IgA, which is primarily involved in mucosal immunity; and IgM, which is primarily involved in acute inflammation. We excluded IgD and IgE, which are present at much lower concentrations. Previous findings showing that the concentration of a specific IgG autoantibody was increased in the peritoneal fluid of patients with endometriosis and that endometrial glandular epithelial staining for both IgG and IgA was significantly increased, suggests that endometriosis may be an autoimmune disease (29, 30). Moreover, the increase in IgG concentration suggests that these patients may have had a precursor condition of endometriosis requiring treatment (31).

We found that inflammatory, innate, and adaptive immune reactions in the peritoneal cavity were integrated, with significant correlations among mRNAs encoding PRRs such as TLRs-1, -2, -4, -5, -6, and -9, and NODs-1 and -2; cytokines such as ILs-1 β , -6, -8, -10, and -12, INF- γ , and TNF- α ; and iNOS and eNOS. These PRRs, cytokines, and NOS are induced by external stimuli, antigens, and pathogens, via distinct pathways; following which they stimulate and facilitate the production of other signaling molecules and/or act synergistically. The manifestations and correlations of PRRs, cytokines, and NOS may be involved in the pathogenesis of endometriosis and in endometriosis-induced immune reactions in the peritoneal cavity.

Our study had several limitations. For ethical reasons, our control group consisted of patients with lesions in the peritoneal cavity, rather than disease

free normal subjects. Although none of the control patients had infections in the peritoneal cavity, the peritoneal lesions may have induced immune reactions. Second, we measured the levels of expression of mRNA instead of protein, since some proteins were not expressed. Third, the response of tissues other than the endometrium was not examined since the study samples consisted only of peritoneal fluid and there have been no comparative studies of serum and endometrial tissue.

We have shown here that mRNAs encoding various PRRs, cytokines, and NOS, as well as IgG, IgA, and IgM, were expressed in the peritoneal cavities of patients with endometriosis, with this expression due to diverse inflammatory and immune reactions. The expression of several PRRs and NOS increased significantly, suggesting that the cooperative interaction of PRRs, cytokines, and NOS in innate immune responses in the peritoneal fluid may be associated with endometriosis.

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Competing Interests

The authors have declared that no competing interest exists.

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