Glycoprotein (gp) 96 expression: induced during differentiation of intestinal macrophages but impaired in Crohn’s disease

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Background: The glycoprotein (gp) 96 links the adaptive with the innate immune system. It is a chaperone with a binding domain for peptides generated by proteasomal degradation. During cellular stress, peptide loaded gp96 can be released and presented to T cells by antigen presenting cells (APCs).

Methods: mRNAs from in vitro differentiated macrophages (iv mac) and normal intestinal macrophages (IMACs) were compared by subtractive hybridisation and Affymetrix GeneChip analysis. Differentiation induced expression of gp96 was investigated in the multicellular spheroid (MCS) model. In vivo gp96 protein expression was detected by double labelling immunohistochemistry of human colon and in the CD4+ CD62L+ T cell transfer mouse model.

Results: Five of 76 clones obtained by subtractive hybridisation revealed >99% sequence homology to gp96. Affymetrix GeneChip analysis confirmed induction of gp96 in IMACs. Gp96 mRNA was detected in IMACs from normal and intestinal bowel disease mucosa. Induction of gp96 protein was observed after seven days in the MCS model of IMAC differentiation. Immunohistochemistry confirmed the presence of gp96 protein in IMACs in normal mucosa as well as in mucosa from patients with ulcerative colitis and diverticulitis. In mucosa from Crohn’s disease (CD) patients, gp96 protein was not detectable. In the CD4+ CD62L+ T cell transfer mouse model, gp96 was verifiable in non-activated IMACs.

Conclusion: Gp96 is induced during differentiation of normal IMACs but is not detected in IMACs in CD mucosa. As gp96 has been described as having a role in tolerance induction, this may be relevant for loss of tolerance against luminal bacteria found in CD patients.
An in vitro model for the differentiation of IMACs is culture of monocyes with HT-29 spheroids. In this model monoocytes are seeded onto spheroids of HT-29 cells and invade them followed by a differentiation process which results in a phenotype similar to IMACs in normal intestinal mucosa with lack of CD14, CD16, CD11b, and CD11c surface expression.

By subtractive hybridisation we found that gp96 was expressed in IMACs from normal mucosa but not in in vitro differentiated macrophages (iv mac). Therefore, we further investigated gp96 expression in normal and inflamed mucosa. The MCS model was used to study gp96 expression during differentiation of IMACs. Finally, we studied regulation of gp96 expression under inflammatory conditions in an animal model of colitis.

**MATERIALS AND METHODS**

**Patients**

Tissue samples were obtained from intestinal mucosa of 14 patients with Crohn’s disease (CD), five patients with ulcerative colitis (UC), three patients with diverticulitis, and 14 control patients with no intestinal inflammation who underwent surgery for other reasons (for example, colon cancer). The degree of inflammation was graded microscopically by determination of the inflammatory infiltrate of neutrophils, eosinophils, and lymphocytes: 0 = no infiltration, 1 = low degree of infiltration, 2 = severe infiltration. All control specimens were negative for inflammatory infiltrate and the CD, UC, and diverticulitis specimens all had a moderate to severe inflammatory infiltrate.

The study was approved by the University of Regensburg Ethics Committee.

**Isolation of human lamina propria mononuclear cells (LPMNCs)**

Surgical specimens from inflamed and normal mucosa were obtained by surgery. Specimens were obtained from the colon of patients with CD and UC after receiving informed consent. Control specimens were taken from patients who underwent surgery for other reasons (colonic carcinomas). On average, specimens were approximately 20 cm² and we obtained about 1×10⁶ IMACs. LPMNCs were isolated as described previously. In brief, mucosa was incubated in calcium and magnesium-free Hank’s balanced salt solution with 1 mmol/l ethylenediaminetetraacetic acid for 15 minutes at 37°C to remove intestinal epithelial cells. Intestinal epithelial cell depleted specimens were incubated for 30 minutes in 2 ml phosphate buffered saline (PBS) with 1 mg/ml collagenase type I (= 336 U/ml), 0.3 mg/ml deoxyribonuclease (DNase I; Boehringer, Mannheim, Germany), and 0.2 mg/ml hyaluronidase without fetal calf serum at 37°C. Cells were filtered through a 70 µm nylon mesh and the CD, UC, and diverticulitis specimens all had a moderate to severe inflammatory infiltrate.

**Isolation and purification of IMACs**

IMACs in isolated LPMNCs were labelled with immunomagnetic MicroBeads, armed with CD33 antibody, and purified twice with the help of type LS separation columns (Miltenyi Biotec, Bergisch Gladbach, Germany), as described recently.

**Generation of in vitro differentiated macrophages**

After leucapheresis, 1×10⁶ monocytes were seeded in Teflon bags and cultured at 37°C with 7% CO₂. By addition of 2% human AB serum for seven days, macrophages were generated out of the monocytes. After incubation at 4°C, cells detached from the Teflon bag.

**Subtractive hybridisation**

Subtractive hybridisation was performed, subtracting cDNA from iv mac from normal IMACs using the Clontech PCR-Select cDNA subtraction kit (Clontech, Palo Alto, California, USA), as previously described. Two hybridisations were performed. In the first, excess cDNA from iv mac was added. Single strand molecules from normal IMACs were significantly enriched for differentially expressed sequences. The entire population of molecules was subjected to polymerase chain reaction (PCR) to amplify the desired differentially expressed sequences. Only molecules with two different adaptors were amplified exponentially. A secondary PCR amplification was performed using nested primers to further reduce any background PCR products and enrich for differentially expressed sequences.

With the clones obtained, a similarity search was performed at http://www.ncbi.nlm.nih.gov/index.html. The BLAST program was used to perform DNA database searches for sequence similarities.

**Affymetrix oligonucleotide array analysis**

Affymetrix GeneChip analysis was performed with help of the Kompetenzzentrum für fluoreszenz Bioanaytik (KFB) in Regensburg (Germany). mRNA was isolated from IMACs from non-inflamed mucosa and from iv mac using polyT magnetic beads (Dynal, Oslo, Norway). In each case, RNA from three patients was pooled to avoid individual differences being responsible for variation.

Double stranded cDNA was synthesised by a Superscript II kit (Life Technologies, Karlsruhe, Germany) using oligo(dT) as primer. Biotin labelled cRNA was prepared by in vitro transcription reaction using an Enzo BioArray HighYield RNA Transcript Labelling Kit (Affymetrix, P/N 900182) based on the manufacturer's protocol. Biotin labelled cRNA was fragmented and, after addition of Control Oligonucleotide B2 (3 nM) and 2× Eukaryotic Hybridisation Controls (bioB, bioC, bioD, cre), hybridised to Affymetrix HG-U133A GeneChips (Affymetrix, Santa Clara, California, USA), scanned on the Affymetrix array scanner, and data analysis performed using the Affymetrix statistical data analysis software, Affymetrix Microarray Suite (version 5.0).

**Reverse transcription-polymerase chain reaction (RT-PCR) for gp96**

Poly(A)-RNA was isolated by polyT magnetic beads (Dynal) from CD3⁺ cells according to the manufacturer’s protocol. Oligonucleotides for gp96 PCR were chosen using the BTI software Gene Tool lite (version 1.0.0.1). For PCR, primers were used as follows: gp96 upstream, 5’ TGG TGT GGT GGA 3’; gp96 downstream, 5’ GGT GCC AGA CCA TCC GTA CT 3’. To test cDNAs for representation and full

**Figure 1**

As a result of a similarity search, five of 76 clones obtained by subtractive hybridisation of intestinal macrophages and in vitro differentiated macrophages were identified as glycoprotein 96 (gp96). Thereby, two different parts of the cDNA were identified. Nucleotides 598–1271 were obtained three times and nucleotides 2105–2713 twice.
MCS were generated according to the liquid overlay culture. Generation of multicellular spheroids (MCS) using the Checker Kit (Invitrogen, Leek, Netherlands) was used. 6K clathrin, and 2K clathrin primer set from the Gene. mRNA levels were higher in macrophages from four donors for iv mac, and 11 monocyte donors. (B) Real-time PCR for gp96 and GAPDH. mRNA levels were higher in macrophages from patients with UC, CD, and NIH mucosa than in monocytes and iv mac. For probe set 200598_at, 11-fold induction and for 200599_at, threefold induction was observed. For statistical analysis the t-test was used. On average, induction of glycoprotein 96 mRNA expression in IMACs from normal mucosa compared with iv mac was sevenfold in the Affymetrix analysis.

length genes, RT-PCR with a 5’ β-actin, 3’ β-actin, GAPDH, 6K clathrin, and 2K clathrin primer set from the Gene Checker Kit (Invitrogen, Leek, Netherlands) was used.

Generation of multicellular spheroids (MCS)

MCS were generated according to the liquid overlay culture technique: 4 x 10^3 cells suspended in 0.2 ml of medium/well were seeded in agarose coated wells of 96 well plates and cultured under static conditions, as described. After seven days of culture, MCS had formed and were used for experiments. MCS supernatants (0.1 ml each) were replaced by freshly isolated monocytes in 0.1 ml of medium supplemented with 2% human AB serum. Cocultures of MCS with monocytes were harvested after 24 hours, three days, and seven days for immunohistochemical analysis.

Antibodies

The following antibodies were used for immunohistochemical identification of gp96: rat anti-gp96 monoclonal antibody (clone 9G10, IgG2a; Stressgen Biotechnologies, Canada), rabbit anti-gp94 polyclonal antibody (Stressgen Biotechnologies), and biotin conjugated rabbit antirat and goat antirabbit secondary antibody (Sigma, Deisenhofen, Germany).

The following monoclonal antibodies were used for immunohistochemical identification of macrophages: mouse antihuman macrophage CD68 (clone KP1; Dako, Hamburg, Germany) and biotin conjugated goat antimouse secondary antibody (IgG, monoclonal; Sigma) for detection of mouse macrophages; and biotinylated rat anti-mac-3 (IgG1k, monoclonal; Cederlane, Canada) and biotinylated rat anti-F4/80 (IgG2a; Serotec, Germany). The monoclonal antibody mouse antihuman macrophage CD33 MicroBeads (Miltenyi Biotec, Bergisch Gladbach, Germany) was used for isolation of macrophages.

For immunofluorescence the following antibodies were used as secondary antibodies: Alexa Fluor 488 chicken antimouse IgG (H+L) (Molecular Probes Europe BV, the Netherlands) and Alexa Fluor 594 conjugated goat antirat IgG (Molecular Probes Europe BV).

Immunohistochemistry/immunofluorescence

For peroxidase staining, frozen sections were cut (5 μm), air dried, and fixed in acetone. Slides were rehydrated with PBS (pH 7.4) and incubated for 30 minutes with 0.3% H2O2 in PBS to quench endogenous peroxidase. Slides were washed with PBS and incubated for 60 minutes with 1% bovine serum albumin (Biomol, Hamburg, Germany) in PBS (blocking buffer) with goat serum (dilution 1:100; Dako, Denmark) for CD68 staining and rabbit immunoglobulin fraction (0.2 mg/ml total protein concentration; Dako) for gp96 staining.

To identify the cell types expressing gp96, an immunohistochemical procedure for sequential double antigen localisation was applied. Firstly, slides were incubated with either rat anti-gp96 (final concentration 5 μg/ml in blocking buffer) or mouse antihuman macrophage CD68 (0.33 μg/ml) primary antisera in blocking buffer for 60 minutes. Secondly, slides were incubated for 30 minutes with 1:500 diluted biotin conjugated antirat or antimouse IgG secondary antibody in blocking buffer. After washing, tissue was incubated with a freshly prepared solution of Vector NovaRED (Vector, Burlingame, California, USA) containing 0.01% hydrogen peroxide. Slides were incubated for 30 minutes with 0.3% hydrogen peroxide in PBS buffer and incubated with the primary and secondary antibodies for the second staining. Sections were preincubated for eight minutes in 0.01% benzidine dihydrochloride (BDHC; Sigma) with 0.03% sodium nitroprusside (Sigma). This was followed by incubation in the reaction medium (0.01% BDHC, 0.005% hydrogen peroxide, and 0.03% sodium nitroprusside) for blue staining. Staining was interrupted with water.

For immunofluorescence, all tissue specimens were fixed in 4% buffered formalin and embedded in paraffin. Specimens were dewaxed and pretreated in a water bath at 70°C for 40 minutes in 0.1% sodium citrate, blocked for 20 minutes with 20% goat serum at room temperature, and
then incubated with the two primary antibodies at 4°C overnight (antibody dilution 1:100). Slides were washed in PBS and further incubated for one hour at 37°C with secondary antibodies (dilution 1:200) conjugated with fluorescein isothiocyanate or Texas red in the dark. Slides were washed and mounted in Vectashield mounting medium with 1.5 μg/ml 4',6-diamidino-2-phenylindole (DAPI; Vector Laboratories). Microscopy was done with a Leitz DMRXE microscope, mostly with 400× magnification. Detailed information on magnification is given in the figure legends.

**CD4⁺CD62L⁺ T cell transfer model of colitis**

Splenic CD4⁺CD62L⁺ T cells from BALB/c mice were isolated as described previously with slight modifications. In brief, CD4⁺ T cells were purified from spleen mononuclear cells of healthy mice by negative depletion of other cell populations using anti-CD8, anti-MHC-II, anti-B220, and anti-CD11b antibodies (purchased from Pharmingen, San Diego, California, USA) followed by anti-rat IgG immunomagnetic microbeads (Miltenyi Biotech, Bergisch Gladbach, Germany). CD4⁺ T cells were further separated by immunomagnetic beads into CD62L⁺ and CD62L⁻ T cells. The former cells (purity >95%) showed high expression of CD45RB by FACS analysis. CD62L⁺ CD4⁺ T cells (0.25 × 10⁶) were resuspended in 200 μl of sterile PBS and injected intraperitoneally into recipient CB-17 SCID mice. Colitis activity was monitored by weight change and histology analysis, as specified below.

**RESULTS**

**Subtractive hybridisation, RT-PCR, and Affymetrix GeneChip analysis**

To analyse whether specific proteins are induced during differentiation of IMACs, subtractive hybridisation of mRNA from iv mac and IMACs from normal mucosa was performed as described above. Evolving clones from this screening represented genes that are expressed in IMACs but not in iv mac.

Overall, 76 different clones were obtained. All cDNA fragments were cloned and sequenced. Some genes were obtained several times as sequenced clones contained different parts of the same gene. As genes obtained several times are considered more likely to be truly upregulated we focused on these clones. A search for the expressed sequence tag database showed that five of 76 subtracted products had >99% homology with mRNA of gp96 (reference sequence in NCBI: X15187). We obtained two different fragments of gp96 (nucleotides 598–1271 and 2105–2713) from which one occurred three and the other two times (fig. 1).

To confirm the subtractive hybridisation data, we used Affymetrix GeneChip analysis. mRNA from three control patients and three donors for iv mac were pooled. mRNA in each case was hybridised onto two chips. For both probe sets for gp96 significantly higher mRNA expression was found in IMACs compared with iv mac. For each probe set we averaged the two chips. For one probe set nearly threefold and for the other 11-fold higher gp96 mRNA expression in IMACs compared with iv mac was found (fig 2).

To further demonstrate the reliability of subtractive hybridisation and the Affymetrix GeneChip analysis, we performed RT-PCR for gp96 with mRNAs from CD-IMACs, UC-IMACs, and control-IMACs from monocytes and from iv mac. In macrophages from CD, UC, and non-inflamed mucosa, but not from blood monocytes, gp96 cDNA was amplified. Also, almost no gp96 cDNA was amplified in iv mac (fig 3A). The integrity of the mRNA was verified by the Gene Checker kit (fig 3A, only GAPDH shown). Additionally,
we used the MCS model of IMACs differentiation to induction of gp96 during differentiation of IMACs. Therefore, based on the results for mRNA expression, we assumed that gp96 expression was detected in intestinal macrophages in Crohn’s disease (B). No NovoRED reaction product was detected using rat IgG2a isotype antibody in the first staining step. No BDHC reaction product was detected using mouse isotype control in the second staining step (C, D). The broken line indicates the crypt border. Original magnification ×400.

**Immunohistochemical analysis of gp96 expression in MCS**

Based on the results for mRNA expression, we assumed induction of gp96 during differentiation of IMACs. Therefore, we used the MCS model of IMACs differentiation to investigate whether gp96 is differentiation specific in IMACs.

Twenty four hours and three days after coculture of HT-29 cells with monocytes, no gp96 was detected in MCS (fig 4A, B). After seven days of coculture, gp96 was detected in high amounts in MCS in a pattern typical of invaded monocytes/macrophages (fig 4C, D). This confirmed induction of gp96 during IMAC differentiation.

**Gp96 protein expression in human intestinal mucosa**

Double labelling immunohistochemistry with specimens from nine patients without intestinal inflammation and seven patients with CD was performed. As shown in fig 5, gp96 was detected in IMACs from patients without intestinal inflammation (fig 5A) but not in macrophages from patients with CD (fig 5B). To exclude the possibility that peptide fragments bound to gp96 mask the epitope for the monoclonal anti-gp96 antibody, additional immunohistochemistry with a polyclonal antibody was performed, revealing identical results (fig 6).

Identical results were obtained by immunofluorescence. As shown in fig 7A–C (arrows), gp96 was expressed in the vast majority of IMACs in normal non-inflamed mucosa. Gp96 negatively stained cells accumulated subepithelially and were preferentially detected close to crypts. Positively stained cells were mainly found in the deeper layers of the lamina propria. IMACs in the mucosa of patients with CD were mostly negative for gp96 (fig 7D–F, green arrows). Only a small number of IMACs stained positive for gp96 (fig 7D–F, white arrows). No gp96 protein expression was detected in other cells in the mucosa in all samples (figs 5–7). To determine whether the absence of gp96 found in CD-IMAC is specific for CD or whether it is an epiphenomenon of mucosal inflammation, we performed immunohistochemistry and immunofluorescence on sections of patients with diverticulitis and UC. As shown in fig 8, during diverticulitis (fig 8A–F) and UC (fig 8G–L), gp96 was detected in large amounts in IMACs similar to non-inflamed controls (figs 5–7, arrows). The distribution of gp96 positive IMACs in the lamina propria of patients with diverticulitis was the same as in patients without inflammation.

**Gp96 expression in the transfer colitis mouse model**

As the function of gp96 for mucosal tolerance and inflammation is difficult to study in humans, we further investigated whether gp96 expression and regulation is similar in an...
animal model of colitis. CD4+ CD62L+ cells were isolated from BALB/c mice and transferred into SCID mice. After seven weeks SCID mice developed colitis and frozen sections from the intestine of seven mice with colitis and from five healthy mice without colitis were stained immunohistochemically to determine gp96 protein expression. As there is no typical general marker for macrophages in mice, we used two different antibodies (Mac-3 and F4/80) to detect macrophages in the intestinal mucosa. In mucosa from mice without colitis, gp96/Mac-3 positive macrophages were detected (fig 9A). In contrast, no gp96 was detected in the mucosa of mice with colitis (fig 9B, F). There were also no Mac-3 positive macrophages detected in the mucosa of mice with colitis (fig 9B). However, in Mac-3 positive macrophages that were located in lymph follicles of mice with colitis, gp96 protein was expressed (fig 9C, D). The majority of IMACs in healthy mice were negative for another macrophage marker, F4/80, which is assumed to be expressed in mature and activated macrophages. Only very few F4/80 positive IMACs were detected in the non-inflamed intestinal mucosa. These IMACs were negative for gp96 (fig 9E). In contrast, in the inflamed mucosa a high number F4/80 positive macrophages were detected which were gp96 negative (fig 9F). Taken together these data indicate that regulation of gp96 protein in this model of colitis is very similar to that found in human CD.

**DISCUSSION**

In the present study we demonstrated induction of gp96 expression during differentiation of IMACs at the mRNA and protein levels. Immunohistochemistry clearly showed gp96 expression in IMACs of the lamina propria from patients with normal non-inflamed mucosa, diverticulitis, and UC. Almost no gp96 protein expression could be found in the intestinal mucosa of patients with CD, raising the possibility that absence of gp96 is involved in CD specific pathophysiology. In mice, gp96 was found to be expressed in mac-3 positive IMACs in the non-inflamed mucosa but was absent in mature F4/80 positive IMACs in the inflamed mucosa of mice with colitis, indicating regulation similar to that found in humans.

Previously, gp96 expression was mainly studied in the context of malignant diseases, such as colorectal cancer or malignant mesothelioma. In both, gp96 expression was enhanced in tumour tissue compared with normal healthy tissue, which indicates that gp96 is involved in antigen presentation by peptide transfer to MHC class I. As a role for gp96 in antigen presentation and induction of tolerance is likely, the presence of this protein in IMACs is interesting. Our data indicate that gp96 is induced during specific differentiation of intestinal APCs. This points to a specific role for this protein in the mucosal immune system.

Gp96 is detected in high amounts in IMACs under conditions of mucosal tolerance and is decreased in CD, associated with loss of tolerance for its own bacterial flora. The role of gp96 during chronic inflammation has not been studied; in contrast, expression of Hsp 90, the cytosolic homologue of gp96, was investigated in patients with CD and UC. No difference in Hsp 90 expression between inflamed and non-inflamed mucosa was observed. However, Ludwig et al showed enhanced expression of Hsp 70 in epithelial cells in CD and UC, independent of the degree of inflammation. Peetermans et al demonstrated enhanced Hsp 60 expression in B7 positive mononuclear cells in the mucosa of patients with intestinal bowel disease (IBD) and enhanced Hsp 60 staining was reported in epithelial cells in UC. These investigations did not provide evidence for a role of these Hsps in chronic mucosal inflammation.

Our study is the first to investigate expression of gp96 in the intestinal mucosa in the context of IBD. In contrast with other studies which showed unchanged or enhanced expression of Hsps during IBD, we demonstrated here that gp96 was downregulated or absent in IMAC-CD. There are several reasons why gp96 may disappear in active CD. The first

**Figure 7** Immunofluorescence for glycoprotein 96 (gp96) expression in the intestinal mucosa of patients with no inflammation and with Crohn’s disease (CD). Paraffin embedded sections were cut and gp96 was detected with a monoclonal antibody. In a second step, a secondary Alexa Fluor 594 conjugated goat antirat IgG antibody was used (red). CD68 was also detected with a monoclonal antibody and in a second step an Alexa Fluor 488 chicken antimouse IgG (H+L) was used (green). Counterstain was done with 4',6-diamidino-2-phenylindole (blue). In non-inflamed mucosa (A–C), most of the intestinal macrophages (A, green, arrows) were gp96 positive (B, red; C, merge, white arrows). No fluorescence was detected using rat IgG2a or mouse IgG1 isotype control (G, H). Original magnification ×200.
possibility is that gp96 protein is secreted by activated IMACs in inflamed CD mucosa in high amounts and is rapidly degraded. Our data from the Affymetrix GeneChip analysis showed that in IMACs from healthy non-inflamed mucosa, proteasome associated proteins were downregulated in comparison with iv mac (data not shown). In IMACs from inflamed mucosa, these genes can be upregulated again, so that protein degradation is enhanced and gp96 is degraded intracellularly. In qualitative PCR in both IMACs from normal and from inflamed mucosal tissue, gp96 mRNA was detected. Because of the low number of cells available after purification and a consecutive low amount of mRNA (under the level of detection), a high number (40) of cycles have to be done in PCR. Therefore, no conclusion about mRNA quantity can be made. Taqman data confirmed the results of subtractive hybridisation and Affymetrix GeneChip analysis which clearly showed enhanced gp96 expression in IMACs compared with monocytes and iv mac.

We showed that gp96 was induced in intestinal macrophages during differentiation of monocytes into an intestinal macrophage-like phenotype in the MCS model. During inflammation, monocytes invade the intestinal mucosa. Invading monocytes do not express gp96. This could explain, to some extent, why no gp96 protein was found in CD patients. However, mRNA expression was induced, as demonstrated. In addition, in UC, monocytes invade the lamina propria and finally express gp96 protein. Therefore, invasion of monocytes does not sufficiently explain the absence of gp96 protein in CD mucosa. An alternative explanation for downregulation of gp96 protein in CD intestinal macrophages could be the occurrence of post transcriptional or post translational regulation. This needs to be evaluated further.

As noted previously, downregulation of gp96 in IMACs of CD mucosa may play a role in loss of tolerance against luminal antigens. Chandawarkar et al found that immunisation with a 5–10-fold higher than optimal dose of gp96 downregulated antitumour immune responses. The authors showed recently that this effect was dependent on CD4+ T cells. These cells have to be further characterised but it is possible that CD25+ regulatory T cells are generated when gp96 is present in high amounts. It has been shown that gp96 cell surface expression in transgenic mice leads to a lupus-like autoimmune disease. Extracellular gp96 that has not bound processed peptides could serve as an endogenous APC activator. Chronic activation of APCs by gp96 could cause

Figure 8 Immunohistochemical detection of glycoprotein 96 (gp96) on paraffin embedded and frozen sections from patients with diverticulitis (A–F) and ulcerative colitis (UC) (G–L). Paraffin embedded sections (A–D and G–J) were cut and gp96 was detected with a monoclonal antibody. In a second step, a secondary Alexa Fluor 594 conjugated goat antirat IgG antibody was used (red). CD68 was also detected with a monoclonal antibody, and in a second step an Alexa Fluor 488 chicken antimouse IgG (H+L) was used (green). Counterstain was done with 4',6-diamidino-2-phenylindole (blue). Frozen sections (E, F, K, L) were cut and fixed in acetone for peroxidase staining. Gp96 was immunohistochemically detected in a first step with Vector NovaRED (red). In a second step, the cellular marker for macrophages (CD68) was detected with benzidine dihydrochloride (BDHC) (dark blue, granular). In patients with diverticulitis (A–E) as well as in patients with UC (G–K), most intestinal macrophages were gp96 positive (arrows). No fluorescence or NovaRED reaction product was detected using rat IgG2a in the first staining step. No fluorescence or BDHC reaction product was detected using mouse isotype control in the second staining step (D, F, J, L). The broken line indicates the crypt border. Original magnification A–D and G–J ×200; E, F, K, L ×400.
A similar mechanism could be responsible for maintaining the inflammation in the intestine during CD. Recently, an immunomodulatory role for another related glycoprotein (HC gp39) was reported. In ex vivo assays, HC gp39 directed immune response was capable of suppressing cytotoxic T cell responses, indicating that the HC gp39 directed immune response in healthy individuals tends towards a regulatory phenotype. CD4\(^+\) T cell lines directed against HC gp39 expressed CD25, GITR, CTLA-4, and Foxp3 molecules and were capable of suppressing other immune responses. It could be speculated that gp96 has a similar effect in the intestinal mucosa. In fact, autoantibodies against gp96 can be detected in healthy mice. This interesting aspect warrants further study.

Very recently Doody et al demonstrated that gp96 can chaperone MHC II restricted epitopes for in vivo presentation to CD4\(^+\) T helper cells. CD4\(^+\) T cells can proliferate but are unable to express effector cytokines such as interferon \(\gamma\) or interleukin 4.

In conclusion, we have demonstrated that gp96 is induced during differentiation of monocytes into IMACs in humans and mice. We propose that gp96 functions as a tolerance mediating molecule. In CD mucosa and mouse colitis, gp96 is not detected. Further studies on the functional role of gp96 in the context of CD are under way.

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REFERENCES
An unusual cause of rectal bleeding in a young woman

Clinical presentation
A 24 year old woman presented with bloody rectal discharge of three months’ duration. She had a medical history of herpes progenitalis and candida vaginitis. On examination, there was no tenderness in the abdomen. Routine stool culture and blood tests were negative. Colonoscopy with indigo carmine dye revealed friable and erythematous nodular mucosa in the lower rectum (fig 1).

Question
What is the diagnosis?
See page 949 for answer

Editor’s Quiz: GI Snapshot

Robin Spiller, Editor

Figure 1
Colonoscopy with indigo carmine dye of the lower rectum.

References