Involvement of the fractalkine pathway in the pathogenesis of childhood hemolytic uremic syndrome

María Victoria Ramos,1 Gabriela C. Fernández,1 Natasha Patey,4 Pablo Schierloh,1 Ramón Exeni,2 Irene Grimoldi,2 Graciela Vallejo,3 Christian Elías-Costa,3 María del Carmen Sasain,1 Howard Trachtman,7 Christophe Combadière,6 François Proulx,3 and Marina S. Palermo1

1Division of Immunology, Institute of Hematological Investigations, Academia Nacional de Medicina, Buenos Aires, Argentina; 2Department of Nephrology, Hospital Municipal del Niño, San Justo, Argentina; 3Department of Nephrology, Hospital de Niños “Ricardo Gutierrez,” Buenos Aires, Argentina; 4Department of Pediatrics, Division of Pediatric Pathology, Hôpital Necker Enfants-Malades, Paris, France; 5Department of Pediatrics, Division of Pediatric Intensive Care, Ste-Justine Hospital, Montreal, QC, Canada; 6INSERM (Institut National de la Santé et de la Recherche Médicale) U543, Université Pierre et Marie Curie, Hôpital Pitié-Salpêtrière, Paris, France; 7Department of Pediatrics, Division of Pediatric Nephrology, Schneider Children’s Hospital, New York, NY

Thrombotic microangiopathy and acute renal failure are cardinal features of post-diarrheal hemolytic uremic syndrome (HUS). These conditions are related to endothelial and epithelial cell damage induced by Shiga toxin (Stx) through the interaction with its globotriaosyl ceramide receptor. However, inflammatory processes contribute to the pathogenesis of HUS by sensitizing cells to Stx fractalkine (FKN), a CXC transmembrane chemokine expressed on epithelial and endothelial cells upon activation, involved in the selective migration and adhesion of specific leukocyte subsets to tissues. Here, we demonstrated a selective depletion of circulating mononuclear leukocytes expressing the receptor for FKN (CX3CR1) in patients with HUS. We found a unique phenotype in children with HUS distinct from that seen in healthy, uremic, or infected controls, in which monocytes lost CX3CR1, down-modulated CD62L, and increased CD16. In addition, the CD56dim natural killer (NK) cell subpopulation was decreased, leading to an altered peripheral CD56dim/CD56bright ratio from 10.0 to 4.5. It is noteworthy that a negative correlation existed between the percentage of circulating CX3CR1+ leukocytes and the severity of renal failure. Finally, CX3CR1+ leukocytes were observed in renal biopsies from patients with HUS. We suggest that the interaction of CX3CR1+ cells with FKN present on activated endothelial cells may contribute to renal injury in HUS. (Blood. 2007; 109:2438-2445)

Introduction

The postdiarrheal form of hemolytic uremic syndrome (HUS) is characterized by hemolytic anemia, thrombocytopenia, and acute renal failure,1,2 and has been associated with enterohemorrhagic infections caused by Shiga toxin (Stx)-producing Escherichia coli (STEC).3,4 Endothelial dysfunction induced by Stx is a central event to the development of thrombotic microangiopathy observed in children with HUS. However, clinical and experimental data suggest that the host’s inflammatory response to other bacterial virulence factors plays a pivotal role in the pathogenesis of the disease. Lipopolysaccharide (LPS) O157 has been shown to activate platelets,5 and several other studies demonstrated activation of myeloid cells by Stx and LPS.6-8 Moreover, we have previously demonstrated that depletion of hepatic and splenic macrophages diminished Stx cytotoxicity in mice.9 We have also demonstrated that circulating levels of C-C chemokines for monocytes (Mo’s) were associated with the severity of postdiarrheal HUS.8

The migration and interaction of leukocytes with blood vessels involve the elaboration of cytokines and chemokines by injured cells.10,11 Leukocyte attachment and rolling on endothelial cells is promoted by vasodilatation and expression of selectins, followed by integrin-mediated cellular adhesion.10,11 Chemokines can be divided broadly into inflammatory and homeostatic chemokines,12 However, the pattern of chemokine expression and function of vascular endothelium are completely modified once activated. Fractalkine (FKN; CX3CL1) is the only member of a novel class of chemokines (CXC). In contrast to all other known chemokines, FKN is a type I transmembrane protein, consisting of a chemokine head tethered to the cell surface by a mucin stalk and a short cytoplasmic tail.13 Metalloproteases lead to the proteolytic cleavage of FKN under constitutive or inflammatory conditions.14 FKN may thus function as an adhesion molecule (membrane-anchored form) as well as a potent soluble (s) chemotactrant.15 FKN has been detected on endothelial15,16 and epithelial cells,17,18 whereas the FKN receptor (CX3CR1) is expressed on Mo's, natural killer (NK) cells, and a subset of CD8+ T cells.19,20 CX3CR1 is absent on neutrophils and B cells.13,14,19 whereas both receptor and ligand have been noted on dendritic cells.21,22 Under conditions of physiologic flow, FKN mediates adhesion of monocytes and NK cells, independently of integrin15,19 and G-coupled protein receptors.21,23 In a murine model of glomerulonephritis, treatment with an anti-CX3CR1 antibody dramatically reduced leukocyte infiltration within glomeruli and improved renal function.25,26 Evidence of CX3CR1-expressing cells within human kidneys suggested a role
for FKN in the pathogenesis of glomerulonephritis as well as in renal transplant rejection.27,28

Accordingly, we hypothesized that the interaction of CX3CR1-carrying cells with FKN present on activated endothelial cells could enhance renal damage during HUS. In the present study, we analyzed CX3CR1-expressing circulating leukocyte populations in children with HUS during the acute phase of the illness. We also evaluated serum concentrations of (s)FKN, and the extent of renal infiltration by CD68 or CX3CR1 evaluated serum concentrations of (s)FKN, and the extent of renal infiltration by CD68 or CX3CR1.

Patients and materials, and methods

Patients and samples

The study was approved by the Hospital Ethical Committees: the Comité de Bioética del Hospital Municipal del Niño de San Justo, San Justo, Buenos Aires, Argentina; the Comité de Bioética del Hospital Ricardo Gutiérrez, Capital Federal, Buenos Aires, Argentina; and that of Ste-Justine Hospital, Montreal, Canada. All patients were enrolled after informed consent was obtained from their parents. The Argentinean group included 31 children who were studied in the acute period of HUS. The diagnostic criteria for HUS were microangiopathic hemolytic anemia with schizontes, thrombocytopenia (platelet count < 150 × 10^9/L), and acute renal failure (serum creatinine level > 106.0 μM). All patients developed HUS after a prodrome of gastroenteritis with bloody diarrhea. Half of the children showed evidence of Stx-producing E. coli O157 by stool culture and/or the presence of atypical HUS due to factor H deficiency (n = 9), healthy control children of similar age (n = 9), and centrifuged at 400g for 30 minutes. Erythrocytes were then treated with FACS Lysing solution (BD Biosciences), washed, and resuspended in 0.2 μL of ISOFLOW (International Link SA, Buenos Aires, Argentina). In all cases, isotype-matched antibodies were assayed in parallel, and the fluorescence was measured on 100 000 events using the Cell Quest program (CellQuest, San Jose, CA) on a Becton Dickinson FACScan. Mo’s were identified and gated according to their forward- and light-scattering (FSC/SSC) dotplot profiles and positive staining for CD14.

PBMC isolation

Blood was diluted 1:2 with saline, layered on a Ficol-Hypaque cushion (Ficoll Pharmacia, Uppsala, Sweden; Hypaque, Winthrop Products, Buenos Aires, Argentina) and centrifuged at 400g for 30 minutes. Erythrocytes were subsequently washed with FCS and antibiotics. Viability of PBMCs was more than 96% as determined by trypan blue exclusion test.

Influence of plasma or Stx on apoptosis of NK cells

PBMCs (1 × 10^6) were incubated for 18 hours in the presence or not of plasma from patients included in the different clinical groups (80%) at 37°C in 5% CO₂, or in medium containing 10^-6 to 10^-3 M Stx type 2 (Stx2) (kindly provided by Dr Sugiyama Junichi, Denka Seiken Co Ltd, Nigata, Japan). After overnight incubation, cells were stained for surface CD3 and CD56 expression, and washed. Apoptosis was assessed based on the annexin V (AV)–FITC protein-binding assay according to manufacturer’s instructions without the addition of propidium iodide (APO-AF kit; Sigma-Aldrich, St Louis, MO). Results were expressed as a percentage of AV-positive cells among CD3^-CD56^- (NK) cells. CD56high and CD56dim were distinguished by the level of CD56 expression.

Immunohistochemistry

We evaluated kidney biopsy specimens from children with epidemic HUS (D + HUS; n = 9), healthy control children of similar age (n = 4), children with atypical HUS due to factor H deficiency (n = 7), acute renal rejection (n = 1), systemic lupus class IV (International Society of Nephrology/ Renal Pathology Society [ISN/RPS] 2003 classification; n = 2), or Henoch–Schönlein purpura (n = 2). Tissue sections (3 μm) were cut on super-frost slides from alcohol formalin acid acetate– or Bouin fluid–fixed, paraffin-embedded blocks and immunostained with mAb anti-iCD68 antigen (macrophage; clone KP1. IgG1; k: 1:1500; DAKO SA, Glostrup, Denmark) and anti-CXCR1 antibody (rabbit polyclonal antibody, AB 1891, 1:20; Chemicon International, Temecula, CA). Sections were dewaxed in a microwave oven in citrate buffer (pH 6) at medium power. After microwave irradiation (2 times for 5 minutes for biopsy specimens fixed in alcohol formalin acid acetate, and 3 times for 5 minutes for specimens fixed in Bouin fluid), slides were allowed to cool at room temperature and were then incubated with the
Monocytes, NK bright, NK dim, Lymphocytes, anti-CX3CR1–FITC mAb on whole blood to avoid variations in its profile after whole-blood analysis by flow cytometry (Figure 1A). Circulating leukocytes were identified by their SSC-versus-FSC staining reaction with 3-3 primary antibody, followed by an indirect streptavidin-biotin method and staining reaction with 3-3 diaminobenzidine (DAB; DAKO ChemMate Detection Kit). Finally, sections were counterstained with hematoxylin. Images were captured from paraffin sections by a Leica DMLB 100 microscope with a Leica DC 300F camera (Leica, Heidelberg, Germany).

Statistic analysis

Results are expressed as means ± SD. Comparative analyses were performed using a nonparametric, unpaired, 2-tailed Mann-Whitney U test. P values less than .05 were considered significant. Correlations between immunologic and clinical data were done with the nonparametric Spearman rank-correlation test.

Results

Definition of leukocyte populations by CX3CR1 expression

Circulating leukocytes were identified by their SSC-versus-FSC profile after whole-blood analysis by flow cytometry (Figure 1A). Therefore, comparative analysis of CX3CR1 expression on leukocytes was performed by multiparameter flow cytometry using the anti-CX3CR1–FITC mAb on whole blood to avoid variations in its cellular expression during isolation of leukocyte subpopulations. As shown in Figure 1B, several discrete CX3CR1+ mononuclear populations were identified in HC, whereas polymorphonuclear neutrophils (PMNs) were negative for CX3CR1, as previously described.13,14 In children with HUS the percentage of CX3CR1+ leukocytes was significantly reduced (Figure 1C-D), whereas all other groups had a similar expression (Figure 1D).

To investigate CX3CR1 expression on circulating mononuclear subpopulations from HC and children with HUS, we stained human peripheral blood cells with the corresponding phenotypic markers.

CX3CR1 expression on peripheral Mo’s from patients with HUS

Mo heterogeneity has been recently demonstrated by flow cytometry according to the expression of CD14 and CD16 molecules.30-32 To investigate CX3CR1 expression on circulating mononuclear subpopulations, we stained human peripheral blood cells with antibodies directed against the corresponding phenotypic markers. As shown in Figure 2A, we could define 4 subsets of CD14+ Mo’s: CX3CR1+/CD16−, CX3CR1−/CD16−, and CX3CR1−/CD16+ (double negative). Most peripheral Mo’s from HC corresponded to CX3CR1+/CD16− (approximately 78%), whereas the other subsets represented less than 15% of cells. Children with HUS showed a significant decrease in the percentages of the circulating CX3CR1+/CD16− “classical monocyte” subset (approximately 45%), with a simultaneous increase in all other subsets (Figure 2B).

Similarly, the percentage of CX3CR1+/CD16− Mo’s was abnormally decreased in IC (approximately 62%), with a simultaneous increase in double-negative or double-positive cells, whereas the

Table 2. Absolute numbers of circulating leukocytes in different clinical groups

<table>
<thead>
<tr>
<th></th>
<th>WBC count, × 10^9/L</th>
<th>No. patients</th>
</tr>
</thead>
<tbody>
<tr>
<td>HC</td>
<td>8.55 ± 2.71</td>
<td>21</td>
</tr>
<tr>
<td>HUS</td>
<td>17.06 ± 6.65*</td>
<td>31</td>
</tr>
<tr>
<td>UC</td>
<td>12.35 ± 5.42</td>
<td>20</td>
</tr>
<tr>
<td>IC</td>
<td>20.56 ± 11.87*</td>
<td>17</td>
</tr>
<tr>
<td>NK cells, × 10^9/L</td>
<td>0.40 ± 0.23</td>
<td></td>
</tr>
<tr>
<td>NK(2/3), × 10^9/L</td>
<td>3.26 ± 1.22</td>
<td></td>
</tr>
<tr>
<td>NK(3/4), × 10^9/L</td>
<td>0.18 ± 0.12</td>
<td></td>
</tr>
</tbody>
</table>

WBC indicates white blood cells. *P < .001 compared with HC. †P < .01 compared with HUS. §P < .005 compared with HC. ‡P < .005 compared with HUS.

Data are expressed as means ± SD. NK cell studies were performed on a lesser number of patients: HC (n = 18), HUS (n = 16), UC (n = 9), and IC (n = 8).

Figure 1. CX3CR1 expression in whole blood. Whole blood from HC (n = 10), patients with HUS (n = 19), IC (n = 10), and UC (n = 14) was stained with FITC-labeled anti-CX3CR1 and analyzed by flow cytometry as described in “Patients, materials, and methods.” (A) Representative dotplot of FSC versus FITC fluorescence of (B) 1 healthy child and (C) 1 child with HUS, showing CX3CR1 expression in mononuclear populations. The corresponding isotype control is also presented. (D) Percentage of CX3CR1 cells among mononuclear cells was calculated according the gates indicated in panels B and C for the different clinical groups. Results are expressed as means ± SEM. **P < .001 compared with all clinical groups using the Mann-Whitney U test.
relative proportions of Mo subsets were similar in HC and UC. Nevertheless, abnormalities present in the IC group were less striking than among children with HUS, making them significantly different (Figure 2B). Interestingly, children with HUS presented a unique phenotype consisting of significantly reduced percentages of circulating Mo’s CX3CR1 (CX3CR1Mo: HC = 84.6 ± 2.7%, n = 22; HUS = 55.5 ± 4.0%, n = 31; UC = 79.0 ± 4.3%, n = 20; and IC = 83.3 ± 1.9%, n = 17; *P < .001 compared with all other clinical groups).

Figure 2C shows that the distribution among the 4 subsets of circulating Mo’s was also different in absolute numbers. IC and children with HUS had the total count of Mo’s increased (Table 2). However, the number of regular Mo’s (CX3CR1+/CD16−) was significantly enhanced in IC, but was conserved in children with HUS, who showed, alternatively, an enhancement in double-positive and double-negative subsets (Figure 2C).

Finally, in agreement with previous reports among healthy adults, we noted that all groups showed a higher expression of CX3CR1 on CD16+ than CD16− Mo’s (P < .001). However, Mo’s from children with HUS showed an abnormally decreased CX3CR1 mean fluorescence intensity (MFI) on the CD16+ subset (CX3CR1 MFI: HC = 115.2 ± 11.1, n = 10; HUS = 71.3 ± 9.3, n = 19; P < .005), as well as on the CD16− subset (HC = 48.1 ± 5.2; HUS = 38.1 ± 3.9; P < .05).

Since Mo subpopulations are distinguished also by the expression of the adhesion receptor CD62L, we further characterized Mo subpopulations by measuring CD62L together with CX3CR1 on CD14+ Mo’s. Most circulating Mo’s in HC (CX3CR1+/CD16− Mo’s) also express CD62L. Comparative analysis showed a significant reduction in the expression of the double-positive (CX3CR1+/CD62L+) population in children with HUS (Figure 3A-B). This was mainly due to the loss of CX3CR1, since the total percentage of CD62L+ cells was conserved (HC = 77.0 ± 5.0%, n = 11; HUS = 86.9 ± 1.6%, n = 13; nonsignificant), and even more, the CX3CR1−/CD62L+ subset was significantly increased (Figure 3B). In this case, the UC and IC groups showed a Mo subset distribution similar to that of HC, and the percentages of all Mo subsets in the IC group were statistically different from those found in children with HUS (Figure 3B).

Figure 3. CD62L/CX3CR1 expression on peripheral Mo’s from different clinical groups. Whole blood from HC (n = 11), patients with HUS (n = 13), UC (n = 6), and IC (n = 11) was stained with PECy5-labeled anti-CD14, PE-labeled anti-CD62L, and FITC-labeled anti-CX3CR1, and analyzed by flow cytometry as previously described. Mo subpopulations were defined according to their CD14 expression, and those of CX3CR1 and CD62L were depicted. (A) Representative dotplots from 1 child within each clinical group showing CX3CR1 and CD62L expression on CD14+ Mo’s. Dot plot quadrants are defined as: CX3CR1+/CD62L+, top right; CX3CR1−/CD62L+, top left; and CX3CR1+/CD62L−, bottom right. The percentages of the different subpopulations are shown within each quadrant. (B) According to CX3CR1 and CD62L membrane expression, 4 subpopulations were defined. Each bar represents the mean ± SEM of the absolute number of cells for each Mo subset within the different clinical groups. *P < .05; **P < .01; ***P < .001 statistically different compared with HC. #P < .05; ##P < .01; ###P < .001 compared with children with HUS.
The MFI of CD62L was abnormally reduced in HUS patients on double-positive (CX3CR1+/CD62L+) cells as well as on CX3CR1−/CD62L+ cells (Figure 3C), indicating that CD62L was either shed or downmodulated.

**CX3CR1 expression on NK cells from patients with HUS**

As the NK cell population (CD3−/CD56+) expresses CX3CR1, we determined by 3-color flow cytometry its percentage among lymphoid cells using anti-CD3-PECy5, anti-CD56-PE, and anti-CX3CR1-FITC mAbs. As shown in Figure 4, children with HUS presented a significant decrease in the percentage of NK cells (CD3−CD56+) relative to lymphocytes. *P < .005 compared with HC.

![Figure 4. Quantification of NK cells in whole blood from different clinical groups. Whole blood from HC (n = 16), patients with HUS (n = 14), UC (n = 5), and IC (n = 6) was stained with PECy5-labeled anti-CD3 and PE-labeled anti-CD56 and analyzed by flow cytometry as described in "Patients, materials, and methods." (A) Representative dotplots from 1 child within each clinical group showing NK (CD3−CD56+) and NKT (CD3+/CD56+) subpopulations in the lymphocyte gate. Dot plot quadrants are defined as: CD3+/CD56+, top left; and CD3+/CD56−, bottom right. The percentages of the different subpopulations are shown within each quadrant. (B) Each bar represents the mean ± SEM of the percentage of NK cells (CD3−CD56+) relative to lymphocytes. *P < .005 compared with HC.

2 show that percentages and absolute counts of the more abundant CD56dim subset were significantly reduced in HUS compared with that of all other clinical groups, without variations in CD56bright cells. The selective decrease in the CD56dim subset could also be observed by analyzing the CD56dim/CD56bright ratio (HC = 10.0 ± 2.0, n = 14; HUS = 4.5 ± 1.3, n = 7; UC = 10.8 ± 2.5, n = 6; and IC = 11.0 ± 4.3, n = 6; **P < .005, HUS compared with HC). Finally, a significant reduction in the NKT population was noted only in children with HUS, both in the relative (HC = 2.0% ± 0.4%, n = 14; HUS = 0.2% ± 0.8%, n = 9; P < .001) and in the absolute number (HC = 61.7 ± 8.7 × 10^6 cells/L, n = 8; HUS = 16.0 ± 5.7 × 10^6 cells/L, n = 8; P < .001). Moreover, the percentage of CX3CR1+ cells among NKT cells was also reduced in patients with HUS (HC = 57.0% ± 5.1%, n = 14; HUS = 33.6% ± 7.7%, n = 9; P < .05).

We then evaluated whether reduction in the NK CD56dim and NKT populations may have been due to specific cell sensitivity to some hypothetical death signals present within circulating blood of children with HUS. We isolated peripheral blood Mo’s from healthy adult donors that were incubated with 80% plasma from HC or from patients with HUS. After overnight incubation, apoptosis of the specific subpopulations was assayed by flow cytometry using anti-CD56-PE and anti-CD3-PECy5 antibodies and AV-FITC, as previously described.35 We did not observe any significant enhancement in the apoptotic rate of either cell types after exposure to HUS-derived plasmas (data not shown). Similarly, no change in apoptosis of these cells was noted after exposure to 10−6 M or 10−8 M of Stx2 (data not shown).

**Immunohistochemistry**

Single labeling for the macrophage marker macrosialin (CD68) and for CX3CR1 was performed on kidney sections from children with postdiarreal and atypical HUS. We noted glomerular infiltration by CD68+ macrophages and CX3CR1+ cells in both forms of HUS (Figure 6). The presence of CD68− and CX3CR1− cells, probably corresponding to neutrophils within the glomeruli, and CX3CR1+ macrophages and/or lymphocytes within tubules of children with either form of HUS, were also observed. The number of infiltrating CD68+ and CX3CR1+ macrophages per glomerulus is shown in Table 3.

**Correlations**

Retrospectively, we analyzed the existence of correlations between the immunologic alterations found in patients with HUS and the severity of HUS, classified according to Gianantonio et al’s criteria.36 We found a weak negative correlation between severity and the percentage of total circulating CX3CR1+ leukocytes.
Among Mo subsets, the severity of HUS was positively correlated with CX3CR1^+ monocytes (r = .66, P = .027, n = 13), and negatively correlated with CX3CR1^+/CD62L^+ (r = -.67, P < .001, n = 13). Moreover, the expression of CD62L on both CX3CR1^+ and CX3CR1^+ monocytes negatively correlated with severity (r = -.62, P = .022, n = 13; and r = -.56, P = .045, n = 13, respectively). In addition, there was no correlation between severity and CX3CR1^+/CD56^+ cells.

Discussion

The main finding of this study was the dramatic decrease of circulating CX3CR1^+ monocytes and NK cells in children with HUS. This observation could be related to a unique phenomenon that involves all leukocyte populations or could be due to different effector and/or regulatory mechanisms for each individual cell type. Two general processes may explain the decrease of a specific leukocyte subset defined by a particular membrane receptor (CX3CR1): (1) regulatory mechanisms at the cellular level, which imply down-regulation of its membrane exposure or synthesis; or (2) disappearance from the circulation by specific interaction with its ligand (FKN, CX3CL1). Different mechanisms could underlie both processes. Modulatory cell pathways include enzymatic cleavage and/or impairment of the biosynthetic machinery via enhancement of the degradation rate, and/or enhanced internalization. On the other hand, specific recruitment of CX3CR1^+ cells by soluble or membrane-bound FKN in tissues may explain a specific cell disappearance from circulation. Although further experimental models or in vitro studies must be designed to ascertain precisely whether one or more of these mechanisms are operating during HUS, valuable information can be obtained from the present study.

The remarkable changes observed in Mo’s from patients with HUS can be summarized as a significant reduction in CX3CR1^+ Mo’s and a down-modulation of CD62L together with an enhanced expression of the CD16 molecule. However, the most distinctive feature of Mo’s from HUS is the loss of the CX3CR1. In fact, Mo subset classification according to CX3CR1 and CD62L expression clearly showed that children with HUS present a unique Mo phenotype distinctive from all other clinical groups. We have demonstrated that children with HUS showed leukocytosis based on increased absolute counts of neutrophils and monocytes (Table 2) and, furthermore, that the increase of total Mo’s results from the enhancement in the relative and absolute number of CX3CR1^− Mo subsets. This altered distribution within Mo subsets observed in patients with HUS could be due to differentiation or modulatory changes of “classical Mo’s” (CX3CR1^+ /CD16^− /CD62L^−) through the acquisition of CD16 and the loss of CD62L and CX3CR1. In this regard, enhancement in the percentage of CD16^+ Mo’s has been demonstrated after in vitro stimulation of human peripheral blood Mo’s with IL-10, GM-CSF plus IL-4, and plasma from patients with HUS. The selectin CD62L (L-selectin), which mediates the initial attachment of leukocytes to endothelial cells, is shed following cell activation and its down-regulation has been proposed to participate

Table 3. Immunohistochemistry from renal biopsies

<table>
<thead>
<tr>
<th>Age, y</th>
<th>Change in time, d</th>
<th>No. cells/glomerulus</th>
</tr>
</thead>
<tbody>
<tr>
<td>Normal kidney</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TBD</td>
<td>5</td>
<td>—</td>
</tr>
<tr>
<td>TBD</td>
<td>8</td>
<td>—</td>
</tr>
<tr>
<td>TBD</td>
<td>20</td>
<td>—</td>
</tr>
<tr>
<td>TBD</td>
<td>25</td>
<td>—</td>
</tr>
<tr>
<td>Postdiarrheal HUS</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TBD</td>
<td>0.75</td>
<td>30</td>
</tr>
<tr>
<td>TBD</td>
<td>2</td>
<td>&lt; 7†</td>
</tr>
<tr>
<td>TBD</td>
<td>2.5</td>
<td>300</td>
</tr>
<tr>
<td>TBD</td>
<td>3</td>
<td>21</td>
</tr>
<tr>
<td>TBD</td>
<td>3</td>
<td>7</td>
</tr>
<tr>
<td>TBD</td>
<td>4</td>
<td>21</td>
</tr>
<tr>
<td>TBD</td>
<td>6</td>
<td>NA</td>
</tr>
<tr>
<td>TBD</td>
<td>6</td>
<td>30</td>
</tr>
<tr>
<td>Atypical HUS</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TBD</td>
<td>1</td>
<td>NA</td>
</tr>
<tr>
<td>TBD</td>
<td>6</td>
<td>NA</td>
</tr>
<tr>
<td>TBD</td>
<td>7</td>
<td>Chronic</td>
</tr>
<tr>
<td>TBD</td>
<td>8</td>
<td>Chronic</td>
</tr>
<tr>
<td>TBD</td>
<td>8</td>
<td>&lt; 7</td>
</tr>
<tr>
<td>TBD</td>
<td>8</td>
<td>&lt; 7</td>
</tr>
<tr>
<td>TBD</td>
<td>16</td>
<td>Chronic‡</td>
</tr>
<tr>
<td>Acute rejection</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TBD</td>
<td>3</td>
<td>—</td>
</tr>
<tr>
<td>Systemic lupus</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TBD</td>
<td>12</td>
<td>—</td>
</tr>
<tr>
<td>TBD</td>
<td>16</td>
<td>—</td>
</tr>
<tr>
<td>Henoch—Schoenlein purpura</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TBD</td>
<td>6</td>
<td>—</td>
</tr>
<tr>
<td>TBD</td>
<td>8</td>
<td>—</td>
</tr>
</tbody>
</table>

Each row beneath the boldfaced headings indicates a different child within that category.

*NA indicates not available; and —, not applicable.
†Cortical necrosis.
‡Recurrence after transplantation.
in the occurrence of leukocytosis.\textsuperscript{31} Down-modulation of CX\textsubscript{3}CR1 in monocytes has not been previously reported.

Alternatively, CX\textsubscript{3}CR1\textsuperscript{+} Mo’s could be retained in inflamed tissues that express high levels of the CX\textsubscript{3}CR1 ligand, FKN, or CX\textsubscript{3}CL1. In this regard, immunohistochemistry detection of CD68\textsuperscript{+} macrophages and CX\textsubscript{3}CR1\textsuperscript{+} leukocytes in kidney sections from patients with HUS supports the concept that these cells could be locally retained. Bazan et al have also reported that FKN induces the migration of Mo’s,\textsuperscript{13} and recently, Chapman et al\textsuperscript{44} and Ancuta et al\textsuperscript{45} reported that CX\textsubscript{3}CR1\textsuperscript{+} Mo’s undergo both transendothelial migration in response to (s)FKN and efficient binding to FKN-expressing cells.

As a component of the innate immune system in circulation, the other CX\textsubscript{3}CR1-carrying population, NK cells, may be also attracted to areas of inflammation.\textsuperscript{34} In particular, CD56\textsuperscript{high} NK cells are strongly attracted by IL-8 and soluble CX\textsubscript{3}CL1, since they express high levels of CXCR1 and CX\textsubscript{3}CR1.\textsuperscript{34} On the contrary, CD56\textsuperscript{bright} NK cells show little or no response to these chemokines, supporting a preferential modulatory role for this NK subset during the early stages of an immune response.\textsuperscript{53} Besides, circulating NKT cells are a more heterogeneous population, and nearly 50\% of them express CX\textsubscript{3}CR1.\textsuperscript{34}

The reduction observed specifically in the percentage of CD56\textsuperscript{dim} NK cells and NKT cells in children with HUS, without a specific induction of the apoptotic rate of these populations, suggests that these cells are being retained on the endothelium or within tissues. The contribution of this phenomenon to endothelial and/or epithelial damage is still uncertain. However, systemic toxicity in patients treated with IL-2 has been related to the selective disappearance of circulating NK and NKT cells upon massive adherence on activated endothelium.\textsuperscript{46-48} Moreover, adult HUS has been reported as secondary to IL-2 treatment of metastatic melanoma.\textsuperscript{49} Although autologous endothelial cells are considered to be protected from NK cell–mediated lytic attack by the expression of MHC class I molecules, viruses, bacteria, or IL-2 administration may directly or indirectly activate endothelial cells, rendering them more susceptible to NK lysis.\textsuperscript{50} During STEC infections, the direct action of Stx and/or inflammatory mediators released by Mo’s and NK cells activates endothelial cells.\textsuperscript{51,52} In this scenario, the expression of FKN could collaborate with the adherence of CX\textsubscript{3}CR1-expressing cells, Mo’s, and NK cells, which in turn would produce cytotoxic factors, enhancing endothelial damage. Moreover, FKN expression is significantly enhanced in response to TNF, IL-1, and IFN-\gamma.\textsuperscript{13,15,53,54} and circulating levels of many cytokines are abnormally increased in patients with HUS.\textsuperscript{55,56}

Finally, we could not rule out that soluble mediators may be inducing a downmodulation or shedding of CX\textsubscript{3}CR1 from the membrane of Mo’s and NK cells in patients with HUS, as shown for NK cells upon activation by CD16 engagement\textsuperscript{14,57} or IL-15.\textsuperscript{58} We have measured (s)FKN in the plasma of children with HUS enrolled in a previous study\textsuperscript{b} (data not shown). Since there were large interindividual variations among healthy children or adults (range, 0-7498 pM), we could not obtain any meaningful interpretation.

Typical HUS occurs after bacterial enteritis; however, distinctive changes occurred in these patients compared with the group of IC. Although IC showed an expanded CD16\textsuperscript{+} Mo subset, the percentage of CX\textsubscript{3}CR1\textsuperscript{+} leukocytes and Mo’s, NK and NKT populations, and the mean expression of CD62L were within the normal range. On the other hand, since UC presented Mo and NK phenotypes similar to that of HC, it is unlikely that the changes noted in HUS resulted from the azotemia and metabolic perturbations per se associated with acute renal failure.

Most important, correlation studies indicated that a lower percentage of circulating CX\textsubscript{3}CR1\textsuperscript{+} leukocytes or CX\textsubscript{3}CR1\textsuperscript{+} Mo’s was associated with a longer duration of anuria and a poorer prognosis. Similarly, a lowered expression of CX\textsubscript{3}CR1\textsuperscript{+} NK cells has been associated with an increased disease activity in patients with multiple sclerosis.\textsuperscript{59} Based on the background discussed, we speculate that early Stx-mediated endothelial injury activates leukocytes with a high cytotoxic potential, and leads to a close leukocyte/endothelial cell interaction through the FKN pathway. This process will determine the specific inflammatory phenotype observed in patients with HUS, and the exacerbation of endothelial damage by cells of the innate immune system.

Acknowledgments

The authors thank Marta Felippo, Nora Galassi, and Norma Riera for their excellent technical assistance. The authors also thank Fundación de la Hemofilia and Academia Nacional de Medicina for the use of the FACSflow cytometer, and the Department of Hemotherapy of Centro de Educación Médica e Investigaciones Clínicas “Norberto Quiró” (CEMIC) for normal blood samples. The authors thank Anna Proietti, research nurse, for her assistance. The authors also want to thank le Comité de Recherché Cliniques, Ste-Justine Hospital Research Centre.

Supported by grants from Fundación Alberto J. Roemers, and Agencia Nacional de Promoción Científica y Tecnológica, Argentina (M.S.P.).

Authorship

Author contributions: M.V.R. and G.C.F. performed phenotypic studies on circulating leukocytes from patients and control groups; R.E., I.G., C.E.C., and G.V. selected patients and control children, obtained the informed consent from their parents and the approval from ethics committee, collected blood samples, and followed clinical and renal evolution; N.P. collected renal samples and performed immunohistochemistry on renal tissues; P.S. and M.C.S. collaborated in phenotypic studies on natural killer cells; H.T. collected urine samples for soluble fractalkine measurements as PI of the Synsorb-Pk clinical trial; C.C. performed ELISA for measurements of soluble fractalkine; F.P. codeigned this study with M.S.P., obtained approval from ethics committee, and collected samples for measurements of soluble fractalkine in plasma; M.S.P. designed and conducted immunologic and phenotypic studies and wrote the manuscript.

Conflict-of-interest statement: The authors declare no competing financial interests.

Correspondence: Marina S. Palermo, Academia Nacional de Medicina, Pacheco de Melo 3081, (C1425AUM), Buenos Aires, Argentina; e-mail: mspalermo@hematologia.anm.edu.ar.

References


Involvement of the fractalkine pathway in the pathogenesis of childhood hemolytic uremic syndrome

María Victoria Ramos, Gabriela C. Fernández, Natasha Patey, Pablo Schierloh, Ramón Exeni, Irene Grimoldi, Graciela Vallejo, Christian Elías-Costa, María del Carmen Sasaiain, Howard Trachtman, Christophe Combadière, François Proulx and Marina S. Palermo