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Cutting Edge: IFN-γ Is a Negative Regulator of IL-23 in Murine Macrophages and Experimental Colitis

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Abstract

IL-23 regulation is a central event in the pathogenesis of the inflammatory bowel diseases. We demonstrate that IFN- γ has anti-inflammatory properties in the initiation phase of IL-23–mediated experimental colitis. IFN- γ attenuates LPS-mediated IL-23 expression in murine macrophages. Mechanistically, IFN- γ inhibits *Il23a* promoter activation through altering NF- κ B binding and histone modification. Moreover, intestinal inflammation is inhibited by IFN- γ signaling through attenuation of *Il23a* gene expression. In germ-free wild-type mice colonized with enteric microbiota, inhibition of colonic *Il23a* temporally correlates with induction of IFN- γ . IFN- γ R1/IL-10 double-deficient mice demonstrate markedly increased colonic inflammation and *IL23a* expression compared with those of IL-10^{-/-} mice. Colonic CD11b⁺ cells are the primary source of IL-23 and a target for IFN- γ . This study describes an important anti-inflammatory role for IFN- γ through inhibition of IL-23. Converging genetic and functional findings suggest that IL-23 and IFN- γ are important pathogenic molecules in human inflammatory bowel disease.

The inflammatory bowel diseases (IBDs) result from inappropriately directed inflammatory responses to the enteric microbiota in a genetically susceptible host. Key participants in the innate immune response to the enteric microbiota are macrophages (1). Of the inflammatory genes induced in macrophages, IL-12 family members play a central role in mediating intestinal inflammation. IL-12 and IL-23 are heterodimeric cytokines composed of a common p40 subunit (*II12b*) and a p35 and p19 (*II23a*) subunit, respectively (2). Recently IL-23 has been strongly implicated in the pathogenesis of human IBD (3).

IL-23, unlike IL-12, promotes a distinct CD4⁺ T cell phenotype characterized by the production of the cytokine IL-17, denoted Th17 cells. IL-23 enhances Th17 function and survival by acting on differentiated Th17 cells that express the IL-23 receptor. Development of Th1, Th2, and Th17 cells are mutually exclusive because differentiation of one subset is inhibited by the presence of another (4). Indeed, IFN- γ , the signature Th1 cytokine induced by IL-12, inhibits Th17 development (4). IFN- γ strongly synergizes with bacterial products to activate and sustain production of IL-12 by dendritic cells and macrophages (5). Importantly, IFN- γ as a proinflammatory cytokine has been implicated in the pathogenesis of multiple chronic

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Disclosures

The authors have no financial conflicts of interest.

inflammatory conditions, including IBD (1). In this study, we demonstrate that IFN- γ is a negative regulator of IL-23 in murine macrophages and experimental colitis.

Materials and Methods

Mice

Wild-type (WT), IL-10^{-/-}, and IFN- γ R1^{-/-} mice on the C57BL/6 background were matched for age in all experiments. IFN- γ R1/IL-10^{-/-} mice were obtained by crossing IL-10^{-/-} and IFN- γ R1^{-/-} mice. Heterozygous offspring were then bred to obtain homozygous IFN- γ R1/ IL-10^{-/-} mice. Littermates were used as controls. 129S6/SvEv germ-free (GF) mice (WT and IL-10^{-/-}) were Cesarian-derived and maintained at the Gnotobiotic Facility at the University of North Carolina. Mice were housed in accordance with guidelines from the American Association for Laboratory Animal Care and Research Protocols, and experiments were approved by the Institutional Animal Care and Use Committee of the University of North Carolina.

General reagents and methods

Murine IL-12 p40, IL-12 p70, IFN- γ (R&D Systems, Minneapolis, MN) and IL-23 (eBioscience, San Diego, CA) ELISA kits were used according to the manufacturers' instructions. The *ll23a* 1.8 kb luciferase reporter plasmid was provided Y.H. Chen (University of Pennsylvania School of Medicine) (6). Quantitative real-time PCR was performed as described previously (7).

Bone marrow-derived macrophages

Bone marrow-derived macrophages (BMMs) were harvested and cultured as described (7).

Transient transfections

BMMs were transiently transfected using AMAXA Nucleofector Technology (AMAXA, Walkersville, MD) using the protocol for murine macrophages. Transfection efficiencies of >50% are routinely obtained (data not shown). Luciferase activity was determined as described (7).

Colonic tissue explant cultures

Colonic sections were processed as described previously (7). Supernatants were collected after 24 h for cytokine ELISAs.

Colonic macrophages

Lamina propria mononuclear cells were isolated from mouse colon by an enzymatic method and density gradient centrifugation, as described previously (8). Lamina propria mononuclear cells were further separated into CD11b⁺ cells using anti-CD11b microbeads (Miltenyi Biotec, Auburn, CA). Purity was >90% by flow cytometric analysis (data not shown).

Histology

Colitis scoring was performed as described (7). Histological scores were determined by a pathologist (T.R.) blinded to experimental protocols.

Chromatin immunoprecipitation assays

Chromatin immunoprecipitation (ChIP) was performed with ChIP-IT Express kit (Active Motif, Carlsbad, CA) according to the manufacturer's protocol. DNA–protein complexes were immunoprecipitated with Abs (RelA, p50, rabbit polyclonal IgG, Santa Cruz Biotechnology,

Santa Cruz, CA; acetylated core histone H4 [H4Act], Upstate Biotechnology, Lake Placid, NY). Real-time PCR primers for the *ll23a* promoter and the Nuc1 region of *ll12b* promoter were used to amplify immunoprecipitated and input DNA (diluted 10-fold).

Statistical analysis

Statistical significance from experiments in cells was determined using Wilcoxon signed-rank tests. Results were considered statistically significant if p < 0.05. Statistical significance for in vivo data was assessed by the Mann-Whitney U test (SPSS, Chicago, IL) with Bonferroni correction.

Results and Discussion

IFN-γ inhibits LPS-induced IL-23 in murine macrophages

In BMMs from C57BL/6 mice, LPS-induced IL-23 protein secretion and IFN- γ significantly inhibited LPS-stimulated IL-23 (Fig. 1*A*). IFN- γ -mediated inhibition of IL-23 was in notable contrast to enhancement of LPS-induced IL-12 p40 and IL-12 p70 by IFN- γ (Supplemental Fig. 1*A*, 1*B*). Consequently, in LPS-activated BMMs from IFN- γ R1^{-/-} mice, a loss of IFN- γ -mediated IL-23 inhibition (Fig. 1*A*) and IL-12 (Supplemental Fig. 1*A*, 1*B*) induction were observed.

Il23a (IL-23 p19) mRNA expression was rapidly induced by LPS by 1 h. IFN- γ inhibited LPS-activated *Il23a* (Fig. 1*B*) and augmented *Il12b* (IL-12 p40) expression (Supplemental Fig. 1*C*). *Il23a* expression levels returned to baseline by 6 h, whereas LPS plus IFN- γ -induced *Il12b* continued to rise (Supplemental Fig. 1*C*).

Whether effects of IFN- γ on IL-23 expression were dependent on the production of IL-10 was next determined. IL-10^{-/-} macrophages demonstrate significantly enhanced LPS-induced IL-23 secretion compared with that of WT BMMs. IFN- γ -inhibited IL-23 expression in IL-10^{-/-} BMMs, and inhibition was abrogated in IFN- γ R1/IL-10^{-/-} BMMs (Fig. 1*C*). IFN- γ -mediated synergistic induction of IL-12 p40 and IL-12 p70 (Supplemental Fig. 1*D*, 1*E*) was also abrogated in IL-10/IFN- γ R1^{-/-} BMMs.

Primary response genes have promoters that exist in an open chromatin structure and/or undergo rapid nucleosome remodeling. In contrast, secondary response genes have delayed induction kinetics, requiring ATP-dependent nucleosome remodeling and new protein synthesis prior to transcription initiation. In the presence of the protein synthesis inhibitor cyclohexamide (CHX), *Il23a* was rapidly induced by LPS in BMMs (Fig. 1*D*), characteristic of a primary response gene. CHX alone induced *Il23a* compared with vehicle-treated BMMs, as reported for other primary response genes (9). Unlike *Il23a*, *Il12b* induction is reduced in the absence of new protein synthesis, as described (9). In the presence of CHX, IFN-γ-mediated inhibition of LPS-induced *Il23a* is preserved. In contrast, *Il12b* induction by LPS and IFN-γ remains inhibited in the absence of new protein synthesis (Fig. 1*D*).

IFN-y prevents ReIA binding to the II23a promoter

Two NF- κ B sites have been reported to mediate LPS-induced *Il23a* promoter activity in murine macrophages (6). A 1.8 kb promoter luciferase reporter plasmid containing the NF- κ B sites (κ 1 –82 and κ 2 –618 from the transcription initiation site) was transiently transfected into BMMs. LPS strongly induced and IFN- γ inhibited *Il23a* promoter activity (Fig. 2*A*). Inhibition of NF- κ B transactivation by IFN- γ likely occurs at the *Il23a* promoter because IFN- γ fails to block phosphorylation of NF- κ B p65 and degradation of I κ B α (Supplemental Fig. 2).

Next, BMMs were cultured with LPS plus IFN- γ , and occupancy of RelA on the distal (κ 2) *Il23a* NF- κ B binding site was analyzed by ChIP using PCR primers that span the *Il23a* promoter sequence from -549 to -680. RelA promoter occupancy was demonstrated 1 h after LPS stimulation. IFN- γ inhibited LPS-induced RelA recruitment to the *Il23a* promoter and enhanced the recruitment of NF- κ B p50 (Fig. 2*B*, upper panels).

Histone acetylation is associated with transcriptionally active chromatin. The core histone H4 was acetylated 1 h after LPS stimulation at the distal *II23a* NF- κ B binding site. IFN- γ inhibited LPS-induced histone H4 acetylation (Fig. 2*B*, *lower panels*). In contrast, LPS plus IFN- γ stimulation was associated with histone H4 acetylation at an NF- κ B site in the *II12b* proximal promoter (Fig. 2*B*, *lower panels*). Therefore, IFN- γ may limit ReIA access to the *II23a* promoter by altering the dynamics of NF- κ B subunit recruitment and by regulating covalent histone modifications.

These results provide new insights into transcriptional inhibition of *Il23a. Il23a* expression has markedly different kinetics of induction and is regulated through notably divergent mechanisms compared with those of another NF- κ B-dependent gene, *Il12b*. Where IFN- γ potently synergizes with bacterial products for optimal induction of *Il12b* gene expression (10), IFN- γ inhibits LPS-mediated *Il23a* expression, surprisingly, through effects on NF- κ B DNA binding and histone acetylation.

Enteric microbiota induce colonic IL-23 expression in experimental colitis

IL- $10^{-/-}$ mice develop chronic intestinal inflammation mediated by IL-23 (11) and dependent on the presence of the enteric microbiota (12). We investigated the role of the enteric microbiota in the regulation of mucosal IL-23 in WT and IL- $10^{-/-}$ mice raised germ-free (GF) and transitioned to a conventionalized (CNV) specific pathogen free microbiota at 8 wk of age. Two weeks after transition, colonic explants from CNV IL- $10^{-/-}$ mice secreted significantly more IL-23 (Fig. 3A) than GF WT, GF IL- $10^{-/-}$, and CNV WT mice.

WT mice transitioned to a CNV microbiota revealed an increase in colonic *Il23a* mRNA after 3 d (Fig. 3*B*, *top panel*) that returned to baseline levels by day 14. Downregulation of colonic *Il23a* temporally correlated with increased colonic IFN- γ (*ifng*) mRNA (Fig. 3*B*, *bottom panel*). These results suggest that IFN- γ expression is a homeostatic checkpoint controlling the initiation of mucosal innate immune responses to the enteric microbiota. Consistent with these results, the enteric microbiota was recently shown to inhibit expression of IL-23 with subsequent effects on expansion and survival of Th17 cells in the colon (13).

In IL-10^{-/-} mice, 2 wk after transition to a CNV microbiota, an increase in colonic *Il23a* and *ifng* mRNA was detected (Supplemental Fig. 3A). Increased colonic expression of *Il23a* correlated with the development of intestinal inflammation (Supplemental Fig. 3B). Therefore, in IL-10^{-/-} mice, IFN- γ was insufficient to completely inhibit colonic *Il23a* expression.

IFN-γ inhibits *II23a* expression in colonic CD11b⁺ lamina propria cells from IL-10^{-/-} mice

Colonic CD11b⁺ lamina propria mononuclear cells (LPMCs) were the primary source of *Il23a* (Fig. 3*C*). IFN- γ inhibited heat-killed *Escherichia coli*-induced expression of *Il23a* in colonic CD11b⁺ WT and IL-10^{-/-} LPMCs (Fig. 3*C*), whereas *Il12b* expression was not inhibited (Supplemental Fig. 3*C*). Thus, IFN- γ and IL-10 are negative regulators of IL-23 in colonic macrophages.

Increased mucosal expression of IL-23 correlates with severity of colonic inflammation in IFN- γ R1/IL-10^{-/-} mice

To understand functional consequences of IFN- γ deficiency in the development of colitis, colonic inflammation and IL-23 expression were determined in IL-10^{-/-} and IFN- γ R1/IL-10^{-/-} mice. Eight-week-old IL-10^{-/-} mice demonstrated minimal or no inflammatory changes. However, age-matched littermate IFN- γ R1/IL-10^{-/-} mice developed significant colonic inflammation (Fig. 4*A*). Severity of colonic inflammation correlated with increased colonic *Il23a* expression (Fig. 4*B*) and IL-23 secretion (Fig. 4*C*) in colon explant cultures. There were no significant differences in colonic *Il12b* and *Il12a* expression between IL-10^{-/-} and IFN- γ R1/IL-10^{-/-} mice (Supplemental Fig. 4*A*, 4*B*). Moreover, IFN- γ /IL-10^{-/-} colonic CD11b⁺ LPMCs demonstrated increased heat-killed *E. coli*-activated *Il23a* induction compared with that of IL-10^{-/-} CD11b⁺LPMCs (Fig. 4*D*). These results suggest that a primary defect in *Il23a* inhibition in colonic macrophages may mediate the development of severe IBD in IFN- γ R1/IL-10^{-/-} mice.

These studies identify protective effects of IFN- γ in two models of experimental colitis. IL-10^{-/-} mice on a C57BL/6 background raised in a CNV environment are relatively resistant to spontaneous colitis (14). In the absence of IFN- γ signaling, development of colitis is accelerated. To specifically test the role of the enteric microbiota in colitis initiation events, GF IL-10^{-/-} mice colonized with microbiota were used. In this model, colitis is rapidly induced upon introduction of the enteric microbiota (15). A robust increase in colonic *Il23a* was observed in GF IL-10^{-/-} mice upon colonization that correlated with colitis severity. Protective properties of IFN- γ have been described in other chronic inflammatory disease models. IFN- γ gene deletion or administration of anti–IFN- γ Abs leads to increased severity of experimental autoimmune encephalomyelitis and collagen-induced arthritis (16–18). A number of mechanisms have been proposed to explain protection afforded by IFN- γ in autoimmunity. For example, IFN- γ directly inhibits Th17 differentiation (19). Beyond T cell responses, little has been described about homeostatic effects of IFN- γ on innate immunity in chronic inflammation.

This study focused on events during the initiation of colitis. By utilizing GF mice transitioned to a CNV microbiota and through studies in IL- $10^{-/-}$, IFN- γ R1/IL- $10^{-/-}$, and IRF-1/IL- $10^{-/-}$ mice, we have clarified mechanisms that may be operative at disease onset. It is possible that with long-standing inflammation, as in human IBD, other mechanisms become more relevant, and in fact IFN- γ may promote inflammation (20).

Regulation of *Il23a* is an important in vivo checkpoint to determine the subsequent T cell response. Hypothetically, as Th1 and Th17 responses are counterregulatory, IFN- γ may act upon the macrophage to attenuate Th17 responses through inhibition of IL-23. We also further implicate macrophage-derived IL-23 in the initiation of experimental colitis, highlighting the protective effects of IFN- γ signaling in IL-10^{-/-} mice. Recently, genome-wide association studies in human IBD have elucidated contributions of single nucleotide polymorphisms located in relevant genomic loci, including the *IL23R*, *IL12B* (3), and *IFNG* (IFN- γ) genes (21). Thus, converging genetic and functional findings suggest that IL-23 and IFN- γ may be important therapeutic targets in human IBD.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Abbreviations used in this paper

BMM	bone marrow-derived macrophage
ChIP	chromatin immunoprecipitation
CHX	cyclohexamide
CNV	conventionalized
GF	germ-free
H4Act	acetylated core histone H4
IBD	inflammatory bowel disease
LPMC	lamina propria mononuclear cell
WT	wild-type

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FIGURE 1.

IFN-γ negatively regulates LPS-mediated IL-23 expression in macrophages. *A*, BMMs from WT and IFN-γR1^{-/-} mice were cultured in the presence of LPS (100 ng/ml, black bars) with or without IFN-γ (10 ng/ml, white bars). Supernatants were analyzed for IL-23 by ELISA. Results are expressed as mean ± SEM of three independent experiments. *p < 0.05 versus LPS-stimulated BMMs. *B*, WT BMMs were stimulated with LPS (100 ng/ml, solid line) and IFN-γ (10 ng/ml, dash line). *Il23a* mRNA expression was quantified by real-time RT-PCR. Results are expressed as fold induction normalized to β-actin. Error bars represent mean ± SEM of three independent experiments. p < 0.05 versus LPS-stimulated BMMs. *C*, BMMs from WT, IL-10^{-/-}, and IFN-γR1/IL-10/^{-/-} mice were cultured with LPS (black bars, 100 ng/ml) plus IFN-γ (gray bars, 10 ng/ml). Supernatants were analyzed for IL-23 by ELISA. Results are expressed as mean ± SEM of three independent experiments. *p < 0.05 versus LPS-stimulated BMMs. *C*, BMMs from WT, IL-10^{-/-}, and IFN-γR1/IL-10/^{-/-} mice were cultured with LPS (black bars, 100 ng/ml) plus IFN-γ (gray bars, 10 ng/ml). Supernatants were analyzed for IL-23 by ELISA. Results are expressed as mean ± SEM of three independent experiments. *p < 0.05 versus LPS-stimulated BMMs. *D*, WT BMMs were incubated for 30 min with DMSO or CHX (5 µg/ml) and then stimulated with LPS (100 ng/ml, black bars) ± IFN-γ (10 ng/ml, grey bars). *Il23a (upper panel*) and *Il12b (bottom panel*) mRNA was analyzed by real-time RT-PCR after 1 h. Results

are expressed as fold induction normalized to β -actin. Error bars represent mean \pm SEM of three independent experiments. *p < 0.05 versus LPS-stimulated BMMs.



FIGURE 2.

IFN- γ inhibits *II23a* promoter activity. *A*, BMMs were transfected with an *II23a* promoter luciferase reporter plasmid and cultured with LPS (100 ng/ml) ± IFN- γ (10 ng/ml) for 18 h. Reporter activity is represented as luciferase units normalized to heat shock protein promoter β -galactosidase activity. Data represent mean ± SEM of three independent experiments. *p <0.05 versus LPS-stimulated *II23a* promoter. *B*, Binding of RelA, p50, and H4Act to the distal NF- κ B site on the endogenous WT *II23a* promoter and H4Act at the nucleosome 1 position of the *II12b* promoter was assessed by ChIP 1 h after incubation with LPS (100 ng/ml) ± IFN- γ (10 ng/ml). Real-time PCR was performed on anti-RelA, anti-p50, and anti-H4Act precipitated DNA samples, respectively. Results are presented as enrichment (percentage input) of RelA, p50, or H4Act DNA binding. Error bars represent mean ± SEM of three independent chromatin preparations from three independent experiments. *p < 0.05 versus LPS-stimulated BMMs



FIGURE 3.

Enteric microbiota induce colonic IL-23 expression in experimental colitis. WT and IL-10^{-/-} mice raised in GF conditions were colonized with enteric microbiota from CNV mice at 8 wk of age. *A*, IL-23 secretion at day 14 postcolonization in colonic explant cultures was determined by ELISA. *B*, Colonic *Il23a (upper panel)* and *ifng* mRNA (*bottom panel*) expression was detected by real time RT-PCR in GF WT mice (day 0) and at days 3 and 14 postcolonization. Each time point includes three individual mouse colons and is representative of three independent experiments. *C*, CD11b⁺ LPMCs were isolated from WT and IL-10^{-/-} mouse colons. CD11b⁺ LPMCs were activated with heat-killed *E. coli* ± IFN- γ (10 ng/ml). *Il23a* and β -actin mRNA expression was detected by real-time RT-PCR. Results are expressed as fold induction normalized to β -actin. Error bars represent mean ± SEM of three independent experiments. *p < 0.05 versus heat-killed *E. coli*-stimulated IL-10^{-/-} CD11b⁺ LPMCs.



FIGURE 4.

Increased colonic inflammation and expression of IL-23 in IFN γ R1/IL-10^{-/-} mice. *A*, Colitis scores of WT, IFN- γ R1^{-/-}, IL-10^{-/-}, and IFN- γ R1/IL-10^{-/-} mice at 8 wk of age. *B*, Colonic *Il23a* mRNA was examined by real-time RT-PCR. *C*, IL-23 protein in supernatants from colon explants cultures were analyzed using cytokine-specific ELISA and from WT, IFN- γ R1^{-/-}, IL-10^{-/-}, and IFN- γ R1/IL-10^{-/-} mice. Results are expressed as mean ± SEM from four to six mice per group. **p* < 0.05 versus IL-10^{-/-} mice. *D*, CD11b⁺ LPMCs isolated from WT, IFN- γ R1^{-/-}, IL-10^{-/-}, and IFN- γ R1/IL-10^{-/-} mice at 8 wk of age were activated with heat-killed *E. coli*, multiplicity of infection (10:1). *Il23a* mRNA expression was detected by realtime RT-PCR. Results are expressed as fold induction relative to unstimulated CD11b⁺ LPMCs normalized to β -actin and represent mean ± SEM of three independent experiments. **p* < 0.05 versus heat-killed *E. coli*-stimulated IL-10^{-/-} CD11b⁺ LPMCs.