Osteopontin Bridging Innate and Adaptive Immunity in Autoimmune Diseases

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Received 6 August 2016; Revised 2 October 2016; Accepted 19 October 2016

Academic Editor: Manoj K. Mishra

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Osteopontin (OPN) regulates the immune response at multiple levels. Physiologically, it regulates the host response to infections by driving T helper (Th) polarization and acting on both innate and adaptive immunity; pathologically, it contributes to the development of immune-mediated and inflammatory diseases. In some cases, the mechanisms of these effects have been described, but many aspects of the OPN function remain elusive. This is in part ascribable to the fact that OPN is a complex molecule with several posttranslational modifications and it may act as either an immobilized protein of the extracellular matrix or a soluble cytokine or an intracytoplasmic molecule by binding to a wide variety of molecules including crystals of calcium phosphate, several cell surface receptors, and intracytoplasmic molecules. This review describes the OPN structure, isoforms, and functions and its role in regulating the crosstalk between innate and adaptive immunity in autoimmune diseases.

1. Introduction

OPN is an acidic glycoprotein that, depending on its intracellular (iOPN) or extracellular (OPN) localization, is involved in inflammation by inducing cell adhesion and migration, regulating the differentiation of proinflammatory lymphocytes, and inhibiting the apoptosis of inflammatory cells. It was initially described as a bone-specific sialoprotein [1] and then as a molecule expressed in activated T cells, consequently being named the “early T cell activated gene” (ETA-1) [2, 3].

OPN is produced by a variety of cell types, such as B and T cells, natural killer (NK) cells, NKT cells, macrophages, neutrophils, dendritic cells (DC), bone cells (osteoblasts and osteocytes), breast epithelial cells, and neurons, and high expression is detected in the bone, lung, liver, brain, joints, adipose tissue, and body fluids such as blood, urine, and milk [4–6].

1.1. OPN Gene. OPN is encoded by an 8 kb gene mapping on chromosome 4q13 and composed of 7 exons; the first exon is untranslated while exons 2–7 contain the coding sequences (Figure 1). Genetic variations of the OPN gene have been described in the 5′ flanking region, exons, introns and the 3′ untranslated region (3′UTR) [7–9]. Some of these variations are associated with development and/or disease activity of several autoimmune diseases [10–13] and some of them influence OPN expression [14]. For instance, the four single nucleotide polymorphisms (SNPs) +282T>C (exon VI: rs4754), +750C>T (exon VII; rs1126616), +1083A>G (3′UTR; rs1126772), and +1239A>C (3′UTR; rs9138) are associated with three haplotypic combinations, that is, 282T-750C-1083A-1239A (haplotype A), 282C-750T-1083A-1239C (haplotype B), and 282C-750T-1083G-1239C (haplotype C), and carriers of haplotype B and haplotype C display higher OPN
serum levels and higher risk of developing several autoim-
mune diseases than haplotype A homozygotes. This effect
may be related to the higher stability of the mRNA coded by
haplotype B and haplotype C compared to that coded by hap-
lotype A [15]. Interindividual differences of OPN expression
may be also influenced by variations in the promoter region,
such as −66T>G (rs28357094) and −156G>GG (rs7687316)
SNPs, which may modulate the transcriptional activity of
the gene [15–18]. Table 1 summarizes several associations
reported between OPN SNPs and autoimmune diseases
(http://www.ncbi.nlm.nih.gov/projects/SNP/) [10, 13–16, 19–
25].

1.2. OPN Structure. The protein is composed of 314 amino
acids, rich in aspartate, glutamate, and serine residues, and
it contains functional domains for calcium binding [26].
Its molecular weight ranges from 44 to 75 kDa, depending
on alternative splicing and postranslation modifications.
Given its composition, OPN is a highly negatively charged
protein lacking extensive secondary structure that displays
8 α-helices and 6 β-sheets. Besides the full-length variant,
named OPN-a and containing all exons, two splicing variants
are OPN-b and OPN-c lacking exon 5 and exon 4, respectively
[27]. All of them use exon 2, which contains the signal
peptide, and are expected to be secreted. Another form
derives from alternative initiation of transcription, lacks the
signal peptide, and is expressed as an intracellular protein
(iOPN) (Figure 1). As both OPN and iOPN are generated
from the same mRNA, selective silencing of only one of
these isoforms is not possible. OPN deficient mice lack both
forms, while administration of OPN neutralizing antibodies
or aptamers selectively blocks only OPN [28]. Other variants
depend on several postranslational modifications, includ-
ing phosphorylation, O-linked glycosylation, sialylation, and
tyrosine sulfation [29–34].

1.3. OPN Functions. OPN functions as a free cytokine in body
fluids or as an immobilized extracellular matrix molecule in
mineralized tissue. Its pleiotropic effects are partly due to its
capacity to interact with multiple ligands including several
cell surface receptors, intracellular signaling molecules, cal-
cium, and heparin.

The binding sites to cell surface receptors include a
RGD (arginine–glycine–aspartate) motif interacting with
integrins αvβ1, αvβ3, αvβ5, αvβ6, α8β1, and a5β1 [35–37],
and a binding site for CD44, in particular for the isoform
CD44v6−v7. Moreover, thrombin cleaves OPN at a conserved
site (162RS169) and exposes a cryptic 162SVVYGLR168 motif

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**Figure 1:** Genomic, transcriptional, and protein features of OPN. The figure shows in the upper panel the genomic organization of the SPP1 gene and the relevant single nucleotide polymorphisms. OPN is transcribed with 3 splicing variants: variant a contains exons 2–7 while variants b and c lack exons 5 and 4, respectively (middle panel). OPN transcripts have also two starting points generating a secreted or/and intracellular form. Several postranslational modifications are also shown (lower panel) including phosphorylation (asterisks), glycosylation, and sulfation sites. Proteases (thrombin and matrix metalloproteinase, MMP) cleavage sites are also depicted.
interacting with the integrins α9β1, α4β1, and α4β7 [38, 39]. The RGD and the cryptic sites are located in the N-terminal fragment of OPN produced by thrombin cleavage (OPN-N), whereas the CD44 binding site is located in the corresponding C-terminal fragment (OPN-C).

The role of OPN in the crosstalk between innate and adaptive immunity (iOPN is described later) is clearly highlighted in the development of proinflammatory T helper (Th) type-1 and Th17 cells (Figure 2). By acting on macrophages, OPN upregulates interleukin- (IL-) 12 production and enhances Th1 development. By acting on Th cells, OPN induces production of IL-17 by triggering αvβ3 integrin and inhibits secretion of IL-10 by triggering CD44 [40]. By interacting with CD44 in Th cells, OPN induces hypomethylation of interferon- (IFN-) γ and IL-17a genes enhancing differentiation of Th1 and Th17 cells. In contrast, CD44 deficiency promotes hypermethylation of IFN-γ and IL-17a and hypomethylation of IL-4 gene, leading to Th2 cell differentiation [41].

We have recently demonstrated that OPN-N and OPN-C generated by thrombin-mediated cleavage display distinct functions. In T cells, OPN-N promotes IL-17 secretion, whereas OPN-C inhibits IL-10 secretion. In monocytes, secretion of IL-6 is induced mainly by OPN-N. In several cell types, including vascular endothelial cells and tumor cells, OPN-N induces migration whereas OPN-C induces adhesion. By contrast, both fragments similarly induce IFN-γ secretion in T cells and tubulogenesis in vascular endothelial cells and inhibit activation-induced cell death in lymphocytes [42].

Human OPN has two strong heparin-binding domains associated with internalization signals, which suggests that it rapidly binds to surface heparan sulfate proteoglycans to be internalized. Interestingly, the thrombin cleavage site is located close to one of these heparin-binding domain. Thus, heparin binding to OPN blocks the access of thrombin and maintains OPN in the full-length form. Furthermore, the N-terminal of OPN is dominated by acidic, negatively charged amino acids, whereas all of the positively charged heparin-binding domains are on the C-terminal part [43, 44].

The biological activity of OPN can be modulated also by cleavage mediated by several matrix metalloproteinases (MMPs), including MMP-1, MMP-2, MMP-3, MMP-8, MMP-9, MMP-10, MMP-11, MMP-12, MMP-13, MMP-14, and MMP-25. These are increased in biological samples of patients with autoimmune diseases, and the majority of them have a detrimental role since their activity inhibit cell adhesion and/or the migration driven by OPN [45, 46]. Human OPN contains three cleavage sites for MMPs located between the amino acids Gly146-Leu167, Ala201-Tyr202, and Asp210-Leu211. Interestingly, MMP-12 processes OPN into a less-inflammatory form, or, alternatively, it generates OPN peptides with anti-inflammatory properties [47].

Other mechanisms controlling OPN functions are post-translational modifications, including O-linked glycosylation, sialylation, phosphorylation, and tyrosine sulfation, which influence each other and complicate the functional study on OPN and its variants [48]. For instance, O-GlcNAcylation antagonizes phosphorylation in terms of abundance, protein distribution, and activity of the protein [49]. Moreover, reduction of sialylation may prevent OPN from binding to cell surface receptors [31].

OPN displays substantial O-glycosylations, while the level of N-glycosylation seems to be low. Accurate analysis of O-glycosylation on human OPN detected multiple complex and heterogeneous glycans and α (2-3) sialic acids and a novel O-glycosylation site (SI46) in the SVVYGLR domain which might regulate the interaction of OPN with integrins [50, 51]. Additionally, steric hindrance of glycan structures on SI46 may impaire the OPN cleavage mediated by MMPs. Other O-glycosylation sites are located in the C-terminus of OPN, which binds to CD44 and may regulate various cellular events, including binding interactions [52].

OPN displays 26 phosphorylation sites which may interplay with the O-glycosylation sites [53]. Nevertheless, one new phosphorylation site, Y209, was detected. Interestingly, among the identified O-glycopeptides, aa205–225, aa234–252, and aa286–298 were also identified as phosphorylation sites, indicating that interplay between O-glycosylation and phosphorylation on these sites may occur in OPN.

OPN phosphorylation seems to play regulatory roles in bone mineralization, OPN-receptor interactions, and tumor metastasis [54].

OPN regulates bone mineralization and has diverse effects on hydroxypatite (HA) formation and growth depending on the extent of phosphorylation, since phosphorylated OPN can bind to Ca2+ and Mg2+ which are essential bone components. Phosphorylated OPN from bone inhibits hydroxypatite mineral deposition in both cell-free systems [55–58] and in cell cultures [59–62]. In contrast the highly phosphorylated milk OPN promotes mineralization in solution [63]. Moreover, in OPN-KO mice, mineralization is enhanced in bone, calcified cartilage [64], vasculature [65],

| SNP        | rs       | Located | Autoimmune disease                | References |
|------------|----------|---------|-----------------------------------|------------|
| −156 G>GG  | Rs7687316| Promoter| MS, TID, SLE                      | [15, 16, 19]|
| −66 T>G    | Rs28357094|         | TID, CD, SLE                      | [10, 19, 20]|
| +282 T>C (8090TH) | Rs4754 | Exon VI | MS, SLE, ALPS/DALD, RA, CD   | [10, 14, 21]|
| +750 C>T (+707; 9250TH) | Rs1126661 | Exon VII | SLE^6, PBC, MS, CD, ALPS/DALD | [10, 14, 21–23, 25]|
| +1083 A>G (9583RD) | Rs1126772 | 3’UTR | BD, SLE^*, MS, ALPS/DALD, CD | [10, 13, 14, 16, 21, 24, 25]|
| +1239 A>C  | Rs9138   | 3’UTR   | CD, TDM1, MS, SLE, ALPS/DALD    | [10, 13–16]|

^4 A recent meta-analysis found an association only in Asian but not in European SLE patients (4) and no association with SLE patients (4) [25].

^5 SNPs may have different numbering depending on the transcript analysed and the assembly used.
iOPN in T follicular helper (TFH) cells, since inducible T cell costimulator (ICOS) signaling induces iOPN interaction with the PI3K p85α regulatory subunit, followed by translocation into the nucleus and binding to B cell lymphoma (Bcl-6) (involved in TFH differentiation) protecting it from proteasome-mediated degradation [73]. Moreover, at perimembrane regions, iOPN colocalizes with CD44-ERM (ezrin-radixin-moesin) actin complexes, and it is involved in cell motility [74]; this activity seems to play a key role in the osteoclast function [75, 76].

2. OPN and Autoimmune Diseases

In the past thirty years, OPN has attracted attention following observations that high levels of OPN can be detected in several autoimmune diseases, like systemic lupus erythematosus (SLE) [77, 78], multiple sclerosis (MS) [79], rheumatoid arthritis (RA) [80], and others [14, 81]. In line with these observations, transgenic overexpression of OPN in a nonautoimmune background causes accumulation of B cells, hypergammaglobulinemia, and production of anti-DNA antibodies, which is typical of SLE [82]. Moreover, OPN deficient mice are relatively protected against MS [83], RA [84], type 1 diabetes mellitus (TIDM) [85], autoimmune uveitis [86], autoimmune hepatitis [87, 88], intestinal bowel disease (IBD) [89], and Sjögren’s syndrome (SS) [90, 91].

The detrimental activity of OPN in autoimmune diseases may involve its ability to promote secretion of IL-17 and IFN-γ in T cells and IL-6 in monocytes, to promote lymphocyte adhesion and migration, to inhibit activation-induced cell death that is involved in the switching off the immune response and to support TFH differentiation (Figure 2).

Genetic investigations, including genome-wide association studies (GWAS), have identified numerous, replicable,
2.1. Systemic Lupus Erythematosus. SLE is a complex autoimmune disease characterized by production of autoantibodies (autoAbs) against nuclear, cytoplasmic, and cell surface molecules that transcend organ-specific boundaries. Tissue deposition of antibodies or immune complexes induces inflammation and subsequent injury of multiple organs and finally results in clinical manifestations of SLE, including glomerulonephritis, dermatitis, thrombosis, vasculitis, and arthritis [96, 97]. The first evidence of a relationship between OPN and SLE was reported in MRL\(^{lpr/lpr}\) mice, developing a disease partly resembling SLE [98]. MRL\(^{lpr/lpr}\) mice were found to carry a loss-of-function mutation of the FAS gene, which was followed by identification of the corresponding human disease, which was named autoimmune lymphoproliferative syndrome (ALPS). MRL\(^{lpr/lpr}\) mice and ALPS patients are characterized by accumulation of polyclonal lymphocytes in the secondary lymphoid organs, expansion of a peculiar subset of T cells expressing TCRαβ but not CD4 or CD8 (named double negative T cells) and autoimmune manifestations. MRL\(^{lpr/lpr}\) mice, but not ALPS patients, produce anti-DNA autoAbs and develop vasculitis, arthritis, and glomerulonephritis causing fatal renal failure [99–101]. In both MRL\(^{lpr/lpr}\) mice and ALPS patients, the disease is due to decreased function of the proapoptotic FAS receptor involved in switching off the immune response. Most ALPS patients carry an inherited or somatic loss-of-function mutation of FAS. Rare ALPS patients carry mutations of FASLG coding for FAS-ligand, or CASP10 coding for caspase 10 involved in FAS signaling: a mutation of FASLG is carried also by MRL\(^{gld/gld}\) mice showing a disease similar to that displayed by MRL\(^{lpr/lpr}\) mice. However, a substantial proportion of ALPS patients display a defective function of FAS in the absence of known mutations, which is a pattern shown also by patients with Dianzani autoimmune lymphoproliferative disease (DALD) displaying lymphoproliferation and autoimmune manifestations but lacking DN T cell expansion [102, 103].

In mice, CD4\(^+\) CD8\(^−\) T cells expressed high levels of OPN.Overexpression of OPN in MRL\(^{lpr/lpr}\) mice induces B cell activation and IgG and IgM production, elevated autoantibodies levels (including autoAbs to double-stranded (ds) DNA), and increased cytokine expression (TNF-α, IFN-γ, and IL-1β) [12, 82, 97]. In MRL\(^{lpr/lpr}\) mice, OPN upregulation begins at the onset of the autoimmune response and positively correlates with the symptom severity [86], which is also influenced by allelic differences in OPN gene [12, 104], since development of glomerulonephritis is favored by the OPN variant carried by MRL strain but not by the one carried by the C3H strain. These allelic variants correlate with different levels of antibody production, tumor necrosis factor- (TNF-) α, IL-1β and IFN-γ expression, and macrophage activation [12]. Similar findings are reported for patients with ALPS or DALD, who show increased levels of serum OPN and an increased risk of developing the diseases in subjects carrying the OPN haplotype B or haplotype C causing production of high levels of OPN. We proposed that high levels of OPN may contribute to the disease by inhibiting lymphocyte apoptosis, worsening the defect due to defective FAS function [14]. In particular, OPN directly inhibits activation-induced lymphocyte apoptosis and promotes secretion of TIMP1 [105] and IL-17 [106], which, in turn, inhibit both Fas-induced and activation-induced lymphocyte apoptosis.

Increased levels of OPN have been reported also in the sera and plasma of SLE patients [16, 84, 107], and their use has been suggested in monitoring SLE severity [107]. In line with these observations, a prospectivestudy suggested that high plasma levels of OPN may be predictors of poor outcome and are associated with the presence of autoAbs anti-ds-DNA and elevated IFN-α levels in the serum [108]. Moreover, high OPN levels in the serum and glomeruli are associated with renal damage, possibly mediated by the OPN’s ability to support secretion of Th1 and Th17 proinflammatory cytokines and inflammatory cell migration and activation [107, 109, 110].

Recently a meta-analysis revealed that the OPN level was significantly higher in SLE patients and particularly in those with renal disease [25]. Moreover, it showed a trend of positive correlation between OPN levels and the systemic lupus erythematosus disease activity index (SLEDAI) [25]. The relationship between OPN and SLE has been confirmed by genetic studies showing correlations between OPN polymorphisms and development of SLE and suggesting that OPN may participate in a complex network of gene-gene and gene-environment interactions accounting for the SLE clinical heterogeneity. In line with this view, OPN polymorphisms have been correlated with specific clinical features of the disease, such as thrombocytopenia and hemolytic anemia, renal disease and opportunistic infections, lymphadenopathy, and high serum levels of IgE. Moreover, a correlation has been found with high serum levels of IFN-α, which is a key cytokine in SLE pathogenesis [16, 20, 22, 111, 112]. Some studies have detected a predominant influence of some polymorphisms on SLE development in males, who account for about 10% of SLE patients, which suggests that the OPN effect may be highlighted in the absence of strong SLE-promoting factors acting on females [113]. However, other reports have
detected the same polymorphism as SLE-promoting factors in females and particularly in the young ones [16, 111], which may be ascribed to OPN's role in long bone remodeling during adolescence [114]. Other studies investigated OPN 9250C/T (rs1126616) polymorphism as SLE susceptibility variants, in association with OPN levels and clinical outcome. Authors found that the frequency of TT genotypes was higher in SLE patients with nephritis compared to controls, suggesting that the CT and TT genotypes could be risk factors for SLE [115]. Recently, Lee and Song conducted a meta-analysis on the role of OPN in SLE. They found that elevated OPN levels positively correlated with SLE activity and demonstrated a significant association between OPN 1239C/A, 9250C/T polymorphisms, and SLE susceptibility [25].

Interestingly, OPN in kidneys may also be secreted by senescent myofibroblasts and drive glomerular fibrosis [116]. OPN is required for myofibroblasts differentiation, and it regulates the molecules involved in tissue fibrosis. High levels of glomerular OPN are associated with macrophage accumulation and progressive renal injury in an antiglomerular basement membrane (GBM) glomerulonephritis (GN) model [116, 117]. OPN may modulate also angiotensin-II- (AII-) induced inflammation, oxidative stress, and fibrosis of the kidney [118]. Moreover, studies showed a localization of OPN at the origin of the fibrotic lesion in Bowman's capsule and the OPN deposition colocalized to the fibrotic lesion [116].

OPN also has an important function in vascular inflammation. It acts through modulation of the proliferation, migration, and accumulation of smooth muscle and endothelial cells [116, 119]. Under injury conditions, OPN plays a regulating role in arterial mineral deposition and in atherosclerotic lesions. OPN levels are high in human atherosclerotic lesions, and in lesions of ApoE−/− mice, a model of atherosclerosis and aneurysm formation especially associated with macrophage and foam cells [120]. In vivo the function of OPN in atherosclerotic plaque formation has been proven by backcrossing ApoE−/− mice with OPN−/− mice. Double knockout mice highlighted the role of OPN in recruiting leukocytes, in macrophage apoptosis, and in reduction of AII-induced aortic aneurysm formation and MMP-2 and MMP-9 activity. These data support the idea that OPN and MMP have a role in regulation and vessel rupture [120].

2.2. Multiple Sclerosis. MS is a chronic inflammatory disease of the Central Nervous System (CNS). At onset, approximately 15% of patients display a primary progressive course (PP), whereas the others display a relapsing-remitting (RR) course, which mostly switches to a secondary progressive (SP) course within 10–15 years [121]. Analysis of the transcriptome has identified more than 50 genes highly overexpressed in MS lesions, and they included OPN [122]. High OPN levels have been reported in the serum, plasma, and cerebrospinal fluid (CFS) of MS patients and levels are higher in RR MS than in PP and SP MS, especially during the relapses. Moreover, OPN is expressed in reactive astrocytes and microglial cells in patients with RR-MS, particularly during the relapses [123–128]. The high OPN levels are positively correlated with the levels of IL-17 [129, 130]. Genetic analyses have associated variations of the OPN gene with MS [21, 131]. In this regard, we found that the variants of the OPN gene coding for a mRNA with increased stability (haplotype B and haplotype C) were associated with increased risk of MS, severe disease course, and rapid evolution of disability [132]. Furthermore, we detected a correlation between the 156G>GG SNP in the OPN promoter region and timing of disability progression and switch to the SP form [15, 132]. In Japanese patients, an OPN SNP (i.e., 9583A>G) has been associated with age of onset of disease [21]. However, other studies have not found associations between OPN variants and MS development or course [131].

Experimental autoimmune encephalomyelitis (EAE), the animal model of MS, is mediated by myelin specific autoreactive Th1 and Th17 cells. OPN deficient mice develop an attenuated form of EAE with a single relapse and lack of exacerbations or progression [79]. From the immunological point of view, they show a shift toward a Th2 cytokine profile, a reduction of proinflammatory cytokines, such as IFN-γ, and TNF-α, and increased numbers of apoptotic immune cells in the CNS lesions. Moreover, daily treatment with recombinant OPN worsened EAE in OPN+/+ mice and the effect was even more striking in OPN deficient mice. These results suggest that OPN influences the disease course not only by supporting expression of proinflammatory cytokines, but also by inhibiting apoptosis of autoreactive immune [133]. Treatment with IFN-β suppressed the production of IL-17 and OPN, through activation of signal transducer and activator of transcription (STAT)1 and suppression of STAT3 activity, and decreased spinal cord infiltration of cells secreting OPN and IL-17 [130].

Proteomic studies performed on MS lesions have detected the expression of several molecules of the coagulation cascade and administration of coagulation inhibitors, such as hirudin, decreased disease severity, and suppressed production of Th1 and Th17 type cytokines [79, 133, 134]. These effects may involve OPN, since thrombin cleavage unmasks the OPN cryptic domain interacting with α4β1 integrin. This plays a key role in the recruitment of autoimmune T cells in MS lesions and is targeted by natalizumab, a humanized monoclonal antibody active in the treatment of RR MS [135]. Moreover, α4β1-OPN interaction prevents nuclear translocation of the transcription factor forkhead box O3A (FOXO3A), blocking transcription of proapoptotic genes such as Bim, Bcl-2 homologous antagonist killer (Bak), and Bcl-2-associated X protein (Bax), and promotes degradation of Ik-B with activation of NF-kB and upregulation of antiapoptotic genes and Th1 and Th17 cytokines [136].

A modulatory effect on OPN activity may also be exerted by OPN cleavage by MMPs whose levels are increased in several autoimmune diseases [45]. In line with this view, MMP-12-deficient mice develop more severe EAE than wild type mice, and this effect disappears in OPN/MMP-12 double-deficient mice [46, 47].

It has been reported that, after spinal cord injury, OPN is expressed by microglia and correlated with cell infiltration. OPN plays a major role in attracting inflammatory cells to the injury site [137] and, for a long time, most authors highlighted
the role of OPN in exacerbating tissue damage after spinal cord injury. By contrast, Hashimoto et al. observed that the expression of Bcl-2, TNF-α, IL-1β, and IL-6 is downregulated in OPN knockout spinal cord after spinal cord injury that may result from a deficiency of OPN’s proinflammatory activity [138]. Lower cytokine expression is accompanied by reduction of the number and activity of microglia/macrophages. Moreover, KO mice showed lower Basso Mouse Scale (BMS) scores than in wild type mice. These findings suggest that OPN is beneficial for recovery from spinal cord injury and plays a neuroprotective role during inflammatory response to spinal cord injury. This is also supported by research showing an upregulation of OPN, secreted by astrocytes and microglial cells, during the demyelination and remyelination, at the site of spinal cord injury [139]. A stroke model, too, in which focal cerebral ischemia is induced by photothrombosis, has shown a neuroprotective and regenerative function of OPN, especially when it is cleaved by thrombin [140].

2.3. Rheumatoid Arthritis. RA is a chronic, disabling autoimmune disease characterized by an inflammatory attack of the joint space, leading to the invasive growth of the synovial tissue and progressive destruction of the articular cartilage and bone [141]. RA patients have a shortened life span, and they suffer cardiovascular diseases caused by accelerated atherosclerosis which are a common cause of death in these patients. In the diseased joints, several cytokines are highly expressed, including OPN together with IL-1 and TNF-α [142].

The first evidence of a relationship between OPN and RA was provided in OPN−/− mice that are protected from joint destruction in collagen-antibody-induced arthritis, an RA animal model (CAIA) [84]. High levels of OPN have been reported in the plasma and synovial fluid of RA patients [143] and have been associated with clinical severity indexes [144]. Furthermore, OPN mRNA is highly expressed in CD4+ synovial T cells and correlates with coexpression of selected OPN receptors, including αv and β1 integrin chains and CD44 [145]. Moreover, OPN plasma levels decrease after treatment with biologics or immunosuppressive drugs [145].

The effects of OPN on bone resorption are mainly ascribed to the interaction with CD44 and αvβ3 integrin expressed by osteoclasts in the site of bone erosion [146]. However, other receptors may also be involved, since blocking αvβ3 integrin by means of anti-β3 antibodies or other specific antagonists, such as SB273005 or cyclic RGD peptides, inhibits bone resorption only partly in animal models of RA [147].

Integrins α4 and β9 chains are expressed in arthritic joints and mAbs against the cryptic domain of OPN ameliorate collagen-induced arthritis by decreasing infiltration of inflammatory cells, proliferation of synovium, and development of bone erosions [148]. In line with a role of the cryptic domain, high levels of thrombin, OPN-N, and OPN-C have been detected in the synovial fluid of patients with RA. The levels of OPN-C and OPN-N correlate with the disease’s severity, and the thrombin inhibitor Hirudin ameliorates collagen-induced arthritis [149]. Moreover, the activity of OPN-N may be inhibited by thrombin-activatable carboxypeptidase B (CPB), which cleaves the C-terminal arginine from the cryptic site [150]. Interestingly, RA synovocytes express a modified form of OPN, forming a complex with fibronectin and thus exposing the cryptic domain that promotes secretion of IL-6 in B cells [151].

In RA, OPN may play a crucial role by promoting differentiation of Th17 and Th1 cells, whose levels are elevated in rheumatoid synovium and correlate with several parameters of inflammation in RA patients [152]. Genetic studies did not detect univocal associations of OPN variants with RA. Some studies did not find any association between OPN polymorphisms and susceptibility to RA nor any correlation between OPN levels and OPN genetic polymorphisms [153, 154]. By contrast, an Italian study associated a SNP in the OPN promoter region (−156G>GG) with RA susceptibility [155]. These differences may be ascribed to the different influence of other concurrent genetic or environmental factors in the different populations of patients.

A recent prospective study, conducted in biologic-naïve patients with RA, pointed out that low OPN serum levels at baseline predict clinical remission one year after initiating tocilizumab treatment but not infliximab treatment [156].

OPN is also a crucial regulator involved in osteoarthritis (OA) progression [157]. OA is the most common form of arthritis, which mainly affects older people. OPN is highly expressed in the joints of OA patients, and its levels correlate with the severity of joint lesion and inflammatory status in the OA patients [158]. Abnormal mechanical load to chondrocytes alters the composition and metabolism of articular cartilage [159], which induces the release of OPN, and the enhanced level of OPN in cartilage leads to the induction of MMP-13 [160]. MMP-13 is thought to be involved in the degradation of cartilage matrix components of type II collagen and the release of proteoglycan from cartilage tissue that promotes the development of OA [161]. Elevated levels of OPN in cartilage also activate the transcription factor NF-κB pathway involved in the production of many inflammatory factors, including chemokines and cytokines (e.g., IL-1, IL-6, IL-8, CXCL1, and CCL2) in cartilage, which leads to the spontaneous production of nitric oxide (NO), prostaglandin E2 (PGE2), IL-1β, IL-6, and IL-8. The overproduction of these cytokines and mediators exerts injurious effects on chondrocyte functions which lead to an imbalance of cartilage homeostasis resulting in progressive articular degeneration [162].

2.4. Other Autoimmune Diseases

2.4.1. Type 1 Diabetes Mellitus (T1DM). In T1DM, the autoimmune process is marked by the presence of antilglutamic acid decarboxylase (GAD), anti-islet cell, or anti-insulin antibodies, but the disease is mainly due to cell-mediated destruction of insulin-producing pancreatic β-cells [163]. This destruction is preceded by insulinitis, a massive invasion of the islets by a mixed population of lymphocytes and macrophages. OPN may play a role in this inflammation since immune responses are increased in T1DM and in diabetic patients correlated with high systolic and diastolic blood pressure, body mass index, low high density lipoprotein, diagnosis of retinopathy, and microalbuminuria.
High OPN concentrations are associated with an unfavorable metabolic profile in these patients and are strong predictors of incident diabetic nephropathy [164, 165]. SNPs analysis of the OPN gene showed that +1239C carriers displayed a significantly higher risk of T1DM than +1239A homozygotes [13]. Another SNP that can have a susceptibility role in T1DM development is located in the position 66. The G allele had significantly higher frequency in controls than T1DM patients. Interestingly, case-control comparison in males showed no significant association, whereas the association was confirmed in females [19]. Intriguingly, a screening of random peptide libraries with sera of young T1D patients detected an epitope of human OPN as an autoantigen expressed in the somatostatin cells of human islets [167].

2.4.2. Sjögren’s Syndrome (SS). SS is an autoimmune disease characterized by lymphocyte infiltration of exocrine glands but can also involve the lungs, kidneys, and nervous system. Moreover, patients with SS are predisposed to develop non-Hodgkin’s B cell lymphoma at a substantially higher rate than the general population. Recent studies suggested a role for OPN in SS pathogenesis [168, 169] since OPN levels are increased in the serum and salivary glands of patients with SS. Transgenic mice expressing OPN under the immunoglobulin enhancer/SV40 promoter spontaneously develop SS and display increased OPN levels both in the salivary glands and systemically. In tissues, OPN colocalizes with the inflammatory infiltration and is associated with reduced saliva production and increased autoantibody levels. These data have been confirmed in both the serum and submandibular salivary gland tissue using the well-characterized NOD/ShiLtJ mice, developing spontaneous SS disease in a highly predictable time frame [170]. Since OPN transgenic mice showed elevated OPN levels especially in B cell, B cell derived OPN has been speculated to play a role in SS development [90], even if a role may be played also by T cell derived OPN, and iOPN in TH1 cells [73].

2.4.3. Inflammatory Bowel Diseases (IBD). IBDs are immune-mediated diseases typically resulting from abnormal mucosal T cell response to commensal bacteria in intestine and involve chronic intestinal inflammation, mucosal damage, and epithelial barrier dysfunction. Plasma concentration of OPN is elevated in patients with ulcerative colitis (UC) and correlate with clinical activity [81]. Moreover, Crohn’s disease (CD) patients show elevated OPN expression in the terminal ileum and elevated plasma OPN levels correlate with disease activity [171–175]. In particular, in patients with active disease, the plasma OPN levels were increased compared with the quiescent disease and reduced after infliximab treatment. Genetic studies also found association of OPN haplotypes with CD susceptibility, and the combined effects of certain OPN variants may modulate IL-22 secretion [10]. However, preclinical studies showed that, in the acute phase of colitis, OPN-KO mice showed more extensive colonic ulcerations and mucosal destruction than wild type mice and the clinical phenotype was ameliorated by delivery of OPN. By contrast, in chronic dextran sulfate sodium- (DSS-) induced colitis, in which a Th1 response of the lamina propria infiltrates played a pivotal role, OPN-KO mice were protected from mucosal inflammation and produced less serum levels of IL-12 than wild type mice. Furthermore, neutralization of OPN was therapeutic in these mice. These data suggest a dual function of OPN in intestinal inflammation: during acute inflammation it activates innate immunity, reducing tissue damage and initiating mucosal repair; during chronic inflammation, it activates adaptive Th1 response, reducing inflammation [176].

2.5. Targeting OPN in Autoimmune Diseases. Patients with RA or OA spontaneously produce anti-OPN autoAbs, and their serum levels are inversely correlated with markers of disease activity [177]. These data are in line with a large body of data showing that autoAbs against inflammatory cytokines can be detected in several inflammatory diseases and suggesting that they are a physiological mechanism to counteract the pathological effects of these cytokines [178]. In line with this model, induction of EAE promotes the production of anti-OPN autoAbs, and remission occurs when their titers peak. Furthermore, DNA vaccination with a plasmid encoding OPN before EAE induction boosts the production of these autoAbs and ameliorates the course of the disease [179]. In MS, we found increased levels of anti-OPN autoAbs in RR MS patients especially in the early phases of the disease and during the remission phase. Moreover, high levels of anti-OPN autoAbs at diagnosis correlate with relented development of disability in the RR MS patients treated with immunomodulating therapy.

These data suggest that the use of antagonists of OPN may be effective in the treatment of autoimmune diseases. In line with this possibility, injection of anti-OPN antibodies ameliorates the disease in primate and mouse RA [84]. These experiments showed that this cryptic epitope is involved in leukocytes migration, cell adhesion, cytokine production, and progression of arthritis [180].Moreover, in a rat model of antiglomerular basement membrane glomerulonephritis, induced by immunizing mice with the nephritogenic T cell epitope pCol(28–40) derived from α3 chain of type IV collagen, treatment with neutralizing antibodies to OPN inhibited development of glomerular fibrosis [117].

These results obtained in animal models have prompted a recent study on safety, tolerability, pharmacokinetics, pharmacodynamic, and efficacy of a monoclonal antibody against OPN in patients with RA [180]. Patients with RA were divided into two random groups, that is, treated with a placebo or a humanized IgG1 monoclonal antibody (ASK8007) directed to the cryptic site of OPN and inhibiting both RGD- and α9β1 integrin-dependent cell binding to human OPN [180, 181]. Overall ASK8007 administration appeared safe and well tolerated up to a highest dose (20 mg/kg), but it did not induce any improvement in joint inflammation and destruction. This can be the consequence of the clinical trial design (early time of evaluation, low power of the study, and aggressive disease compared to the mild preclinical one), low affinity of the mAb for the human OPN, or the methodology used targeting OPN but not iOPN. To overcome this problem, silencing OPN expression might be a good approach since mice with collagen-induced arthritis ameliorated the inflammatory
response and bone destruction (articular swelling and cartilage erosion) in the ankle joint upon inhibition of OPN expression, by mean of lentiviral OPN short hairpin RNA [182]. Moreover, preclinical data on experimental autoimmune uveitis, in which targeting of OPN has been obtained by mean of a small interfering RNA (siRNA) [183], are encouraging.

Results showed that plasma levels of OPN and the clinical and histopathological scores of disease were lower in the siRNA-treated group than that in the control group.

3. Conclusions

OPN is multifaceted protein exerting several roles in inflammation, adaptive immunity, tissue repair, bone formation, and cell signaling. These heterogeneous activities may be ascribed to the multiple variants of OPN including those due to transcriptional, posttranscriptional, and posttranslational modifications. These variants may be variably involved in the pathogenesis of different autoimmune diseases and clarification of the role of each variants is crucial to tailor appropriate therapeutic approaches targeting this complex molecule.

Competing Interests

The authors declare that they have no competing interests.

Acknowledgments

This research was supported by the Fondazione Italiana Sclerosi Multipla (FISM, Genova 2010/R12-2011/R11), the Associazione Italiana Ricerca sul Cancro (IG 14430, AIRC, Milan), Fondazione Amici di Jean (Torino), and Fondazione Cassa di Risparmio di Cuneo (Cuneo).

References

[1] F. P. Ross, J. Chappel, J. I. Alvarez et al., “Interactions between the bone matrix proteins osteopontin and bone sialoprotein and the osteoclast integrin avf3 potentiate bone resorption,” Journal of Biological Chemistry, vol. 268, no. 13, pp. 9901–9907, 1993.
[2] A. Oldberg, A. Franzen, and D. Heinegard, “Cloning and sequence analysis of rat bone sialoprotein (osteopontin) cDNA reveals an Arg-Gly-Asp cell-binding sequence,” Proceedings of the National Academy of Sciences of the United States of America, vol. 83, no. 23, pp. 8819–8823, 1986.
[3] R. Patarca, G. J. Freeman, R. P. Singh et al., “Structural and functional studies of the early T lymphocyte activation 1 (Eta-1) gene. Definition of a novel T cell-dependent response associated with genetic resistance to bacterial infection,” The Journal of Experimental Medicine, vol. 170, no. 1, pp. 145–161, 1989.
[4] D. T. Denhardt and M. Noda, “Osteopontin expression and function: role in bone remodeling,” Journal of Cellular Biochemistry. Supplement, vol. 30–31, pp. 92–102, 1998.
[5] S. K. Ramaiah and S. Rittling, “Role of osteopontin in regulating hepatic inflammatory responses and toxic liver injury,” Expert Opinion on Drug Metabolism & Toxicology, vol. 3, no. 4, pp. 519–526, 2007.
[6] T. Uede, “Osteopontin, intrinsic tissue regulator of intractable inflammatory diseases,” Pathology International, vol. 61, no. 5, pp. 265–280, 2011.
[7] F. Giacopelli, N. Rosatto, M. T. Divizia, R. Cusano, G. Caridi, and R. Ravazzolo, “The first intron of the human osteopontin gene contains a C/EBP-beta-responsive enhancer,” Gene Expression, vol. 11, no. 2, pp. 95–104, 2003.
[8] H. Iwasaki, Y. Shinohara, Y. Ezura et al., “Thirteen single-nucleotide polymorphisms in the human osteopontin gene identified by sequencing of the entire gene in Japanese individuals,” Journal of Human Genetics, vol. 46, no. 9, pp. 544–546, 2001.
[9] S. Mochida, M. Hashimoto, A. Matsu et al., “Genetic polymorphisms in promoter region of osteopontin gene may be a marker reflecting hepatitis activity in chronic hepatitis C patients,” Biochemical and Biophysical Research Communications, vol. 313, no. 4, pp. 1079–1085, 2004.
[10] J. Glas, J. Seidner, C. Bayrle et al., “The role of osteopontin (OPN/SPP1) haplotypes in the susceptibility to Crohn’s disease,” PLoS ONE, vol. 6, no. 12, Article ID e29309, 2011.
[11] N. Barizzone, M. Marchini, F. Cappiello et al., “Association of osteopontin regulatory polymorphisms with systemic sclerosis,” Human Immunology, vol. 72, no. 10, pp. 930–934, 2011.
[12] T. Miyazaki, M. Ono, W.-M. Qu et al., “Implication of allelic polymorphism of osteopontin in the development of lupus nephritis in MRL/lpr mice,” European Journal of Immunology, vol. 35, no. 5. pp. 1510–1520, 2005.
[13] A. Chiocchetti, E. Orilieri, G. Cappellano et al., “The osteopontin gene +1239A/C single nucleotide polymorphism is associated with type 1 diabetes mellitus in the Italian population,” International Journal of Immunopathology and Pharmacology, vol. 23, no. 1, pp. 263–269, 2010.
[14] A. Chiocchetti, M. Indelicato, T. Bensi et al., “High levels of osteopontin associated with polymorphisms in its gene are a risk factor for development of autoimmunity/lupus proliferations,” Blood, vol. 103, no. 4, pp. 1376–1382, 2004.
[15] C. Comi, G. Cappellano, A. Chiocchetti et al., “The impact of osteopontin gene variations on multiple sclerosis development and progression,” Clinical and Developmental Immunology, vol. 2012, Article ID 212893, 6 pages, 2012.
[16] S. D’Alfonso, N. Barizzone, M. Giordano et al., “Two single-nucleotide polymorphisms in the 5’ and 3’ ends of the osteopontin gene contribute to susceptibility to systemic lupus erythematosus,” Arthritis and Rheumatism, vol. 52, no. 2, pp. 539–547, 2005.
[17] S. Mochida, T. Yoshimoto, S. Mimura et al., “Transgenic mice expressing osteopontin in hepatocytes as a model of autoimmune hepatitis,” Biochemical and Biophysical Research Communications, vol. 317, no. 1, pp. 114–120, 2004.
[18] F. Giacopelli, R. Marciano, A. Pistorio et al., “Polymorphisms in the osteopontin promoter affect its transcriptional activity,” Physiological Genomics, vol. 15, no. 1, pp. 87–96, 2004.
[19] R. Marciano, G. D’Annunzio, N. Minuto et al., “Association of alleles at polymorphic sites in the osteopontin encoding gene in young type I diabetic patients,” Clinical Immunology, vol. 131, no. 1, pp. 84–91, 2009.
[20] T. Trivedi, B. S. Franek, S. L. Green et al., “Osteopontin alleles are associated with clinical characteristics in systemic lupus erythematosus,” Journal of Biomedicine and Biotechnology, vol. 2011, Article ID 802581, 2011.
[21] M. Niino, S. Kikuchi, T. Fukazawa, I. Yabe, and K. Tashiro, “Genetic polymorphisms of osteopontin in association with
multiple sclerosis in Japanese patients," *Journal of Neuroimmunology*, vol. 136, no. 1-2, pp. 125–129, 2003.

[22] A. C. Forton, M. A. Petri, D. Goldman, and K. E. Sullivan, "An osteopontin (SPP1) polymorphism is associated with systemic lupus erythematosus," *Human Mutation*, vol. 19, no. 4, p. 459, 2002.

[23] K. Kikuchi, A. Tanaka, H. Miyakawa et al., "Eta-1/osteopontin genetic polymorphism and primary biliary cirrhosis," *Hepatology Research*, vol. 26, no. 2, pp. 87–90, 2003.

[24] Y. H. Lee and G. G. Song, "Correlation between circulating osteopontin level in systemic lupus erythematosus and disease activity and associations between osteopontin polymorphisms and disease susceptibility: a meta-analysis," *Lupus*, 2016.

[25] E. Källning, B. Christensen, E. S. Sørensen, T. Vorup-Jensen, and J. K. Jensen, "Osteopontin binds multiple calcium ions with high affinity and independently of phosphorylation status," *Bone*, vol. 66, pp. 90–95, 2014.

[26] M. F. Young, J. M. Kerr, J. D. Termine et al., "cDNA cloning, mRNA distribution and heterogeneity, chromosomal location, and RFLP analysis of human osteopontin (OPN)," *Genomics*, vol. 7, no. 4, pp. 491–502, 1990.

[27] J. D. Coombes, M. Swiderska-Syn, L. Dollé et al., "Osteopontin neutralisation abrogates the liver progenitor cell response and fibrogenesis in mice," *Gut*, vol. 64, pp. 1120–1131, 2014.

[28] M. Nemir, M. W. DeVouge, and B. B. Mukherjee, "Normal rat kidney cells secrete both phosphorylated and nonphosphorylated forms of osteopontin showing different physiological properties," *The Journal of Biological Chemistry*, vol. 264, no. 30, pp. 18202–18208, 1989.

[29] G. F. Weber, S. Zawaideh, S. Hikita, V. A. Kumar, H. Cantor, and S. Ashkar, "Phosphorylation-dependent interaction of osteopontin with its receptors regulates macrophage migration and activation," *Journal of Leukocyte Biology*, vol. 72, no. 4, pp. 752–761, 2002.

[30] V. Shanmugam, I. Chackalaparampil, G. C. Kundu, A. B. Mukherjee, and B. B. Mukherjee, "Altered sialylation of osteopontin prevents its receptor-mediated binding on the surface of oncogenically transformed tsB77 cells," *Biochemistry*, vol. 36, no. 19, pp. 5729–5738, 1997.

[31] B. Christensen, C. C. Kazanecki, T. E. Petersen, S. R. Ritting, D. T. Denhardt, and E. S. Sørensen, "Cell type-specific post-translational modifications of mouse osteopontin are associated with different adhesive properties," *The Journal of Biological Chemistry*, vol. 282, no. 27, pp. 19463–19472, 2007.

[32] K. Singh, M. W. DeVouge, and B. B. Mukherjee, "Physiological properties and differential glycosylation of phosphorylated and nonphosphorylated forms of osteopontin secreted by normal rat kidney cells," *The Journal of Biological Chemistry*, vol. 265, no. 30, pp. 18696–18701, 1990.

[33] B. He, M. Mirza, and G. F. Weber, "An osteopontin splice variant induces anchorage independence in human breast cancer cells," *Oncogene*, vol. 25, no. 15, pp. 2192–2202, 2006.

[34] S. Denda, U. Müller, K. L. Crossin, H. P. Erickson, and L. F. Reichardt, "Utilization of a soluble integrin-alkaline phosphate chimera to characterize integrin α 8 β 1 receptor interactions with tenasin: murine α 8 β 1 binds to the RGD site in tenasin-C fragments, but not to native tenasin-C," *Biochemistry*, vol. 37, no. 16, pp. 5464–5474, 1998.

[35] D. D. Hu, J. R. Hoyer, and J. W. Smith, "Ca 2+ suppresses cell adhesion to osteopontin by attenuating binding affinity for integrin αβ3," *The Journal of Biological Chemistry*, vol. 270, no. 17, pp. 9917–9925, 1995.

[36] Y. Yokosaki, K. Tanaka, F. Higashikawa, K. Yamashita, and A. Eboshida, "Distinct structural requirements for binding of the integrins αvβ6, αvβ3, αvβ5, α5β1 and α9β1 to osteopontin," *Matrix Biology*, vol. 24, no. 6, pp. 418–427, 2005.

[37] P. M. Green, S. B. Ludbrook, D. D. Miller, C. M. T. Horgan, and S. T. Barry, "Structural elements of the osteopontin SVVYGLR motif important for the interaction with α6 integrins," *FEBS Letters*, vol. 503, no. 1, pp. 75–79, 2001.

[38] K. Ito, S. Kon, Y. Nakayama et al., "The differential amino acid requirement within osteopontin in α4 and α9 integrin-mediated cell binding and migration," *Matrix Biology*, vol. 28, no. 1, pp. 11–19, 2009.

[39] G. Murugaiyan, A. Mittal, and H. L. Weiner, "Increased osteopontin expression in dendritic cells amplifies IL-17 production by CD4+ T cells in experimental autoimmune encephalomyelitis and in multiple sclerosis," *The Journal of Immunology*, vol. 181, no. 11, pp. 7480–7488, 2008.

[40] H. Guan, P. S. Nagarkatti, and M. Nagarkatti, "Role of CD44 in the differentiation of Th1 and Th2 cells: CD44-deficiency enhances the development of Th2 effectors in response to sheep RBC and chicken ovalbumin," *Journal of Immunology*, vol. 183, no. 1, pp. 172–180, 2009.

[41] E. Boggio, C. Dianzani, C. L. Gigliotti et al., "Thrombin cleavage of osteopontin modulates its activities in human cells in vitro and mouse experimental autoimmune encephalomyelitis in vivo," *Journal of Immunology Research*, vol. 2016, Article ID 9345495, 13 pages, 2016.

[42] G. Platzer, A. Scheldbauer, A. Chemelli et al., "The metastasis-associated extracellular matrix protein osteopontin forms transient structure in ligand interaction sites," *Biochemistry*, vol. 50, no. 27, pp. 6113–6124, 2011.

[43] Metastasis, Osteopontin, Heparin Cooling Inflammation Blog, http://coolinginflammation.blogspot.it/2008/10/metastasis-osteopontin-heparin.html.

[44] R. Agnihotri, H. C. Crawford, H. Haro, L. M. Matrisian, M. C. Havrda, and L. Liaw, "Osteopontin, a novel substrate for matrix metalloproteinase-3 (stromelysin-1) and matrix metalloproteinase-7 (matrilysin)," *The Journal of Biological Chemistry*, vol. 276, no. 30, pp. 28261–28267, 2001.

[45] H. Masuda, Y. Takahashi, S. Asai, and T. Takayama, "Distinct gene expression of osteopontin in patients with ulcerative colitis," *Journal of Surgical Research*, vol. 111, no. 1, pp. 85–90, 2003.

[46] A. G. DaSilva, L. Liaw, and V. W. Yong, "Cleavage of osteopontin by matrix metalloproteinase-12 modulates experimental autoimmune encephalomyelitis disease in C57BL/6 mice," *American Journal of Pathology*, vol. 177, no. 3, pp. 1448–1458, 2010.

[47] P. Beltrao, P. Bork, N. J. Krogan, and V. Van Noort, "Evolution of functional cross-talk of protein post-translational modifications," *Molecular Systems Biology*, vol. 9, no. 1, article 714, 2013.

[48] A. M. Butt, D. Feng, M. Idrees, Y. Tong, and J. Lu, "Computational identification and modeling of crosstalk between phosphorylation, O-β-glycosylation and methylation of FoxO3 and implications for cancer therapeutics," *International Journal of Molecular Sciences*, vol. 13, no. 3, pp. 2918–2938, 2012.
demyelinating disease," *Science*, vol. 294, no. 5547, pp. 1731–1735, 2001.
[80] S. Ohsima, N. Yamaguchi, K. Nishioka et al., "Enhanced local production of osteopontin in rheumatoid joints," *Journal of Rheumatology*, vol. 29, pp. 2061–2067, 2002.
[81] R. Mishima, F. Takeshima, T. Sawai et al., "High plasma osteopontin levels in patients with inflammatory bowel disease," *Journal of Clinical Gastroenterology*, vol. 41, no. 2, pp. 167–172, 2007.
[82] J. Iizuka, Y. Katagiri, N. Tada et al., "Introduction of an osteopontin gene confers the increase in B1 cell population and the production of anti-DNA autoantibodies," *Laboratory Investigation*, vol. 78, no. 12, pp. 1523–1533, 1998.
[83] M. Braitch and C. S. Constantinescu, "The role of osteopontin in experimental autoimmune encephalomyelitis (EAE) and multiple sclerosis (MS)," *Inflammation and Allergy — Drug Targets*, vol. 9, no. 4, pp. 249–256, 2010.
[84] K. Umoto, M. Ishijima, S. R. Ritting et al., "Osteopontin deficiency protects joints against destruction in anti-type II collagen antibody-induced arthritis in mice," *Proceedings of the National Academy of Sciences of the United States of America*, vol. 99, no. 7, pp. 4556–4561, 2002.
[85] T. Nomiyama, D. Perez-Tilve, D. Ogawa et al., "Osteopontin mediates obesity-induced adipose tissue macrophage infiltration and insulin resistance in mice," *Journal of Clinical Investigation*, vol. 117, no. 10, pp. 2877–2888, 2007.
[86] M. Kitamura, K. Iwabuchi, N. Kitaichi et al., "Osteopontin aggravates experimental autoimmune uveoretinitis in mice," *Journal of Immunology*, vol. 178, no. 10, pp. 6567–6572, 2007.
[87] W.-K. Syn, S. S. Choi, E. Liaskou et al., "Osteopontin is induced by hedgehog pathway activation and promotes fibrosis progression in nonalcoholic steatohepatitis," *Hepatology*, vol. 53, no. 1, pp. 106–115, 2011.
[88] O. Morales-Ibanez, M. Dominguez, S. H. Ki et al., "Human and experimental evidence supporting a role for osteopontin in alcoholic hepatitis," *Hepatology*, vol. 58, no. 5, pp. 1742–1756, 2013.
[89] K. Heilmann, U. Hoffmann, E. Witte et al., "Osteopontin as a two-sided mediator of intestinal inflammation," *Journal of Cellular and Molecular Medicine*, vol. 13, no. 6, pp. 1162–1174, 2009.
[90] S. Husain-Krautter, J. M. Kramer, W. Li, B. Guo, and T. L. Rothstein, "The osteopontin transgenic mouse is a new model for Sjögren’s syndrome," *Clinical Immunology*, vol. 157, no. 1, pp. 30–42, 2015.
[91] D. F. Frenzel, L. Borkner, J. Scheurmann, K. Singh, K. Scharffetter-Kochanek, and J. M. Weiss, "Osteopontin deficiency affects imiquimod-induced psoriasis-like murine skin inflammation and lymphocyte distribution in skin, draining lymph nodes and spleen," *Experimental Dermatology*, vol. 24, no. 4, pp. 305–307, 2015.
[92] J. Bentham, D. L. Morris, D. S. Cunnigham Grahame et al., "Genetic association analyses implicate aberrant regulation of innate and adaptive immunity genes in the pathogenesis of systemic lupus erythematosus," *Nature Genetics*, vol. 47, no. 12, pp. 1457–1464, 2015.
[93] E. A. Stahl, S. Raychaudhuri, E. F. Remmers et al., "Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci," *Nature Genetics*, vol. 42, no. 6, pp. 508–514, 2010.
[94] C. Cotsapas, B. F. Voight, E. Rossin et al., "Pervasive sharing of genetic effects in autoimmune disease," *PLoS Genetics*, vol. 7, no. 8, Article ID e1002254, 2011.
serum osteopontin and interferon-α by osteopontin genotype in systemic lupus erythematosus," *Genes & Immunity*, vol. 10, no. 5, pp. 487–494, 2009.

[112] Y. Tanino, N. Hizawa, S. Konno et al., "Sequence variants of the secreted phosphoprotein 1 gene are associated with total serum immunoglobulin E levels in a Japanese population," *Clinical & Experimental Allergy*, vol. 36, no. 2, pp. 219–225, 2006.

[113] S. Han, J. M. Guthridge, I. T. W. Harley et al., "Osteopontin and Systemic Lupus Erythematosus: a probable gene–gender interaction," *PLOS ONE*, vol. 3, no. 3, Article ID e0001757, 2008.

[114] D. T. Denhardt, M. Noda, A. W. O’Regan, D. Pavlin, and J. S. Berman, "Osteopontin as a means to cope with environmental insults: regulation of inflammation, tissue remodeling, and cell survival," *Journal of Clinical Investigation*, vol. 107, no. 9, pp. 1055–1061, 2001.

[115] S. Salimi, M. Noora, S. Nabizadeh et al., "Association of the osteopontin rs126616 polymorphism and a higher serum osteopontin level with lupus nephritis," *Biomedical Reports*, vol. 4, no. 3, pp. 355–360, 2016.

[116] J. Merszei, J. Wu, L. Torres et al., "Osteopontin overproduction is associated with progression of glomerular fibrosis in a rat model of anti-glomerular basement membrane glomerulonephritis," *American Journal of Nephrology*, vol. 32, no. 3, pp. 262–271, 2010.

[117] C. Zhou, J. Wu, L. Torres et al., "Blockade of osteopontin inhibits glomerular fibrosis in a model of anti-glomerular basement membrane glomerulonephritis," *American Journal of Nephrology*, vol. 32, no. 4, pp. 324–331, 2010.

[118] T. Wolak, H. Kim, Y. Ren, J. Kim, N. D. Vaziri, and S. B. Nicholas, "Osteopontin modulates angiostatin II–induced inflammation, oxidative stress, and fibrosis of the kidney," *Kidney International*, vol. 76, no. 1, pp. 32–43, 2009.

[119] C. M. Giachelli, L. Liaw, C. E. Murry, S. M. Schwartz, and M. Almeida, "Osteopontin expression in cardiovascular diseases," *Annals of the New York Academy of Sciences*, vol. 760, pp. 109–126, 1995.

[120] D. Brummer, A. R. Collins, G. Noh et al., "Angiotensin II–accelerated atherosclerosis and atherectomy formation is attenuated in osteopontin–deficient mice," *The Journal of Clinical Investigation*, vol. 112, no. 9, pp. 1318–1331, 2003.

[121] B. G. Weinschenker, "The natural history of multiple sclerosis," *Neurologic Clinics*, vol. 13, no. 1, pp. 119–146, 1995.

[122] S. A. Lund, C. M. Giachelli, and M. Scatena, "The role of osteopontin in inflammatory processes," *Journal of Cell Communication and Signaling*, vol. 3, no. 3–4, pp. 311–322, 2009.

[123] M. H. J. Vogt, L. Lopatinskaya, M. Smits, C. H. Polman, and L. Nagelkerken, "Elevated osteopontin levels in active relapsing–remitting multiple sclerosis," *Annals of Neurology*, vol. 53, no. 6, pp. 819–822, 2003.

[124] C. Sinclair, M. Mirakhor, J. Kirk, M. Farrell, and S. McQuaid, "Up-regulation of osteopontin and αvβ3-crystallin in the normal appearing white matter of multiple sclerosis: an immunohistochemical study utilizing tissue microarrays," *Neuropathology and Applied Neurobiology*, vol. 31, no. 3, pp. 292–303, 2005.

[125] M. Comabella, I. Pericot, R. Goertsches et al., "Plasma osteopontin levels in multiple sclerosis," *Journal of Neuroimmunology*, vol. 158, no. 1–2, pp. 231–239, 2005.

[126] L. Börnsten, M. Khademi, T. Olsson, P. S. Sørensen, and F. Sellebjerg, "Osteopontin concentrations are increased in cerebrospinal fluid during attacks of multiple sclerosis," *Multiple Sclerosis Journal*, vol. 17, no. 1, pp. 32–42, 2011.

[127] M. Carecchio and C. Comi, "The role of osteopontin in neurodegenerative diseases," *Journal of Alzheimer's Disease*, vol. 25, no. 2, pp. 179–185, 2011.

[128] E. M. Hur, S. Youssef, M. E. Haws, S. Y. Zhang, R. A. Sobel, and L. Steinman, "Osteopontin-induced relapse and progression of autoimmune brain disease through enhanced survival of activated T cells," *Nature Immunology*, vol. 8, no. 1, pp. 74–83, 2007.

[129] R. K. Han, Y. F. Cheng, S. S. Zhou et al., "Increased circulating Th17 cell populations and elevated CSF osteopontin and IL-17 concentrations in patients with Guillain-Barré syndrome," *Journal of Clinical Immunology*, vol. 34, no. 1, pp. 94–103, 2014.

[130] M. Chen, G. Chen, H. Nie et al., "Regulatory effects of IFN-β on production of osteopontin and IL-17 by CD4+ T Cells in MS," *European Journal of Immunology*, vol. 39, no. 9, pp. 2525–2536, 2009.

[131] S. Caliler, L. F. Baranzini, A. Swerdlin et al., "Multiple Sclerosis Genetics Group. Osteopontin polymorphisms and disease course in multiple sclerosis," *Genes and Immunity*, vol. 4, no. 4, pp. 312–315, 2003.

[132] A. Chiocchetti, C. Comi, M. Indelicato et al., "Osteopontin gene haplotypes correlate with multiple sclerosis development and progression," *Journal of Neuroimmunology*, vol. 163, no. 1–2, pp. 172–178, 2005.

[133] M. Jansson, V. Panoutsakopoulou, J. Baker, L. Klein, and H. Cantor, "Cutting edge: attenuated experimental autoimmune encephalomyelitis in Eta-1/osteopontin-deficient mice," *Journal of Immunology*, vol. 168, no. 5, pp. 2096–2092, 2002.

[134] H. Neumann, I. M. Medana, J. Bauer, and H. Lassmann, "Cytotoxic T lymphocytes in autoimmune and degenerative CNS diseases," *Trends in Neurosciences*, vol. 25, no. 6, pp. 313–319, 2002.

[135] L. Steinman, "Blocking adhesion molecules as therapy for multiple sclerosis:natalizumab," *Nature Reviews Drug Discovery*, vol. 4, no. 6, pp. 510–518, 2005.

[136] L. Steinman, "Some misconceptions about understanding autoimmunity through experiments with knockouts," *Journal of Experimental Medicine*, vol. 185, no. 12, pp. 2039–2041, 1997.

[137] A. O’Regan and J. S. Berman, "Osteopontin: a key cytokine in cell-mediated and granulomatous inflammation," *International Journal of Experimental Pathology*, vol. 81, no. 6, pp. 373–390, 2000.

[138] M. Hashimoto, D. Sun, S. R. Ritting, D. T. Denhardt, and W. Young, "Osteopontin-deficient mice exhibit less inflammation, greater tissue damage, and impaired locomotor recovery from spinal cord injury compared with wild-type controls," *Journal of Neuroscience*, vol. 27, no. 13, pp. 3603–3611, 2007.

[139] R. Selvaraju, L. Bernasconi, C. Losberger et al., "Osteopontin is upregulated during in vivo demyelination and remyelination and enhances myelin formation in vitro," *Molecular and Cellular Neuroscience*, vol. 25, no. 4, pp. 707–721, 2004.

[140] M. Rabenstein, J. Hucklenbroich, A. Willuweit et al., "Osteopontin mediates survival, proliferation and migration of neural stem cells through the chemokine receptor CXCR4," *Stem Cell Research & Therapy*, vol. 6, article99, 2015.

[141] M. D. McKee and A. Nanci, "Osteopontin at mineralized tissue interfaces in bone, teeth, and osseointegrated implants: ultrastructural distribution and implications for mineralized tissue formation, turnover, and repair," *Microscopy Research and Technique*, vol. 33, no. 2, pp. 141–164, 1996.
[142] M. G. Attur, M. N. Dave, S. Stuchin et al., “Osteopontin: an intrinsic inhibitor of inflammation in cartilage,” Arthritis and Rheumatism, vol. 44, no. 3, pp. 578–584, 2001.

[143] F. Zhang, W. Luo, Y. Li, S. Gao, and G. Lei, “Role of osteopontin in rheumatoid arthritis,” Rheumatology International, vol. 35, no. 4, pp. 589–595, 2015.

[144] S. G. Gao, K. H. Li, K. B. Zeng, M. Tu, M. Xu, and G. H. Lei, “Elevated osteopontin level of synovial fluid and articular cartilage is associated with disease severity in knee osteoarthritis patients,” Osteoarthritis and Cartilage, vol. 18, no. 1, pp. 82–87, 2010.

[145] K. Suzuki, B. Zhu, S. R. Ritting et al., “Colocalization of intraacellular osteopontin with CD44 is associated with migration, cell fusion, and resorption in osteoclasts,” Journal of Bone and Mineral Research, vol. 17, no. 8, pp. 1486–1497, 2002.

[146] S. Oshahima, H. Kobayashi, N. Yamaguchi et al., “Expression of osteopontin at sites of bone erosion in a murine experimental arthritis model of collagen-induced arthritis: possible involvement of osteopontin in bone destruction in arthritis,” Arthritis & Rheumatism, vol. 46, no. 4, pp. 1094–1101, 2002.

[147] A. M. Badger, S. Blake, R. Kapadia et al., “Disease-modifying activity of SB 273005, an orally active, nonpeptide avf3 (vitronectin receptor) antagonist, in rat adjuvant-induced arthritis,” Arthritis & Rheumatism, vol. 44, no. 1, pp. 128–137, 2001.

[148] N. Yamamoto, F. Sakai, S. Kon et al., “Essential role of the cryptic epitope SLAYGLR within osteopontin in a murine model of rheumatoid arthritis,” The Journal of Clinical Investigation, vol. 112, no. 2, pp. 181–188, 2003.

[149] M. Yamaga, K. Tsuji, K. Miyatake et al., “Osteopontin level in synovial fluid is associated with the severity of joint pain and cartilage degradation after anterior cruciate ligament rupture,” PLoS ONE, vol. 7, no. 11, Article ID e49014, 2012.

[150] S. A. Sharif, X. Du, T. Myles et al., “Thrombin-activatable carboxypeptidase B cleavage of osteopontin regulates neutrophil survival and synovioctye binding in rheumatoid arthritis,” Arthritis and Rheumatism, vol. 60, no. 10, pp. 2902–2912, 2009.

[151] Y. Take, K. Nakata, J. Hashimoto et al., “Specifically modified osteopontin in rheumatoid arthritis fibroblast-like synovioctyes supports interaction with B cells and enhances production of interleukin-6,” Arthritis & Rheumatism, vol. 60, no. 12, pp. 3591–3601, 2009.

[152] G. Chen, X. Zhang, R. Li et al., “Role of osteopontin in synovial Th17 differentiation in rheumatoid arthritis,” Arthritis and Rheumatism, vol. 62, no. 10, pp. 2900–2908, 2010.

[153] E. Urcelay, A. Martinez, A. Mas-Fontao et al., “Osteopontin gene polymorphisms in Spanish patients with rheumatoid arthritis,” The Journal of Rheumatology, vol. 32, no. 3, pp. 405–409, 2005.

[154] G. Xu, W. Sun, D. He et al., “Overexpression of osteopontin in rheumatoid synovial mononuclear cells is associated with joint inflammation, not with genetic polymorphism,” The Journal of Rheumatology, vol. 32, no. 3, pp. 410–416, 2005.

[155] F. Ceccarelli, S. D’Alfonso, C. Perricone et al., “The role of eight polymorphisms in three candidate genes in determining the susceptibility, phenotype, and response to anti-TNF therapy in patients with rheumatoid arthritis,” Clinical and Experimental Rheumatology, vol. 30, no. 6, pp. 939–942, 2012.

[156] K. Izumi, Y. Kaneko, M. Hashizume, K. Yoshimoto, and T. Takeuchi, “Baseline serum osteopontin levels predict the clinical effectiveness of tocilizumab but not infliximab in biologic-Naive patients with rheumatoid arthritis: a single-center prospective study at 1 year (the Keio First-Bio Cohort study),” PLOS ONE, vol. 10, no. 12, Article ID e0145468, 2015.

[157] C. Cheng, S. Gao, and G. Lei, “Association of osteopontin with osteoarthritis,” Rheumatology International, vol. 34, no. 12, pp. 1627–1632, 2014.

[158] S. Honsawek, A. Tanavalee, M. Sakdinakiatkoon, M. Chayanapatkul, and P. Yukanandana, “Correlation of plasma and synovial fluid osteopontin with disease severity in knee osteoarthritis,” Clinical Biochemistry, vol. 42, no. 9, pp. 808–812, 2009.

[159] T. M. Griffin and F. Guilak, “The role of mechanical loading in the onset and progression of osteoarthritis,” Exercise and Sport Sciences Reviews, vol. 33, no. 4, pp. 195–200, 2005.

[160] F.-J. Zhang, W.-B. Yu, W. Luo, S.-G. Gao, Y.-S. Li, and G.-H. Lei, “Effect of osteopontin on TIMP-1 and TIMP-2 mRNA in chondrocytes of human knee osteoarthritis in vitro,” Experimental and Therapeutic Medicine, vol. 8, no. 2, pp. 391–394, 2014.

[161] R. C. Billinghurst, L. Dahlberg, M. Ionescu et al., “Enhanced cleavage of type II collagen by collagenases in osteoarthritic articular cartilage,” The Journal of Clinical Investigation, vol. 99, no. 7, pp. 1534–1545, 1997.

[162] M. G. Attur, I. R. Patel, R. N. Patel, S. B. Abramson, and A. R. Amin, “Autocrine production of IL-1β by human osteoarthritis-affected cartilage and differential regulation of endogenous nitric oxide, IL-6, prostaglandin E2, and IL-8,” Proceedings of the Association of American Physicians, vol. 110, no. 1, pp. 65–72, 1998.

[163] D. Daneman, “Type 1 diabetes,” The Lancet, vol. 367, no. 9513, pp. 847–858, 2006.

[164] I. Barchetta, C. Alessandri, L. Bertocci et al., “Increased circulating osteopontin levels in adult patients with type 1 diabetes mellitus and association with dysmetabolic profile,” European Journal of Endocrinology, vol. 174, no. 2, pp. 187–192, 2016.

[165] M. Talat, L. Sherief, H. El-Saadany, A. Ahmed, R. Salah, and M. sakr, “The role of osteopontin in the pathogenesis and complication of type 1 diabetes mellitus in children and adolescents,” Journal of Clinical Research in Pediatric Endocrinology, 2016.

[166] D. Gordin, C. Forsblom, N. M. Panduru et al., “Osteopontin is a strong predictor of incipient diabetic nephropathy, cardiovascular disease, and all-cause mortality in patients with type 1 diabetes,” Diabetes Care, vol. 37, no. 9, pp. 2593–2600, 2014.

[167] A. Fierabracci, P. A. Biro, Y. Yiannou et al., “Osteopontin is an autoantigen of the somatostatin cells in human islets: identification by screening random peptide libraries with sera of patients with insulin-dependent diabetes mellitus,” Vaccine, vol. 18, no. 1–4, pp. 342–354, 1999.

[168] Y. H. Wang, M. S. Zhao, and Y. Zhao, “Expression of osteopontin in labial glands of patients with primary Sjögren’s syndrome,” Beijing Da Xue Xue Bao, vol. 44, no. 2, pp. 236–239, 2012.

[169] G.-L. Wu, N.-Y. Wu, T.-Y. Li, Y.-S. Fan, G.-Y. Yu, and W.-W. Lu, “Effect of Chinese herbal medicines for nourishing yin, supplementing qi, and activating blood on reproductive endocrine activity and immune functions in patients with primary Sjögren’s syndrome,” Chinese Journal of Integrative Medicine, vol. 21, no. 10, pp. 778–783, 2015.

[170] S. Cha, A. B. Peck, and M. G. Humphreys-Beher, “Progress in understanding autoimmune exocrinopathy using the non-obese diabetic mouse: an update,” Critical Reviews in Oral Biology and Medicine, vol. 13, no. 1, pp. 5–16, 2002.

[171] N. Gassler, F. Autschbach, S. Gauer et al., “Expression of osteopontin (Eta-1) in Crohn disease of the terminal ileum,” Scandinavian Journal of Gastroenterology, vol. 37, no. 11, pp. 1286–1295, 2002.

[172] J. Agnholt, J. Kelsen, L. Schack, C. L. Hvas, J. F. Dahlerup, and E. S. Sorensen, “Osteopontin, a protein with cytokine-like
properties, is associated with inflammation in Crohn's disease," *Scandinavian Journal of Immunology*, vol. 65, no. 5, pp. 453–460, 2007.

[173] J. N. Gordon and T. T. MacDonald, "Osteopontin: a new addition to the constellation of cytokines which drive T helper cell type 1 responses in Crohn's disease," *Gut*, vol. 54, no. 9, pp. 1213–1215, 2005.

[174] T. Sato, T. Nakai, N. Tamura et al., "Osteopontin/Eta-1 upregulated in Crohn's disease regulates the Th1 immune response," *Gut*, vol. 54, no. 9, pp. 1254–1262, 2005.

[175] S. Komine-Aizawa, H. Masuda, T. Mazaki, M. Shiono, S. Hayakawa, and T. Takayama, "Plasma osteopontin predicts inflammatory bowel disease activities," *International Surgery*, vol. 100, no. 1, pp. 38–43, 2015.

[176] K. Heilmann, U. Hoffmann, E. Witte et al., "Osteopontin as a two-sided mediator of intestinal inflammation," *Journal of Cellular and Molecular Medicine*, vol. 13, no. 6, pp. 1162–1174, 2009.

[177] M. Sakata, J.-I. Tsuruha, K. Masuko-Hongo et al., "Autoantibodies to osteopontin in patients with osteoarthritis and rheumatoid arthritis," *Journal of Rheumatology*, vol. 28, no. 7, pp. 1492–1495, 2001.

[178] G. Cappellano, E. Orilieri, A. D. Woldetsadik et al., "Anticytokine autoantibodies in autoimmune diseases," *American Journal of Clinical and Experimental Immunology*, vol. 1, no. 2, pp. 136–146, 2012.

[179] L. Steinman, S. Youssef, N. Van Venrooij et al., "Response to Comment on 'The influence of the proinflammatory cytokine, osteopontin, on autoimmune demyelinating disease'" *Science*, vol. 299, article 1845, 2003.

[180] M. J. H. Boumans, J. G. A. Houbiers, P. Verschueren et al., "Safety, tolerability, pharmacokinetics, pharmacodynamics and efficacy of the monoclonal antibody ASK8007 blocking osteopontin in patients with rheumatoid arthritis: a randomised, placebo controlled, proof-of-concept study," *Annals of the Rheumatic Diseases*, vol. 71, no. 2, pp. 180–185, 2012.

[181] S. Kon, Y. Yokosaki, M. Maeda et al., "Mapping of functional epitopes of osteopontin by monoclonal antibodies raised against defined internal sequences," *Journal of Cellular Biochemistry*, vol. 84, no. 2, pp. 420–432, 2002.

[182] C.-M. Su, Y.-C. Chiang, C.-Y. Huang, C.-I. Hsu, Y.-C. Fong, and C.-H. Tang, "Osteopontin promotes oncostatin M production in human osteoblasts: implication of rheumatoid arthritis therapy," *The Journal of Immunology*, vol. 195, no. 7, pp. 3355–3364, 2015.

[183] D. Iwata, M. Kitamura, N. Kitaichi et al., "Prevention of experimental autoimmune uveoretinitis by blockade of osteopontin with small interfering RNA," *Experimental Eye Research*, vol. 90, no. 1, pp. 41–48, 2010.