The Many Faces of a Monogenic Autoinflammatory Disease: Adenosine Deaminase 2 Deficiency

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Abstract
Purpose of Review We aim to describe the pathophysiology, clinical findings, diagnosis, and treatment of deficiency of adenosine deaminase 2 (DADA2).

Recent Findings DADA2 is a multi-organ disease of children and less often adults, which can present with wide-ranging manifestations including strokes, medium vessel vasculitis, hematologic disease, and immunodeficiency. Diagnosis is through detection of reduced activity level of the adenosine deaminase 2 (ADA2) enzyme and/or identification of bi-allelic mutations in the ADA2 gene. Outside of high-dose glucocorticoids, conventional immunosuppression has been largely ineffective in treating this relapsing and remitting disease. Vasculitic-predominant manifestations respond extremely well to tumor necrosis factor-α inhibition. Hematopoietic stem cell transplantation can lead to normalization of enzyme activity, as well as resolution of vasculitic, hematologic, and immunologic manifestations, although treatment-related adverse effects are not uncommon.

Summary Early detection of this disease across multiple disciplines could prevent devastating clinical outcomes, especially in genetically pre-disposed populations.

Keywords Adenosine · Adenosine deaminase 2 deficiency · Vasculitis · Polyarteritis nodosa · Monogenic disease

Introduction
Deficiency of adenosine deaminase 2 (DADA2) is a monogenic autoinflammatory disorder that affects multiple organ systems with highly variable clinical presentations. First described in 2014 by two separate groups as a mimic of polyarteritis nodosa (PAN), there have been over 260 cases of DADA2 identified to date [1••, 2••]. Abnormalities in adenosine breakdown play a key role in the pathogenesis.

Pathophysiology of DADA2
Adenosine is formed by the breakdown of adenine nucleotides, with increased concentrations seen in the setting of cellular damage [3, 4]. The effect of adenosine is mediated by four G protein-coupled cell surface receptors (A₁, A₂A, A₂B, and A₃), present on many different cell types [5••]. Receptor activation triggers either a decrease (A₁ and A₃) or increase (A₂A and A₂B) in intracellular cyclic AMP, which then mediates a change in cellular activation [3]. The varying effect of adenosine on cellular activity and local inflammation seems to be determined by the interplay between the local adenosine concentration, degree of cellular receptor expression, receptor type, and the receptor affinity; most of which can be affected by environmental factors [5••]. Drugs that manipulate the adenosine pathway can be effective in inflammatory diseases [4].

The adenosine deaminases catalyze the deamination of adenosine to inosine, and deoxyadenosine to deoxyinosine, a key part of the purinergic pathway [6••] (Fig. 1). There are two primary isoforms of adenosine deaminase in humans, adenosine deaminase (ADA) 1 and 2. The major adenosine deaminase in humans is ADA1, a 40-kDa monomer present in
almost all cells, functioning to reduce levels of adenosine in the intracellular space, as high levels are cytotoxic [6]. Known as severe combined immunodeficiency (SCID), loss-of-function mutations in the ADA1 gene lead to increased apoptosis of T and B cells, resulting in recurrent infections [7]. ADA2 is present only at low levels in the physiologic state, increasing during periods of stress [6]. Unlike ADA1, ADA2 is a 57-kDa homodimer, secreted into the extracellular space, with low affinity for adenosine in physiologic conditions, although this can change based on temperature and pH [6]. Structurally, while ADA1 and ADA2 have somewhat similar catalytic domains, ADA2 has additional domains allowing it to mediate protein dimerization and cell surface binding in the extracellular space [8]. The protein dimerization domain is structurally homologous with a family of adenosine deaminase growth factors [6], which have been shown to be critical in the development of frog [9] and fly [10] embryos. ADA2 has been proposed to have a role in endothelial cell and hematopoietic cell development, although this enzyme is not present within human endothelial cells [2, 6, 8]. Endothelial cell instability in patients with DADA2 is felt to predispose to an underlying vasculitic phenotype [1, 2]. ADA2 is predominantly expressed by monocytes undergoing T cell-dependent differentiation into macrophages and dendritic cells [1]. ADA2 seems to also have cytokine-like growth factor properties, namely by activating CD4+ T cells and monocytes via cell surface binding and formation of the immunological synapse [11]. ADA2 is also felt to have autocrine-type growth factor properties by stimulating macrophage proliferation and inducing T cell-dependent differentiation of monocytes to macrophages [11]. ADA2 binds to different cell types via cell surface proteoglycans, but can bind specifically to T cells via the adenosine receptors [11]. ADA2 binds preferentially to specific lymphocyte subsets without receptors more specific to ADA1 [5].

Neutrophils have adenosine receptors. A chronic upregulation of neutrophil activity has been proposed to also be behind the vasculopathy component of DADA2 based on findings of upregulated interferon-stimulated gene transcripts in peripheral blood and overexpression of neutrophil-derived genes in two DADA2 patients. Neutrophil activation could contribute to reduced endothelial cell integrity [12]. Changes in the adenosine metabolism pathways cause dysregulation of neutrophil extracellular trap formation (i.e., NETosis) in patients with DADA2 [13]. Increased NETosis has been identified in mesenteric artery tissue with myeloid-cell predominant inflammation [13]. Increased circulation of low-density granulocytes, a subset of neutrophils with propensity for NETosis, was also increased in patients with DADA2 [13]. NETosis also increases macrophage release of TNF-α [13].
Dysregulation of the NETosis can also be seen in other autoimmune diseases, specifically systemic lupus erythematosus and ANCA-associated vasculitis [14, 15] (Fig. 1).

DADA2 patients have demonstrated a shift in differentiated myeloid cells toward a prominence of pro-inflammatory M1 macrophages, likely tied to ADA2’s physiologic role in maintaining the balance of myeloid lineage cells and their differentiation [2••]. This polarization of the monocyte-macrophage population is felt to cause a release of pro-inflammatory cytokines and lead to downstream tissue damage [2••, 16].

Upregulation of several key pro-inflammatory cytokines has been identified in patients with DADA2. Increased TNF-α has been seen both peripherally and in affected tissues [1••, 17•]. TNF-α secretion is thought to come predominantly from pro-inflammatory macrophages and chronically activated neutrophils [2••, 12]. The increased TNF-α production in these patients is felt to be a large part of the reason for the efficacy of TNF-α inhibition in DADA2 patients, especially those with vasculitic phenotypes. Increased TNF-α levels has also been proposed as a mechanism behind hematologic manifestations of the disease, as TNF-α may have a role in bone marrow failure in patients with aplastic anemia [18]. Patients with DADA2 also have upregulation of type 1 interferon-stimulated genes [12, 19, 20•, 21, 22]. Type I interferons decreases after treatment [19] and may be useful as a biomarker for DADA2 activity. Additional elevated peripheral cytokines have included IL-6 [23–25, 26•], IL-8 [25], and IL-10 [27, 28] (Fig. 1).

In the physiologic state, ADA2 binds to neutrophils, monocytes, NK cells, and the CD16 subset of macrophages. A decrease in NK, NKT, and CD16-positive monocytes was found to be common in many DADA2 patients [5•••]. Several studies have found depletion of memory B cells, while naïve B cell subsets are present in increased or normal levels [2••, 29••, 30], suggestive that aberrant B cell development and differentiation is also part of the pathogenesis of DADA2 (Fig. 1).

Clinical Manifestations of DADA2

Since its discovery, the phenotype of DADA2 has significantly broadened to include not only vasculitis but also immunodeficiency and hematologic manifestations. Ocular involvement is highly variable, with many symptoms likely secondary to underlying cerebrovascular disease, such as vision loss and oculomotor palsies [20•, 31•], including internuclear ophthalmoplegia. Additional ocular involvement has included strabismus [27, 32•], conjunctivitis [33], uveitis, papillitis [20•], and optic neuritis [17•, 32•, 34]. Vasculitis of the temporal artery has been described in a 6-year-old girl with histologic changes mimicking giant cell arteritis [35]. Hearing loss, including neurosensory hearing loss [17•, 36], has been described as well.

Neurologic involvement is estimated to occur in 50–77% of patients [32•, 37•, 38, 39, 40•]. A key characteristic neurologic manifestation has been central nervous system (CNS) arterial infarctions in pediatric patients with PAN-like manifestations, in contrast with classic childhood PAN, in which CNS disease is less common [41•]. Both ischemic and hemorrhagic CNS infarctions have been described. In a cohort of 12 Turkish patients with DADA2 with neuroimaging, the most common lesions were acute and/or chronic lacunar infarcts in the brainstem and/or deep gray matter (75%), with up to 50% revealing findings compatible with recurrent ischemic infarcts [34]. Several cases have suggested that cerebral MRI may be normal early on or under-report the true burden of disease [22, 42]. Additionally, conventional cerebral angiography of the CNS has been found normal in patients with ischemic lesions on MRI, again suggestive that there may be a predominance of CNS small vessel vasculitis in this subset of patients [43–45]. Alternatively, vasospasm may cause ischemia and/or reduced vascular integrity in the absence of inflammation, contributing to strokes with a paucity of findings on conventional angiogram, but consistent clinical and MRI findings [45]. DADA2 can also mimic Sneddon syndrome when presenting with livedo reticularis and neurologic manifestations [30]. Meningitis and encephalitis have been reported [31•]. Up to 53% of patients with DADA2 may have peripheral nervous system (PNS) involvement which can include mononeuritis multiplex, cranial neuropathies, and polyneuropathy (sensory or motor) [17•, 31•, 38, 39, 46].

Direct cardiopulmonary involvement is less common. Cardiac involvement specifically has included pericarditis, myocarditis, cardiomyopathy, long QT syndrome, and aortic root enlargement [17•, 40•, 47, 48]. Cavitary lung lesions, ARDS, and pleuritis have been described, in addition to recurrent pneumonias in the setting of immunodeficiency and cytopenias [20•, 32•, 37•, 49].

Visceral organ and gastrointestinal manifestations can include colicky abdominal pain, mesenteric ischemia, mesenteric artery aneurysms, hepatic artery aneurysms, gastrointestinal perforation, pancreatitis, portal hypertension, hepatoportal sclerosis, hepatosplenomegaly, and intestinal amyloidosis [31•, 32•, 40•, 42, 50]. By both imaging and histology, mesenteric, and renal aneurysms are largely indistinguishable from PAN. A 17-year-old male with a homozygous ADA2 mutation (G47A) and heterozygous for a Familial Mediterranean Fever (FMF) genetic mutation (V276A) was found to have aneurysmal changes of the abdominal veins [51]. Isolated hepatic involvement in the form of hepatomegaly ranges from 19 to 27% [17•, 29••, 52], and hepatitis was found to be present in 36% of patients [29••]. Nodular regenerative hyperplasia has been seen most frequently on liver biopsy [42]. Notably, CVID patients have also been shown...
Gene Mutations within the ADA2 Gene

ADA2 was purified in 2005 and determined to be encoded by the ADA2 gene (formally CECR1), located on chromosome 22q11.1 [6, 23]. A variety of types of mutations have since been reported including missense, nonsense, splice site mutations, frameshift mutations, deletions, and copy number variations [16, 61, 70]. Between 2014 and 2020, at least an additional 67 known pathogenic mutations were documented, with a total of 98 known variants, many not classified, of uncertain significance, or potentially benign [72].

Several genotype-phenotype correlations have been proposed, however, such correlations are complicated by newly discovered phenotypic manifestations, epigenetics, incomplete penetrance, environmental factors affecting gene expression, and compound heterozygosity for two different mutations. Vasculitis-predominant phenotypes more commonly had missense mutations and at least some residual enzymatic activity of ADA2 (around 3%), while those with pure red cell aplasia and bone marrow failure were more likely to have mutations leading to complete loss of function of the enzyme and/or minimal residual enzyme activity, including insertions/deletions, nonsense, or missense mutations [73]. While there are some genotypes that

to have nodular regenerative hyperplasia and autoimmune hepatitis, suggesting a potential similar mechanism in these cohorts [42]. Splenic artery irregularities have also been demonstrated [50]. Hypertension has been well described in DADA2 patients, even in those with hematologic phenotypes [27, 40]. Reports of renal involvement have included renal artery aneurysms, segmental glomerulosclerosis (collapsing variant), and renal amyloidosis [32, 50, 53]. One Turkish pediatric patient with renal amyloidosis was concurrently found to be heterozygous for the FMF gene (Met694Val mutation) and responsive to canakinumab [53].

Cutaneous manifestations are common, ranging from livedo reticularis/racemosa, nodular lesions, soft tissue/subcutaneous edema, ulcers, and erythema-multiforme-like lesions [31, 54, 55]. The prevalence of livedo racemosa has been reported to be as high as 73% of patients [39]. Skin biopsies can show medium vessel vasculitis, as seen in PAN, or leukocytoclastic vasculitis [17].

Less specific symptoms, including recurrent fevers, myalgias, arthralgias, arthritis, and oral aphthosis, are well described. There have been isolated cases of myositis [17, 20, 29, 31, 56, 57, 58].

Hematologic and immunodeficiency phenotypes were recognized later, with manifestations appearing to be closely linked. Pure red cell aplasia (PRCA), hemolytic anemia, autoimmune neutropenia, isolated refractory thrombocytopenia, and pancytopenia have all been reported [26, 27, 28, 31, 37, 49, 57, 59, 60]. Bone marrow biopsy in one DADA2 patient with bicytopenia revealed myelofibrosis [32]. In a separate cohort of five patients, two with PRCA had bone marrow studies consistent with PRCA, while one with hemolytic anemia had a hypercellular marrow, and the fourth who presented with recurrent fevers and pancytopenia had a normocellular marrow [37].

Cytopenia-predominant disease has been closely linked to immune deficiency, with findings of memory B cell deficiency and decreased B cell differentiation in these patients [2, 29, 30, 61]. The concurrence of DADA2 and lymphoproliferative diseases has been reported in several cases. Childhood onset Hodgkin’s lymphoma has been described in three patients, two of which were siblings [42, 62]. Two patients have been described with T cell large granular lymphocytic lymphoproliferation [49]. One patient presented with a syndrome resembling multicentric Castleman disease, responding well to anti-IL6 therapy [24].

Immunodeficiency can be mild (more commonly seen in vasculitis-predominant disease), or more severe (more commonly seen in hematologic-predominant disease). Varying degrees of antibody deficiencies can be seen including pure antibody deficiency or pan-hypogammaglobulinemia as in CVID [17, 29, 32, 49, 63, 64, 65, 66]. Verrucosis, herpes virus infections, and an increased susceptibility to infection with dsDNA viruses have been reported [26, 49, 61, 67, 68]. In a study screening 181 adolescent and adult patients with antibody deficiency or CVID, 11 patients were identified with mutations in the ADA2 gene. Within this cohort, upper and lower respiratory infections were slightly more common than herpes, intestinal, or urinary tract infections [29].

Overlap with other inherited disorders has been noted, often incidentally. A Turkish cohort of 196 patients with systemic autoinflammatory diseases screened for ADA2 mutations found two patients with pathogenic mutations and four with mutations of undetermined significance, three of which also had mutations in the FMF gene as well [69]. FMF gene mutations in patients with DADA2 have been seen by other groups as well [53]. Both FMF and DADA2 are known to have increased allele frequency in the Turkish population, possibly explaining their co-occurrence in this population. Noonan Syndrome-like disorder with loose anagen hair [70] and X-linked recessive nephropathisis (i.e., Dent’s disease) [71] have been described. Two siblings with chronic mucocutaneous candidiasis, retinal vasculitis, elevated IgG, and neutropenia were found to have a 770-kb deletion on chromosome 22q11.1 encompassing both the IL17 receptor gene and ADA2 [33] (Table 1).
seem to correlate with one of the three proposed categories, Lee and colleagues propose that the phenotypes of DADA2 represent a spectrum rather than distinct entities, as demonstrated by the R169Q missense variant, which was found in all the phenotypic categories [73].

**Epidemiology of DADA2**

The estimated prevalence of DADA2 could be as high as 4:100,000, based on allele frequencies of in silico-predicted ADA2 damaging variants [74]. The carrier frequency for the specific mutation, Gly47Arg, is felt to be as high as 1:10 in Georgian-Jewish populations and 1:500 in Turkish populations [74]. The Arg169Gln mutation is estimated to have a carrier frequency of 1:500 in northern European populations (Finnish and Dutch) [36, 49, 74, 75], and 1:2100 in the general population [41]. This mutation has also been reported in other European populations including a German cohort with immunodeficiency and/or antibody deficiency and has been proposed to potentially have an association with a lymphoproliferative phenotype [29, 49]. An Italian variant, Thr360Ala,
has been identified as well [17•]. These subsets of increased genetic frequency is felt to not only reflect founder population effects but also relative isolation of smaller populations, population bottlenecks, and population expansion that could lead to expansion of variants with deleterious effects [49]. Overall, patients with DADA2 have been identified globally, with reported ethnicities including Japan, Singapore, China, Morocco, Brazil, Middle East, Europe, and North America.

**Diagnosis of and Screening for DADA2**

To confirm the diagnosis, ADA2 activity level and/or genetic testing have been used. Enzyme activity testing level is largely based on the degree of inosine production from adenosine abstracts of patient plasma. Some study groups use the dried plasma spot method, where plasma is prepared from blood samples and applied to Guthrie filter cards [64•]. Filter cards are then dried and enzyme activity is determined via high-performance liquid chromatography [2•]. Caorsi and colleagues based ADA2 enzyme activity by measuring concentration of inosine and adenosine levels from adenosine-stimulated monocytes with or without an ADA1 inhibitor. Lower enzyme activity has correlated with severity of disease in many cases [17•]. Heterozygotes have been shown to have lower levels than normal controls, with variable levels of clinical symptoms [17•].

Genetic testing, whether this is gene-targeted or comprehensive genomic testing, can help confirm the diagnosis. Many studies continue to use Sanger sequencing for targeted gene sequencing of the ADA2 gene (e.g., of all 10 exons of the gene) and/or to confirm the exact mutation within the gene after larger sequencing has been performed [36]. Other studies have used next-generation sequencing as part of a broader approach, including whole exome (WES) or whole genome sequencing (WGS) [76]. Case studies with patients with low ADA2 activity and classic phenotypes but without identifiable mutations raise the question of mutations in introns or other non-coding parts of the gene [17•]. Comprehensive genomic testing also allows for detection of other concurrent genetic mutations that could be confounding the diagnosis. Ram and colleagues proposed a diagnostic algorithm for advanced genetic testing in those with a high clinical suspicion of DADA2. Patients with features of inflammation (e.g., fever and elevated CRP), cutaneous or neurologic features of vasculitis, and recurrent or chronic disease course in adults would benefit from further targeted Sanger sequencing of the ADA2 gene [20•]. Somez and colleagues recommended the addition of hematologic manifestations and parental consanguinity to this algorithm [77].

Currently, the decision to screen for DADA2 is largely based on physician discretion. Some have proposed screening PAN patients with early disease (under age 10), other family members with PAN, certain manifestations (skin, neurologic, cerebral, or bleeding) or resistant disease [78]. A North American cohort screening of 117 adult-onset PAN patients showed that over 3% of patients were homozygous or compound heterozygous for ADA2 gene mutations [79••]. While the prevalence of DADA2 is low in adult-onset PAN, consideration could be made for screening those with a familial predisposition for PAN, arterial strokes, hematologic manifestations, or immunologic manifestations.

**Treatment of DADA2**

Early treatment is important to prevent potentially devastating complications. It has become apparent that specific hematologic and immunologic phenotypes may not respond to the same therapies as those with vasculitic phenotypes [73••]. While in the acute phase, high-dose systemic glucocorticoids can reduce inflammation, most patients have refractory or relapsing disease upon attempts to taper [17•]. For patients with a vasculitic-predominant phenotype, TNF-α inhibition has been shown to be highly efficacious [16, 78, 80]. Anti-TNF-α therapy has strong protective effects against stroke [40•, 80••]. There has not been enough data to support one TNF-α inhibitor over another, with most receiving etanercept, adalimumab, or infliximab [1••, 2••, 32•, 39•]. TNF-α inhibitors have no apparent effect on ADA2 enzyme activity levels [80••]. Long-term, possibly lifelong, use of TNF-α inhibitors has been proposed because of the risk of relapse with therapy cessation, especially in those with vasculitis and CNS manifestations [80••].

Patients with immunodeficiency and cytopenia-driven phenotypes tend to be less responsive to TNF-α inhibition. These patients may be candidates for hematopoietic stem cell transplantation (HSCT). Successful HSCT has been reported in twenty-one patients with DADA2 to date [25, 26•, 28, 31•, 63, 67, 68••]. All patients had resolution of disease manifestations (including cytopenias, immune cell deficiencies, and vasculitis), normalization of ADA2 levels, and reduction of key cytokines (TNF-α, IFN-α, and IL-6) [68••]. Several patients required more than one transplant; two of whom had received their initial transplant from a healthy sibling later found to have ADA2 gene mutations. Thus, matched alloge- neic donors may be preferred, as healthy siblings are often heterozygotes and could have subclinical disease or delayed-onset disease. Three of the total twenty-one patients required additional stem cell boosts following decreased donor chimerism with their initial transplant [67, 68•••]. Hashem and colleagues have successfully used a reduced conditioning regimen, raising consideration that these DADA2 patients’ baseline immunodeficiency may reduce risk for rejection [81••]. This also raises the question that graft lymphocytes may also...
have survival advantage over ADA2-deficient cells; therefore, nonmyeloablative regimens may be successful.

Reported complications from the HSCTs included acute/chronic graft-versus-host disease, viral reactivation, other severe post-HSCT infections, post-HSCT autoimmune phenomena, and pineal gland hemorrhage [26•, 28, 63, 68••]. The potential for a heightened state of inflammation in DADA2 during the peri-transplant period could affect engraftment and the overall success of the transplant [26•]. This heightened inflammatory state could also increase risk for CVA in the pre- and post-transplant periods [26•]. Two patients had veno-occlusive disease complicating the transplant, requiring acute fluid restriction [68••]. The decreased endothelial integrity in DADA2 patients possibly increases the risk for veno-occlusive disease in these patients undergoing HSCT.

A number of other therapies have been trialed, many of which are used as first-line therapies for PAN. Nonspecific anti-inflammatory medications like NSAIDs and colchicine have been used, especially when the patient is initially suspected to have concurrent FMF. Stronger immunosuppressive agents including mycophenolate mofetil, azathioprine, cyclosporine, sirolimus, tacrolimus, methotrexate, cyclophosphamide, and rituximab have also been trialed in isolated cases with success [50, 71]. Thalidomide achieved complete response in a small number of Italian patients; however, its potential toxicities, including fetal toxicity and peripheral neuropathy, must be considered [17•]. Other than in a single patient mimicking multicentric Castleman disease, IL-6 blockade has not been successful in many cases [24, 25, 55]. Anakinra was not found to be effective in an Italian patient with vasculitis-predominant manifestations [17•]. Canakinumab has been effective in a patient with renal amyloidosis and heterozygous for the FMF gene [53]. Cyclosporine was used for cytopenia-predominant disease with efficacy; however, it was ultimately stopped for treatment-related adverse effects with a rapid return of cytopenias [57]. Pentoxifylline was used for a patient with mononeuritis multiplex and cutaneous manifestations, however with incomplete control of livedo reticularis [46]. IVIG has been used for hypogammaglobulinemia and recurrent infections, but was not enough to reduce other disease manifestations [64•]. Monthly fresh frozen plasma infusions replace ADA2 levels, but the effect is too transient for long-term effectiveness [32•, 40•, 80••].

While anti-platelet therapy has been used successfully in Kawasaki’s disease, there is a theoretical potential for these therapies to worsen hemorrhagic infarcts or lead to hemorrhagic conversion of ischemic infarcts in patients with DADA2.

Future therapies that could change the course of the disease include recombinant ADA2. Gene therapy or gene editing has been proposed, although models have not been developed. Treatment targets including modulation of adenosine receptors, modulation of adenosine-specific NET formation, or the interaction between NET mediators and macrophages are also potential treatment targets [13••].

There are no guidelines about when to treat patients, especially given the broad spectrum of disease and disease severity. Healthy heterozygotes, often healthy family members of index cases, and asymptomatic genetically proven DADA2 patients are probably safe to monitor for the development of any DADA2 symptoms. For these patients, starting a TNF-α inhibitor at early onset of disease could prevent devastating complications.

**Conclusions**

DADA2 demonstrates how single mutations in the gene encoding a monogenic metabolic pathway can result in highly variable, multi-organ disease with potential for high morbidity and mortality. Early identification and diagnosis of this phenotype has significant treatment implications and can dramatically change the disease course. Allele frequency is not limited to endogamous cultures, although it is higher in these populations, and this disease is likely more prevalent than what is currently reported. Awareness of this disease by pediatricians, internists, hematologists, immunologists, infectious disease physicians, hepatologists, and rheumatologists is essential in detection and early management of these patients. A low threshold for screening is important for adults and children with PAN, especially those with affected family members and those who do not respond to conventional therapies. A comprehensive approach to treatment, from both the medical and the patient community, may be the key to developing ongoing effective therapies for this disease.

**Compliance with Ethical Standards**

**Conflict of Interest**  The authors declare that they have no conflicts of interest.

**Human and Animal Rights and Informed Consent**  This article does not contain any studies with human or animal subjects performed by any of the authors.

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