

Adenosine Deaminase (ADA)-Deficient Severe Combined Immune Deficiency (SCID): Molecular Pathogenesis and Clinical Manifestations

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Abstract Deficiency of adenosine deaminase (ADA, EC3.5.4.4), a housekeeping enzyme of purine metabolism encoded by the *Ada* gene, is a cause of human severe combined immune deficiency (SCID). Numerous deleterious mutations occurring in the ADA gene have been found in patients with profound lymphopenia ($T^+ B^- NK^-$), thus underscoring the importance of functional purine metabolism for the development of the immune defense. While untreated ADA SCID is a fatal disorder, there are multiple life-saving therapeutic modalities to restore ADA activity and reconstitute protective immunity, including enzyme replacement therapy (ERT), allogeneic hematopoietic stem cell transplantation (HSCT) and gene therapy (GT) with autologous gene-corrected hematopoietic stem cells (HSC). We review the pathogenic mechanisms and clinical manifestations of ADA SCID.

Keywords Adenosine Deaminase · Purine metabolism · SCID · Lymphopenia · Gene therapy · Clinical trials

Abbreviations

ADA Adenosine deaminase
Ado Adenosine

AdoHCy	S-Adenosyl-L-homocysteine
BMT	Bone marrow transplantation
d-Ado	2'-Deoxyadenosine
d-ATP	Deoxyadenosine triphosphate
DFSP	Dermatofibrosarcoma protuberans
d-Ino	Deoxyinosine
dCydK	Deoxycytidine kinase
ERT	Enzyme replacement therapy
GALT	Gut-associated lymphoid tissue
GI	Gastrointestinal
HSCT	Hematopoietic stem cell transplantation
Ino	Inosine
PEG-ADA	Polyethylene glycol conjugated adenosine deaminase
PJP	Pneumocystis jiroveci pneumonia
SCID	Severe combined immune deficiency

Introduction

Severe combined immunodeficiency (SCID) arises from profound defects of immune system development and function, and affected individuals are susceptible to severe and recurrent infections. The condition is genetically heterogeneous and to date approximately 20 different genetic causes have been identified [1]. The identification of ADA deficiency as a cause of SCID occurred through serendipitous recognition of the absence of ADA enzyme from a few SCID patients, when ADA isozymes were being analyzed to compare compatibility among potential family donors and were the first form of SCID for which an underlying cause was determined [2]. ADA deficiency accounts for 10–20% of human SCID, with an estimated incidence of ~ 1/500,000, based on a large population-based newborn screening ascertainment [3]. ADA-deficient SCID is unique among the different genetic

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causes of human SCID in that it is not lymphocyte-specific, but rather a systemic purine metabolic disorder causing SCID and potentially extra-immune complications in the pulmonary, hematologic, GI, neurologic, or skeletal systems. [4]

The pathophysiology connecting adenosine deaminase deficiency and SCID has been intensively studied (Table 1) and most of the mechanisms of the lymphotoxicity are understood (Fig. 1) (Table 2). The mechanisms for the other manifestations of ADA deficiency are not fully understood, although there are some insights, as discussed below. The development of a mouse model by ADA gene knockout greatly increased understanding of pathogenic mechanisms and processes in the disease. Fortunately, many of these manifestations are prevented by early restoration of ADA activity, by either ERT, allogeneic HSCT, or gene therapy. The management of ADA deficiency is not discussed in this review but is the subject of a recent review by the same authors [5].

Features of the *Ada* Gene and Enzyme

Genetics

The *Ada* gene was isolated in 1983 and is localized on the long arm of human chromosome 20 (locus 20q13.11) and has been well characterized [6, 7]. It is 32 kb long and has 12 exons separated by 11 introns. The promoter region of the gene is 135 bases long, lacks the “TATA” and “CAAT” sequences that are usually present in eukaryotic promoters, and is extremely rich in G/C residues (82%) [8]. It also contains three inverted repeats that allow the formation of cruciform structures, two direct repeats of 10 and 16 bp, and five G/C-rich motifs (GGGCGGG) disposed in a strikingly symmetrical fashion [9]. Northern blot analysis detected a major 1.6 kb mRNA transcript and a minor 5.8 kb transcript that may be a precursor of the mature 1.6 kb message [10].

A large number of mutations have been identified in children with ADA SCID [11]. Studies expressing the defective

Table 1 Pathophysiology of adenosine deaminase (ADA)-deficient SCID

Bi-allelic mutations in the adenosine deaminase (<i>ADA</i>) gene
Absence of adenosine deaminase enzyme (<i>ADA</i>)
Adenosine and deoxyadenosine nucleosides not catabolized to inosine
Adenosine and deoxyadenosine accumulate systemically
Lymphocytes phosphorylate and trap the adenine nucleotides, esp. dATP
High levels of dATP are cytotoxic by multiple mechanisms causing death of T/B/NK lymphocytes
Pan-lymphopenia causes SCID, with opportunistic bacterial, viral and fungal infections possible
Systemic purine metabolic disorder may also affect liver, lungs, GI, and possibly CNS

ADA enzyme from known ADA SCID patient mutations in bacterial systems revealed that the genotype is an important determinant of enzymatic activity [12]. The mutations could be categorized into those that resulted in (i) single amino acid substitutions [13], (ii) premature stop codons [14], (iii) RNA splicing errors [15, 16], (iv) defects in post-translational modification of the ADA protein [17], and (v) insertions or deletions [2, 18–20] of a DNA segment carrying the ADA gene. A number of specific mutations are seen more frequently in ADA patients and some of these are associated with certain ethnic groups most probably as a result of a founder effect [21]. In some cases, subtle changes in the configuration of the protein render the ADA enzyme unstable and/or catalytically inactive and thus prone to a rapid degradation [22–25]. In other cases, the mutation was found in a regulatory region controlling expression of the ADA gene [26, 27].

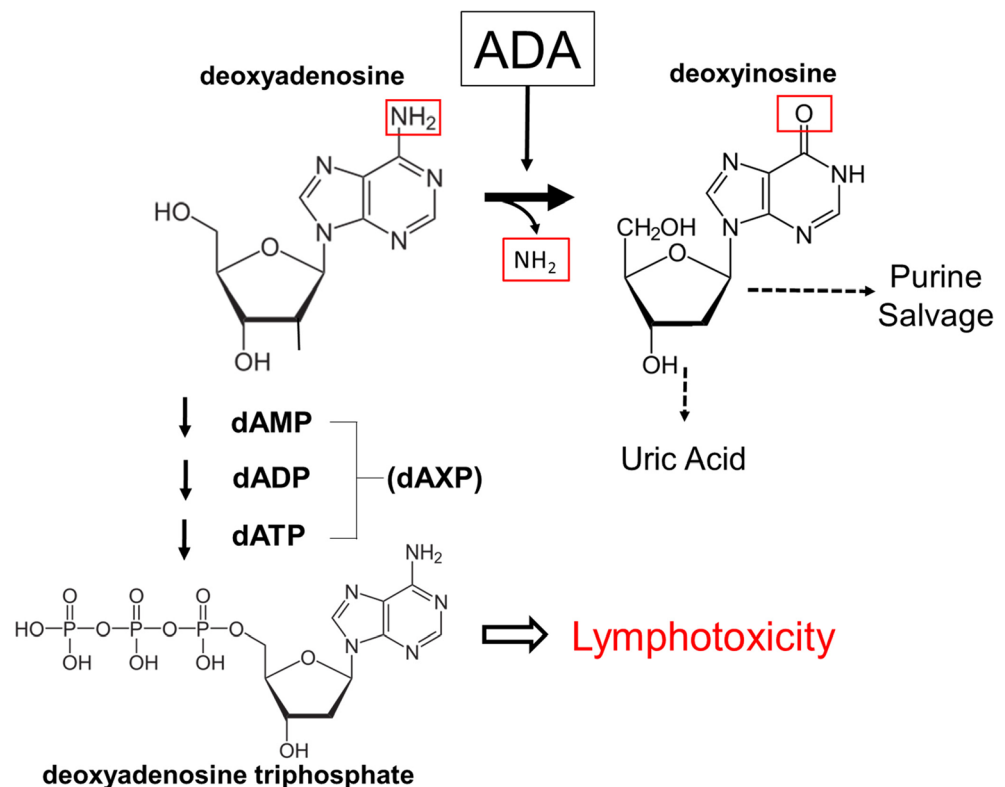
Molecular Features

The complete ADA amino acid sequence, deduced from complementary DNA (cDNA) and protein sequencing, consists of 363 amino acids, including the initiator Met. The enzyme consists of a single polypeptide chain with an estimated molecular weight of approximately 41 kDa, which contains carbohydrate and additional post-translational modifications [10, 28]. The ADA amino acid sequence is highly conserved from bacteria to humans [29]. The sequences of human and murine ADA are ~83% identical [30]; those of human and bovine ADA are ~93% identical [31]. The homology between human and bovine ADA is of particular relevance since enzyme replacement therapy for ADA SCID is derived from bovine ADA [5].

Expression Pattern

The ADA enzyme has a large phylogenetic distribution and has been found in a wide variety of organisms, from microorganisms [29, 32] to plants [33], from invertebrate [34] to vertebrate [35], from lower mammals [31, 36] to humans [37]. In addition to expression in virtually all mammalian cells and tissues, ADA plays a unique and essential role in the differentiation and maturation of the immune system. ADA activity in the thymus is much higher than that in any other organ [38]. Intermediate ADA activity is observed in the GI tract, spleen, lymph nodes, and skin while a low level of activity is observed in all non-lymphoid organs such as the brain, liver, and kidney [39]. Among lymphocytes, ADA activity in cortical thymocytes is higher than that in medullary thymocytes and peripheral lymphocytes [40, 41], while activity in T cells is higher than that in B cells [42, 43]. ADA immunoreactive protein and translatable ADA mRNA were indeed found to be six to eight times higher in T-lymphoblast lines than in B-lymphoblast lines, which corresponded to increased ADA catalytic activity

Fig. 1 Biochemistry of adenosine deaminase-deficient severe combined immune deficiency (ADA SCID). The nucleoside deoxyadenosine is deaminated by the enzyme adenosine deaminase (ADA) to produce deoxyinosine. Deoxyinosine may either be further catabolized down to uric acid or may be salvaged for further use in purine metabolic pathways. Lymphocytes especially have high levels of kinases which can successively phosphorylate deoxyadenosine to the nucleotides dAMP, dADP and dATP (collectively referred to as dAXP). dATP is the major metabolite that produces the pan-lymphotoxicity that leads to SCID



and protein in T cells compared to B cells [7]. These observations suggest that fundamental differences in the rate of degradation of the ADA protein and in nucleotide metabolism may exist between T cells and B cells, and among members of the thymocyte lineage at various stages of maturation. The expression pattern of ADA highlights its importance in different organ systems and also why pathology in ADA SCID is not just confined to the immune system but also has severe systemic consequences in the brain and elsewhere.

Table 2 Mechanisms of lymphotoxicity in ADA SCID [133]

Thymus
Widespread thymocyte apoptosis
Abnormal thymocyte distribution with reduced CD4+CD8+ double-positive cells and increased double-negative cells
Periphery, including lymph nodes, spleen, and circulating cells
Accumulation of adenosine and 2'-deoxyadenosine, which are toxic to lymphocytes, through incompletely understood mechanisms, resulting in decreased peripheral survival of T, B, and NK cells
Abnormal cell surface markers on T cells
T cell receptor (TCR) activation defects resulting in defective activation and expansion
Inhibition of TCR-induced signaling by adenosine
Exacerbation of 2'-deoxyadenosine spontaneous and TCR-mediated apoptosis of lymphocytes
Alterations in peripheral B cell distribution in spleen zones

Researchers have demonstrated that ADA is not only a cytosolic enzyme, but it can be also found as an ecto-enzyme on the external side of the plasma membrane of hematopoietic cells [44], which suggests an obvious role of the enzyme in degrading extracellular adenosine to inosine [45]. However, ecto-ADA, which seems to be identical to intracellular ADA and has a globular structure, does not directly interact with membranes but rather with membrane proteins [46, 47]. So far, two types of surface anchoring proteins for ecto-ADA have been described. The first type, with only one member, is CD26 (dipeptidyl-peptidase IV, DPP4), a multifunctional protein of 110 kDa strongly expressed on epithelial and endothelial cells as well as fibroblasts and leukocytes [48]. The second type of ecto-ADA-binding proteins includes the adenosine receptors (AR) A1 (A1R) [49] and A2B (A2BR) [50]. The functional role of ecto-ADA/CD26 complex is still unclear, although CD26 was described as a proteolytic enzyme with a wide tissue distribution almost 30 years ago [51]. However, it has been shown that ecto-ADA acts as a costimulatory molecule which facilitates a variety of specific signaling events in different cell types [52, 53]. The associated ecto-ADA/CD26 has been proposed to have a costimulatory function during T cell receptor (TCR)/CD3 complex engagement [54, 55] and to mediate costimulatory signals in the immunological synapse leading to the stimulation of the T helper 1 (Th1) and pro-inflammatory cytokines IFN- γ , TNF- α , and IL-6 [56]. A further role of ecto-ADA is to facilitate the signal transduction via A1R. Binding of ADA increases A1R affinity for the ligand, indicating that ADA is needed for an effective coupling between A1R and

heterotrimeric G proteins [49]. Binding of ADA to A2BR regulates agonist binding to A2BRs by increasing the ligand-binding affinity and the 5'-*N*-ethylcarboxamidoadenosine (NECA)-induced second messenger production. Therefore, in lymphocytes, cell surface ADA, apart from degrading extracellular adenosine, regulates those actions of adenosine that are mediated via adenosine receptors of the A2B subtype [50].

Pathogenesis of ADA Deficiency

Adenosine deaminase is a key enzyme of the purine-metabolism pathway which catalyzes the irreversible deamination of adenosine (Ado) and 2'-deoxyadenosine (d-Ado) into inosine (Ino) and 2'-deoxyinosine (d-Ino), respectively [57, 58]. Adenosine and 2'-deoxyadenosine are purine nucleosides that are intermediates in the pathway of purine nucleotide degradation, and their turnover in cells involves a complicated series of regulated reactions [59]. A number of disorders of purine metabolism have been shown to lead to immunodeficiencies [60].

In non-affected individuals, levels of purine nucleosides and deoxy-nucleosides in blood and urine are extremely low or undetectable. Alternatively, in patients with ADA SCID, marked elevation of both d-Ado and Ado in fluids and cellular components has been repeatedly noted and is considered the main cause of lymphotoxicity and other immunological and systemic pathologies [61, 62].

Toxic Effect Caused by d-Ado Buildup

2'-Deoxyadenosine is a component of DNA and primarily derives from its degradation. Thus, lymphoid tissues such as the bone marrow and thymus, where lymphocytes undergo apoptosis during the maturation and differentiation process, are predisposed to generate and retain the highest amount of d-Ado [39, 63]. In the absence of ADA activity, d-Ado accumulates in extracellular compartments and within cells, where it is converted by the enzyme deoxycytidine kinase (dCydK) to deoxyadenosine triphosphate (d-ATP) [64]. Retained d-ATP has the potential to (i) generate DNA strand breaks [65], (ii) inhibit ribonucleotide reductase, an enzyme that participates in DNA synthesis and repair [66], (iii) induce apoptosis in developing thymocytes [67–69], and (iv) interfere with terminal deoxynucleotidyl transferase (TdT) activity, thus limiting V(D)J recombination and antigen receptor diversity [70]. Additionally, d-Ado irreversibly inactivates the enzyme *S*-adenosyl-L-homocysteine hydrolase (SAHH) which catalyzes the reversible reaction: adenosine + L-homocysteine ↔ *S*-adenosyl-L-homocysteine (AdoHCy). The SAHH inactivation prevents hydrolysis of AdoHCy, and as a consequence, its accumulation results in inhibition of transmethylation reactions necessary for effective lymphocyte

activation [71–73]. The cytotoxic buildup of d-Ado, d-ATP, and AdoHCy in lymphocytes is considered as the primary cause of lymphotoxicity. This is most apparent in the thymus where there is a very rate of thymocyte turnover during negative and positive selection and where the need for ADA expression is highest. The consequence of ADA deficiency and substrate accumulation is therefore to significantly disrupt thymopoiesis so leading to the lymphopaenia seen in ADA deficiency.

Toxic Effect Caused by Ado Buildup

Adenosine is a purine nucleoside composed of a molecule of adenine attached to a ribose sugar molecule moiety and Ado is a component of adenine nucleotides including ATP and RNA [74]. ADA deficiency leads to both intracellular and extracellular accumulation of adenosine. Under normal physiological conditions, adenosine functions as an extracellular signaling molecule playing a role in fine-tuning the immune response via activation of plasma membrane G protein-coupled receptors (GPCR) of target cells. However, this balanced regulation may be disrupted when adenosine concentrations are elevated, as in ADA deficiency [39, 75]. It has been shown that elevated levels of Ado may contribute to thymic apoptosis and inhibition of T cell activation and expansion causing severe T lymphopenia in mice and humans [76, 77]. For example, activation of the A_{2A} adenosine receptor by extracellular adenosine generates intracellular production of cAMP as a consequence of tissue damage-associated deep hypoxia. Elevated levels of cAMP inhibit and trigger “off signaling pathways” in activated immune cells in a delayed negative feedback manner in order to prevent additional tissue damage [78, 79]. Apart from T cell receptor signaling, adenosine is involved in the control of blood pressure and heart rate [80, 81], renal function [82], pulmonary inflammation [62, 83, 84], and in neurotransmission [85].

Clinical Manifestations

Immune Deficiency Many groups have previously reported the lymphocytotoxic effects of the accumulated adenine nucleotides, namely adenosine and deoxyadenosine, and their metabolites resulting in profound pan-lymphopenia (T, B and NK lymphocytes) in most ADA-deficient patients [86, 87]. Furthermore, these lymphocytes have been characterized by failed proliferation to mitogens, antigens, or allogeneic blood mononuclear cells [88]. Patients with missense mutations and some minimal residual ADA enzyme activity may present at older ages (e.g. beyond 1 year) and have demonstrated less severe lymphopenia [89]. In addition to impaired cellular immunity, ADA SCID patients have shown impaired humoral immunity as demonstrated by hypogammaglobulinemia and poor response to vaccination. Assessments in the murine model have shown absent germinal

center formation, leading some groups to suggest that splenic dysfunction, previously noted via decreased mass and damaged structure in ADA SCID models, inhibits B cell antigen-dependent maturation [90]. Indeed, live vaccines are contraindicated in this population, as with other immunodeficient groups, given the risk of reactivation and disseminated disease. Poliani et al. noted thymic atrophy and disrupted thymic architecture in tissue analysis of ADA SCID patients [91]. ADA SCID patients have been noted to present with a spectrum of opportunistic infections, including *Pneumocystis jiroveci* pneumonia (PJP), candidiasis, otitis media, and other upper respiratory infections, atypical mycobacterium, herpes viruses, and respiratory viruses, as well as live vaccination agents, including BCG, rotavirus, MMR, or varicella [92].

Myeloid Abnormalities Features of myeloid dysplasia and bone marrow hypocellularity in ADA SCID have previously been reported [93]. It has also been noted that prior to enzyme replacement therapy, absolute neutrophil counts were inversely proportional to plasma deoxynucleotide levels, suggesting that neutropenia occurs as a result of metabolic toxicity. In an analysis of 13 patients with active disease, the majority of the analyzed patients had hypogranular or vacuolated neutrophils and most had hyperlobular neutrophils. Of note, two cases demonstrated pseudo-Pelger-Huet neutrophils. Ubiquitously, atypical eosinophils were seen and characterized by cytoplasmic vacuoles, inconsistent granulation, or hyperlobular nuclei; these findings were independent of the presence of absence of absolute eosinophilia [93].

Furthermore, patients with ADA SCID have demonstrated increased incidence of antibiotic-induced myelotoxicity and prolonged neutropenia following nonmyeloablative chemotherapy [94]. In one report, 3 of 4 patients undergoing gene therapy with low dose Busulfan preconditioning had prolonged neutropenia. Notably, these three patients also developed prolonged neutropenia following antibiotic therapy with trimethoprim-sulfamethoxazole, β -lactam antibiotics, and vancomycin [93].

Autoimmunity An interesting phenomenon of autoimmunity has been described in patients with ADA SCID and recapitulated in some mouse models. Clinical manifestations of autoimmunity may include autoimmune hypothyroidism, diabetes mellitus, hemolytic anemia, and immune thrombocytopenia seen in mild or late-onset forms of ADA SCID [95]. Notably, classical forms of ADA SCID in humans and untreated ADA^{-/-} mice do not develop clinical or laboratory features of autoimmunity, prior to treatment, perhaps because of the severe lymphopenia [96]. Cases of autoimmunity, including hemolytic anemia and thyroiditis, have been reported in at least nine patients on long-term polyethylene glycol-conjugated adenosine deaminase (PEG-ADA) therapy [95]. It has been postulated that PEG-ADA therapy interferes with Treg function via increased adenosine

turnover due to abnormally high levels of ADA in the extracellular environment and results in breakdown of tolerance. Treg generation of extracellular adenosine has previously been described as one of the mechanisms of Treg-mediated suppression of immune response, which maintains peripheral self-tolerance and prevents autoimmunity. Untreated ADA SCID mice showed excessively high extracellular adenosine concentrations, interfering with T cell function, including that of Tregs, while excessive conversion of extracellular adenosine by PEG-ADA in treated mice appeared to block the Treg suppressor function [96]. Additionally, T cell and B cell receptor (TCR and BCR, respectively) signaling are critical in the development of a functional repertoire of immune cells in addition to destruction of auto-reactive cells. In the disease state, excessive extracellular adenosine increases intracellular cAMP which inhibits downstream intracellular signaling following TCR or BCR activation, resulting in attenuated response of the antigen-bound cell [95]. Sauer et al. further posited that this downregulation of intracellular signaling could contribute to the survival of auto-reactive cells as the excessively strong TCR or BCR signals are dampened, when they would otherwise trigger apoptosis of the auto-reactive cell [95]. ADA SCID patients who have undergone HSCT and gene therapy modalities have demonstrated development of physiologic levels and function of Tregs, although it is worth noting that autoimmunity has also been reported in these populations as well [97–99].

Neurodevelopment Significant neurological or developmental problems have now been reported in several series of ADA SCID patients [100–102], although the exact frequency within the ADA-deficient population is not known. Cognitive ability, as assessed by standard intelligence tests, was significantly lower in a group of ADA SCID patients with respect to age-matched controls, though it was not significantly lower than the group of non-ADA SCID controls who underwent prior BMT [100]. Notably, there appeared to be an inverse correlation between deoxyadenosine triphosphate (dATP) levels at diagnosis and IQ [100]. Additionally, behavior testing has demonstrated behavioral abnormalities in these patients that were statistically significant, with respect to non-ADA SCID controls who received prior BMT [100, 102] as well as those treated with long-term PEG-ADA replacement [102]. Recent imaging data revealed white matter and ventricular changes on magnetic resonance imaging as well as EEG differences with respect to age-controlled unaffected patients [102]. Audiometric data has shown bilateral high-frequency sensorineural deafness in the majority of ADA SCID patients who underwent audiometric assessments post BMT. This hearing loss has been linked to the multisystem pathology of ADA SCID, rather than BMT conditioning regimens or ototoxic anti-infective medications given to assessed patients [102–104].

Mechanisms of CNS toxicity in ADA SCID are incompletely understood. Neurons or glial cells may be directly

damaged by the metabolic abnormalities seen in ADA deficiency, or the pathologies may be secondary to insults, such as hypoxia related to pneumonia. Additionally, it is not known if the CNS defects develop in utero and are then static or if they progress postnatally. Normally, ADA expression is relatively high in the brain as previously measured, and it has been postulated that the absence of this enzyme results in an accumulation of dATP and resultant neuronal cell dysfunction in a similar manner to that seen in lymphocytes [100]. Chick embryonal neurons studied in vitro and in vivo have provided further support for this hypothesis, by demonstrating sympathetic ganglia and optic tectum cell loss with inhibition of ADA. Interestingly, the severity of cell loss was dependent upon the embryonal stage of the chick, with inhibition of ADA earlier in the gestation resulting in greater volume loss, with the authors positing that the ADA inhibition resulted in halting cell proliferation [105]. Because the degree, if any, of neuronal loss in the prenatal and postnatal periods is not known, it is difficult to determine whether early interventions, either ERT, HSCT, or gene therapy prevent or reverse them, although the clinical data suggests that these modalities do not result in improvement of neurological and audiological defects [100, 101, 103, 104].

Pulmonary The accumulation of dATP has previously been shown in mouse models to cause pulmonary insufficiency and has been a major cause of mortality in these models [106, 107]. Histologically, affected mice demonstrated accumulated activated alveolar macrophages and eosinophils, with abnormal alveolar formation. Furthermore, these histologic findings resolved with ERT [106].

Previous case reports of patients with ADA SCID have noted frequent occurrence of reactive airway diseases, even after successful treatment with ERT and BMT; these diseases include allergic rhinitis and asthma [107]. Furthermore, eczema has also been reported in these cases, supporting a predilection toward atopy in affected patients [108]. Extracellular adenosine is involved in purinergic signaling and modulates inflammation, and adenosine is elevated in bronchoalveolar lavage fluid from patients with asthma. Adenosine inhalation in an allergic mouse model sensitized with ragweed, resulting in a dose-dependent bronchoconstriction. Bronchoalveolar lavage and tissue testing of these adenosine-treated mice demonstrated airway inflammation as shown by elevations of eosinophils, lymphocytes, and neutrophils, in addition to decreased macrophages [109]. ADA SCID patients have a relative adenosine excess in the pulmonary system, which likely accounts for the increased incidence of reactive airway disease complications noted [110].

Booth et al. examined cohorts of patients with ADA SCID and X-linked SCID with respiratory disease characterized by similar clinical and radiographic findings. They were able to isolate an infectious etiology significantly more frequently in

the latter group, suggesting that lung pathology in ADA SCID may often be non-infectious. The non-infectious pulmonary disease found in patients with ADA SCID is thought to be due to metabolic injury, which is not seen in X-linked SCID [107].

Pulmonary alveolar proteinosis (PAP) represents a clinically significant non-infectious etiology of pulmonary disease in untreated ADA SCID patients [107, 111]. In these cases, patients exhibit hypoxemia, tachypnea, and infiltrates on chest X-ray. Lung biopsies have been found to contain proteinaceous material in the alveoli, and BAL fluid has shown abundant PAS+ foamy macrophages. This clinical presentation is quite similar to that of PJP or other pneumonias, and these infectious causes need to be ruled-out to allow PAP to be attributed as the cause. Additionally, the rate of PAP recognized in patients with ADA SCID greatly exceeds that noted in patients with SCID from another mechanism [111]. Notably, PAP was reversed with ERT or transplantation [111].

Gastrointestinal Prior studies with untreated ADA-deficient patients have demonstrated impaired development of gut-associated lymphoid tissue (GALT) [112]. Assessed patients did not have functional lymphoid cells and the Peyer's patches lacked germinal centers.

With regard to liver function, ADA-deficient mice have severe hepatic abnormalities resulting in rapid hepatocellular degeneration, which proves fatal to them in the perinatal period [113]. Although there are case reports of humans with ADA deficiency with hepatic pathology, by and large, patients with ADA SCID do not typically demonstrate clinically significant hepatic disease in early infancy or childhood, although they may have persistent mildly elevated liver enzymes on laboratory analysis [114].

Of note, with respect to overall intestinal motility in ADA SCID patients, there are reports of affected children with varying degrees of dysmotility and resultant challenges with feeding [16]. ADA expression has previously been shown to be quite elevated in the gastrointestinal tract relative to other tissues [115]. Taken together with the clinically significant gastrointestinal deficits observed in ADA-deficient patients, the finding of relative overexpression of ADA in the gut suggests that adenosine plays a fundamentally important role in the development and function of the gastrointestinal tract. Adenosine or its metabolites may act as an intestinal neurotransmitter or as a regulator of intestinal vasodilatation. In support of the latter possibility, a prior analysis of jejunal tissue undergoing absorption following suffusion with a nutrient solution showed hyperemia in the control versus treatment with ADA, suggesting that adenosine becomes elevated in response to nutrients to cause vasodilation and this may be blunted by exogenous ADA. Additionally, these authors also demonstrated increased arteriolar diameter and resultant intestinal blood flow in a dose-dependent manner when treated with topical adenosine [116]. ADA metabolizes adenosine to inosine, which has been established as a much weaker vasodilator relative

to adenosine. Other disease processes with chronic intestinal vascular congestion, such as failing Fontan procedures or portal hypertension seen in hepatic failure, have well-established malabsorptive complications, including protein-losing enteropathy as well as fat and carbohydrate malabsorption. We speculate that abnormalities of adenosine in the congenital absence of ADA may play a role in the clinical symptoms of feeding intolerance, chronic diarrhea, and failure to thrive often seen in untreated ADA SCID infants.

Skeletal Children affected by ADA SCID have previously demonstrated a combination of unique skeletal abnormalities, especially prominent bulbous ends of the costochondral junctions (Rachitic Rosary) that may be seen on chest X-ray [117]. They may have impaired linear growth by the time of diagnosis. Microscopic analysis of bones from patients with ADA who died without any treatment showed variable short growth plates with diminished numbers of functional chondrocytes [118]. Furthermore, the characteristic skeletal phenotype with altered structural and mechanical properties of bones in ADA gene knockout mice has been linked to imbalance in the receptor activator of nuclear factor κ B ligand (RANKL)/osteoprotegerin axis that results in impaired osteoblast function and decreased osteoclastogenesis [119]. Similar abnormalities of the RANKL pathway were observed in untreated ADA SCID patients [119].

Dermatologic Dermatofibrosarcoma protuberans (DFSP) is a rare dermatologic soft tissue sarcoma seen with increased frequency in ADA SCID [120, 121]. The underlying chromosomal aberration is a translocation (t[17;22][q22;q13]), which produces a *COL1A1*-platelet-derived growth factor β (*PDGFB*) fusion gene [121]. When associated with ADA SCID, DFSP has demonstrated specific features not characteristically seen in other presentations, including early presentation and multicentricity. DFSP lesions were found in 8 of 12 patients by Kesserwan et al. and were predominantly located on the trunk and extremities [121]. The group posited that the characterized DNA repair defects in patients with ADA SCID resulted in the increased frequency of DFSP in this population [121]. DFSP is slow growing with metastasis rates of < 2% of cases [122]. Interestingly, there have been reports of DFSP occurrence and relapse of DFSP in ADA SCID patients following successful treatment with HSCT and PEG-ADA supplementation, suggesting that once formed in the ADA-deficient state, the DFSP may persist or progress even in an ADA replete environment. Given the predilection for this condition in treated and untreated ADA SCID, high clinical suspicion for the presence of DFSP and skin biopsy of suspicious lesions for diagnosis was propounded [121]. The current standard of care has involved complete excision of the lesion, once diagnosed by skin biopsy, although these cases may become complicated by local relapses [123].

What Has Been Learned from the Mouse Models of ADA SCID?

In attempts to produce a murine model of ADA SCID, the *Ada* gene was knocked out in murine embryos; unexpectedly, the complete absence of ADA enzyme turned out to cause perinatal lethality, due to hepatotoxicity [124, 125]. Similar severe liver problems are not seen in humans with two nonsense mutations of the *Ada* gene, indicating differences in the purine metabolic pathways in humans and mice. However, we have observed that ADA SCID patients, even after successful HSCT or gene therapy, may have persistently mildly elevated serum levels of hepatic transaminase enzymes, and this may be a mild form of the metabolic hepatotoxicity, although it does not seem to have significant clinical consequences.

The same group of investigators had previously defined an enhancer element upstream of the murine *Ada* gene that directed expression of ADA in the placenta and trophoblast [126]. Expression of ADA in these tissues of gestating murine mothers is critical to fetal survival, as evidenced by the inability of *Ada* $-/-$ female mice to bear *Ada* $-/-$ pups after breeding with *Ada* $+/-$ males (necessitating crosses of *Ada* $+/-$ females with $-/-$ males to produce litters with ~ 50% *Ada* $-/-$ mice for studies)(personal observation). They then created a transgenic mouse using the placental/trophoblast enhancer to drive an *Ada* cDNA, leading to expression of ADA during gestation [127]. Crossing the mice with the placental transgene with the *Ada* gene knock-out mice allowed the *Ada* $-/-$ mice to survive beyond birth, but then be severely ADA-deficient post-natally.

The neonatally rescued *Ada* $-/-$ mice were then observed to die in a highly consistent manner at 3 weeks of life [127]. Death in these mice was due to pulmonary insufficiency, with profound airway inflammation seen, with enlarged and foamy macrophages and eosinophils, in the absence of detectable infection [106]. This may be mechanistically similar to PAP that occurs commonly in human ADA SCID infants prior to treatment [111]. Early institution of ADA enzyme replacement therapy, using the clinical-grade-pegylated bovine ADA drug, was shown to rapidly rescue the mice from the pulmonary toxicity, essentially allowing normal murine lifespan with chronic administration [110]. Subsequently, other modes of ADA restoration were also shown to prevent the pulmonary complications, including congenic bone marrow transplants from ADA-replete donors [128], autologous bone marrow transplantation using retroviral [129], or lentiviral vectors [130, 131] to stably introduce an *Ada* cDNA into the stem cells, or in vivo gene therapy by intravenous delivery of the *Ada* cDNA in lentiviral vectors [132]. Adeno-associated virus (AAV) vectors have been used in a version of the *Ada* KO mouse in which foregut ADA expression prevents the lethal phenotype, although efficacy is hard to gauge in this mouse as there is partial immune reconstitution with foregut ADA expression. The in vivo route primarily delivers the *Ada*

gene to the liver, which presumably acts as a metabolic sink to catabolize adenine nucleotides and achieve systemic detoxification and trans-rescue of lymphopoiesis.

Untreated *Ada*^{-/-} mice have a SCID immunophenotype, when studied in the first weeks of life with severe pan-lymphopenia and profoundly reduced central lymphoid tissues. However, any of the life-saving modes that restore ADA activity to prevent pulmonary death also lead to reversal of lymphopenia and the SCID immunophenotype. Unlike other forms of SCID with a cell intrinsic defect that provides an absolute dependence on gene correction for lymphocyte production, ADA deficiency may be improved by cross-correction of ADA-deficient cells by ADA expressed in other cells lowering system ADA substrate levels sufficiently to allow ADA-deficient lymphocytes to develop and survive. Neonatal *Ada*^{-/-} mice transplanted with congenic ADA-replete bone marrow showed donor lymphoid specific chimerism of 5–10% but with normalization of immune cell numbers suggesting that a small number of ADA-replete cells are able to “cross correct” the system and allow recovery of endogenous defective ADA-deficient cells [128].

Another important observation made in the ADA^{-/-} mouse model is on the biodistribution of ADA ERT administered intramuscularly [110]. The ADA was found essentially only in the plasma compartment and not in lymphoid tissues, such as thymus or spleen. The presence of high extracellular levels of ADA activity and low intracellular activity with ADA ERT may lead to consequences such as the predilection to autoimmune complications, as discussed (above) and incomplete correction of thymopoiesis and the subnormal immune reconstitution that occurs in patients.

The murine model has also helped elucidate a distinct neurophenotype present in ADA-deficient SCID. Specifically, mouse models demonstrated decreased activity and increased anxiety-like attributes. Subsequent biochemical studies by the researchers revealed consistent metabolic alterations and anomalous adenosine receptor signaling. Perhaps most notably, treatment with ADA replacement reversed metabolic perturbations but did not influence the signaling abnormalities. The authors noted that PEG-ADA treatment did not increase ADA levels in the brain, thought to be due to inability of medication passage through the blood-brain barrier. Notably, there was a 10-fold decrease in brain levels of adenosine despite the absence of ADA in this tissue, which was attributed to peripheral ADA activity with total body decrement in adenosine levels, as adenosine and other metabolites diffuse readily across the blood-brain barrier [102].

Finally, the *Ada* gene knockout mouse model has been used extensively to model clinical gene therapy approaches, to test efficacy and safety of approaches [131]. There had been concern that ongoing ADA ERT at the time of a gene therapy procedure would diminish the extent of immune restoration, due to rescue of ADA-deficient cells blunting the putative selective advantage of ADA-replete cells that would lead to

reconstitution with gene-corrected cells. In fact, it was found that continuing ERT beyond the time of transplant of *Ada*^{-/-} mice and for the first month after gene therapy did not prevent the development of a robust population of gene-corrected T lymphocytes, which typically do not begin to emerge into the peripheral for several months after transplant [129]. Based on these observations, some current clinical trials of gene therapy for ADA SCID are continuing ADA ERT for the first month beyond the transplant; this maintains lymphocytes that had developed under ERT during the peri-transplant period and minimizes the period of lymphopenia before de novo lymphopoiesis occurs from the gene-corrected graft.

Conclusion

ADA SCID is a multisystem disease with pathology extending well beyond the immune system. The lymphotoxicity caused by accumulation of deoxyadenosine has long been understood to underlie the severe immune deficiency in these patients, and more recently, complications with autoimmunity following initiation of treatment for the disease are coming to light. Intrinsic impairments in neurodevelopment are often exacerbated by long-hospital stays due to infection or malnutrition due to gastrointestinal pathology or infection. Early recognition of disease and initiation of treatment can reverse this pathology, and we would encourage early involvement of a developmental specialist to promote language, motor, and social skills acquisition. Pulmonary structural pathology, namely PAP, independent of infection, has been a well-recognized complication of this disease and predilections toward reactive airway diseases and atopy are increasingly being recognized in this population. Hepatic pathology is inconsistently seen in this population and is much milder than that noted in mouse models. Gastrointestinal disease, widely noted in clinical settings, is incompletely understood at a molecular level. We propose that a constant relative adenosine excess at the intestinal level promotes vasodilation and hyperemia, creating diffuse vascular intestinal congestion and impairing nutrient absorption. This mechanism would be consistent with frequent reports of malnutrition, feeding intolerance, and reflux which are often seen in this population and incompletely accounted for by infection. Skeletal and skin manifestations of the disease are also commonly involved, and practitioners caring for these children should have a high index of suspicion for potential linear growth impairment, impaired skeletal remodeling, and DFSP. Mouse models have been critical in our understanding of the multisystem pathologies encountered in this disease and in the development of current treatment modalities used in the clinical setting.

ADA deficiency remains a complex and challenging condition. Our greater understanding of the mechanisms underlying the immunological defects has led in part to improved

treatment options for this condition such as hematopoietic stem cell gene therapy. Our increased recognition of the multisystem defects may help drive further research that will allow better understanding and further management options.

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Compliance with Ethical Standards

Conflict of Interest DACS is an employee of Orchard Therapeutics Limited and DBK and HBG are members of the Scientific Advisory Board and consultants to Orchard Therapeutics, who are developing a commercial lentiviral vector gene therapy product.

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