also suggest that early initiation of PEG-ADA should be considered for ADA-deficient patients suffering from PAP, particularly when rapid hematopoietic stem cell transplantation cannot be performed. Our study also has several significant limitations. While we focused on AMs, future studies are needed to assess the role of pneumocytes and other lung cell abnormalities in the development of PAP in ADA deficiency. In addition, although we found that phagocytosis of IgG-coated beads by ADA-deficient AMs was normal, surfactant uptake was not directly determined. Moreover, AM function differs between humans and mice; hence, further studies with AMs isolated from ADA-deficient patients or developed from induced pleuri-potent stem cells of ADAdeficient patients will be required.

In conclusion, we establish here that ADA-KO mice develop PAP, in association with adenosine, surfactant, and AM abnormalities, and that early restoration of blood ADA activity can prevent PAP and early lethality in ADA-deficient mice.

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Mendelian inheritance of elevated serum tryptase associated with atopy and connective tissue abnormalities

To the Editor:

The cause of allergic disease is generally considered a complex polygenic process highly influenced by environmental exposures, with mast cells playing a major effector role. Genome-wide association studies and candidate gene approaches have produced an array of allelic variants that impart risk for allergic diatheses.^{E1} Although identification of monogenic allergic disease and inherited mast cell disorders has remained less common, several heritable mutations have been identified that help illuminate some pathophysiologic mechanisms underlying the development of atopy in the general population. These include serine peptidase inhibitor, Kazal type 5 (SPINK5) mutations in patients with Netherton syndrome¹; dominant negative signal transducer and activator of transcription 3 (STAT3) mutations in patients with a form of hyper-IgE syndrome²; and phospholipase C, gamma 2 (PLCG2) mutations in patients with a form of familial cold urticaria.³ Within the mast cell compartment, both germline and somatic mutations in KIT have been identified in patients with rare familial presentations of mastocytosis.^{4,5}

While seeking to characterize patients with unique, inherited allergic phenotypes among patients referred for evaluation of severe allergic skin, airway, or gastrointestinal disease or systemic mastocytosis (SM), we identified 9 atopic subjects with persistent increases in serum basal tryptase levels in the absence of evidence for a clonal mast cell disorder. Thorough clinical evaluations were undertaken, total and fractionated serum tryptase levels were obtained, in vitro basophil activation was assayed, and bone marrow biopsies were performed in 5 of these index patients. For a complete description, see the Methods section in this article's Online Repository at www.jacionline. org. Increased basal serum tryptase levels were found to segregate with distinct clinical features, and it was noted that multiple family members of each index patient shared elements of this phenotype. Serum tryptase levels were obtained in all available family members, and an autosomal dominant inheritance pattern of increased basal total serum tryptase levels was revealed in all 9 families (Fig 1). The mean basal total serum tryptase level measured during a resting state was 21.6 \pm 1.4 ng/mL among subjects with inherited tryptasemia (n = 33). Five family members' tryptase levels were unable to be obtained; these data were excluded from statistical analysis but are included with data from unaffected family members in Table E1 in this article's Online Repository at www.jacionline.org. Tryptase fractionation in a subset of 18 affected patients from 8 of 9 families further revealed that mature tryptase in serum was undetectable (<1 ng/mL, see Table E2 in this article's Online Repository at www.jacionline.org).

Prominent among features segregating with basal serum tryptase level increases were symptoms consistent with chronic and episodic mast cell degranulation, with 26 of 33 subjects

	Family 1	Family 2	Family 3	Family 4	Family 5	Family 6	Family 7	Family 8	Family 9
) • • • •				▎°┬■ □┬◆、◆		
Ages	8, 10, 44, 70	11, 39, 69	6, 10,13, 41	21, 55, 55	17, 22, 23, 48, 57, 78	24, 43, 50	2, 26, 29, 53	12, 20, 44, 68	9, 46
Symptoms No. affected									
Cutaneous	4	1	4	3	5	2	2	2	2
Flushing	1	-	2	3	4	1	1	1	-
Urticaria	4	-	3	-	5	2	2	1	1
Angioedema			1		3	-	-	1	-
Pruritis	4	1	2	1	5	1	1	-	2
Connective Tissue	3	3	4 4	3	0	2 2	3 3	3	2
Hypermobility Retained PrimaryTeeth	2	3	4	1	-	2	3	2	-
Arched Palate	1	1	2	1	-	1	1	1	2
Bone/Vascular abnormali		-	-	3	_	-	î	2	-
Atopy	4	3	4	3	6	3	2	4	2
Anaphylaxis		2	1	$2(1^{\dagger})$	3	-	ī	1	
Eczema	3	1	3	-	3	-	1	2	2
Asthma	4	2	2	2	4	1	2	1	1
Environmental Allergy	4	2	2	2	4	3	1	3	2
Food Allergy	3	2	1	1	4	-	1	1	-
Drug Allergy	3	1	2 ⁵	1	5	2	2	1	-
Gastrointestinal	4	3	4	3	5	2	3	4	0
Episodic Pain	2	-	2	3	3	2	3	2	-
IBS/Episodic Diarrhea	1	14	3	-	4	2	2	1	-
GERD Eosinophilic Esophagitis	4	1 2	3	2	3	1	2	2	-
Failure to Thrive in Infan		2	- 3	1	-	-	-	-	-
Neuropsychiatric	4	1	3	2	3	2	4	3	2
Dysautonomia ²	4	-	3	-	2	1	-	-	-
Anxiety/Depression	4	1	-	1	2	-	-	2	-
Behavioral Disorder	2	1	2	-	1	1	1	-	-
Chronic Pain ³	2	-	1	2	2	1	3	1	-
Developmental Delay	1	1	-	-	-	-	1	-	2

Pedigrees, ages, and clinical phenotypes of affected individuals

FIG 1. Pedigrees, ages, and clinical phenotypes of affected subjects. *GERD*, Gastroesophageal reflux disease; *IBS*, irritable bowel syndrome. †Hymenoptera hypersensitivity.¹Abormalities include pectus excavatum, plagiocephaly/brachycephaly, torticollis, femoral anteversion, tibial torsion, congenital mitral valve insufficiency, tortuous/ectatic intracranial vessels, and scoliosis.²Principally manifesting as postural orthostatic tachycardic syndrome.³Arthralgia, fibromyalgia, and vulvodynia.⁴Encopresis.⁵Immediate hypersensitivity reaction to radiocontrast media in 1 patient.

reporting episodic urticaria, flushing, and/or cramping abdominal pain frequently associated with urgency, diarrhea, or both. These symptoms could be unprovoked or triggered by heat, exercise, vibration, emotional stress, nonspecific foods, or minor physical trauma. A history of anaphylaxis was reported in 10 of 33 subjects from 6 different families, with 7 occurring following exposure to foods, 2 caused by insect stings, and another without an identified trigger.

Additional triggers in 3 subjects with food-mediated anaphylaxis included stress/heat/exercise (n = 2) and anesthesia (n = 1, Fig 1). One subject (family 1) with a normal serum tryptase level (6.6 ng/mL) had a history of anaphylaxis; this reaction was due to Hymenoptera envenomation (see Table E1).

Gastrointestinal manifestations, whether chronic or episodic, were another prominent feature seen in 28 of 33 subjects with increased tryptase levels; these included eosinophilic gastrointestinal disease, gastroesophageal reflux disease, tenesmus, fecal urgency, irritable bowel syndrome, and diarrhea. Atopic symptoms were present in 31 of 33 tryptasemic subjects, with environmental allergies and asthma being reported among 28 of 33 affected subjects. Connective tissue abnormalities were present in 23 of 33 subjects with increased tryptase levels from 8 of 9 families, and chronic musculoskeletal pain was present among 11 of 33 tryptasemic subjects from 6 of 9 families. Autonomic dysfunction, manifesting as postural orthostatic tachycardia syndrome, was reported in 10 of 33 affected subjects, and a neuropsychiatric diagnosis was present in 17 of 33 affected subjects (Fig 1). Each of these 5 clusters of clinical characteristics (cutaneous, connective tissue, gastrointestinal, atopic, and neuropsychiatric) were independently and significantly associated with increased tryptase levels (Fig 2, *A*). For additional clinical characterization of the families, see the Results section in this article's Online Repository at www.jacionline.org.

Because some of the reported symptoms in tryptasemic subjects could have been consistent with the familial presentation of a clonal mast cell disorder, index patients in 5 of 9 families underwent bone marrow biopsy. A significant increase in mast cell numbers was observed (P < .05; mean, 9.6 cells/high-power field [hpf]; range, 4-19.4 cells/hpf) compared with that seen in healthy volunteers (mean, 2.5 cells/hpf; range, 1-5 cells/hpf; see Fig E1 in this article's Online Repository at www.jacionline. org). However, none of the patients met the World Health Organization established criteria for a diagnosis of SM or monoclonal mast cell activation.^{E2} Mast cell aggregates were not present, aberrant expression of CD2/CD25 was absent, and KIT D816V mutation test results were negative in all 5 patients. Spindled-shaped mast cells (>25%) were observed in a single patient. In the 4 families in which a bone marrow biopsy was not performed, the clinical phenotype aligned with the cohort in which SM was excluded.

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