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Atopic Dermatitis Phenotypes Impact Expression of Atopic Diseases Despite Similar Mononuclear Cell Cytokine Responses

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1 **Atopic Dermatitis Phenotypes Impact Expression of Atopic Diseases Despite Similar**
2 **Mononuclear Cell Cytokine Responses**

3

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34

35

36 **Abstract**

37 **Background:** The atopic march refers to the co-expression and progression of atopic diseases in
38 childhood, often beginning with atopic dermatitis, although children may not “progress” through
39 each atopic disease.

40 **Objective:** We hypothesized that future atopic disease expression is modified by atopic
41 dermatitis phenotype, and that these differences result from underlying dysregulation of cytokine
42 signaling.

43 **Methods:** Children (n=285) were enrolled into the Childhood Origins of ASThma birth cohort
44 and followed prospectively. Rates of atopic dermatitis, food allergy, allergic rhinitis, and asthma
45 were assessed longitudinally from birth to 18 years of age. Associations between atopic
46 dermatitis phenotype and food allergy, allergic rhinitis, asthma, allergic sensitization, exhaled
47 nitric oxide, and lung function were determined. Peripheral blood mononuclear cell responses
48 (IL-5, IL-10, IL-13, IFN- γ) to dust mite, phytohemagglutinin, Staphylococcus aureus Cowan I,
49 and tetanus toxoid were compared among atopic dermatitis phenotypes.

50 **Results:** Atopic dermatitis at year 1 was associated with an increased risk of food allergy
51 (p=0.004). Both persistent and late-onset atopic dermatitis were associated with an increased risk
52 of asthma (p<0.001), rhinitis (p<0.001), elevated total IgE (p<0.001), percentage of
53 aeroallergens with detectable IgE (p<0.001), and elevated exhaled nitric oxide (p=0.002).
54 Longitudinal analyses did not reveal consistent differences in PBMC responses among dermatitis
55 phenotypes.

56 **Conclusion:** Atopic dermatitis phenotype is associated with differential expression of other
57 atopic diseases. Our findings suggest peripheral blood cytokine dysregulation is not a mechanism

58 underlying this process, and immune dysregulation may be mediated at mucosal surfaces or in
59 secondary lymphoid organs.

60

61 **Clinical Implications:** Identifying clinical factors and immune mechanisms that underlie the
62 association between atopic dermatitis in early childhood and subsequent allergic disease may
63 lead to personalized strategies towards allergic disease prevention.

64

65 **Capsule Summary** Early persistent atopic dermatitis is associated with risk for food allergy,
66 rhinitis, and asthma. AD onset after age 3 years is associated with asthma and rhinitis risk.
67 These findings suggest that timing of skin disease is important for allergic disease expression.

68

69 **Key Words:** Atopic dermatitis, atopic march, atopic dermatitis phenotypes, allergic
70 sensitization, progression of atopic disease

71

72 **Abbreviations:** Atopic dermatitis (AD), interleukin (IL), peripheral blood mononuclear cell
73 (PBMC)

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75

76

77 INTRODUCTION

78 Atopic dermatitis (AD) is the most common chronic pediatric inflammatory skin
79 condition, affecting approximately 11-20% of children in the United States(1–4). Prevalence of
80 childhood AD is increasing worldwide(5). Most cases of AD manifest before 5 years of age, with
81 a variable natural course that is difficult to predict. In some children, symptoms improve over
82 time, but approximately 50% of children with AD will have persistent symptoms into
83 adulthood(6).

84 AD is associated with allergic sensitization and the development of further atopic
85 diseases in childhood. AD, food allergy, asthma, and allergic rhinitis are traditionally considered
86 to be atopic diseases. These diseases occur together more often than would be expected if there
87 was no association between them, and they share the common feature of atopy. However, the age
88 of onset, organ involvement, clinical presentation, and comorbidities of allergic diseases vary
89 widely between individuals.

90 This co-expression and progression of AD to other allergic diseases has been termed the
91 atopic march. The atopic march posits a sequential progression of allergic diseases, starting with
92 AD in childhood and culminating in development of asthma and rhinitis later in childhood.
93 Temporally, food allergy is often co-expressed with AD, most often in early life. Asthma and
94 allergic rhinitis are considered later manifestations of the “atopic march” process. Allergic
95 sensitization to aeroallergens tends to develop after food allergens, and the delayed timing of
96 sensitization to inhaled allergens and subsequent allergic airway inflammation may contribute to
97 the later age of onset for these diseases(7–9). AD is a well-established risk factor for
98 development of asthma and rhinitis later in childhood, and multiple longitudinal and cross-
99 sectional investigations have confirmed the association between childhood AD and asthma(10–

100 14). Additionally, AD manifesting before 2 years of age has been associated with more severe
101 and persistent asthma(15,16).

102 However, only a small subset of children follow the “classic” atopic march. We have
103 previously described the use of latent class modeling analysis to identify 3 AD phenotypes in the
104 Childhood Origins of Asthma (COAST) study(17), a high-risk birth cohort composed of children
105 with parental histories of asthma and/or allergies. To investigate the role of AD phenotypes in
106 atopic disease co-expression and progression, we conducted longitudinal analyses comparing
107 incidence and expression of atopic diseases and biomarkers of allergic disease from birth until 18
108 years of age among the 3 AD groups. To determine if systemic immune dysregulation
109 contributes, we also measured and compared stimulated peripheral blood mononuclear cell
110 (PBMC) cytokine response patterns of each AD phenotype group longitudinally. The
111 identification of at-risk infants would provide valuable opportunities for early intervention by
112 evaluating for sensitization, implementing allergen avoidance measures, or initiating controller
113 therapies as indicated.

114

115

116 **Methods**

117 The Childhood Origins of ASThma (COAST) cohort is comprised of 285 children
118 enrolled from November 1998 through May 2000(18). To qualify, at least one parent was
119 required to have history of physician-diagnosed asthma and/or respiratory allergies. Participants
120 were enrolled at birth and followed prospectively. Questionnaires for both parents and children
121 were periodically administered and included questions regarding health histories, with a focus on
122 atopic diseases and environmental exposures. Physical examinations were performed at regularly
123 scheduled visits.

124 AD was defined as having an Eczema Area and Severity Index score (completed by study
125 team) greater than or equal to 1, physician diagnosis of AD at study visit, or parental report of
126 physician-diagnosed AD. Greater than 99% of yearly AD diagnoses were made by EASI score or
127 physician diagnosis at study visit(17). Food allergy was defined by using allergen-specific IgE
128 test results and historical reports of clinical reactions from parents or documentation in the
129 medical record. Asthma was defined as the documented(19,20). Rhinitis was defined as having
130 perennial or seasonal frequent sneezing, itchy nose, and/or rhinorrhea, and was ascertained by
131 affirmative response on historical questionnaires as previously described(21).

132 Blood was routinely collected and specific IgE to dog, cat, cockroach, ragweed, birch,
133 timothy grass, *Alternaria alternata*, *Dermatophagoides farinae*, and *Dermatophagoides*
134 *pteronyssinus*, were measured by using automated fluoroenzyme immunoassays as previously
135 described¹⁶. Allergen-specific IgE values of 0.35 kU/L or greater were considered positive.
136 Additional studies performed at annual routine clinic visits included total IgE level measurement,
137 peripheral blood eosinophil count measurement, exhaled nitric oxide measurement (FeNO), and
138 spirometry.

139 Blood samples from birth (cord blood), 1, 3, 6, 8, and 11 years of age was used for
140 cytokine analysis. Cord blood mononuclear cells (MNCs) and peripheral blood mononuclear
141 cells (PBMCs) were stimulated with dust mite, PHA, *Staphylococcus aureus* cowan, and tetanus
142 toxoid, and levels of IFN- γ , IL-5, IL-10, and IL-13 in culture supernatants were evaluated by
143 means of ELISA as previously described(22).

144

145 **Statistical analysis**

146 Latent class analysis (LCA) was previously performed using the observed pattern of AD
147 in a child during the first 6 years of life(17). Briefly, the Schwarz or Bayesian information
148 criterion was used to select the number of latent classes; a 3-class model was selected. After
149 considering co-variables, the 3-class model was selected by Bayesian information criterion.
150 Classifications from the 3-class LCA model with covariates were used for all subsequent
151 analyses.

152 Comparisons of the demographic characteristics between AD phenotypes were completed
153 using data from 1 year. Categorical variables were compared using Fisher's exact test and
154 continuous measures were compared using a Kruskal-Wallis Rank Sum test.

155 Repeated measures analyses were performed with the "repeated" statement in SAS
156 GLIMMIX using the link function appropriate for the outcome. All measures that represent
157 presence/absence of the outcome (food allergy, allergic rhinitis, asthma, aeroallergen
158 sensitization) were analyzed with logistic regression (using the logit link) and represented
159 graphically as the percent at each year with the outcome present. All other outcomes were
160 analyzed with linear regression (using the identity link) and represented graphically as the mean
161 at each age. The model included a term for age (up to 18 levels), AD phenotype (3 levels) and

162 the interaction. The interaction p-value is reported and represents the test of whether there are
163 any ages where the outcome differs by AD status. Where the overall term is significant, we did
164 pairwise comparisons of the AD phenotypes.

165 Since the numbers in the late-onset AD group are small, we did secondary analyses
166 where those with late-onset AD were grouped either with the none/intermittent AD or with the
167 persistent AD and re-evaluated the models.

168 Cytokines were analyzed after log transformation. SAS v9.4 (Cary, NC) was used for all
169 analyses.

170

171 **Results**

172 **Characteristics of study population**

173 Three AD phenotypes were identified as previously described(17) by latent class
174 analysis: (1) “none/intermittent” group was comprised of children who never had AD or had an
175 intermittent course (n=180, 63%); “late onset” group had no AD early in life with onset later in
176 childhood, between 4-6 years of age (n=38, 13%), and (3) “persistent” group developed AD in
177 infancy and had persistent symptoms throughout the observation period (n=67, 24%). Further
178 details regarding the phenotypes have been previously published(17). Participant demographics
179 are shown in Supplement Table 1.

180

181 AD phenotype and the association of other atopic diseases in childhood

182 To evaluate the influence of each AD phenotype on the risk of development of
183 subsequent allergic disease, we first compared rates of food allergy in our cohort longitudinally,
184 stratified by AD phenotype. At 1 year of age, the persistent AD group already has a significantly
185 higher proportion of food allergy (15%) compared to 5% in the none/intermittent and late-onset
186 AD groups (Figure 1A). This association continued throughout childhood; the persistent AD
187 group consistently 2-3 fold higher incidence of food allergy compared to the none/intermittent
188 and late-onset groups. Overall, the differences were not significant (Table I) despite these
189 apparently consistent differences. Since the late-onset group appeared very similar to the
190 none/intermittent group, we did a secondary analysis where late-onset group was combined with
191 none/intermittent group and compared to persistent AD. This analysis showed an overall
192 significant association ($p=0.004$).

193 We next examined rates of asthma in each AD group longitudinally from 6-18 years of
194 age. Both persistent and late-onset AD phenotypes had double the rate of asthma compared to the
195 none/intermittent phenotype at 6 years of age, and quadruple the rate throughout the remainder
196 of childhood (Figure 1B). Compared to children with none/intermittent AD, those with either
197 persistent or late-onset AD had significantly higher rates of asthma longitudinally throughout
198 childhood ($p < 0.001$). There was no significant difference in rates of asthma between the
199 persistent and late-onset AD groups (Table I).

200 Both persistent and late-onset AD were associated with higher rates of rhinitis compared
201 to none/intermittent AD (Figure 1C). All 3 AD groups had higher rates of rhinitis at 1 year of age
202 that decreased sharply by 2 years of age. Rates of rhinitis then increased steadily throughout

203 childhood. There was no significant difference in rates of rhinitis between the persistent and late-
204 onset AD groups (Table I). Specific IgE confirmed rhinitis is shown in Supplemental Table 1.

205 When we compare AD diagnosis at year 1 (persistent) with children that do not have AD
206 at year 1 (none/intermittent and late-onset), food allergy ($p=0.004$), asthma ($p<0.001$) and self-
207 reported rhinitis ($p<0.001$) are more common throughout childhood (Figure 2)

208

209 **AD phenotypes and biologic markers of atopy**

210 We next compared biologic markers of Type 2 inflammation between the 3 AD groups
211 longitudinally from birth to 18 years of age. Mean total IgE levels for all 3 groups were similar at
212 1 year of age; by 5 years of age the persistent AD group had significantly higher total IgE levels
213 compared to the other phenotypes, which persisted across childhood (Figure 3A). Total IgE level
214 was not significantly different between the none/intermittent and late-onset groups (Table I).

215 Both persistent and late-onset AD were associated with higher peripheral eosinophil
216 counts longitudinally throughout the study period compared to the none/intermittent AD
217 phenotype (Figure 3B). There was no significant difference in eosinophil counts between the
218 persistent and late-onset groups (Table I).

219 Allergic sensitization, defined as detectable specific IgE to at least one of the allergens
220 tested, was not significantly different between the 3 groups, although the persistent AD trended
221 higher from 3-9 years of age (Figure 3C). Quantitative assessment of the degree of allergic
222 sensitization comparing the percentage of allergens tested with detectable IgE revealed a
223 significantly higher percentage of positive allergens in the persistent AD group compared to both
224 late-onset and transient AD phenotypes (Figure 3D). In addition, late-onset AD had significantly
225 higher percentage of positive allergens compared with the none/intermittent group (Table I).

226 When looking at food sensitization, food sensitization to both egg and peanut was more common

227 with persistent AD, but this was not statically significant (Figure 1). Similar to food allergy
228 diagnosis, when we compare persistent AD compared to the other phenoytes, we see a significant
229 difference in both allergens, peanut =0.05, egg p=0.007) (Figure 2)

230

231

232 **AD phenotypes and physiologic indicators of atopy**

233 We measured FeNO (Figure 4A) and spirometry (Figure 4B) from 6 to 17 years of age.
234 Longitudinal analyses revealed an association between both persistent and late-onset AD (but not
235 none/intermittent AD) with higher FeNO (Table I). In contrast, no significant association was
236 found between AD phenotypes and FEV1/FVC ratio (Table I).

237

238 **AD phenotypes and cytokine responses**

239 To evaluate for evidence of underlying cytokine dysregulation, PBMC cytokine
240 responses after stimulation were compared between AD phenotypes. There was variability in
241 ages with data available; all available data was analyzed. We compared levels of IL-10 and IL-13
242 after stimulation with dust mite at 1, 3, and 6 years of age; levels of IFN- γ , IL-5, IL-10, and IL-
243 13 after stimulation with PHA at birth, 1, 3, 6, 7, and 11 years of age; levels of IFN- γ after
244 simulation with *Staphylococcus aureus* at birth, 3, and 6 years of age; levels of IL-10 after
245 stimulation with *Staphylococcus aureus* at birth, 1, 3, and 6 years of age.; levels of IFN- γ after
246 stimulation with tetanus toxoid at birth, 1, 3, and 6 years of age; levels of IL-10, IL-13, and IL-5
247 after stimulation with tetanus toxoid at 1, 3, and 6 years of age. No significant associations
248 between AD phenotype and PBMC cytokine response profiles were identified (Table II).

249

250 Discussion

251 The increasing prevalence of atopic diseases(23–27) has intensified focus on
252 characterizing the mechanisms underlying the development of these disorders in childhood.
253 Additionally, allergic diseases once thought of as single entities actually represent several related
254 endotypes and/or phenotypes, and elucidating these differences is crucial to refine our
255 understanding of allergic disease co-expression and mechanisms of disease. We have previously
256 utilized latent class analysis to identify 3 distinct AD phenotypes in a high-risk birth cohort –
257 none/intermittent, late-onset, and persistent AD. In this study, we found that AD phenotype,
258 based on age of onset and persistence of AD, is differentially associated with other atopic disease
259 expression and progression in childhood.

260 This study is unique in that we were able to perform a longitudinal analysis with follow-
261 up to 18 years, accompanied by robust clinical, laboratory, and immunologic data. This provides
262 a more complete picture of immunologic development and changes that occur throughout
263 childhood. Although only persistent AD was associated with food allergy, both persistent and
264 late-onset AD were associated with asthma and rhinitis. This likely reflects the different
265 pathogenesis of these disorders, and how age of onset effects disease expression.

266 Early onset AD was associated with food allergy, while other AD phenotypes were not.
267 The dual-allergen exposure hypothesis(28,29) suggests that allergic sensitization can occur
268 through cutaneous exposure, while early oral consumption of food protein induces tolerance. Our
269 study illustrates that AD that manifests in infancy is most crucial in perturbing the development
270 of oral tolerance, and that associated impairment of the skin barrier *at this time in the lifespan* is
271 most important for food allergy. In food allergy, timing, and balance of cutaneous versus oral

272 exposure predominantly influence whether food allergy or tolerance occurs. In contrast, asthma
273 and rhinitis have overlapping but differential risk factors.

274 Sensitization to inhaled allergens is a well-established risk factor for asthma and rhinitis,
275 and is more likely to occur later in childhood compared to food allergen sensitization(7). The
276 ORCA Study characterized the natural history of sensitization during the first 6 years of life of
277 229 children with moderate-to-severe AD. The results of that study showed food sensitization
278 decreasing over time, from 58% to 34%, while aeroallergen sensitization increased from 17% to
279 67% over the study period(30). Another study showed that sensitization to food allergens is more
280 common in children from 0-4 years of age, while environmental allergies are more common after
281 4 years of age(31). This may help explain why late onset AD is associated with an increased risk
282 for asthma and rhinitis, but not food allergy.

283 We also found associations between persistent AD and increased total IgE, higher
284 percentage of allergens with detectable specific IgE, and higher blood eosinophils. We found no
285 associations between AD and single allergen sensitization. These findings likely reflect the high-
286 risk nature of the cohort, whereby the children in the cohort have a higher risk for allergic
287 sensitization compared to the general population. Thus, there are high levels of sensitization
288 throughout the cohort, regardless of AD phenotype. However, there were significant differences
289 in the percentage of aeroallergens children were sensitized to between the 3 AD phenotypes,
290 with persistent AD having the highest percentage of positive aeroallergens. Indeed, by 11 years
291 of age at least 80% of the cohort was sensitized to at least one allergen. The increased in Type 2
292 biomarkers in persistent AD is consistent with the understanding that atopic dermatitis is
293 associated with Type 2 inflammation. Th2 polarizaiton is often initiated with increased
294 expression of epithelial cytokines, such as TSLP. Thymic stromal lymphopoiectin (TSLP) is an

295 alarmin that can stimulate Th2 cell polarization and cytokine responses(34). TSLP is
296 undetectable in normal skin, but highly expressed in lesional skin of AD patients(35). Expression
297 of TSLP, as well as IL-33 and IL25 promote Th2 inflammation. Indeed, expression of IL-4, IL-5,
298 and IL-13 is upregulated in the lesional skin in AD patients compared to healthy controls(32,33).
299 CCL17, CCL18, CCL22 and CCL26 are type 2 chemokines and have been found to be
300 overexpressed in AD lesional skin(36). We also found an association between persistent and late-
301 onset AD and elevated FENO, but not with decreased FEV1/FVC ratio. This is likely because
302 FENO is a better marker of allergic inflammation than FEV1/FVC ratio. For example, children
303 with non-allergic asthma would be expected have a decreased FEV1/FVC ratio, but not an
304 elevated FENO.

305 We hypothesized that patient with persistent AD (with more atopic manifestations),
306 would have evidence of underlying peripheral blood cytokine dysfunction. We found no
307 associations between AD phenotypes and PBMC cytokine signaling patterns, and these negative
308 findings suggest that the signaling networks underlying allergic disease progression are not
309 demonstrated by peripheral cytokine levels. This raises a potential role for barrier function and
310 local immune responses in atopic disease expression patterns. Indeed, epithelial barrier function
311 is a well-known contributor to allergic inflammation. Further investigation in the role of
312 epithelial cytokine levels (such as TSLP, IL-33 and IL-25), and how barrier function relates to
313 these cytokines are needed. Also, mutations in filaggrin, a skin barrier protein, have been
314 associated with both AD and asthma expression(37,38) likely by a variety of mechanisms, such
315 as increased penetrance of allergens and irritants, and promotion of inflammation. Genetic study
316 in a larger sample size can help clarify this further. Epithelial cells and resident immune cells in

317 the tissue such as dendritic cells may influence cytokine release patterns in regional lymph
318 nodes, and alter the subsequent immune response.(34,39,40)

319 In conclusion, we have shown that both age of onset and persistence of AD differentially
320 associate with expression of atopic disease, and peripheral blood cytokine dysregulation does not
321 appear to be an underlying mechanism. The growing body of evidence demonstrating the
322 importance of multiple phenotypes and disease variance in allergic disease lends evidence
323 against a relatively simple, linear progression through the atopic march. Although many children
324 with AD do indeed co-express other atopic disease(s), only a subset of patients follow the strict
325 and “classic” path. Further longitudinal studies will provide a more complete picture compared
326 to cross-sectional studies, and enable identification of the mechanisms of disease co-expression.
327 A more complete understanding of risk factors for atopic disease progression will enable
328 development of targeted precise interventions.

329

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331

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443 **FIGURE LEGENDS**

444

445 **FIG 1.** Associations between AD phenotypes and atopic diseases. Persistent AD, but not
446 transient or late-onset AD, was associated with higher rates of food allergy (A). In contrast, both
447 persistent and late-onset AD were associated with higher rates of asthma (B) and rhinitis (C).
448 There was no association of AD phenotype with peanut (D) or Egg (E) sensitization. Blue circles
449 = none/intermittent AD, open green triangles = late onset AD, red triangles = persistent AD
450

451 **FIG 2.** Associations between AD diagnosis at year 1 and atopic diseases. AD at year 1
452 (persistent AD) was associated with higher rates of food allergy (A), asthma (B), and self-
453 reported rhinitis (C) in childhood compared to children without AD at year 1 (none/intermittent
454 combined with late onset). The association with peanut sensitization (D) was at the level of
455 significance, and egg sensitization was associated with AD at year 1 (E). Red triangles = AD
456 present at age 1, Blue circles = no AD at year 1.

457

458 **FIG 3.** Associations between AD phenotypes and biologic markers of atopy. Persistent AD
459 group had a significantly higher mean total IgE compared to the other AD groups (A). Persistent
460 and late-onset AD had significantly higher peripheral eosinophil count compared to transient
461 group (B). No significant differences were seen when sensitization rates to at least one
462 aeroallergen were compared (C); when the percentage of allergens with detectable specific IgE
463 was compared, significant differences were seen between all 3 AD groups. Blue circles =
464 none/intermittent AD, open green triangles = late onset AD, red triangles = persistent AD
465

466

467 **FIG 4.** Associations between AD phenotypes and lung function. Persistent and late-onset AD
468 were associated with an elevated FeNO (A), while no significant differences were seen with
469 FEV1/FVC ratio (B). Blue circles = none/intermittent AD, open green triangles = late onset AD,
470 red triangles = persistent AD

471







