

- 1 Evolving Concepts in how Viruses Impact Asthma  
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86

87 **Abstract**

88

89 Over the last decade, there have been substantial advances in our understanding about how  
90 viral infections regulate asthma (Table 1). Important lessons have been learned from birth  
91 cohort studies examining viral infections and subsequent asthma, understanding the  
92 relationships between host genetics and viral infections, the contributions of respiratory viral  
93 infections to patterns of immune development, the impact of environmental exposure on severity  
94 of viral infections, and how the viral genome influences host immune responses to viral  
95 infections. Further, there has been major progress in our knowledge about how bacteria  
96 regulate host immune responses in asthma pathogenesis. In this article, we also examine the  
97 dynamics of respiratory tract bacterial colonization during viral upper respiratory tract infection,  
98 in addition to the relationship of the gut and respiratory microbiomes with respiratory viral  
99 infections. Finally, we focus on potential interventions that could decrease virus-induced  
100 wheezing and asthma. There are emerging therapeutic options to decrease severity of  
101 wheezing exacerbations caused by respiratory viral infections. Primary prevention is a major  
102 goal and a strategy toward this end is considered.

103

104 Key words: virus, asthma, genetics, immune, microbiome

105

106 Abbreviations:

107 Respiratory syncytial virus (RSV)

108 Rhinovirus (RV)

109 Airway responsiveness (AR)

110 Fusion (F)

111 Type 2 (Th2)

112 Cadherin-related family member 3 (*CDHR3*)

113 Genome wide association study (GWAS)

114 Environmental tobacco smoke (ETS)

115 Upper respiratory tract infection (URI)

116 Short chain fatty acids (SCFA)

117 Forced expiratory volume in 0.5 s ( $FEV_{0.5}$ )

118 Wheezing lower respiratory tract illness (WLRI)

119 Inhaled corticosteroids (ICS)

120 Long acting beta agonists (LABA)

121 Lower respiratory tract illness (LRTI)

122 Plasmacytoid dendritic cells (pDC)

123

124 **Introduction**

125

126 Over the last decade, there have been substantial advances in our understanding about how  
127 viral infections regulate asthma (Table 1). Important lessons have been learned from birth  
128 cohort studies examining viral infections and subsequent asthma, understanding the  
129 relationships between host genetics and viral infections, the contributions of respiratory viral  
130 infections to patterns of immune development, the impact of environmental exposure on severity  
131 of viral infections, and how the viral genome influences host immune responses to viral  
132 infections. Further, there has been major progress in our knowledge about how bacteria  
133 regulate host immune responses in asthma pathogenesis. In this article, we also examine the  
134 dynamics of respiratory tract bacterial colonization during viral upper respiratory tract infection,  
135 in addition to the relationship of the gut and respiratory microbiomes with respiratory viral  
136 infections. Finally, we focus on potential interventions that could decrease virus-induced  
137 wheezing and asthma. There are emerging therapeutic options to decrease severity of  
138 wheezing exacerbations caused by respiratory viral infections. Primary prevention is a major  
139 goal and a strategy toward this end is considered.

140

## 141 **The viral genome and how it influences host immune responses to viral infections**

142           Respiratory syncytial virus (RSV) and rhinovirus (RV) are important causes of wheezing  
143 in early life and wheezing illness with these viruses have been associated with increased  
144 asthma risk later in childhood. At age 6, there is an increased risk of asthma if children had  
145 wheezing illness with RSV (odds ratio 2.6), RV (odds ratio 9.8), or both RSV and RV (odds ratio  
146 10.0) in the first 3 years of life. RSV is a negative-sense, single stranded RNA virus that is a  
147 member of the Paramyxoviridae family and is the leading cause of hospitalization each year in  
148 the United States in children under 1 year of age.<sup>1</sup> There are three species of RV in the  
149 enterovirus genus, and all are positive-sense, single-stranded RNA viruses that have protein  
150 capsids. RV are the most frequently detected viruses in wheezing children over the age of 1  
151 year, and from children and adults with acute exacerbations of asthma.

152           The clinical manifestations of a viral infection in the respiratory tract result from a  
153 complex interplay of the host, environment, and virus. To make comparisons between different  
154 immune responses elicited by diverse viruses, host and the environmental conditions must be  
155 held constant in order to prevent the introduction of confounding factors. This requires artificial  
156 conditions, such as the use of human cell lines for *in vitro* infection studies, the infection of  
157 genetically identical animals, such as mice, housed in the same environment, and the use of a  
158 standard viral inoculum. Determining the effect of specific genes within a virus requires that all  
159 other viral genes are identical. Such studies have begun but are still relatively new.

160           Experiments in models of RSV genomes have provided important insights into how the  
161 viral genome influences host immune responses to infection. Three RSV strains commonly  
162 used in pathogenesis studies are A2, line 19, and Long. RSV A2 infection in BALB/c mice  
163 resulted in a predominant IFN- $\gamma$  immune response, no production of the Th2 cytokine IL-13 in  
164 the lung, an absence of airway mucus, and no airway responsiveness (AR) to methacholine.<sup>2</sup>  
165 Infection with RSV Long similarly did not result in host IL-13 production in the lung nor was there  
166 airway mucus.<sup>3</sup> However, line 19 infection in genetically identical mice in the same environment

167 caused the host immune response to produce IL-13, decreased IFN- $\gamma$  compared to A2 infection,  
168 airway mucus, and heightened AR.<sup>2</sup> Sequencing of the A2, line 19, RSV Long strains revealed  
169 six amino acid differences between line 19 and the A2 and Long strains, of which 5 amino acid  
170 differences were in the fusion (F) protein.<sup>3</sup> To determine the contribution of the F gene of each  
171 virus to disease pathogenesis, a reverse genetics approach was undertaken by creating  
172 chimeric viruses whereby an A2 virus was manipulated to replace the A2 F gene with either the  
173 F gene of either line 19 or Long. Infection with the chimeric virus containing the line 19 F gene  
174 caused decreased host IFN- $\alpha$  lung levels, higher viral load in the lungs, greater lung IL-13  
175 protein, augmented airway mucus, and increased AR, compared to the chimeric viruses  
176 containing either A2 or Long F proteins.<sup>3</sup> Therefore, this reverse genetic approach provided the  
177 opportunity to not only discover which genes in RSV line 19 strain were responsible for the lung  
178 IL-13, airway mucus, and AR, but also the identification of the specific amino acids that caused  
179 airway remodeling. These techniques not only provide the knowledge of unique components of  
180 the viral genome that contribute to specific pathogenic features but may also assist in vaccine  
181 and therapeutic strategies aimed at the proteins responsible for specific disease characteristics.

182 Future perspectives: To date, there have not been studies that reveal a relationship between  
183 RSV genotypes and the presence of wheezing in hospitalized children with bronchiolitis or  
184 bronchopneumonia; however, this may be a function of the lack of application of technology to  
185 sequence strains because of cost. Studies relating viral genetics to severity of illness in mice  
186 have demonstrated the intricate interactions between viral genome, viral proteins, host cell  
187 function and metabolism and immune response. Developing a greater understanding of this  
188 chain of events could highlight several therapeutic opportunities, including identification of high-  
189 priority pathogens, inhibition of viral or host proteins that are critical for replication, and  
190 strategies to inhibit virus-induced skewing of immune responses that favors viral replication over  
191 host defense.

192

### 193 **Host genetics and viral respiratory infections**

194 A number of studies have begun to shed light on the relationships among host genetics,  
195 viral infections and acute and long-term respiratory outcomes. Candidate gene approaches  
196 have been utilized to identify associations between genetic polymorphisms and viral respiratory  
197 illness outcomes. Polymorphisms in several antiviral and innate immune genes have been  
198 linked to susceptibility to respiratory viruses, infection severity, and virus-induced asthma  
199 exacerbations, and have been replicated across multiple cohorts (Table 2). These genes  
200 include *STAT4*, *JAK2*, *MX1*, *VDR*, *DDX58*, and *EIF2AK2*.<sup>4</sup> Additionally, whole exome  
201 sequencing has been utilized to identify rare variants in innate immune responses linked to  
202 severe respiratory viral infections. Autosomal recessive *IRF7* deficiency has been observed in  
203 one patient in association with severe influenza infection and acts through impairment of  
204 interferon amplification.<sup>5</sup> Dominant negative loss-of-function variants in *IFIH1*, critical to viral  
205 RNA sensing, have been shown to be a risk for intensive care unit hospitalization due to viral  
206 infections in previously healthy children.<sup>6</sup>

207 A number of polymorphisms have been specifically associated with increased severity of  
208 illnesses associated with hospitalization from RSV infection. A candidate gene approach  
209 identified SNPs in the innate immune genes *VDR*, *IFNA5*, and *NOS2* as risk factors for RSV  
210 bronchiolitis.<sup>7</sup> In order to further elucidate associations of host genetics with RSV illness severity  
211 and asthma risk, a recent review examined overlap amongst genes associated with both  
212 outcomes. This approach identified a number of genes involved with both innate immunity and  
213 type 2 (Th2) inflammatory responses (*ADAM33*, *IL4R*, *CD14*, *TNF*, *IL13* and *IL1RL1*) that are  
214 highly relevant to these outcomes.<sup>8</sup>

215 The most replicated association between host genetics and asthma risk is the 17q21  
216 locus. In two birth cohort studies, variants in this locus, including *ORMDL3* and *GSDMB*, were  
217 also associated with increased risk of wheezing with RV infections in early life.<sup>9</sup> Interestingly,  
218 these variants were only associated with increased risk of subsequent asthma in children who

219 developed RV wheezing in the first 3 years of life. In contrast, early life RSV wheezing was not  
220 linked to 17q21 variants in these cohorts. In addition, farm exposures<sup>10</sup> and pets<sup>11</sup> in the home  
221 lessen the risk of asthma for children with high-risk 17q21 genotypes. In each case, the genetic  
222 risk associated with 17q21 was buffered by protective environmental exposures.

223 RV virulence varies by species; RV-A and RV-C are more likely to cause illnesses,  
224 wheezing and lower respiratory tract infection compared to RV-B,<sup>12</sup> which has a slower rate of  
225 replication and induces muted cytokine and chemokine responses.<sup>13</sup> Whether there are  
226 individual types within RV species that are more virulent is unknown, and difficult to study given  
227 the genetic diversity of these viruses and high mutation rate. A functional polymorphism in  
228 cadherin-related family member 3 (*CDHR3*) has been associated by genome wide association  
229 study (GWAS) with early childhood asthma and severe wheezing episodes.<sup>14</sup> Interestingly,  
230 *CDHR3* is a receptor that enables binding and replication of RV-C, suggesting that this link  
231 between *CDHR3* and asthma risk may be mediated by RV-C infections.<sup>15</sup> In support of this  
232 hypothesis, children with the risk polymorphism in *CDHR3* were recently found to have greater  
233 risk of RV-C illnesses, but not illnesses associated with other viruses.<sup>16</sup>

234 Future perspectives: These genetic associations among respiratory virus susceptibility, infection  
235 severity and subsequent asthma risk may prove to be important to risk stratify populations, and  
236 potentially provide new therapeutic targets for reducing illness severity and subsequent risk.  
237 Further, unbiased approaches have been employed recently to identify pathways of gene  
238 expression in the upper airway that differentiate a viral cold that resolves from one that leads to  
239 an asthma exacerbation.<sup>1</sup> Efforts are ongoing to understand how these gene expression  
240 patterns are regulated in hopes of identifying new personalized therapeutic strategies to prevent  
241 asthma exacerbation. The integration of multiple “omics” approaches holds promise to provide  
242 the ability to unravel these complex relationships.

243

244 **Environmental factors affecting the inception and severity of asthma exacerbations**



245 The exposome, defined as the measure of all exposures that influence the health of an  
246 individual, is an important determinant of asthma risk during the lifespan of an individual.<sup>17</sup> Early  
247 exposures can set in motion pathways that will ultimately define illnesses and symptom  
248 exacerbations, which is especially true when considering the ontogeny of asthma. From birth  
249 through school years, children are frequently exposed to a variety of respiratory pathogens,  
250 allergens, microbes and airway irritants. The pathogenic or beneficial effects of these  
251 exposures and their interactions remain the focus of research to develop new interventions and  
252 preventive therapies.

253 Most of the initial research devoted to the ontogeny of asthma focused on RSV infections  
254 which are frequently detected by culture and tests for RSV antigen in nasal washes from  
255 wheezing infants during the mid-winter months. Studies in the past reported that flares of  
256 wheezing caused by RSV leading to hospitalization during infancy increased the risk for  
257 developing asthma and allergy.<sup>18, 19</sup> However, recent studies indicate that the more severe  
258 episodes (i.e., those requiring hospitalization) of infantile wheezing caused by RSV increase the  
259 risk for subsequent wheeze in infants and toddlers, but it is less certain whether RSV-induced  
260 wheeze influences the development of atopy or asthma as children grow older.<sup>20</sup>

261 In contrast, flares of wheezing caused by RV are more strongly linked to persistent  
262 wheezing and the development of asthma, especially in children who are sensitized to allergens  
263 at an early age.<sup>21-23</sup> In keeping with this, the dominant risk factors for asthma attacks that require  
264 hospital care among children after 3 years of age is the combination of allergic airway  
265 inflammation and RV infection.<sup>24-26</sup> As a result, several host factors should be considered in  
266 efforts to treat asthma exacerbations more effectively and to reduce the risk for asthma  
267 development, For example:

268 1) There may be phenotypes of asthmatic children who would benefit from development of  
269 vaccines to RV or RSV. For example, genetic variations at the 17q21 locus and a coding

270 SNP in *CDHR3* (the receptor for RV-C genotypes) increase the risk for wheezing with  
271 RV during childhood.<sup>9, 14</sup>

272 2) There is current interest in whether the administration of a biologic such as omalizumab  
273 (anti-IgE antibody) during early childhood will have a disease modifying effect after this  
274 intervention is discontinued (i.e., the Preventing Asthma in High Risk Kids (PARK) trial;  
275 NCT02570984).

276 3) There is evidence that the asthmatic airway, especially epithelial cells and innate  
277 lymphoid cells, has a Type 2 bias with enhanced production of TSLP, IL-13, and IL-25 in  
278 parallel with decreased type I and III IFN responses that are needed for effective anti-  
279 viral killing and clearance.<sup>27-34</sup> This bias may increase the susceptibility of allergic  
280 asthmatics to RV infections. Once infected, however, *in vivo* studies have shown that  
281 during RV infections viral loads and clearance are similar among children and young  
282 adults with asthma compared to non-allergic individuals without asthma.<sup>31</sup> At present,  
283 mechanisms to explain these differences are poorly understood. A better understanding  
284 is likely to come from research focused on the cascade of early, innate cellular and  
285 molecular events that follow RV infection of epithelial cells.

286 Future perspectives: Airway inflammation caused by recurrent infections (predominantly with  
287 RV) in the allergic host will continue to be the focus of research designed to develop new  
288 therapies to help children and young adults with asthma. Whether treatments targeting allergic  
289 inflammation will be sufficient to reduce the frequency and severity of exacerbations (e.g., using  
290 new monoclonal antibody-based biologics), or whether additional therapeutics will be needed to  
291 decrease the frequency of RV infections, or enhance viral killing, remains to be determined.  
292 Looking to the future, the evaluation of other interventions such as the administration of  
293 antibiotics to treat secondary bacterial pathogens, or azithromycin to reduce wheezing following  
294 virus infection also deserve further study, along with investigations to determine whether the  
295 administration of vitamin D, probiotics, and dietary modifications (e.g., fish oil) will have benefits.

296 In contrast, the adverse effects of airway irritants such as environmental tobacco smoke (ETS)  
297 and air pollution (e.g., diesel fuel) on the severity and persistence of RV-induced asthma remain  
298 poorly understood.

299

### 300 **Effects of respiratory viral infections on patterns of immune development**

301 Acute wheezing illnesses with respiratory tract viruses in infancy and early childhood  
302 represent an important risk factor for childhood recurrent wheezing and later asthma  
303 development. This link is particularly well-established with RV and RSV, suggesting that these  
304 viruses may have a causative role, and significant research is directed towards understanding  
305 how these viruses can alter immune development to contribute to asthma pathogenesis. That  
306 said, causation remains unproven and asthma prevention strategies targeting viral illnesses do  
307 not currently exist.

308 RV-associated wheezing, in particular, is associated with a higher asthma risk than other  
309 viruses; this has been consistently demonstrated across multiple studies.<sup>21, 23, 35-38</sup> Many of  
310 these studies have linked RV-induced wheezing with other asthma risk factors, in particular  
311 markers of atopy including allergic sensitization, increased eosinophils, and atopic eczema,  
312 suggesting possible additive or synergistic effects in increasing asthma risk. Experimental  
313 models have demonstrated alterations in type-2 immune responses to RV that may account for  
314 this risk (Figure 1). Mouse models have demonstrated that neonatal RV (RV1B) infection results  
315 in persistent airway hyperresponsiveness, mucous cell metaplasia, and IL-13 production that  
316 does not occur in adult mice. Furthermore, knockout of IL4R prevents this response, consistent  
317 with an IL13-dependent process.<sup>39</sup> Subsequent work demonstrated that RV infection leads to  
318 expression of epithelial derived cytokines IL-25, IL-33, and TSLP and an increase in ILC2s as  
319 an important source of airway IL-13; blocking these pathways with anti-IL-25 attenuates  
320 neonatal RV-induced AHR and mucous cell metaplasia.<sup>40, 41</sup> While there is no equivalent human

321 evidence regarding the immune effects of RV in early life, these same pathways are known to  
322 play a key role in the response to RV leading to exacerbations in established asthma.<sup>29</sup>

323 A key question however, is whether underlying Th2 inflammation or RV associated  
324 wheezing comes first. A prospective birth cohort analysis has shown that allergic sensitization  
325 generally precedes RV wheezing but not the other way around, suggesting allergic sensitization  
326 may lead to more severe RV illnesses and the development of asthma.<sup>42</sup> Supporting this  
327 observation, *in vitro* studies have shown that Th2 inflammation can inhibit type I and III  
328 interferon antiviral responses to RV infections,<sup>43, 44</sup> which may increase susceptibility to more  
329 severe RV infections. However in contrast, several human studies have demonstrated  
330 increased IFN signatures in asthmatic children with virus infections, as well as in severe asthma  
331 in adults;<sup>45-47</sup> these might represent different disease states, as a recent report found that early  
332 life exacerbation-prone asthma was correlated with low IFN signatures, while the highest IFN  
333 signatures were associated with later-onset asthma.<sup>48</sup>

334 Allergy is a major risk factor for the progression from wheezing illnesses to asthma, and  
335 this has been a very consistent finding across multiple cohorts.<sup>21, 49, 50</sup> Allergic sensitization  
336 precedes wheezing illnesses in most young children,<sup>42</sup> and allergic inflammation can impair  
337 antiviral responses *in vitro*<sup>51</sup> and *in vivo*.<sup>52</sup> This suggests that allergic airway inflammation can  
338 increase susceptibility to and severity of viral respiratory illnesses. Allergic sensitization in early  
339 childhood may also modify the relationship between microbial colonization and respiratory  
340 outcomes. In preschool children whose airways were colonized with pathogen-dominated  
341 microbiomes, sensitized children were at increased risk for chronic asthma while non-sensitized  
342 children were likely to have transient wheeze that resolved by age 4 years.<sup>53</sup>

343 It is well established that hospitalization for RSV bronchiolitis in the first year of life is  
344 associated with later development of asthma.<sup>38, 54-56</sup> RSV induces a broad innate immune  
345 response in infants including systemic interferon, neutrophil, and inflammatory pathways, and  
346 distinct RSV strains and concomitant airway bacteria can influence the severity of infection.<sup>57, 58</sup>

347 The risk for more severe RSV-illnesses has also been linked to polymorphisms in several  
348 immune regulatory genes,<sup>59</sup> many of which also can influence asthma risk. However, whether  
349 RSV is causal remains a subject of debate with two large cohort studies showing different  
350 conclusions,<sup>59, 60</sup> one suggesting causation and the other an underlying genetic predisposition.  
351 Notably, two prevention studies using palivizumab (a monoclonal antibody directed against  
352 RSV) in high-risk infants found that prevention of more severe RSV-illnesses decreased the risk  
353 of childhood recurrent wheezing but not asthma development.<sup>61, 62</sup> Ultimately RSV infection  
354 appears to have the greatest impact on asthma risk during a critical window of lung  
355 development for infants born during the fall (in the Northern hemisphere) who are at ~4 months  
356 of age during the peak of the winter RSV season. It has been well-established that RSV  
357 infection can induce pathologic Th2 immune responses, especially within the context of  
358 formalin-inactivated RSV vaccination.<sup>63-65</sup> More recently, studies in mice have demonstrated the  
359 ability for RSV-related Pneumonia Virus of Mice as well as human RSV infection to break  
360 tolerance to allergens in neonatal mice.<sup>66</sup> Furthermore, it is now appreciated that RSV triggers  
361 release of epithelial-derived cytokines that promote Th2 responses and can induce ILC2  
362 responses following infection.<sup>67, 68</sup> These same epithelial cytokines have also been implicated in  
363 RV infection, perhaps suggesting a shared innate Th2-skewing mechanism during viral  
364 infection.<sup>27, 40</sup>

365 Future perspectives: Fully understanding the patterns of immune development that lead to  
366 asthma inception, and how such patterns are affected by exogenous exposures including viral  
367 infections, will direct asthma prevention research. Ongoing studies are focused on altering Th2-  
368 skewing in early life including through blocking IgE and through altering microbial exposures. If  
369 effective, decreasing Th2-inflammation in early life may function in part through enhancing  
370 antiviral responses.<sup>51, 69</sup> However, antiviral specific therapies including RV and RSV vaccines,  
371 may also prove to be critical in asthma prevention.

372

373 **Dynamics of respiratory tract bacterial colonization during viral upper respiratory tract**  
374 **infection**

375 Detection of viruses in the upper airway during peak viral seasons can be as high as  
376 90% in prospective studies.<sup>70</sup> However, rates of illness are significantly lower, leading  
377 researchers to question why some patients are more susceptible to increased morbidity when  
378 they have a viral upper respiratory tract infection (URI). One factor that has been shown to  
379 increase upper and lower airway symptoms during viral infections are bacteria.<sup>71</sup>

380 These bacteria collectively constitute the microbiota. The upper airway microbiota  
381 develops over the first year of life with alterations in the natural development associated with  
382 increased risk for URIs during the first few years of life.<sup>72, 73</sup> The most abundant bacteria within  
383 the upper airway of infants and children are *Staphylococcus*, *Streptococcus*, *Moraxella*,  
384 *Haemophilus*, *Dolosigranulum*, and *Corynebacterium*.<sup>71, 72, 74-78</sup>

385 In several infant cross-sectional and cohort studies, the presence of *Streptococcus*,  
386 *Moraxella*, or *Haemophilus* during upper respiratory infection increases the likelihood that the  
387 infant will have lower airway symptoms.<sup>72, 76</sup> Studies examining airway bacteria during RSV  
388 bronchiolitis have reported links between an increased abundance of *Streptococcus*<sup>9</sup> and  
389 *Haemophilus*.<sup>78</sup> In contrast, RV-bronchiolitis is associated with an increased abundance of  
390 *Moraxella* and *Haemophilus*.<sup>9</sup> While these studies suggest that a bacteria-virus interaction  
391 occurs during infancy, only a few studies have examined the association between virus and  
392 bacteria in school-age children. One such study revealed that children with *Streptococcus* or  
393 *Moraxella* present in their airway are more likely to have cold and asthma symptoms during a  
394 naturally occurring RV infection.<sup>71</sup> Collectively, these studies demonstrate that an association  
395 exists between specific bacteria and illness severity.

396 While *Streptococcus*, *Moraxella* and *Haemophilus* are often associated with an increase  
397 in viral-associated symptoms, a higher abundance of *Corynebacterium*, *Staphylococcus* and  
398 *Dolosigranulum* is often present in the airway in the absence of viral detection and clinical

399 symptoms.<sup>75, 77, 78</sup> In addition, when the latter three bacteria are enriched in the upper airway,  
400 infants are less likely to have an acute respiratory illness,<sup>72</sup> and school-age children are less  
401 likely to have a symptomatic illness during RV infection.<sup>75</sup> Furthermore, high abundance of  
402 *Lactobacillus* in the upper airway during RSV illness is associated with a decreased risk of  
403 childhood wheeze,<sup>77</sup> suggesting that bacteria present in the airway during viral illnesses may  
404 contribute to both illness severity and long term sequela.

405 Future perspectives: Because most studies examining airway bacteria during viral infection  
406 have been cross-sectional, observational studies, it remains unclear how airway microbes affect  
407 the epithelium, and whether these interactions contribute to the causation of wheezing illnesses,  
408 asthma development in young children, and exacerbations of established asthma. Greater  
409 insight is needed into metabolic, immunologic and toxic effects of bacteria on epithelial cells that  
410 could contribute to acute illnesses and asthma risk. While many studies have examined  
411 changes in bacteria that occur during viral infection, few have examined how the airway  
412 microbiome influences susceptibility vs. resilience to viral infection. Some bacteria could  
413 promote a “pro-inflammatory” environment thereby making the airway susceptible to viral  
414 infection. The presence of *H. influenzae* in the infant airway prior to viral infection is associated  
415 with increased expression of local inflammatory cytokines suggesting a link between bacteria  
416 and airway inflammation.<sup>79</sup> In contrast, mice receiving intranasal administration of *Lactobacillus*  
417 *rhamnosus* prior to viral infection have enhanced antiviral immune responses,<sup>80</sup> suggesting that  
418 some bacteria protect the airway and reduce the risk of symptomatic viral infection. Greater  
419 understanding of these relationships may lead to new preventive approaches to acute viral-  
420 bacterial illnesses and perhaps the development of childhood asthma.

421

## 422 **The influence of the gut microbiome on viral infections of the respiratory tract**

423 The gut microbiome represents the most abundant and diverse microbial environment in  
424 the human body, comprised of approximately 40 trillion bacteria.<sup>81</sup> These bacteria have

425 coevolved with humans over millennia to contribute to a symbiosis in which humans consume  
426 prebiotic fiber which is metabolized by resident microbes in the gut to create short chain fatty  
427 acids (SCFA) which in turn regulate immune responses.<sup>82, 83</sup> Alterations in this relationship are  
428 occurring in modern times due to practices such as the frequent use of antibiotics, and the  
429 consumption of a high-sugar, low fiber diet. As a consequence, a state of microbial dysbiosis,  
430 or an ecological imbalance, may result, which leads to the loss of metabolic capabilities and  
431 predisposes infants to both the development of atopic diseases as well as an increased  
432 susceptibility to viral infections.<sup>84</sup>

433         Although epidemiologic evidence strongly supports a role of the gut microbiome in the  
434 development of asthma, the mechanisms remain unclear.<sup>85, 86</sup> The most popular theory to  
435 explain these observations is that colonization with certain gut bacteria have a direct anti-  
436 inflammatory effect on the respiratory tract decreasing the likelihood of airway hyperreactivity.<sup>87</sup>  
437 However, there is evidence that certain species of microbiota in the gastrointestinal tract prime  
438 the respiratory immune system to effectively fight viral pathogens. Immunologic factors in early  
439 life such as low blood cell interferon responses<sup>88-91</sup> and attenuated cytokine production<sup>92</sup> have  
440 been associated with increased risk for wheezing in infancy. Furthermore, patterns of  
441 metabolites (which can regulate immune responses<sup>93</sup>) at birth are associated with the risk for  
442 wheezing illnesses.<sup>94</sup> The idea that delayed immune maturation might contribute to wheezing is  
443 supported by studies showing that early life exposures to dogs,<sup>95, 96</sup> farm life,<sup>10, 97</sup> and increased  
444 microbes and allergens<sup>98</sup> are inversely related to the risk of wheezing illnesses. Furthermore,  
445 exposure to these microbes and allergens during the prenatal period or infancy may be  
446 immunostimulatory.<sup>99, 100</sup> A loss, therefore, of these resident microbes may then lead to a  
447 predisposition to viral infections and in turn, the development of asthma.

448         Several studies have proposed mechanisms for the influence of the gut microbiota on  
449 both local and distant immune functions. SCFA have been shown to have a local effect on  
450 immune responses through their influence on mucosal barrier function, and a loss of SCFA-



451 producing bacteria has been implicated in the development of food allergy.<sup>101</sup> Recent advances  
452 have also shown that this symbiosis also influences vital immune responses in other systemic  
453 tissues. For example, in the absence of SCFA, mucosal barrier function can break down and  
454 allow for translocation of gut pathobionts, bacteria that are symbiotic under normal conditions  
455 but pathogenic when removed from their normal environment, which, in turn, can drive  
456 autoimmunity.<sup>102</sup> Similarly, in a murine model intact commensal bacteria in the gut were  
457 required for adaptive immune responses to respiratory influenza virus infection. Specifically,  
458 when mice were treated with antibiotics, they had reduced virus-specific antibody titers, CD4+  
459 T-cell responses, and cytokine secretion which consequently resulted in elevated viral titers post  
460 infection. This impairment, however, was rescued by local or distal injection of Toll-like receptor  
461 ligands.<sup>103</sup> Further, exposure to house dust from homes with dogs enriched the cecal  
462 microbiome in a murine model with *L. johnsonii*, which protected them against infection with  
463 RSV.<sup>104</sup>

464 Future Perspectives: Although the pathways remain incomplete, evidence continues to mount  
465 that the gut microbiome can influence the maturation of the immune system in viral defense and  
466 therefore the development of asthma.<sup>105</sup> Future therapies look to a role of probiotics for the  
467 prevention and treatment of allergic disorders, with recent evidence that atopy risk may be  
468 associated with a dysbiosis of the gut microbiome. Studies have shown that in asthma, MMP9  
469 (members of a family of enzymes that cleave extracellular matrix proteins) levels were  
470 significantly increased and treatment with the probiotic, *L. rhamnosus* GG (LCC), decreased  
471 MMP9 expression in lung tissue and inhibited inflammatory cell infiltration, as well as reducing  
472 exhaled nitric oxide among 4- to 7-year olds in pediatric asthma.<sup>106</sup>

473 In early childhood, total fecal IgE levels appear to be specifically correlated with house dust  
474 mite-specific IgE levels, indicating that fecal IgE levels represent markers of allergic response to  
475 aeroallergens. A significant correlation of fecal IgE levels with *Dorea spp.* and *Clostridium spp.*  
476 related to allergic rhinitis and asthma, respectively, suggest that modulation of particular subsets

477 of gut microbial dysbiosis could contribute to the susceptibility to allergic airway diseases.<sup>107</sup>

478 Future work is required for identification of specific species and functional studies to understand  
479 the strength and mechanism of these associations. In the future, it is critical to understand more  
480 precisely the microbiota composition. Optimized biomarker studies of the microbial taxa and the  
481 metabolites involved in asthma-associated dysbiosis could help identify infants at risk of asthma  
482 before symptoms. This would also provide a scientific rationale for future therapeutic strategies  
483 aimed at restoring an altered infant gut microbiome. Future studies need to revolve around  
484 state-of-the-art methods for the evaluation of the microflora to better define indications, the  
485 probiotic strains and the type of prebiotic to be used.

486

487 **Potential for primary prevention: clinical trials aiming to prevent the development of the**  
488 **episodic wheeze phenotype**

489 The inception of childhood asthma is tightly related to early life events such as  
490 respiratory infections and the development of aeroallergens sensitization. Other co-factors (e.g.,  
491 vitamin D) may modulate asthma inception pathways. Previous and on-going clinical trials,  
492 geared for asthma prevention, have targeted these pathways and co-factors.

493 Early life respiratory infections are significant determinants of childhood asthma<sup>108</sup>. In  
494 young toddlers, prevention of severe RSV bronchiolitis may reduce the risk of episodic  
495 wheeze/asthma development<sup>109, 110</sup>. In preterm infants (33-35 wks), palivizumab treatment  
496 during the RSV season resulted in a 73% reduction in the number of wheezing days during the  
497 first year of life, and outside of the RSV season<sup>109</sup>. A follow up study from the same cohort,  
498 revealed that at the age of 6 years the intervention resulted in a 41% relative risk reduction in  
499 parent-reported asthma, but the forced expiratory volume in 0.5 s (FEV<sub>0.5</sub>) percentage predicted  
500 values, which was an additional primary outcome, were similar between the palivizumab and  
501 placebo treated infants<sup>110</sup>.

502            Since early life respiratory infections cannot be completely prevented, attenuation of the  
503 immune/inflammatory processes during these infections may be another pathway for asthma  
504 prevention. This concept is illustrated by the results of a proof-of concept clinical trial in 40  
505 infants hospitalized with RSV bronchiolitis. In this trial, azithromycin treatment for 2 weeks,  
506 during acute RSV bronchiolitis, reduced the likelihood of developing recurrent wheeze during  
507 the subsequent year<sup>111</sup>. Azithromycin effects were attributed to anti-inflammatory properties  
508 and/or its effects on the airway microbiome<sup>112</sup>. A larger confirmatory trial is ongoing (APW-RSV  
509 II; NCT02911935; Table 3).

510            Based on observational studies that linked maternal vitamin D deficiency to childhood  
511 asthma, two clinical trials (VDAART<sup>113</sup>, COPSAC2010<sup>114</sup>) investigated whether maternal vitamin  
512 D supplementation (2400 IU/day<sup>113</sup>, 4000IU/day<sup>114</sup>) during pregnancy would prevent  
513 asthma/recurrent wheeze in their children. A recent meta-analysis that combined these two  
514 trials revealed that this intervention resulted in a 25% significant reduction in asthma/recurrent  
515 wheeze risk during the first 3 years of life<sup>115</sup>. The effect was most profound among women with  
516 sufficient serum vitamin D levels at randomization highlighting the importance of normal pre-  
517 conception vitamin D levels<sup>115</sup>. It was suggested that vitamin D beneficial effects may be related  
518 to enhancement of in-utero lung growth and development and promotion of antimicrobial effects,  
519 thereby reducing early life respiratory infections, and/or providing immune modulation effects<sup>116</sup>.

520            Omega-3 fatty acids were suggested to have anti-inflammatory effects, potentially due to  
521 decreased production of arachidonic acid metabolites. In a recent clinical trial, high dose  
522 Omega-3 fatty acids supplementation (2.4 g daily) to pregnant women, beginning at 24 week of  
523 gestation, resulted in a 30% relative risk reduction of persistent wheeze or asthma at age 3  
524 years<sup>117</sup>. These positive effects were driven by subgroups of children born to mothers with a  
525 variant of the gene encoding fatty acid desaturase, predisposing to low ability to produce  
526 omega-3 fatty acids, and by infants born to mothers with low omega-3 fatty acids baseline blood  
527 levels. These sub-group analyses suggest the plausibility of a precision-medicine approach of

528 this potential future intervention. Nevertheless, it is important to assure that high dose omega-3  
529 fatty acids does not possess any safety issues, before omega-3 fatty acids may be utilized for  
530 asthma prevention.

531 The ongoing ORBEX clinical trial (NCT02148796) is attempting to modulate the infant  
532 immune system by treating high-risk preschool children with Broncho-Vaxom® for 2 years to  
533 prevent/delay the development of wheezing lower respiratory tract illness (WLRI) during a third  
534 observation year. Broncho-Vaxom® contains bacterial lysates and was previously shown to  
535 reduce the rate of respiratory infections<sup>118</sup>. Hence, it is postulated that prevention of early life  
536 WLRI will prevent the development of the recurrent wheeze phenotype. Finally, the ongoing  
537 PARK clinical trial (NCT02570984) is targeting the association between allergic sensitization  
538 and asthma inception. PARK investigates whether treatment of high-risk preschool children with  
539 Omalizumab for 2-years would prevent asthma development, and whether the treatment would  
540 decrease asthma severity among infants who will develop asthma, during an additional 2-year  
541 observation period.

542 Future perspectives: This is an exciting time for all involved in childhood asthma prevention:  
543 recent clinical trials have shown the feasibility of asthma prevention, and multiple clinical trials  
544 are ongoing toward this goal. In addition to targeting type 2 immune responses, new  
545 interventions are needed to inhibit viral replication, either with specific inhibitors or strategies to  
546 boost the development of global antiviral responses in the airways. Finally, studies in farming  
547 environments strongly suggest that environmental exposure can lower the risk of viral  
548 respiratory illness in addition to reducing allergy.<sup>10, 97</sup> Identifying relevant mechanisms is likely to  
549 lead to new preventive approaches to virus-induced wheeze and asthma.

550

551 **New therapeutic options to decrease severity of asthma exacerbations caused by**  
552 **respiratory viral infection**

553           Recent studies have focused on short-term increases in standard asthma therapy,  
554 vitamin D supplementation, azithromycin and anti-IgE therapy. However, mixed efficacy results  
555 limit the widespread application of many of these therapies in clinical practice.

556 Maintenance inhaled corticosteroids (ICS) are effective in reducing the risk of asthma  
557 exacerbations and, when combined with inhaled long acting beta agonists (LABA), this  
558 decreases the risk further. However, exacerbations continue to occur. Attempts to increase  
559 dosing of inhaled steroid with early signs of loss of asthma control with viral infection, termed  
560 the “yellow zone”, to decrease exacerbation risk, have yielded mixed results. GINA guidelines  
561 suggest increasing ICS at onset of symptoms as part of a self-management plan  
562 (<http://www.ginasthma.org>). A Cochrane database review (including five studies in adults and  
563 three studies in children) concluded that current evidence does not support increasing ICS in  
564 mild to moderate asthma patients as part of a self-management plan to treat exacerbations.<sup>119</sup>

565           A clinical trial examined this question further in 254 children aged 5-11 with history of  
566 mild to moderate persistent asthma with at least one previous exacerbation in the past year.<sup>120</sup>  
567 Children were treated for 48 weeks with low dose inhaled steroid and assigned to either  
568 continue this or quintuple the dose for 7 days at onset of loss of asthma control. There was no  
569 significant difference in the rate of severe exacerbations in the groups. The total corticosteroid  
570 exposure in the high dose group was 16% higher (including both inhaled corticosteroid use and  
571 prednisone) and there was an effect on linear growth velocity between the high dose and low  
572 dose group (-0.23 cm/year), suggesting potential risk without identifiable benefit of the therapy.

573           Vitamin D levels have been inversely associated with asthma severity, including  
574 hospitalization for severe infections.<sup>121</sup> A large study aimed at optimizing low Vitamin D levels  
575 through supplementation did not reduce rates of colds or treatment failures in adults with  
576 asthma.<sup>122, 123</sup> In contrast, a meta-analysis of seven randomized trials demonstrated a significant  
577 reduction in asthma exacerbations, with the effect seen only in patients with low vitamin D at

578 baseline.<sup>124</sup> There are ongoing studies in children with asthma examining the possible role of  
579 vitamin D supplementation in preventing asthma exacerbations (Table 3).

580 Current guidelines do not recommend the use of antibiotic treatment for episodes of  
581 asthma-like symptoms in children, yet they are commonly used. A randomized, double-blind,  
582 placebo-controlled trial conducted in the US, evaluated the role of early administration of  
583 azithromycin in prevention of progression to severe lower respiratory tract illness (LRTI)  
584 symptoms.<sup>125</sup> Preschool children, age 12-71 months, with history of recurrent severe wheezing  
585 in the setting of LRTI, were randomized to azithromycin 12 mg/kg for 5 days (307 patients) or  
586 placebo (300 patients). The medications were to be started as soon as the children developed  
587 signs or symptoms that typically preceded the development of a severe LRTI. The primary  
588 outcome measure was the number of respiratory tract infections not progressing to a severe  
589 LRTI. The azithromycin group experienced a lower risk of progression to a severe LRTI than the  
590 placebo group.

591 In the COPSAC<sub>2010</sub> (Copenhagen Prospective Studies on Asthma in Childhood 2010  
592 cohort), children (age 1-3) with recurrent asthma-like symptoms within this cohort were enrolled  
593 in a study to assess the duration of episodes when treated with azithromycin.<sup>126</sup> With each  
594 episode of 3 days of consecutive symptoms (wheeze, cough, dyspnea), children were  
595 randomized to receive 10 mg/kg azithromycin or placebo for 3 days. Seventy-two children from  
596 the recurrent asthma-like symptoms group had 158 episodes. The azithromycin treatment  
597 shortened the days of symptoms, 3.4 days compared with 7.7 days after placebo,  
598 corresponding to a calculated reduction in episode length of 63.3%. More improvement was  
599 seen when the treatment was started earlier in the episode; however, treatment did not  
600 significantly affect the time to next episode of troublesome lung symptoms in children.

601 With these episodes, a hypopharyngeal aspirate was collected and cultured for common  
602 bacterial pathogens and a nasopharyngeal aspirate was collected for viral PCR. Overall, the  
603 presence of any cultured pathogenic bacteria did not significantly alter the treatment effect

604 compared to episodes without bacteria present; however, azithromycin was more effective in  
605 those whose culture grew *H. influenzae*. The treatment effect in these studies is promising;  
606 however, resistance to these antibiotics and eliminating commensal microbes along with  
607 pathogens are concerns with repeated treatment.

608 Birth cohort studies have shown allergic sensitization to be a risk factor for RV-induced  
609 wheeze.<sup>42</sup> Additionally, in one prospective cohort study, the severity of RV-triggered asthma  
610 exacerbation increased as the degree of allergen sensitization increased, with serum IgE levels  
611 (total IgE and allergen specific IgE) increasing from baseline during the exacerbation.<sup>127</sup>  
612 Persistence of asthma by age 13 was most strongly associated with wheezing illness with RV  
613 and aeroallergen sensitization in early life<sup>37</sup> suggesting a role for both viral infection and allergic  
614 sensitization in the development of asthma.

615 A possible mechanism for impaired response to viral infections in allergic asthmatics is a  
616 decreased secretion of IFN in response to viral infection. Purified plasmacytoid dendritic cells  
617 (pDC) from patients with allergic asthma were shown to secrete less IFN- $\alpha$  in response to  
618 exposure with influenza A virus.<sup>51</sup> Increased Fc $\epsilon$ RI $\alpha$  expression and serum IgE levels were  
619 inversely associated with IFN-  $\alpha$  secretion. The increased susceptibility to viral wheeze in atopic  
620 patients and impaired antiviral response in these patients suggests a role for possible  
621 therapeutic intervention to decrease allergic inflammation with the goal of decreasing asthma  
622 exacerbations in response to viral infection.

623 Omalizumab, a humanized monoclonal antibody that selectively binds to IgE, has  
624 recently been studied as an add-on therapy to prevent fall asthma exacerbations in atopic  
625 asthmatics in the Preventative Omalizumab or Step-Up Therapy for Fall Exacerbations  
626 (PROSE) study.<sup>128</sup> The PROSE study included 478 children, age 6-17, with respiratory allergy  
627 and asthma, randomized to either inhaled corticosteroid boost, add on omalizumab or placebo.  
628 All patients had guidelines-based care in addition to the add-on treatment (ICS boost,  
629 omalizumab or placebo). Treatment was begun 4-6 weeks before the participant's school start

630 day and ended 90 days after school start date. Omalizumab treatment significantly decreased  
631 the odds of having at least 1 exacerbation, whereas boosting ICS did not reduce risk.  
632 Omalizumab increased IFN- $\alpha$  responses to RV *ex vivo*. Within the omalizumab group, greater  
633 restoration of IFN- $\alpha$  responses were associated with fewer exacerbations. In this trial,  
634 omalizumab was associated with a decreased frequency of RV illnesses, decreased duration of  
635 RV infection as well as decreased frequency of overall respiratory illness, and reduced peak RV  
636 shedding.<sup>52</sup> Omalizumab reduced expression of Fc $\epsilon$ R1 $\alpha$  on the surface of pDC and this  
637 reduction was associated with lower exacerbation rates and correlated with enhanced IFN- $\alpha$   
638 production, suggestion a possible mechanism for the interaction between allergic sensitization  
639 and virus-induced asthma exacerbations.<sup>69</sup> However, the connection between the pDC type I  
640 IFN production and asthma exacerbation will benefit from further study.

641 In an observational study following children with asthma presenting with an acute  
642 asthma exacerbation triggered by RV, the use of omalizumab for at least 4 weeks prior to  
643 presentation was associated with reduced severity of exacerbation compared with patients  
644 primarily treated with ICS.<sup>129</sup> This suggests a benefit in not only frequency and duration of  
645 asthma exacerbation, but also severity of exacerbation.

646 Another possible mechanism for the interaction between allergic sensitization and virus-  
647 induced asthma exacerbations is the presence of anti-viral IgE in response to infection. In RSV  
648 infection in infants, RSV specific IgE was detected in nasopharyngeal secretions, with  
649 significantly higher titers in subjects with wheezing.<sup>130, 131</sup> Correlation of the peak titers with  
650 degree of hypoxia was also noted. Following known exposure to a specific laboratory strain,  
651 RV-specific IgE could be detected in human sera.<sup>132</sup> While the IgE response to RV and RSV  
652 are associated with infection, the role of IgE in the host response to these infections is not fully  
653 understood. Given the decreased exacerbations with use of omalizumab, further investigation  
654 into the role of anti-viral IgE is indicated.



655 Future perspectives: Given the morbidity of RSV and RV infections in patients with asthma, a  
656 consistent and effective treatment approach is highly desirable. While studies have found  
657 possible benefits to treatment with azithromycin and omalizumab, the widespread use of these  
658 treatment approaches is not currently justified. Further characterization of risk in this patient  
659 population and additional work to delineate the mechanisms by which these drugs are effective  
660 may lead to selection of patients most appropriate for these therapies.

661

662

### 663 **Conclusion**

664

665 There have been important advances in our knowledge of the relationship between viruses and  
666 asthma over the last decade. Advances in scientific methods have provided innovative  
667 opportunities to examine host, environment, and viral interactions that either protect against or  
668 increase vulnerability to asthma development and exacerbations. The exploration of the  
669 contribution of the respiratory and gut microbiome to virally-induced asthma is in its infancy and  
670 we suspect that over the next 5 years there will be major advances in this area. Finally, primary  
671 prevention is a major goal to diminish the morbidity of virally-mediated wheezing, asthma and  
672 exacerbations. Until primary prevention becomes a reality, clinical trials examining the impact of  
673 established medications, as well as novel therapies, will be critical to diminish the impact of viral  
674 infections on wheezing and asthma.

675

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Table 1. Review of most salient points

1. Polymorphisms in several antiviral and innate immune genes have been linked to susceptibility to respiratory viruses, infection severity, and virus-induced asthma exacerbations, and have been replicated across multiple cohorts; these genes include *STAT4*, *JAK2*, *MX1*, *VDR*, *DDX58*, and *EIF2AK2*.
2. Rhinovirus virulence varies by species; RV-A and RV-C are more likely to cause illnesses, wheezing and lower respiratory tract infection compared to RV-B.
3. RV infection leads to expression of epithelial derived cytokines IL-25, IL-33, and TSLP and an increase in ILC2s as an important source of airway IL-13; blocking these pathways with anti-IL-25 attenuates neonatal RV-induced AHR and mucous cell metaplasia in mice.
4. Two prevention studies using palivizumab (a monoclonal antibody directed against RSV) in high-risk infants found that prevention of more severe RSV-illnesses decreased the risk of childhood recurrent wheezing but not asthma development.
5. In several infant cross-sectional and cohort studies, the presence of *Streptococcus*, *Moraxella*, or *Haemophilus* during upper respiratory infection increases the likelihood that the infant will have lower airway symptoms; studies examining airway bacteria during RSV bronchiolitis have reported links between an increased abundance of *Streptococcus* and *Haemophilus*, while in contrast, RV-bronchiolitis is associated with an increased abundance of *Moraxella* and *Haemophilus*.
6. The presence of *H. influenzae* in the infant airway prior to viral infection is associated with increased expression of local inflammatory cytokines suggesting that a link exists between bacteria and airway inflammation; in contrast, mice receiving intranasal administration of *Lactobacillus rhamnosus* prior to viral infection have an enhanced antiviral immune response<sup>80</sup>, suggesting that some bacteria may protect the airway and help prevent viral infection.
7. The gut microbiome also regulate pulmonary anti-viral immunity; in a murine model intact commensal bacteria in the gut were required for adaptive immune responses to respiratory influenza virus infection.
8. Unique components of the viral genome contribute to respiratory illness and knowledge of these factors may also assist in vaccine and therapeutic strategies aimed at the proteins responsible for specific disease characteristics.
9. Omalizumab, a humanized monoclonal antibody that selectively binds to IgE, decreased fall asthma exacerbations in atopic asthmatics and increased IFN- $\alpha$  responses to RV *ex vivo* in the Preventative Omalizumab or Step-Up Therapy for Fall Exacerbations (PROSE) study, whereas boosting ICS did not reduce risk.
10. A recent meta-analysis that combined two clinical trials (VDAART, COPSAC2010) investigated whether maternal vitamin D supplementation (2400 IU/day, 4000IU/day) during pregnancy revealed that this intervention resulted in a 25% significant reduction in asthma/recurrent wheeze risk during the first 3 years of life.

1120 Table 3. Future/ongoing interventional studies examining treatments for viral triggered  
 1121 asthma  
 1122

Study Title	Study Population	Intervention	Primary Outcome Measurement	Estimated Completion Date
Vitamin D In the Prevention of Viral-Induced Asthma in preschoolers	children age 1-<6 with recurrent cold triggered asthma attacks, expected enrollment 865 subjects	Baseline and 3.5 month high dose vitamin D 100,000 IU and daily Vitamin D dose 400 IU OR placebo	Number of courses of rescue oral steroids (OCS) over 7 months	December 2022 <i>enrolling</i>
Azithromycin to Prevent Wheezing Following Severe RSV Bronchiolitis II	children 1-18 months of age, hospitalized due to RSV bronchiolitis, expected enrollment 200 subjects	Azithromycin (10 mg/kg x 7 days followed by 5 mg/kg x 7days) OR placebo	Time to occurrence of a 3rd episode of post-RSV wheezing, observation over 48 months	December 2021 <i>not yet enrolling</i>

1123 Figure 1. Immune responses to virus in the allergic asthmatic host. In the healthy host, anti-viral  
1124 IFN responses control and clear respiratory viral infections. In allergic asthmatics, the release of  
1125 the type 2-skewing cytokines TSLP, IL-25, and IL-33 promote the induction of Th2 cytokines  
1126 and the suppression of IFN responses, in addition to promoting airway hyperreactivity (AHR)  
1127 and increased mucus and IgE production. Furthermore, IgE has the capacity to suppress IFN- $\alpha$   
1128 production by Plasmacytoid DCs (pDCs).  
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Table 1. Review of most salient points

1. Polymorphisms in several antiviral and innate immune genes have been linked to susceptibility to respiratory viruses, infection severity, and virus-induced asthma exacerbations, and have been replicated across multiple cohorts; these genes include *STAT4*, *JAK2*, *MX1*, *VDR*, *DDX58*, and *EIF2AK2*.
2. Rhinovirus virulence varies by species; RV-A and RV-C are more likely to cause illnesses, wheezing and lower respiratory tract infection compared to RV-B.
3. RV infection leads to expression of epithelial derived cytokines IL-25, IL-33, and TSLP and an increase in ILC2s as an important source of airway IL-13; blocking these pathways with anti-IL-25 attenuates neonatal RV-induced AHR and mucous cell metaplasia in mice.
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Table 2. Polymorphisms in several antiviral and innate immune genes have been linked to susceptibility to respiratory viruses, infection severity, and virus-induced asthma exacerbations, and have been replicated across multiple cohorts.

Gene	Function
<i>STAT4</i>	Transcription factor required for IL-12 signaling in the development of Th1 cells from naïve CD4 T cells
<i>JAK2</i>	A non-receptor tyrosine kinase critical for signaling of the GM-CSF, gp130, and single chain receptor families.
<i>MX1</i>	Responsible for the antiviral state against influenza infection
<i>VDR</i>	Vitamin D receptor
<i>DDX58</i>	Involved in antiviral signaling in response to viruses containing a dsDNA genome
<i>EIF2AK2</i>	Innate antiviral immune response to viral infection that can trigger apoptosis via FADD-mediated caspase 8
<i>IRF7</i>	Critical role in the innate immune response against DNA and RNA viruses
<i>IFIH1</i>	Provides instructions for making the MDA5 protein that has a critical role in innate antiviral immunity
<i>IFNA5</i>	One of the type I IFN- $\alpha$ isoforms that has antiviral activities
<i>NOS2</i>	Nitric oxide synthase gene that mediates the antiviral activity of IFN- $\gamma$
<i>ADAM33</i>	Member of the ADAM (a disintegrin and metalloprotease domain) family identified as a major susceptibility gene in asthma
<i>IL4R</i>	Interleukin 4 receptor through which IL-4 and IL-13 signal to induce IgE class switching and airway mucus metaplasia
<i>CD14</i>	Multiple functions, one of which is critical for TLR signaling in host defense
<i>TNF</i>	Tumor necrosis factor that is produced in abundance by mast cells and has roles in cell survival and proliferation
<i>IL13</i>	Important in airway mucous cell metaplasia, airways responsiveness, VCAM expression
<i>IL1RL1</i>	One subunit of the receptor for IL-33, which can activate ILC2 and promote CD4 T cell differentiation toward Th2 phenotype
<i>CDHR3</i>	Cadherin that is the receptor for rhinovirus



Table 3. Future/ongoing interventional studies examining treatments for viral triggered asthma

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