

# **Asthma Exacerbation and Infection:** **Beyond a Simple Trigger**

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# Contents

- Infection in asthma exacerbation
- Viral vs. bacterial mechanisms
- Host immune phenotypes
- Infection-prone asthma
- Therapeutic implications

## Why Does Infection Matter in Asthma?

- Respiratory viral infections are the **leading trigger of exacerbations**
- Identified in **~70–80% of cases**
- Exacerbations → **major driver of morbidity & healthcare burden**
  
- But not all infections lead to exacerbations  
→ Host immune response determines outcome
  
- **Exacerbation is not pathogen-driven alone, but shaped by host immune susceptibility.**

# Isolated Pathogens During Asthma Exacerbation

	Total <i>n</i> = 407*	No obesity <i>n</i> = 236	Obesity <i>n</i> = 171	<i>P</i>
<b>Virus</b>				
Influenza	67/407 (16.5)	41/236 (17.4)	26/171 (15.2)	0.560
Human rhinovirus	37/328 (11.3)	23/196 (11.7)	14/132 (10.6)	0.751
RSV	17/326 (5.2)	10/194 (5.2)	7/132 (5.3)	0.953
Metapneumovirus	14/326 (4.3)	7/195 (3.6)	7/131 (5.3)	0.444
Coronavirus	12/328 (3.7)	9/196 (4.6)	3/132 (2.3)	0.374
Parainfluenza	10/328 (3.0)	6/196 (3.1)	4/132 (3.0)	1.000
Adenovirus	3/328 (0.9)	1/196 (0.5)	2/132 (1.5)	0.567
Enterovirus	1/264 (0.4)	0	1/105 (1.0)	0.398
<b>Bacteria</b>				
<i>S. pneumoniae</i>	25/407 (6.1)	15/236 (6.4)	10/171 (5.8)	0.833
<i>P. aeruginosa</i>	17/407 (4.2)	12/236 (5.1)	5/171 (2.9)	0.282
<i>C. pneumoniae</i>	10/217 (4.6)	9/119 (7.6)	1/98 (1.0)	0.024
<i>K. pneumonia</i>	9/407 (2.2)	5/236 (2.1)	4/171 (2.3)	1.000
<i>M. pneumoniae</i>	9/254 (3.5)	4/138 (2.9)	5/116 (4.3)	0.736
<i>H. influenzae</i>	8/407 (2.0)	4/236 (1.7)	4/171 (2.3)	0.725
<i>E. coli</i>	6/407 (1.5)	4/236 (1.7)	2/171 (1.2)	1.000
MRSA	4/407 (1.0)	2/236 (0.8)	2/171 (1.2)	1.000
<i>M. catarrhalis</i>	3/407 (0.7)	1/236 (0.4)	2/171 (1.2)	0.575
MSSA	2/407 (0.5)	0	2/171 (1.2)	0.176
<i>S. maltophilia</i>	2/407 (0.5)	2/236 (0.8)	0	0.512
<i>B. pertussis</i>	1/126 (0.8)	0	1/54 (1.9)	0.429

## Virus

- Influenza
- Rhinovirus
- RSV
- Metapneumovirus
- Coronavirus
- Parainfluenza

## Bacteria

- *S. pneumoniae*
- *P. aeruginosa*
- *C. pneumoniae*
- *M. pneumoniae*
- *H. influenzae*

# Spectrum of Infection in Asthma

## 1. Viral Infection (Primary Triggers)

다양한 respiratory virus가 천식 악화의 주요 원인으로 작용

### Common viruses

- Rhinovirus (most frequent)
- Respiratory syncytial virus (RSV)
- Influenza / Parainfluenza
- Coronavirus / Adenovirus

## 2. Bacterial Infection / Detection

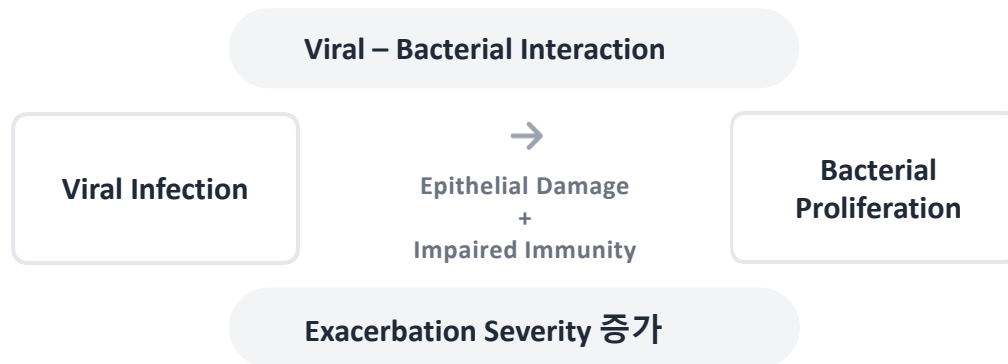
### Key Pathogens

- Haemophilus influenzae*
- Moraxella catarrhalis*
- Streptococcus pneumoniae*
- Pseudomonas aeruginosa*

### Clinical Relevance

- Post-viral bacterial overgrowth
- Chronic detection in severe asthma subset

## 3. Interaction Spectrum



## 4. Temporal Dimension

Pattern	Characteristics
Acute viral infection	Sudden exacerbation trigger
Post-viral bacterial	Delayed worsening / Secondary peak
Persistent colonization	Recurrent exacerbation & inflammation

**Asthma exacerbation reflects a spectrum of infection rather than a single mechanism**

# Viral vs Bacterial Infection in Asthma Exacerbation

	Viral Infection	Bacterial Infection
<b>Common organism</b>	Rhinovirus, RSV	<i>Haemophilus influenzae</i> , <i>Pseudomonas aeruginosa</i>
<b>Trigger role</b>	Primary trigger of exacerbation	Secondary or persistence-driven
<b>Onset</b>	Rapid (acute)	Subacute / chronic
<b>Immune response</b>	IFN-driven antiviral response	Neutrophilic, innate immune activation
<b>Key pathway</b>	↓ IFN (especially in asthma) → viral persistence	Impaired clearance → bacterial colonization
<b>Inflammation type</b>	T2 amplification (eosinophilic)	Non-T2 / neutrophilic
<b>Host defect</b>	Defective antiviral immunity	Impaired innate surveillance (NK, macrophage)
<b>Clinical pattern</b>	Episodic exacerbation	Recurrent / persistent symptoms
<b>Airway status</b>	Relatively reversible	Structural damage (bronchiectasis overlap)
<b>Treatment implication</b>	ICS, biologics (anti-IgE, anti-IL5)	Antibiotics, macrolide, host-directed therapy

# Host Phenotype Modifies Infection-Driven Exacerbation

## T2-High / Atopic Asthma

- Atopy, IgE sensitization
- Type 2 inflammation dominant
- Viral-triggered allergic inflammation

### Mechanism

- Th2 cytokine environment
- Amplified inflammatory response

### Clinical pattern

- Episodic / seasonal exacerbation

## Non-T2 / Infection-Prone Phenotype

- Less atopy, recurrent infection
- Severe / persistent disease
- Infection persists due to impaired clearance

### Mechanism

- Impaired host defense (↓IFN, NK dysfunction)
- Persistent airway inflammation

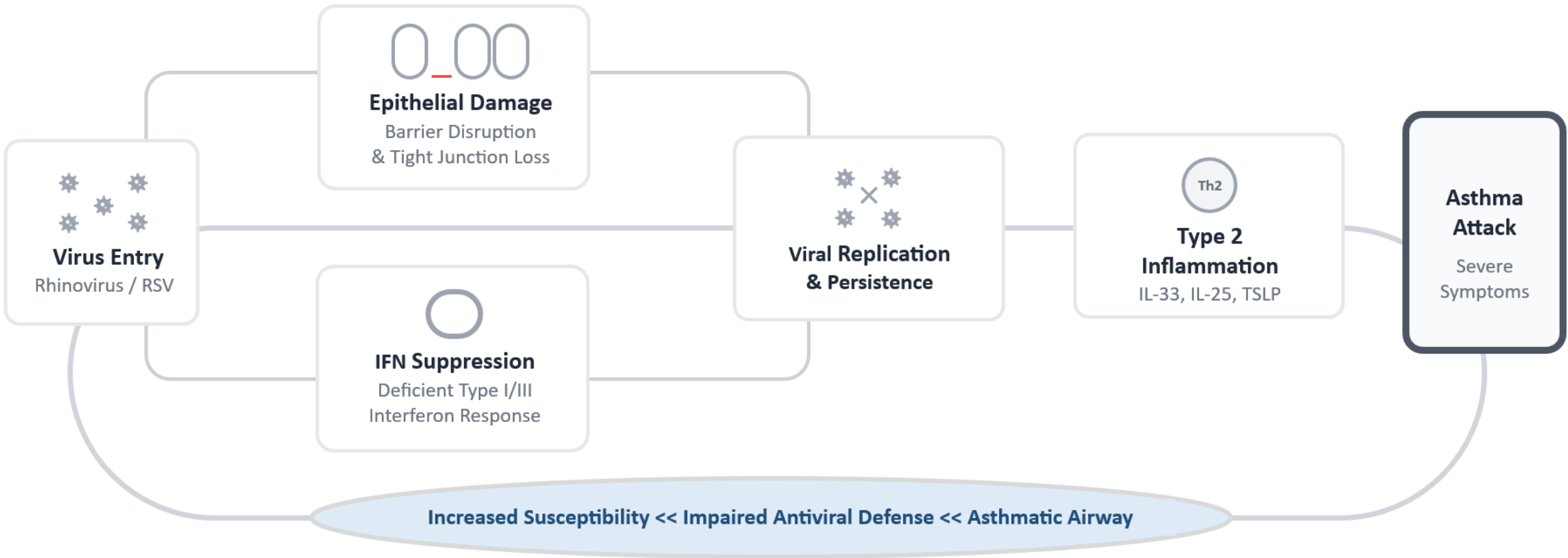
### Clinical pattern

- Recurrent / progressive

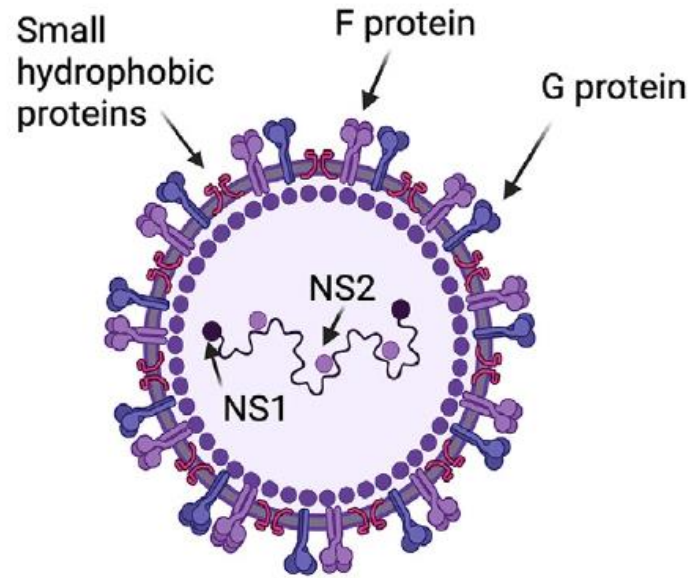
**Infection is a universal trigger,  
but its clinical impact depends on host susceptibility and immune phenotype.**

# Viral Infection Drives Asthma Exacerbation

## via Immune and Barrier Dysregulation



# Respiratory Syncytial Virus (RSV)



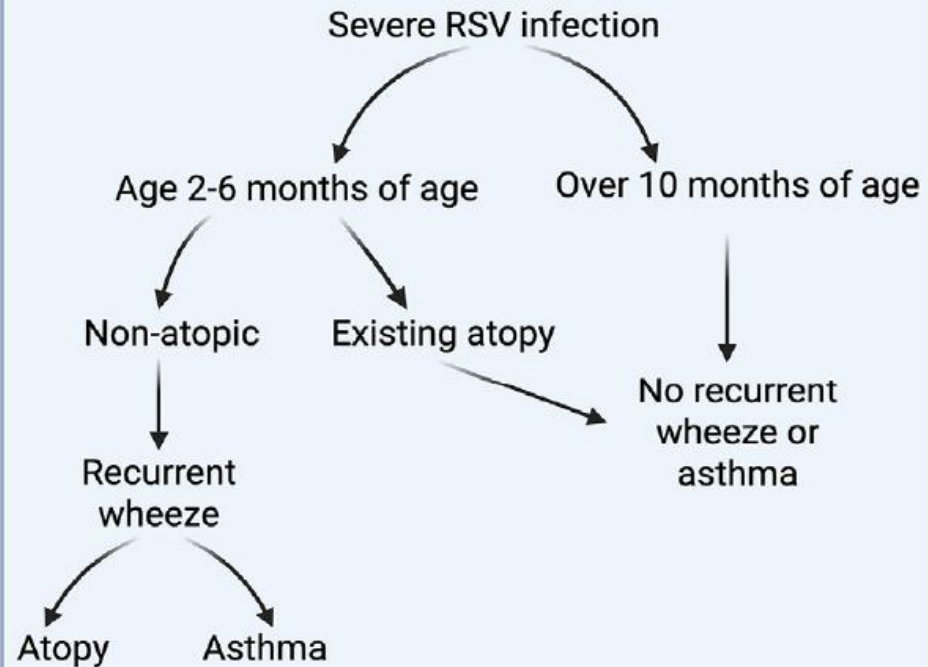
Two subtypes  
**RSV-A**  
 10 genotypes  
**RSV-B**  
 13 genotypes

*Asthma exacerbation*

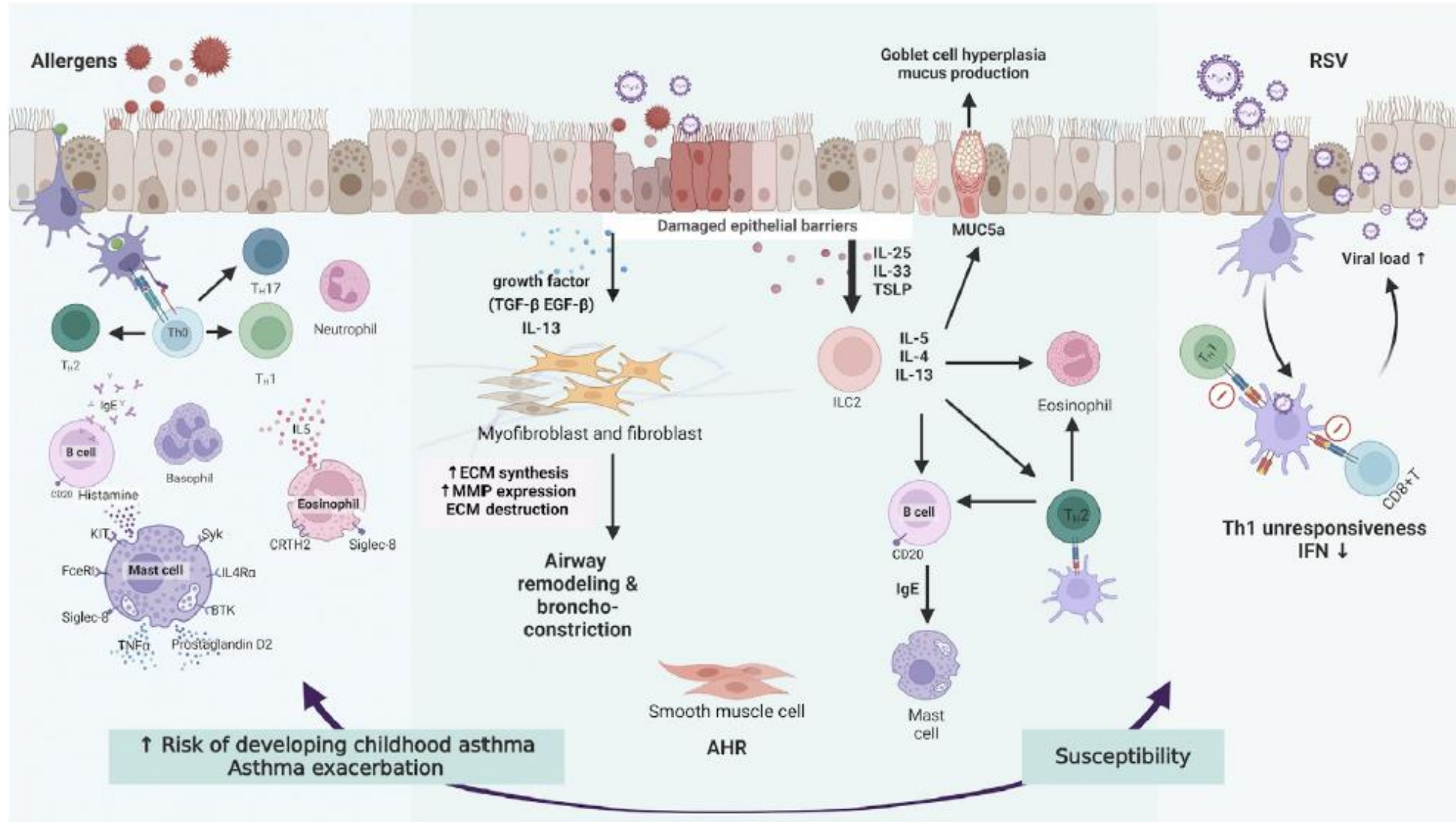
**Minor** cause of asthma exacerbations

- Causes 58,000-80,000 hospitalizations and 100-300 deaths annually in children less than 5 years old
- Circulates between October-April, peaking in December
- Multiple FDA approved vaccines and anti-RSV monoclonal antibodies

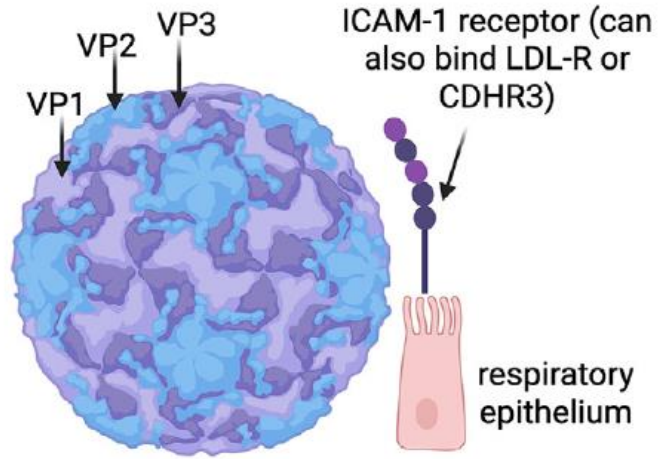
## *Development of asthma*



# RSV Infection Shifts the Immune Response from Th1 → Th2



# Rhinovirus (RV)



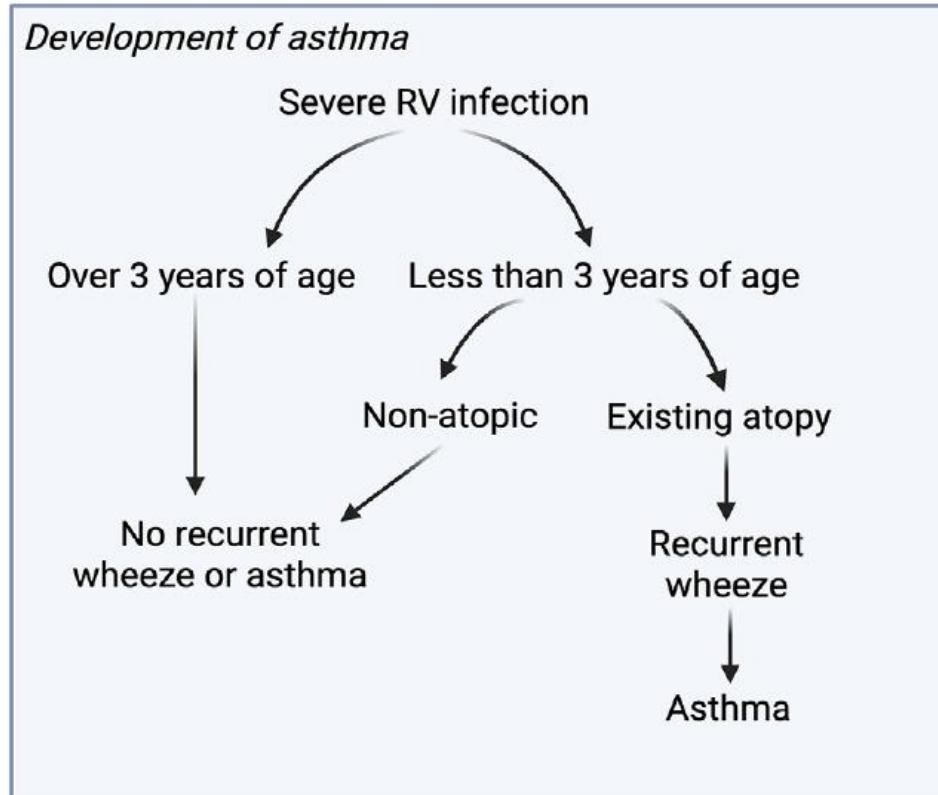
160 antigenically distinct serotypes across three species:

- RV-A
- RV-B
- RV-C

*Asthma exacerbation*

**Major** cause of asthma exacerbations

- Contributes to 30-50% of all respiratory tract infections
- The most frequently detected virus in children
- RV-A and RV-C are the predominant circulating species
- No FDA approved treatments, supportive care only



# Role of Other Respiratory Viruses in Asthma Exacerbation

	Influenza virus	Human coronavirus (SARS-CoV-2)	Human metapneumovirus (hMPV)
Characteristics	<ul style="list-style-type: none"> <li>모든 연령에서 asthma exacerbation 유발</li> </ul>	<ul style="list-style-type: none"> <li>wide clinical spectrum (asymptomatic → severe)</li> </ul>	<ul style="list-style-type: none"> <li>RSV와 유사한 paramyxovirus</li> <li>소아에서 wheezing 흔함</li> </ul>
Clinical significance with asthma	<ul style="list-style-type: none"> <li>exacerbation의 흔한 원인 (약 10–25%)</li> <li>severe AE risk 증가</li> </ul>	<ul style="list-style-type: none"> <li>generally weaker trigger compared to influenza</li> <li>not a strong independent risk factor (early pandemic data)</li> </ul>	<ul style="list-style-type: none"> <li>asthma exacerbation 유발 가능</li> <li>일부 연구에서:               <ul style="list-style-type: none"> <li>recurrent wheezing 증가</li> <li>asthma 위험 증가 가능성</li> </ul> </li> </ul>
Mechanism	<ul style="list-style-type: none"> <li>IL-33 증가</li> <li>ILC2 활성화 → IL-13 증가</li> <li>airway inflammation ↑</li> </ul>	<ul style="list-style-type: none"> <li>ACE2 ↓ (T2-high asthma) → susceptibility modulation</li> <li>IFN dysregulation / epithelial injury</li> <li>일부에서 persistent airway symptoms (long COVID)</li> </ul>	<ul style="list-style-type: none"> <li>airway epithelial infection / injury</li> <li>IL-33 release → ILC2 activation (→ IL-5, IL-13)</li> <li>relatively weak IFN response → delayed viral clearance</li> </ul>

# Respiratory Viruses in Asthma: Key Summary

## Major Viral Contributors

- Rhinovirus & RSV  
→ Primary drivers of asthma exacerbation
- Human metapneumovirus (hMPV)  
→ Associated with recurrent wheezing and possible asthma risk
- Human coronavirus (SARS-CoV-2)  
→ wide clinical spectrum, generally weaker trigger compared to influenza
- Influenza virus  
→ Contributes to exacerbation via IL-33-mediated inflammation

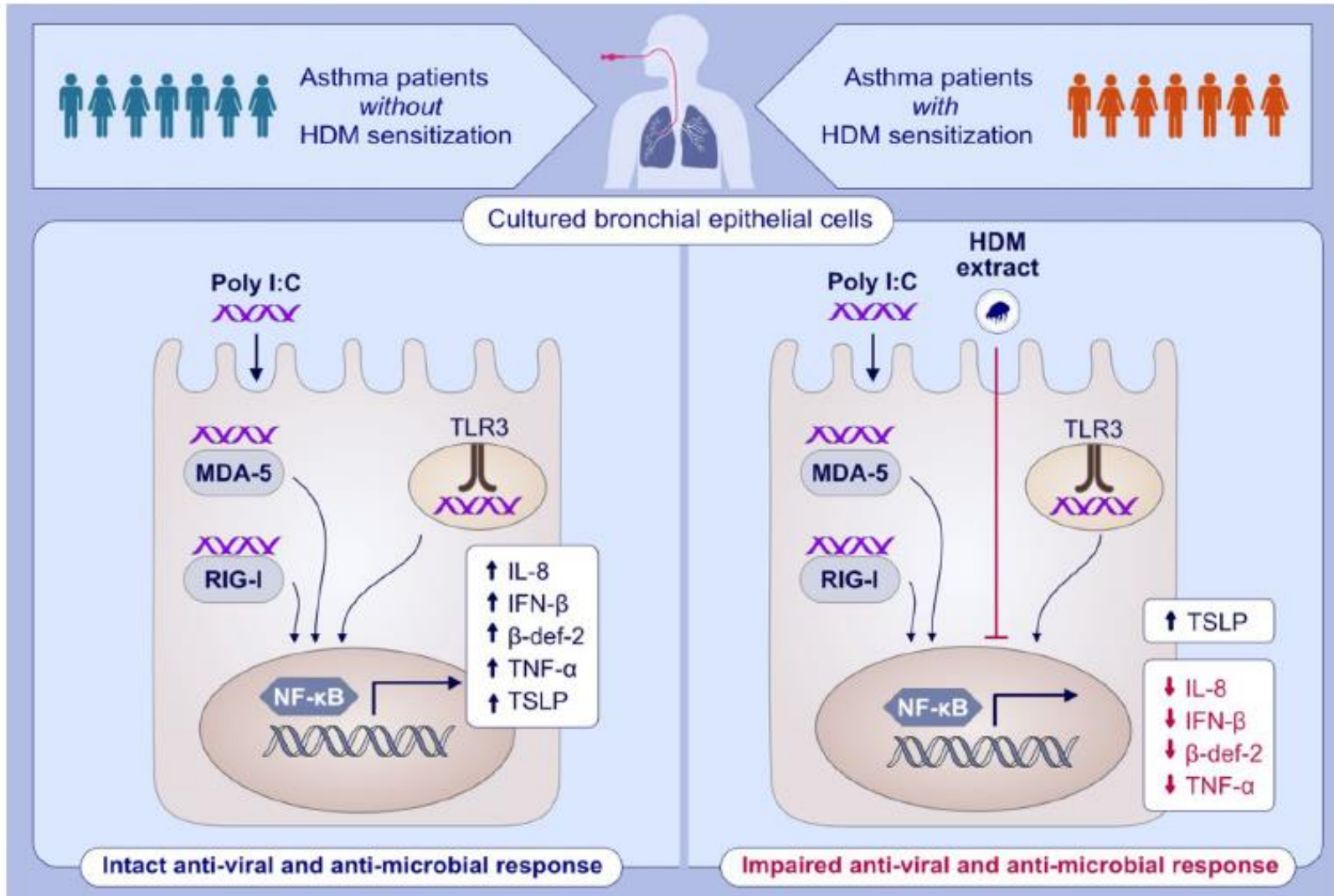
## Shared Mechanisms

- Airway epithelial infection
- Activation of type 2 inflammation
- Increased airway hyperresponsiveness

**Early-life infection burden may be more important than the specific virus type.**

**While multiple viruses contribute to asthma exacerbation, their roles differ, and importantly, the cumulative burden of infection may be more relevant than the specific virus**

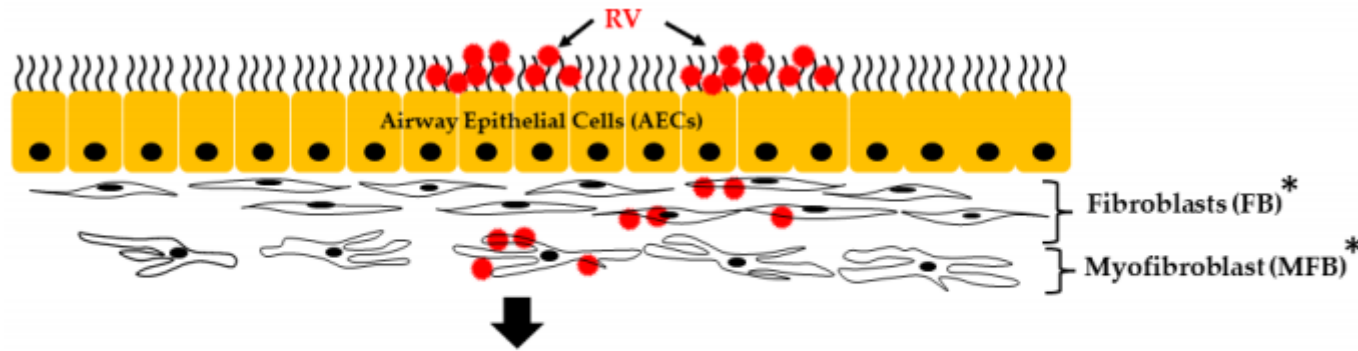
# Allergen Exposure Weakens Infection Defense in Asthma



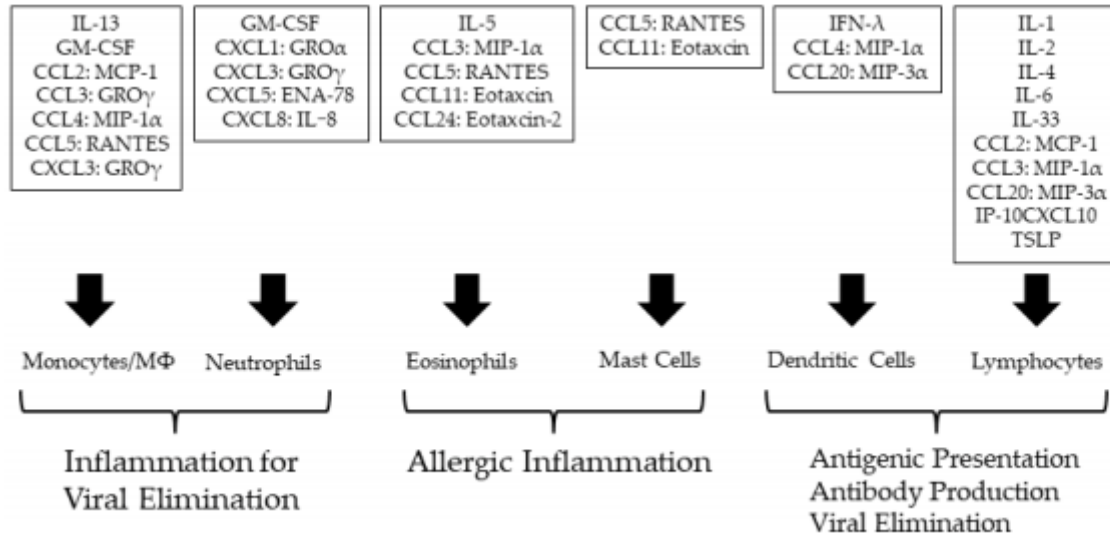
## virus + HDM 에서:

- IFN ↓ → viral clearance 감소
- IL-8/TNF ↓  
→ neutrophil recruitment 감소
- defensin ↓  
→ bacterial defense 감소
- TSLP 유지  
→ T2 inflammation 지속

# Rhinovirus-Induced Asthma: Beyond the Th1/Th2 Paradigm



RV-infection-induced cytokines from AECs, FB, and MFB



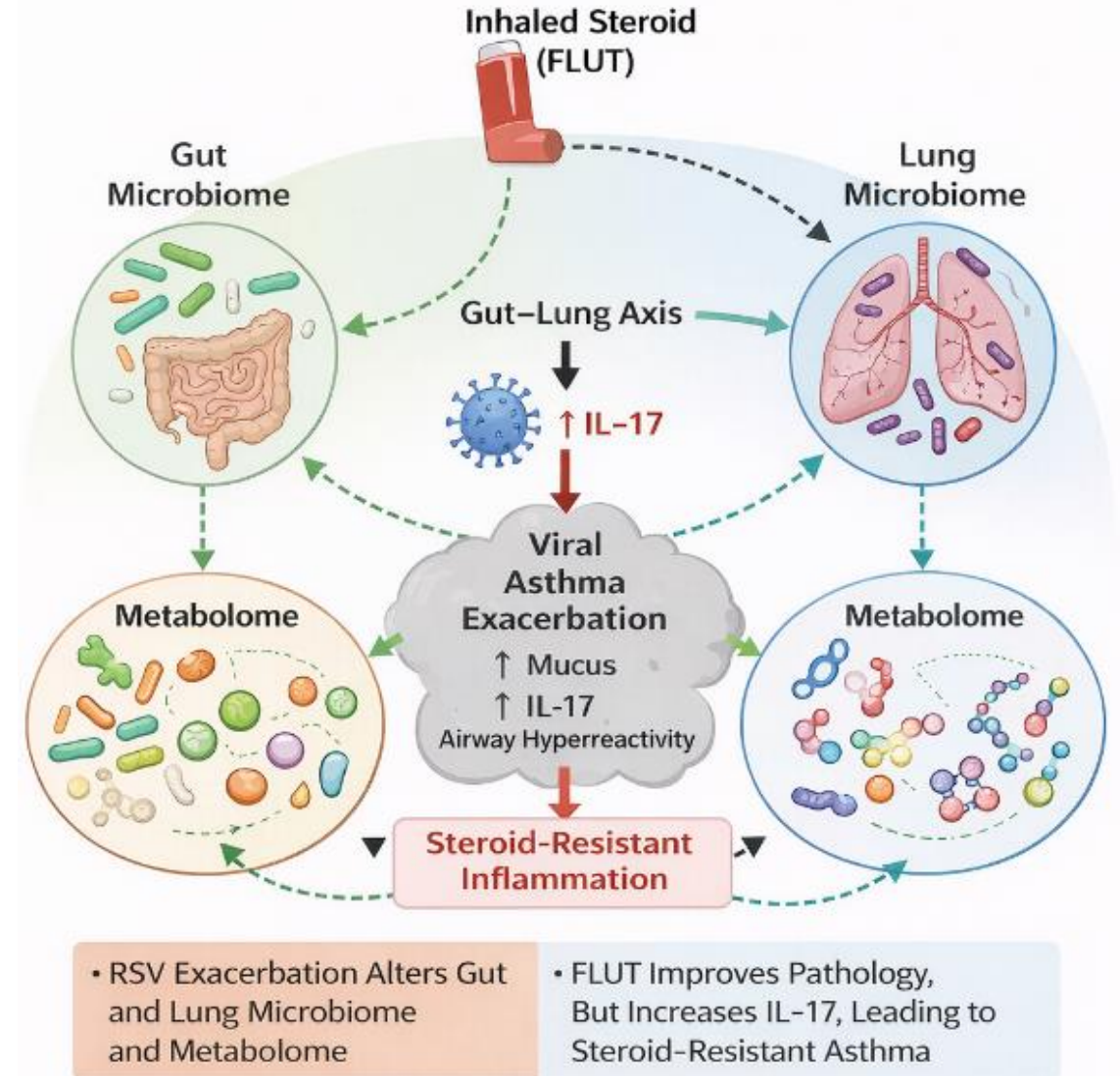
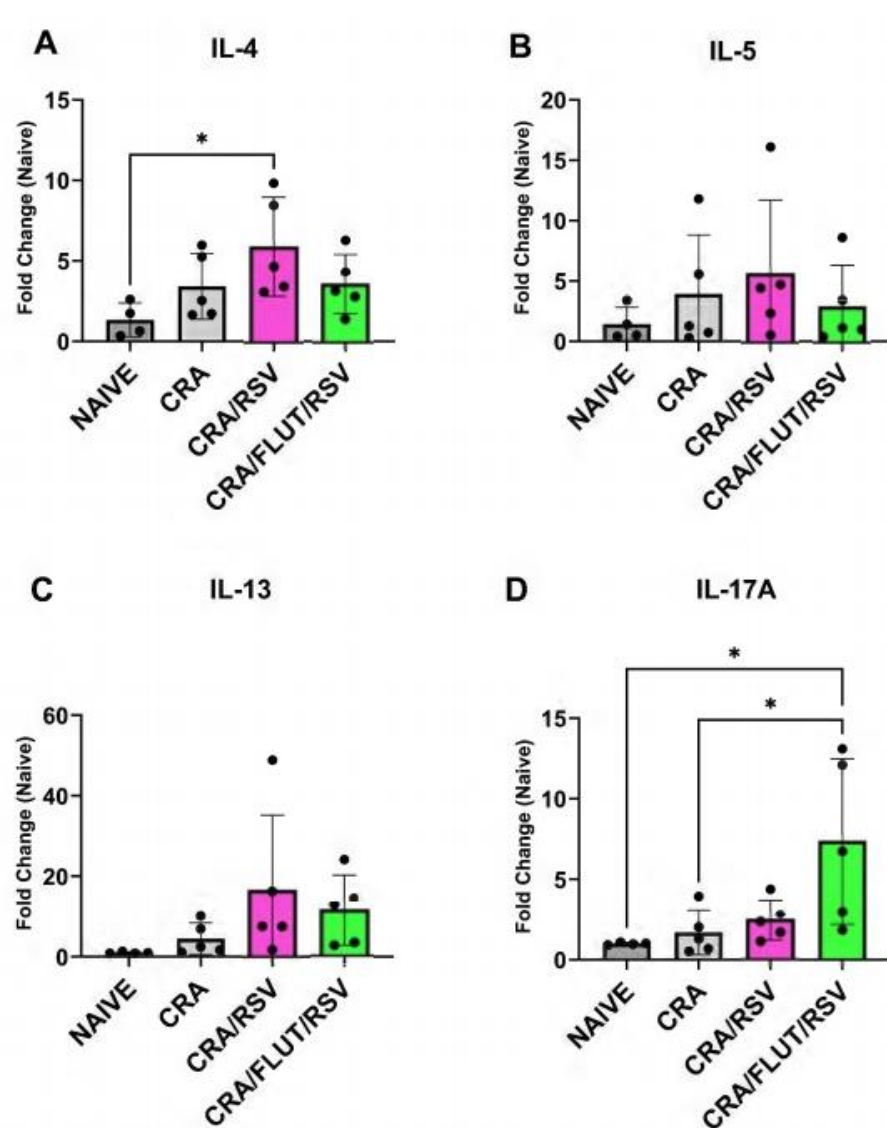
RV infection

- epithelial damage
- PRR activation
- cytokine release (Th1 + Th2)
- immune cell recruitment
- inflammation amplification
- airway remodeling
- asthma exacerbation

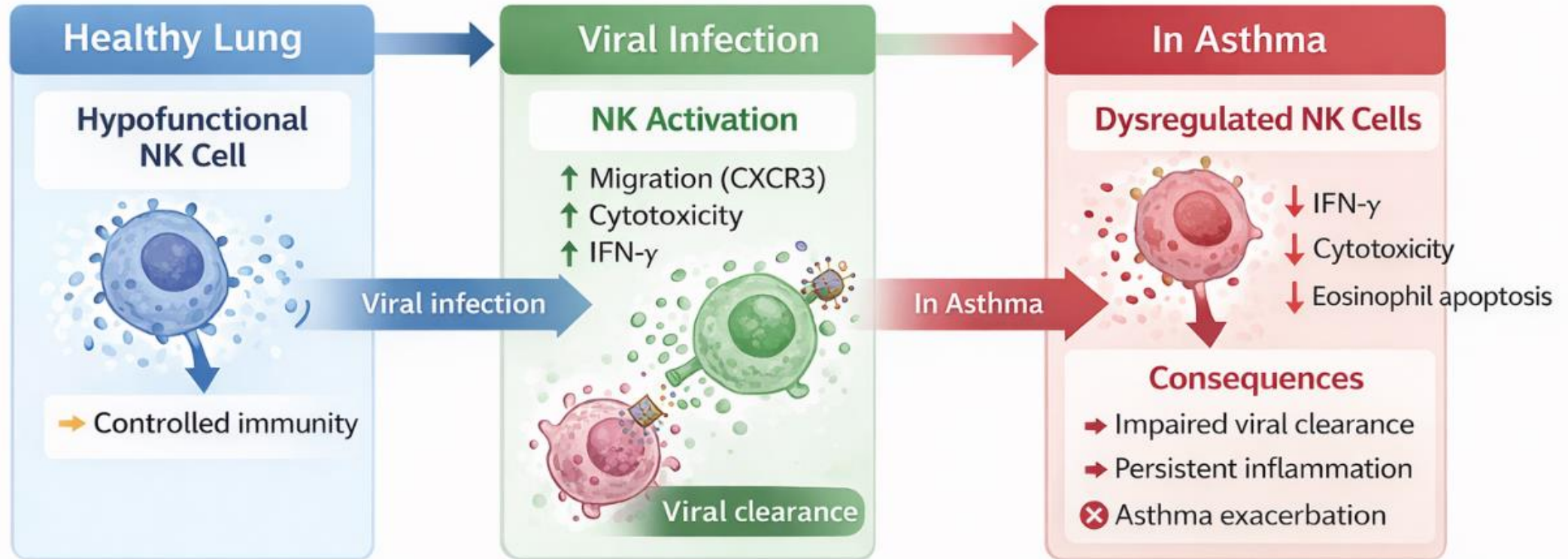
## Clinical Implications

- Virus-induced asthma  $\neq$  simple allergy
- Mixed immune phenotype disease
- Impaired **viral clearance** → **persistent inflammation**
- Target pathways:  
**IFN signaling, epithelial immunity, type 2 inflammation**

# Microbiome–Metabolome Interaction in Virus-Induced Asthma Exacerbation



# Dysregulated NK Cells Link Viral Infection to Asthma Exacerbation



NK cells are key antiviral effectors in the lung  
 Normally tightly suppressed to prevent tissue damage  
 In asthma, NK cells show functional dysregulation  
 This leads to failed viral control + unresolved inflammation

# Infection-Prone Asthma Phenotype: Who are they?

## Clinical Clues

- Frequent antibiotics exposure
- Severe asthma subset
- Bronchiectasis overlap (CT)
- Recurrent exacerbation despite ICS

## Microbiologic Pattern

- Repeated detection of same pathogen
- Chronic airway colonization
- Often subclinical persistence

## Dominant Bacterial Species

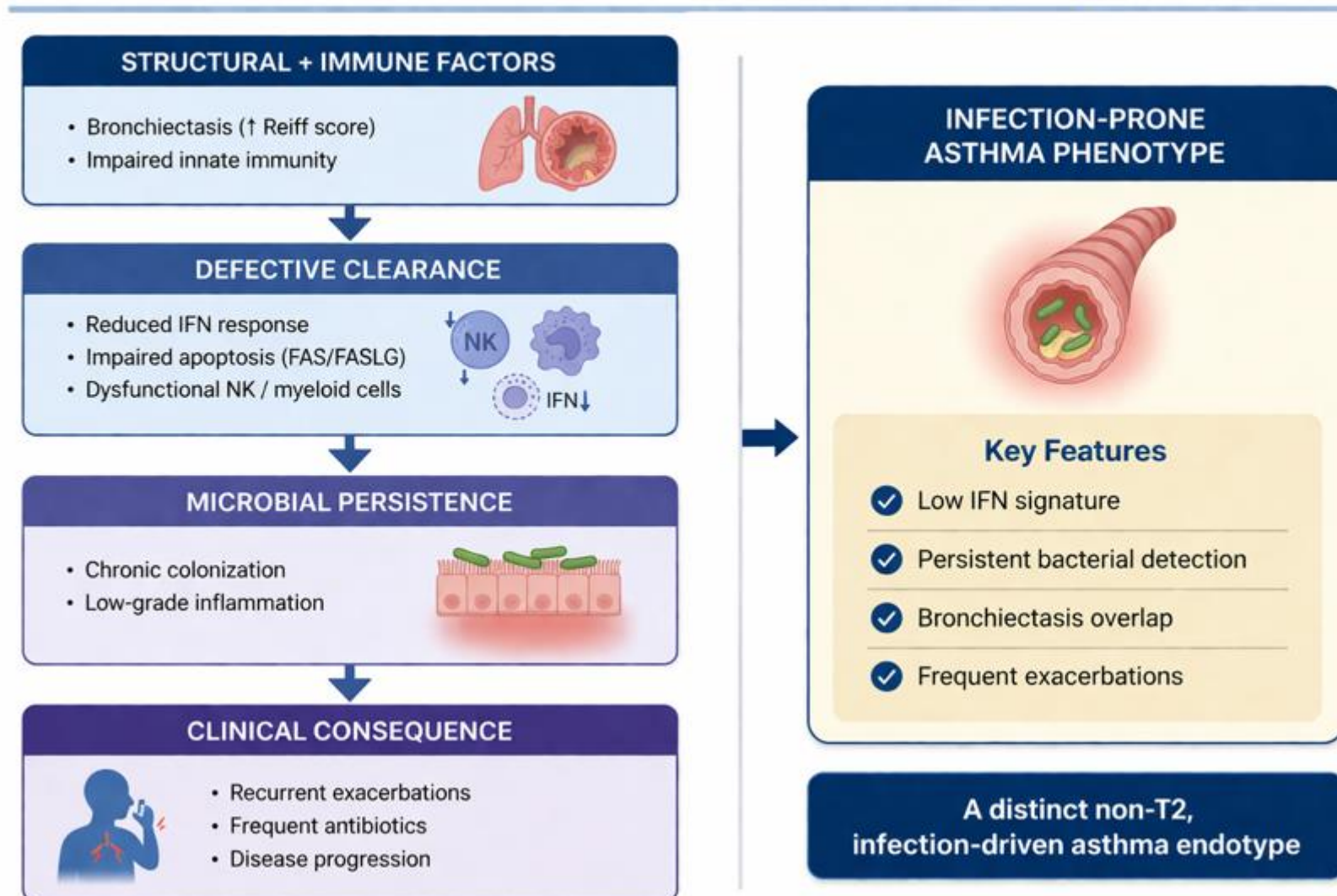
- *Pseudomonas aeruginosa*
- *Haemophilus influenzae*
- *Moraxella catarrhalis*

**Not just infection-prone, but clearance-impaired phenotype**

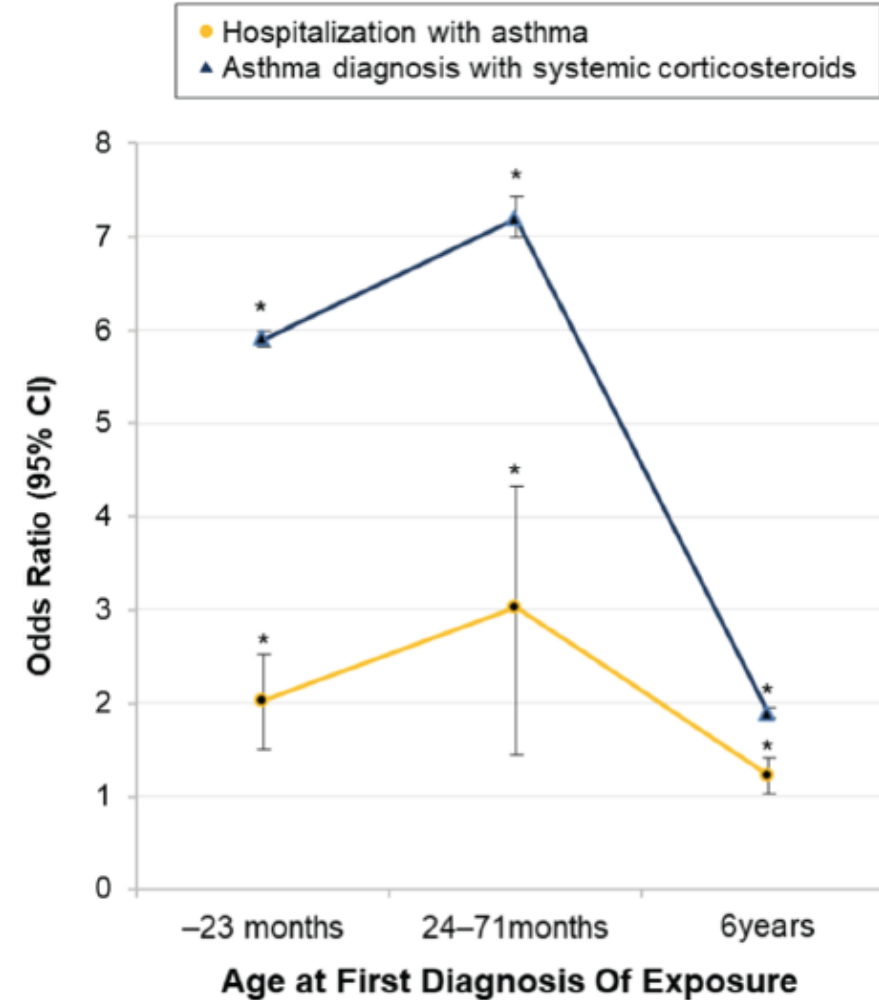
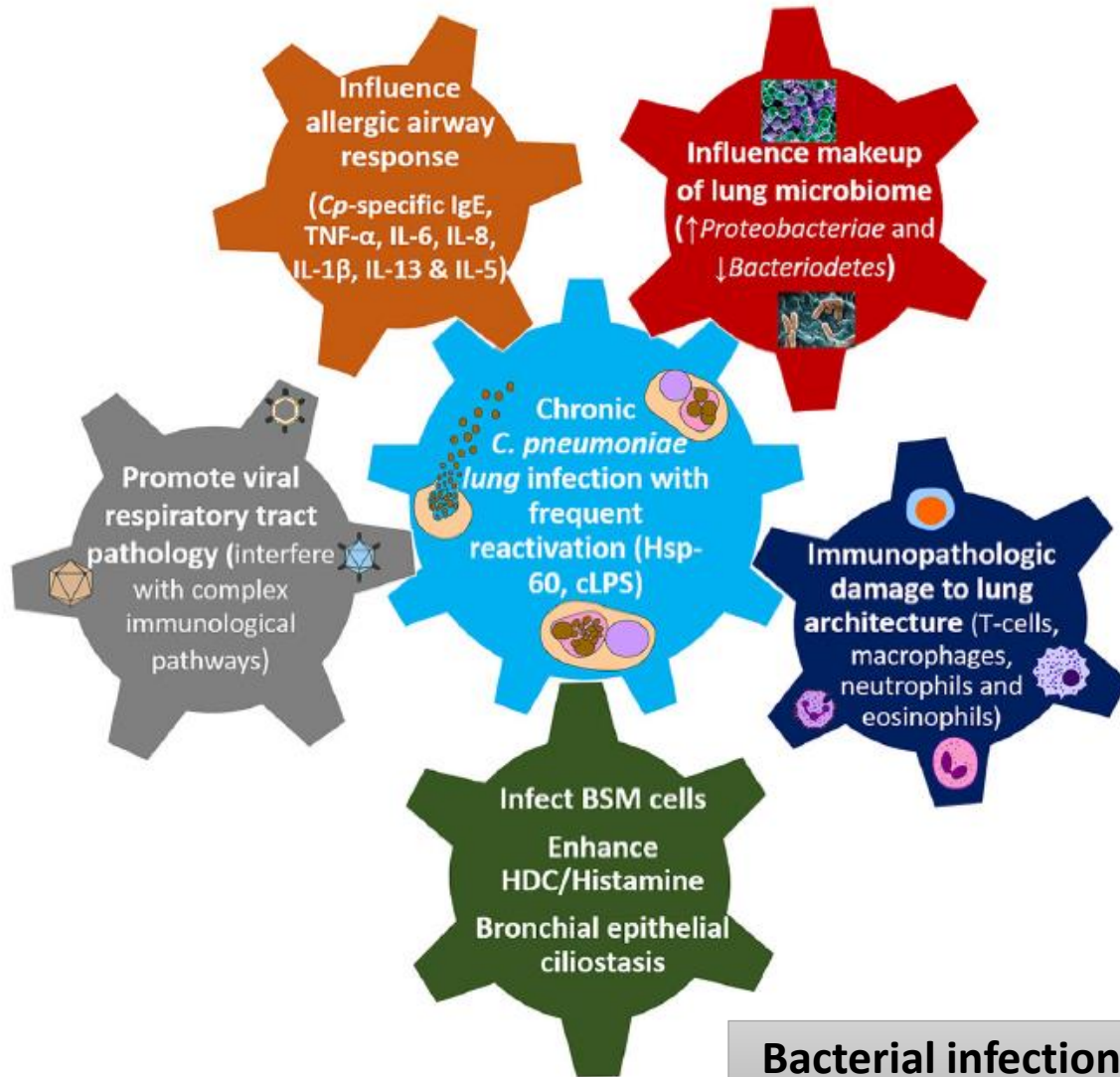
# Mechanistic Model: Infection–Exacerbation Cycle

## Proposed Model: Infection-Driven Exacerbation Loop

Mechanism of the Infection-Prone Asthma Phenotype

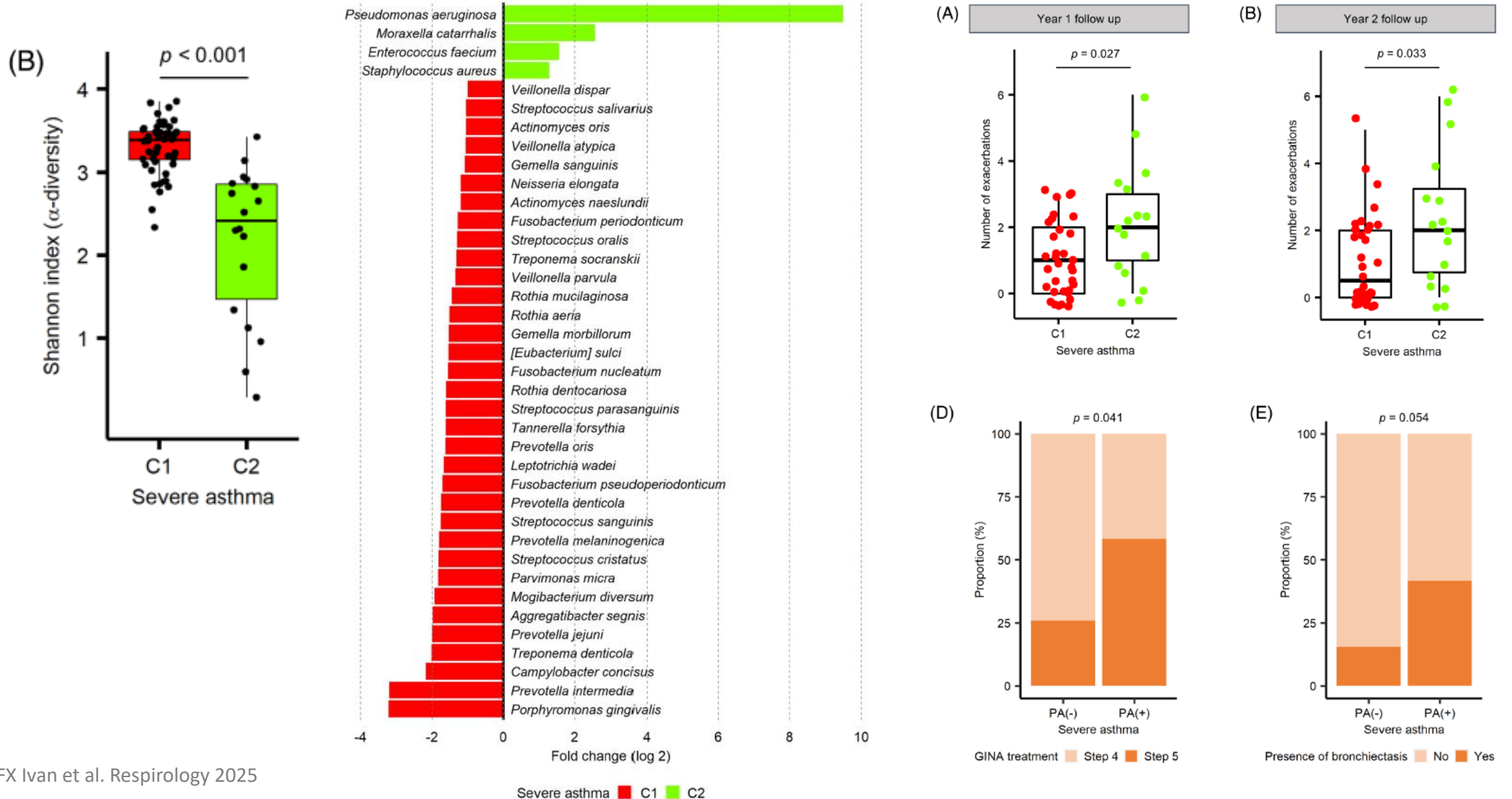


# Bacterial Infection–Driven Asthma Exacerbation



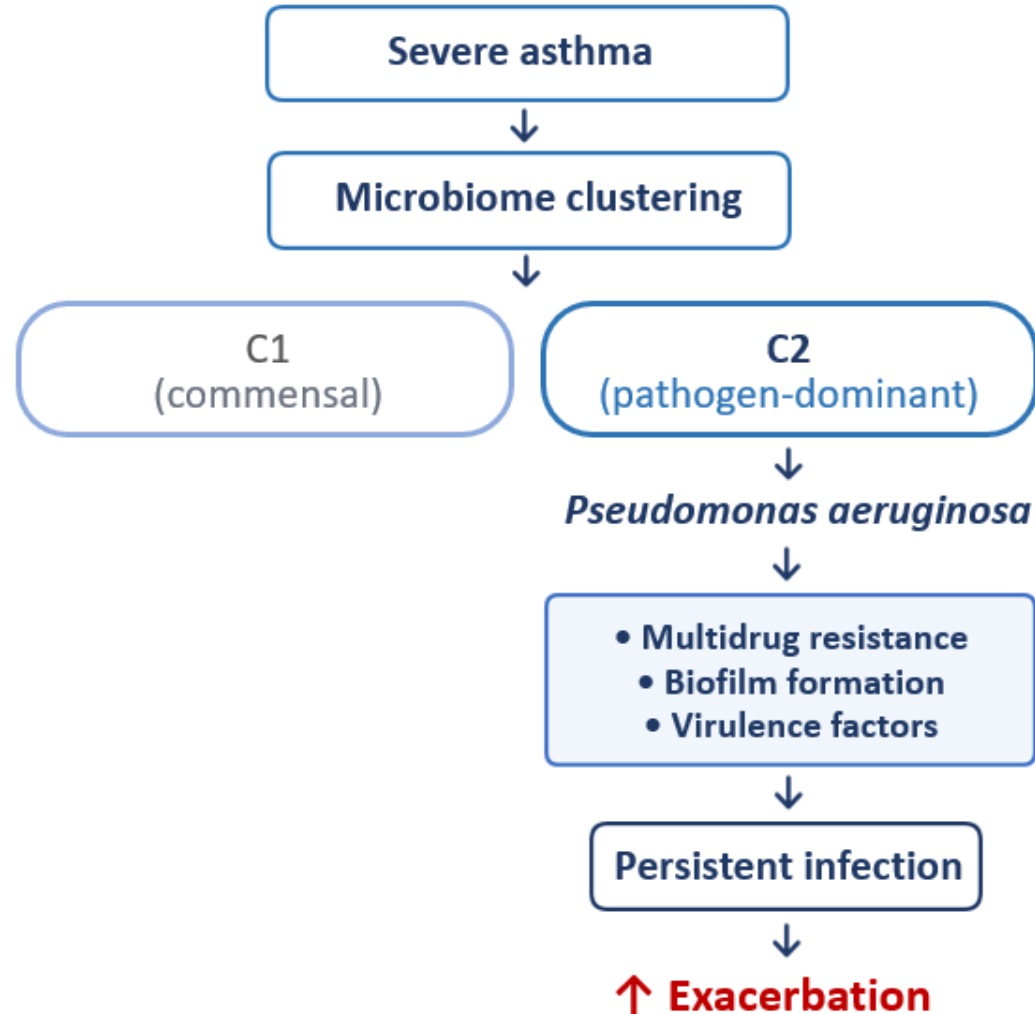
**Bacterial infection is not just a trigger — it is a disease-modifying factor in asthma**

# Pseudomonas-dominant microbiome defines a high-risk severe asthma phenotype



# Pseudomonas-dominant microbiome defines a high-risk severe asthma phenotype

## Mechanism / Concept



## Key Findings

### Analysis Results

↑ **Pseudomonas abundance**  
Significant increase observed

2 **Distinct Clusters**  
C1 (Commensal) vs C2 (Pathogen)

⚠ **Cluster C2 Impact:**  
↑ **Exacerbations (IRR ~2.4)**  
*\* Independent of bronchiectasis*

⌚ ↑ **Multidrug resistance genes**

⚙ ↑ **Virulence (biofilm, secretion)**

# Immunodeficiency-Driven Infection and Asthma Exacerbation

**Table 2.** Variations of selected features after 1 year of IRT in the T2-low group (panel A) and in the T2-high group (panel B)

	Baseline	After IRT	<i>p</i> value
<b>Panel A</b>			
Exacerbations/year	5.5 (1–16)	1.5 (0–7)	<0.01
Respiratory infections	3 (1–14)	1 (0–5)	<0.001
OCS	1/12 (8.3)	1/12 (8.3)	NS
OCS dose, mg	7.5	5	NS
BEC, cells/ $\mu$ L	80 (30–140)	115 (70–520)	<0.05
ACT	16 (10–19)	22 (18–25)	<0.01
ACQ5	2.4 (1.7–3.5)	0.4 (0.1–1.1)	<0.01
	Baseline	After IRT	<i>p</i> value
<b>Panel B</b>			
Exacerbations/year	7 (2–20)	1 (0–6)	<0.001
Respiratory infections	5 (0–20)	1 (0–5)	<0.001
OCS	5/21 (23.8)	2/21 (9.5)	<0.005
OCS dose, mg	17 (5–25)	3.75 (2.5–5)	NS
BEC, cells/ $\mu$ L	200 (50–760)	170 (10–1,400)	NS
ACT	15 (11–18)	23 (18–25)	<0.01
ACQ5	2.6 (1.9–3.3)	0.5 (0.2–1.3)	<0.01

Results:

36.4%가 T2-low phenotype

IVIg 효과

- 감염 & 악화 감소
- infection: 4  $\rightarrow$  1 ( $p < 0.001$ )
- exacerbation: 6  $\rightarrow$  1 ( $p < 0.001$ )

Phenotype 변화

T2-low 환자에서:

- eosinophil 증가 (80  $\rightarrow$  115,  $p < 0.05$ )
- 일부는 T2-high로 이동

# Infection as a Phenotype-Modifying Driver in Asthma

## ① Immunodeficiency



↓ IgG / PAD  
impaired humoral  
immune response  
reduced pathogen  
clearance

## ② Recurrent Infections



Frequent viral (e.g.,  
HRV, RSV) and/or  
bacterial (e.g.,  
*H. influenzae*,  
*S. pneumoniae*)  
exacerbations

## ③ Airway Inflammation & Remodeling



Persistent inflammation  
(Type 2 and/or non-T2)  
Structural changes  
(e.g., goblet cell  
hyperplasia, smooth  
muscle thickening)

## ④ Dysbiosis & Chronic Colonization



Reduced microbial  
diversity  
Emergence of  
pathogenic bacteria  
Creation of a  
pro-inflammatory  
microenvironment

## ⑤ Severe, Treatment-Resistant Asthma



Persistent symptoms  
Frequent, severe  
exacerbations  
Corticosteroid resistance  
Need for biologics or  
other advanced therapies

**RECURRENT INFECTIONS** and the associated immune responses can **DRIVE** the progression to a more **SEVERE, INFLAMMATORY,** and **TREATMENT-RESISTANT** asthma phenotype.

### KEY CORRELATIONS:

Higher infection rate → ↑ FeNO levels  
↑ Sputum neutrophils → ↓ FEV1 % predicted  
Antibiotic use → Altered airway microbiome

# Phenotype-Based Therapeutic Approach in Infection-Associated Asthma

## 1. Universal infection response (all asthma)

- Viral/bacterial trigger → epithelial injury + inflammation
- Standard care: ICS ± LABA Short-term OCS (exacerbation)

## 2. T2-amplified phenotype

- Virus → ↑ IL-33 / TSLP → ↑ IgE / eosinophilia
- Key feature: exaggerated T2 response
- Treatment focus: Anti-IgE (omalizumab), Anti-IL5 / IL5R (mepolizumab, benralizumab), Anti-IL4R (dupilumab)  
Anti-TSLP (Tezepelumab)
- Effect: ↓ exacerbation frequency ↓ virus-induced amplification

## 3. Immune-impaired / infection-prone phenotype

- ↓ IFN response
- impaired NK / macrophage function
- chronic bacterial persistence
- Treatment focus:  
NOT inflammation suppression alone  
→ host defense restoration

# Expanding Treatment Strategies: Beyond Anti-Inflammation

## 1. Macrolide Therapy

(Azithromycin)

### Anti-inflammatory + Antibacterial

↓ Exacerbation (esp. non-T2 / neutrophilic)

### Key Mechanism

- ✓ ↓ IL-8 / neutrophil recruitment
- ✓ Modulation of airway microbiome

### Best Candidates

- ✓ Frequent exacerbators
- ✓ Chronic bacterial detection

## 2. ICS Balance

Critical Issue

### Dual Effect

- ✓ ↓ T2 inflammation
- ⚠ BUT → ↑ Infection risk (bacterial, viral)

### Clinical Implication

- > Overuse in non-T2 → Harmful
- > Need precision dosing

⚠ ICS Overuse Risk

## 3. Biologics

### Primary Benefit

- ★ Strong effect in T2-high asthma
- ⊕ Additional benefit: ↓ Virus-induced exacerbation (e.g., anti-IgE)

### Limitation

⊖ Limited effect in infection-driven / non-T2 phenotype

## 4. IFN Augmentation

Emerging

**Target:** Impaired antiviral response

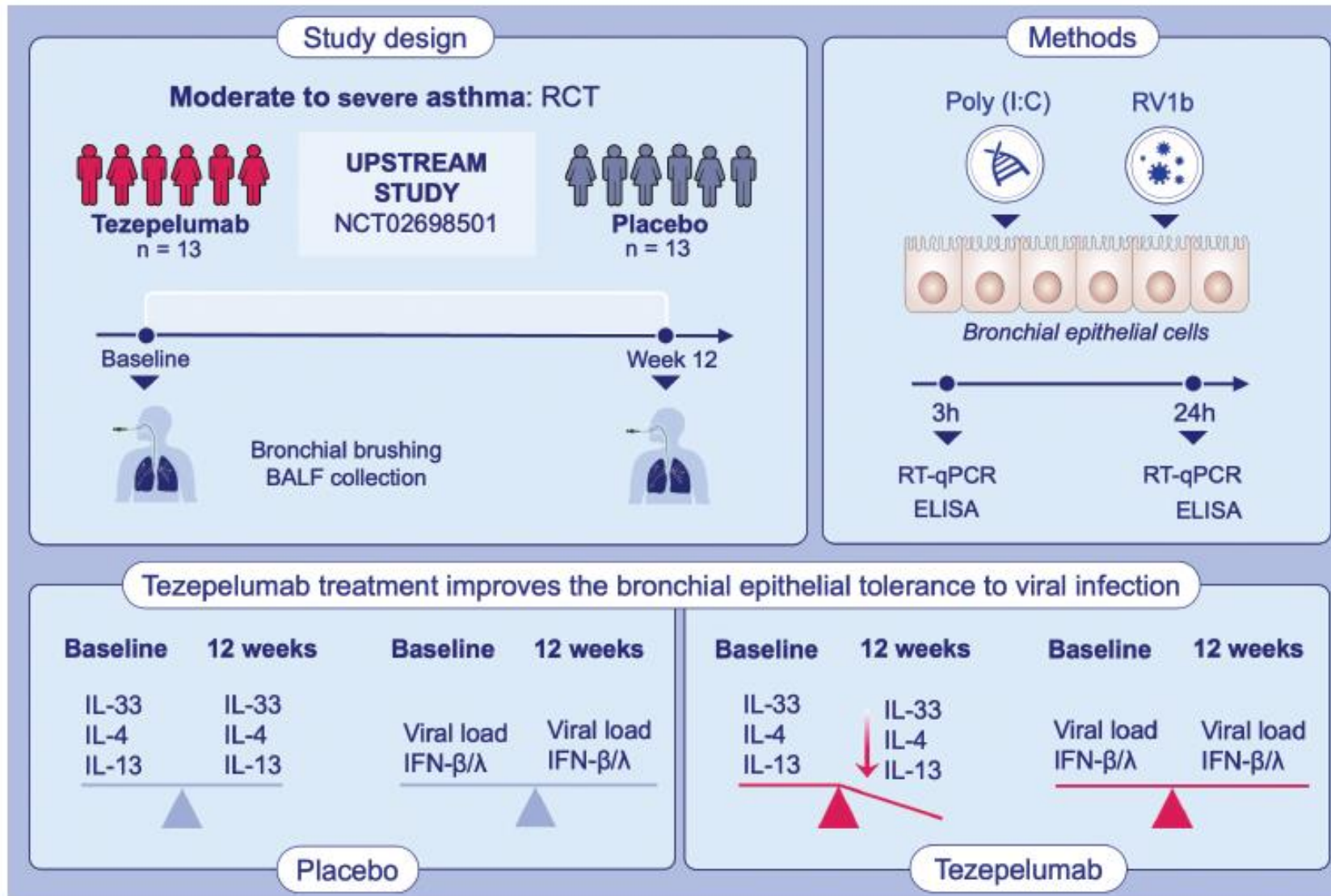
### Examples

- 👤 Inhaled IFN-β (clinical trials)
- ⌘ TLR agonists

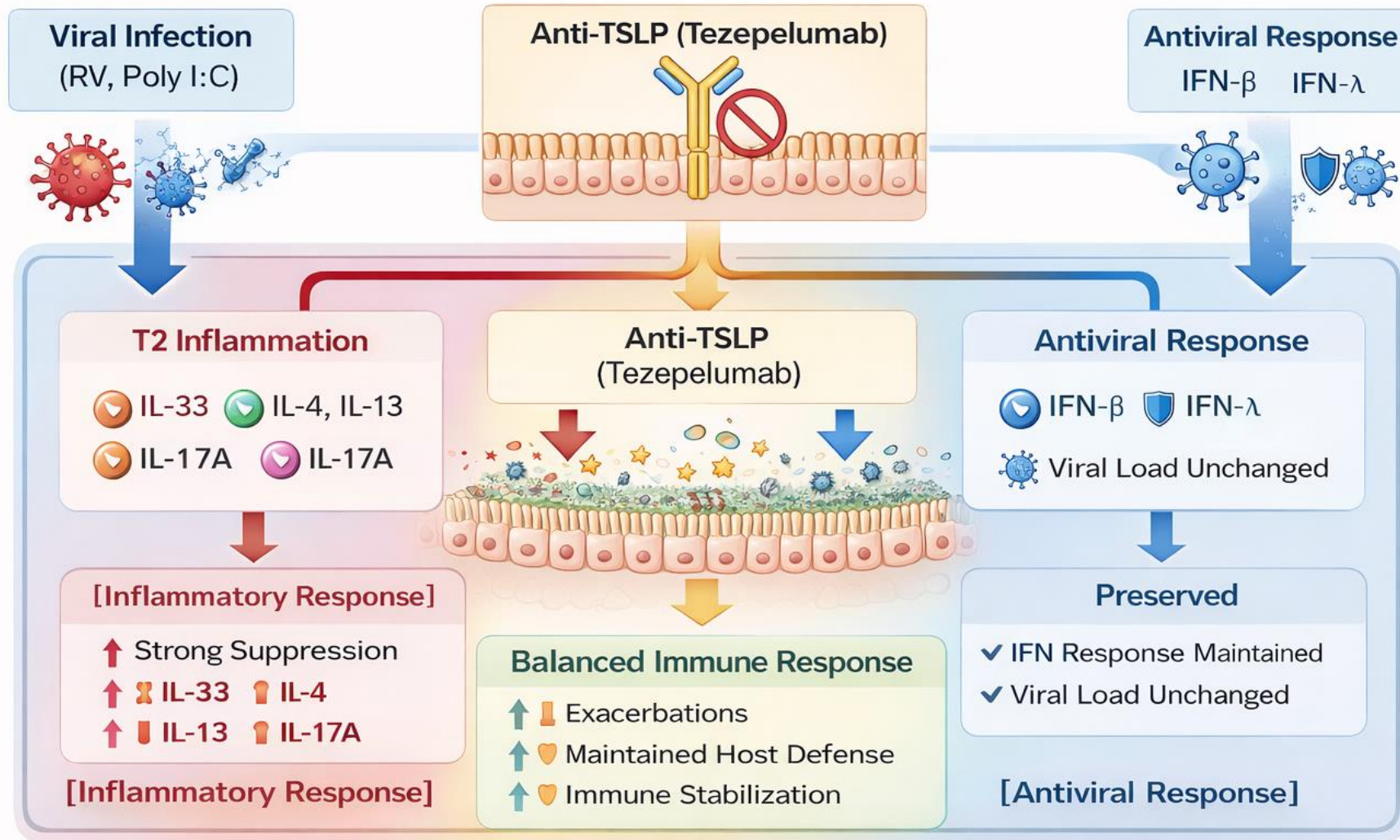
### Potential Role

- 🎯 Infection-prone asthma
- 🕒 Early viral infection phase

# Tezepelumab: Modulating Viral-Induced Inflammation Without Impairing Antiviral Defense



**Tezepelumab: Selective Modulation of Viral-Induced Immune Response**



Tezepelumab suppresses inflammatory amplification while preserving antiviral immunity.

# Toward Precision Therapy in Infection-Prone Asthma

## Candidate biomarkers

- ✓ IFN signature (↓ = infection-prone)
- ✓ Blood transcriptomics
  - immune suppression pattern
  - interferon pathway downregulation
- ✓ NK cell function
  - cytotoxicity ↓
  - impaired viral/bacterial clearance
- ✓ Microbiome profiling
  - chronic colonization pattern
  - dysbiosis

## Future directions

Define infection-prone asthma cohort

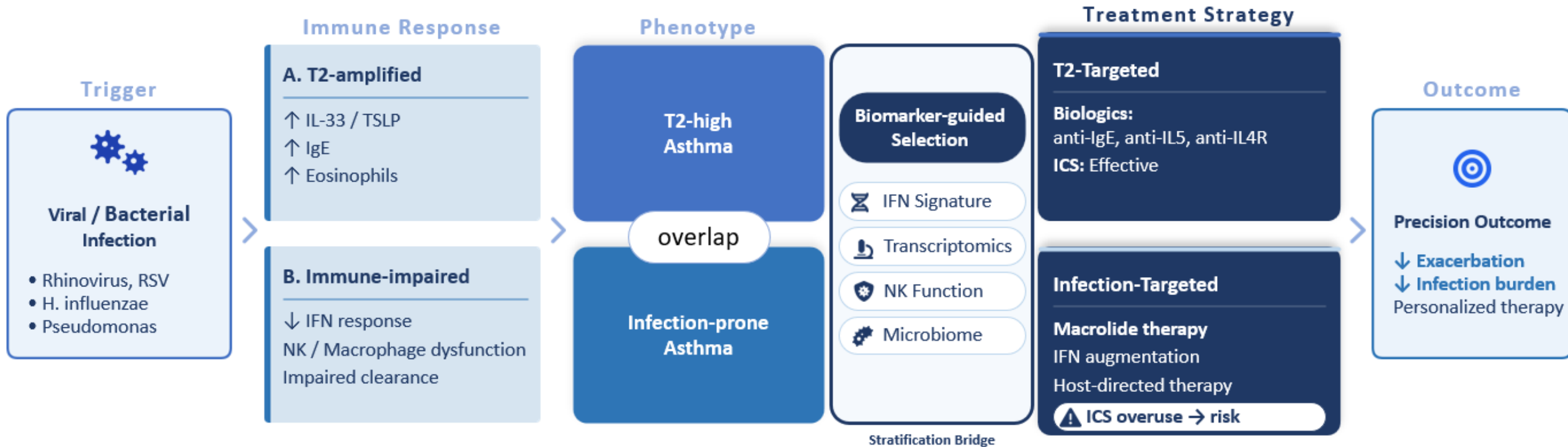
Integrate:

immune profiling  
microbiome  
transcriptomics

Goal:

**Mechanism-based, not symptom-based treatment**

# Precision Treatment in Infection-Associated Asthma



# Summary

- Infection  $\neq$  single mechanism
- Host immunity determines outcome
- T2-high  $\rightarrow$  inflammation amplification
- Infection-prone  $\rightarrow$  impaired clearance
- Bacterial persistence  $\rightarrow$  disease progression
- **$\rightarrow$  Phenotype-based precision therapy is essential**

경청해 주셔서 감사합니다.

