SUPPLEMENTARY MATERIAL

Supplementary figure 1. Comparison of sphingolipid metabolites among the groups studied A. Diacylglycerol; B. Ceramide; C. Sphingomyelin; D. Sphingosine; E. Sphingosine-1-phosphate; F. Sphinganine. The significance of any differences between groups was tested using the Mann-Whitney *U* test. *P*-values were corrected for multiple comparisons using the Benjamini and Hochberg's false discovery rate (FDR).



*Mann-Whitney U test, P-values were corrected for multiple comparisons using the Benjamini and Hochberg's false discovery rate (FDR).

Supplementary figure 2. Correlation between total IgE, egg white/milk-specific IgE, and gut sphingolipid

A. Total IgE; B. Egg white-specific IgE; C. Milk-specific IgE. sphingolipid metabolites, and clinical parameters are log transformed. The data was analyzed by spearman correlation.

Log [18:0/18:2 DAG Log [18:0/20:4 og [18 00 50 r = 0.121 • = 0.191 = 0.171 = 0.179 = 0.182 Log [Total IgE, kU/L] i ż ż Log [Total IgE, kU/L] Log [Total IgE, kU/L] Log [Total IgE, kU/L] Log [Total IgE, kU/L] Log [24:0 Sphingo Log [16:0 Log [24:1 Pog r = 0.182 0.196 Log = 0.267 0.205 0.217 Log [Total IgE, kU/L] Log (Sphingosine 0.138 r = -0.149 Log [Total IgE, kU/L] Log [Total IgE, kU/L]

(A) Total IgE

J Investig Allergol Clin Immunol 2024; Vol. 34(4): 263-265 doi: 10.18176/jiaci.0979

(B) Specific IgE to Egg white



(C) Specific IgE to Milk



J Investig Allergol Clin Immunol 2024; Vol. 34(4): 263-265 doi: 10.18176/jiaci.0979

Supplementary figure 3. Principal component analysis (PCA) score plot of target lipidome data from sample feces and QCs was generated using MetaboAnalyst 5.0 after log transformation of target lipidome data. No filter was applied, and missing values were estimated by replacement with 1/5 of the minimum positive value in the original data.

