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1 Increased periostin associates with greater airflow limitation in patients receiving inhaled corticosteroids $\mathbf{2}$

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48 Abstract

Background: Periostin, an extracellular matrix protein, contributes to subepithelial 49thickening in asthmatic airways, and its serum levels reflect airway eosinophilic 50inflammation. However, the relationship between periostin and the development of airflow 5152limitation, a functional consequence of airway remodeling, remains unknown. 53**Objective:** To determine the relationship between serum periostin levels and pulmonary function decline in asthmatic patients on inhaled corticosteroid (ICS) treatment. 54Methods: 224 asthmatic patients (average age 62.3 years) treated with ICS for at least 554 years were enrolled. Annual changes in forced expiratory volume in one second (FEV₁), 56from at least one year after the initiation of ICS treatment to the time of enrollment or later 57(average 16.2 measurements over 8 years per individual), were assessed. At enrollment, 58clinical indices, biomarkers including serum periostin, and periostin gene polymorphisms 59were examined. Associations between clinical indices or biomarkers and a decline in FEV₁ of 60 $30 \text{ mL} \cdot \text{yr}^{-1}$ or greater were analyzed. 61

Results: High serum periostin levels ($\geq 95 \text{ ng/mL}$) at enrollment, the highest treatment step, higher ICS daily doses, a history of admission due to asthma exacerbation, comorbid or a history of sinusitis, and ex-smoking were associated with a decline in FEV₁ of 30 mL·yr⁻¹ or greater. Multivariate analysis revealed that high serum periostin, the highest treatment step, and ex-smoking were independent risk factors for the decline. Polymorphisms of periostin gene were related to higher serum periostin levels (rs3829365) and a decline in FEV₁ of 30 mL·yr⁻¹ or greater (rs9603226).

- 69 Conclusions: Serum periostin appears to be a useful biomarker for the development of
 70 airflow limitation in asthmatic patients on ICS.
- 71

72 Clinical implications (25 words)

73 Serum periostin levels reflect greater FEV₁ decline in asthmatic patients on inhaled

74	corticosteroid treat	nent. POSTN gen	e polymorphism	s may also be	e helpful for	identifying
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- 75 rapid FEV_1 decliners.
- 76 Key words
- Asthma, inhaled corticosteroids, lung function decline, periostin, *POSTN* gene polymorphism,
- 78 sinusitis, treatment step
- 79

80 Abbreviations

- 81 ACT: asthma control test
- 82 ECP: eosinophil cationic protein
- 83 FAS I: fasciclin I
- 84 FEV₁: forced expiratory volume in one second
- 85 FVC: forced vital capacity
- 86 hsCRP: high sensitivity C-reactive protein
- 87 ICS: inhaled corticosteroids
- 88 IgE: immunoglobulin E
- 89 IL: interleukin
- 90 ROC: receiver operating characteristic
- 91 SNP: single-nucleotide polymorphism
- 92 TGF- β : transforming growth factor beta
- 93
- 94 Total word counts for the text and the abstract are 3800 and 258 words, respectively.

95 **Capsule summary (32 words)**

- 96 This is the first study to identify a relationship between high serum periostin and greater
- 97 annual decline in FEV_1 , which sheds new light on serum periositin as a useful biomarker in
- 98 asthma.

99 Introduction

Airway inflammation and remodeling are key features of asthma that have been 100demonstrated by pathological¹ and radiological findings^{2,3}. Physiologically, patients with 101 asthma show a greater decline in pulmonary function than subjects without asthma⁴. Studies 102that were mostly conducted in the era before inhaled corticosteroids (ICS) demonstrated that 103more severe symptoms or severe exacerbations⁵⁻⁷, long-standing asthma⁸, and smoking 104 history^{4, 8} were moderate to strong risk factors for greater decline in pulmonary function⁵. 105Blood and sputum eosinophilia^{9, 10} and genetic predisposition¹¹⁻¹³ were also potential risk 106 factors. Owing to early intervention with ICS, however, airway inflammation and the degree 107 of annual decline in pulmonary function have been attenuated in a majority of asthmatic 108 patients¹⁴⁻¹⁶. Meanwhile, a subset of patients still show accelerated decline in FEV₁ and 109 develop irreversible airway obstruction despite adequate treatment^{17, 18}. van Veen et al. found 110 that exhaled nitric oxide of 20 ppb or higher is a predictor of accelerated decline in 111pulmonary function in patients with difficult-to-treat asthma¹⁸. However, other biomarkers for 112113greater decline in FEV₁ despite treatment with ICS remain unknown. The airway inflammation of asthma is classically characterized by infiltration and 114 activation of eosinophils, mast cells, and Th2 cells with several mediators and Th2 cytokines, 115such as interleukin (IL)-4, IL-5, and IL-13^{19, 20}. Periostin, a secreted, 90-kDa, extracellular 116 matrix protein that is induced by IL-4 and IL-13, was originally isolated as an osteoblast-117specific factor; it shares structural homology to the insect cell adhesion molecule fasciclin I

specific factor; it shares structural homology to the insect cell adhesion molecule fasciclin I
(FAS I) and binds to fibronectin, tenascin-C, and collagen^{21, 22}. In airway epithelial cells
collected from patients with asthma, periostin is one of the up-regulated genes²³, and its
expression is correlated with thickness of the airway basement membrane²⁴. Takayama et al.
clearly demonstrated that periostin is deposited in the airway subepithelial layer in asthmatic
patients. Moreover, serum periostin is identified as the single best predictor of airway
eosinophilia in patients with severe asthma who remain symptomatic despite maximal ICS

treatment²⁵. Therefore, we hypothesized that periostin would be a novel biomarker of
Th2/eosinophil-driven airway inflammation and greater decline in pulmonary function, a
functional consequence of airway remodeling in patients with asthma.

In this study, the effects of biomarkers and clinical indices on greater annual decline in pulmonary function in asthmatic patients on ICS treatment were examined, with the specific aim of determining the association between serum periostin levels and pulmonary function decline. Polymorphisms of the *POSTN* gene, which encodes periostin, were also examined on the hypothesis that *POSTN* gene polymorphisms may affect serum periostin levels.

134 Methods

135 For full details see Online Repository

136 **Patients**

Patients with asthma were recruited from nine institutions belonging to the Kinki 137Hokuriku Airway disease Conference where asthma specialists manage patients. Asthma was 138diagnosed according to the American Thoracic Society criteria²⁶. From September 2009 to 139140December 2011, patients were enrolled if they had received ICS treatment for 4 years or more, undergone three or more pulmonary function tests when they were stable, and were free from 141exacerbations for at least one month. The first pulmonary function test was performed at least 142143one year after the commencement of ICS treatment and at 25 years of age or older. Patients 144who had smoked more than 10 pack-years, smoked in the past one year, or had other 145pulmonary diseases were excluded. This study was approved by the ethics committee of each participant institution and 146was registered in the UMIN Clinical Trials Registry (Registry ID UMIN000002414). Written 147148informed consent was obtained from all participants. 149150Measurements At enrollment, patients underwent a work-up that included answering a self-151completed questionnaire, spirometry, and blood tests. After enrollment, spirometry was 152repeated at least 6 months later for up to 12 months. 153154Self-completed questionnaire and clinical indices 155The self-completed questionnaire was composed of 4 major items, as presented in 156Table 1. The Asthma Control Test (ACT)TM was also scored. The treatment step at enrollment 157

158 was determined according to the Global Initiative for Asthma 2010 guideline 27 .

160 **Pulmonary function**

161 Spirometry was performed using an electrical spirometer, which was calibrated once a 162 week, at each institution. Spirometry data were obtained only when patients were stable. To 163 determine pulmonary function on daily medications, ICS and other controllers, including 164 long-acting β_2 agonists, leukotriene receptor antagonists, or slow-release theophylline, were 165 not withdrawn before spirometry.

166

167 Measurement of systemic biomarkers

Blood eosinophil and neutrophil counts, and serum levels of total immunoglobulin E
(IgE), specific IgE against common inhaled allergens, eosinophil cationic protein (ECP), high
sensitivity C-reactive protein (hsCRP), and periostin were determined.

Serum periostin levels were measured using an enzyme-linked immunosorbent assay at
Shino-test (Kanagawa, Japan), as described previously²⁸. Pooled serum periostin level data
from 66 healthy subjects [mean (SD), 60.7 (16.7) years old, 40 males]^{28,29} were used for
comparison with those of asthmatic patients.

175

176 Haplotype analysis, DNA extraction, and genotyping of the *POSTN* gene

A total of 47 single-nucleotide polymorphisms (SNPs) in the region of the *POSTN* gene and its upstream, total 39 kb, was captured in the HapMap Japanese data set. Haplotype analysis identified 4 major haplotypes and 2 minor haplotypes. Two minor haplotypes were grouped into the closest major haplotype, and 3 tag SNPs that determined the 4 haplotypes were identified (Figure 1).

Genomic DNA was isolated from blood cells using a QIAamp DNA Blood Mini Kit
(Qiagen, Tokyo, Japan). SNPs were genotyped using a Taqman genotyping assay according
to the manufacturer's instructions (Applied Biosystems, Tokyo, Japan) and analyzed using an
Applied Biosystems 7300 Real-Time PCR System (Applied Biosystems).

186

187 Statistical analysis

Statistical analyses were performed using JMP version 9.0 (SAS Institute Inc., Tokyo, 188 Japan). Annual changes in FEV₁ (Δ FEV₁) were estimated for each subject by fitting a least-189square regression line to all of his/her all available data points. Receiver operating 190 191characteristic (ROC) curve analysis was performed to determine a serum periostin cut-off 192value for asthmatic patients. The effects of serum biomarkers or other indices on ΔFEV_1 were estimated using a generalized linear mixed model with adjustment for sex, height, age at 193enrollment, and FEV₁ at the first measurement. The institutions were included as random 194effects in this model. On univariate analysis of ΔFEV_1 , the adjusted p value, i.e., q value, 195which was a measure of significance in terms of the false discovery rate, was obtained using 196 R and QVALUE software³⁰ to determine spurious significance in multiple testing. The effects 197on the dichotomous data for a decline in FEV₁ of $-30 \text{ mL} \cdot \text{yr}^{-1}$ or greater³¹ were similarly 198estimated using a generalized linear mixed model by IBM SPSS Advanced Statistics 19 199200(SPSS Inc., Tokyo, Japan). Multivariate analysis was performed using variables with p < 0.10201on univariate analysis, except for ICS daily maintenance dose because of its strong correlation with treatment step. On multivariate analysis, the periostin level was considered 202203as a dichotomous variable (high or low) instead of a continuous variable. Correlation coefficients between serum periostin levels and clinical indices were estimated by fitting 204least-square regression lines to data, in which institutions were included as random effects. 205206Unpaired *t*- and Chi-square tests were performed for comparisons of continuous and 207dichotomous variables, respectively. When data were not normally distributed, they were logtransformed. Data are presented as means (SD). P values ≤ 0.05 were considered significant. 208

209 **Results**

210 **Patients' characteristics**

211Initially, 233 patients were enrolled in this study, but 9 patients were excluded: 5 with a 212smoking history of more than 10 pack-years and 4 who did not have enough pulmonary function data available. The demographic data of the remaining 224 patients are presented in 213214Table 2. The mean age at enrollment was 62.3 (13.7) years. Overall, 130 (58%) had onset of 215asthma at 40 years or older. The average number of measurements of FEV₁, follow-up period, and ΔFEV_1 of 224 patients were 16.2 (13.9) times, 8.0 (4.5) years, and -7.8 (34.6) mL \cdot yr⁻¹, 216respectively. The distribution of ΔFEV_1 in this population is shown in Figure E1 in the Online 217218Repository. Within 2 years after diagnosis, 46% of patients started ICS treatment. At enrollment, 82% of patients took controllers such as long-acting β_2 agonists, leukotriene 219220receptor antagonists, or sustained release theophylline to achieve adequate asthma control. Based on a questionnaire, adherence to medication was satisfactory; 49% of the participants 221222never and 38% seldom forgot to take ICS or other medications. Based on ACT scores, 50% 223was totally controlled, and 38% scored from 20 to 24, indicating that they were well 224controlled at enrollment.

Serum periostin levels of asthmatic patients [92.8 (38.4) ng/mL] were significantly 225higher than those of healthy subjects [39.1 (24.5) ng/mL, p < 0.001]. The ROC curve analysis 226 was performed to discriminate patients with asthma who were thought to have refractory Th2 227inflammation despite long-term ICS treatment from healthy subjects. The highest specificity 228229among the 4 cut-off values tested was achieved at 95 ng/mL (0.985) in the comparison study 230of 224 asthmatic patients and 66 healthy subjects. Therefore a cut-off value of 95 ng/mL was 231used to define a high serum periostin group, although it had relatively lesser sensitivity (0.379) (see Figure E2 in the Online Repository). In asthmatic patients, 85 patients (38%) had 232high serum periostin levels (\geq 95 ng/mL). Of the 85 patients, 40 patients (47%) were on 233

treatment step 4, according to the treatment step classification²⁷, and 9 patients (11%) were
on treatment step 5.

236

Associations between serum periostin levels and greater annual decline in FEV₁ and a decline in FEV₁ of 30 mL·yr⁻¹ or greater

In an analysis of continuous values of ΔFEV_1 , greater decline in FEV₁ was associated 239240with higher serum periostin levels at enrollment, treatment step 5, lower ACT scores, incomplete adherence to medications, comorbid or a history of sinusitis, and comorbid 241diabetes mellitus (Table 3). When patients were stratified into two groups according to their 242243serum periostin levels, high serum periostin (\geq 95 ng/mL) was also associated with greater decline in FEV₁ (Table 3). Of these, high serum periostin was significant after controlling for 244multiple testing using the false discovery rate (q = 0.03, data not shown in Table 3).³⁰ 245Multivariate analysis revealed that greater decline of FEV₁ was solely associated with high 246serum periostin (\geq 95 ng/mL) (estimated effect -5.39, 95% confidence interval -10.0 to -0.77, 247248p = 0.02). Fifty-two patients (23%) showed a decline in FEV₁ of 30 mL·yr⁻¹ or greater [mean -24951.8 (18.4) mL \cdot yr⁻¹] and were considered rapid decliners³¹. When adjusted by confounders, 250higher serum periostin levels at enrollment, treatment step 5, a history of admission due to 251asthma exacerbation, higher ICS daily doses, comorbid or a history of sinusitis, and ex-252smoking were associated with a decline in FEV₁ of 30 mL·yr⁻¹ or greater. High serum 253periostin (\geq 95 ng/mL) was also associated with a decline in FEV₁ of 30 mL·yr⁻¹ or greater 254(Table 4). On multivariate analysis, high serum periostin (\geq 95 ng/mL), treatment step 5, and 255ex-smoking were independent risk factors for a decline in FEV_1 of 30 mL·yr⁻¹ or greater 256(Table 4). 257

258 Of the 224 patients, 19 patients were on treatment step 5, and 36 patients took high-259 dose ICS (1,000 μg or higher doses of ICS equivalent to fluticasone propionate daily). When 260 patients were stratified into the high periostin group, the average ΔFEV_1 of patients on

treatment step 5 (n = 9) was -41.0 (49.3) mL \cdot yr⁻¹, and 7 of them (78%) had excess decline;

262 the average ΔFEV_1 of patients on high-dose ICS (n=18) was -34.3 (39.4) mL · yr⁻¹, and 11 of

263 them (61%) had a decline in FEV_1 of 30 mL·yr⁻¹ or greater.

264

265 Serum periostin levels and clinical indices

266In 224 patients, serum periostin levels were weakly associated with blood eosinophil counts (Figure 2), serum IgE (Figure 2) and ECP levels (r = 0.25, p = 0.0005), 267ICS-untreated period, i.e. period between onset of asthma and the initiation of ICS therapy (r 268= 0.16, p = 0.01), daily maintenance doses of ICS at enrollment (r = 0.13, p = 0.05), and a 269history of admission due to asthma exacerbation (r = 0.15, p = 0.03). Serum periostin levels 270271were significantly higher in patients on high-dose ICS ($\geq 1,000 \ \mu g$ daily) than in the remaining patients (110.3 ng/mL vs. 89.5 ng/mL, p = 0.003). Lastly, serum periostin levels 272were higher in patients with sinusitis than in those without sinusitis (103.9 ng/mL vs. 88.3 273274ng/mL, p = 0.007). Serum periostin levels did not show any seasonal variability or 275association with age at onset of asthma (data not shown).

276

277 **POSTN** gene polymorphisms

Associations between polymorphisms of the *POSTN* gene, which encodes periostin, and both serum periostin levels and pulmonary function decline were then investigated. In one patient, DNA quality was insufficient for genotyping; thus, 3 tag SNPs of the *POSTN* gene were analyzed in 223 patients. All genotyped data were in Hardy-Weinberg equilibrium. The frequencies of the 3 tag SNPs and analysis results using dominant and recessive models for serum periostin levels and a decline in FEV_1 of 30 mL·yr⁻¹ or greater are presented in Table 5.

285 Serum periostin levels were higher in patients with the GG genotype of rs3829365 than

in those with the GC/CC genotype (GG 98.7 ng/mL vs. GC/CC 86.1 ng/mL, p = 0.003). 286287rs1028728 was not associated with serum periostin levels or with the frequency of rapid decliners, but patients with the TT genotype of rs1028728, 4 patients only, showed no 288significant decline compared with the AA/AT genotype (AA/AT -8.6 mL \cdot yr⁻¹ vs. TT 29.3 289mL·yr⁻¹, p = 0.03). Rapid decliners were more frequently observed in patients with the minor 290A allele of rs9603226 than in the GG genotype (GG 16% vs. AG/AA 30%, p = 0.02). A 291marked difference in the frequency of rapid decliners was observed when patients were 292stratified into the high periostin group [GG of rs9630226 (n = 37) 19% vs. AG/AA (n = 47) 29345%, p = 0.01]. 294

295 **Discussion**

To the best of our knowledge, this is the first study to identify a relationship between 296greater decline in FEV₁ and higher serum periostin levels, particularly if they were 95 ng/mL 297or more, in asthmatic patients on ICS treatment. It was also shown that high serum periostin, 298together with treatment step 5 and light ex-smoking, was an independent risk factor for a 299decline in FEV₁ of 30 mL·yr⁻¹ or greater. In addition, polymorphisms of the *POSTN* gene, 300 which encodes periostin, were associated with serum periostin levels and a decline in FEV_1 301 of 30 mL·yr⁻¹ or greater in asthmatic patients. These findings suggest that serum periostin 302 may be a useful biomarker for the development of airflow limitation in asthmatic patients on 303 ICS. 304

In this study, despite long-term treatment with ICS with or without other controllers, 30523% of asthmatic patients were rapid decliners who showed a decline in FEV₁ of 30 mL·yr⁻¹ 306 or greater, for which treatment step 5 was an independent risk factor. Adherence to ICS 307treatment and the frequency of early intervention with ICS did not differ between rapid 308 309decliners and non-decliners, although long-term adherence to ICS was undetermined in the 310 present study. In previous studies of patients who were not treated with ICS, severe exacerbation of asthma contributed to greater annual decline of pulmonary function^{6,7}, but the 311exacerbation-related greater annual decline disappeared in an early intervention group with 312ICS treatment in the START study⁶, which might be interpreted to mean that asthmatic 313patients on ICS treatment have little risk of accelerated FEV₁ decline. However, since the 314315START study originally recruited mild persistent asthmatic patients, its results cannot simply be applied to severe asthmatic patients. As observed in the present study, there would be a 316 subset of asthmatic patients still at risk of greater annual decline of pulmonary function 317despite intensive treatment for asthma. 318

Persistent eosinophilic airway inflammation is a key process in irreversible airway
 obstruction¹⁰. Indeed, exhaled nitric oxide of 20 ppb or higher is a risk factor for accelerated

 FEV_1 decline in patients with difficult-to-treat asthma¹⁸. Studies on novel therapies for 321refractory eosinophilic asthma, i.e., anti-IL-5 therapy³² and anti-IL-13 therapy³³, revealed that 322these treatments may reverse airway remodeling when patients are adequately targeted, 323 suggesting the necessity of establishing "companion diagnostics" for this population. 324According to the most recent study, serum periostin is the single best biomarker reflecting 325sputum and tissue eosinophilia among several biomarkers, including blood eosinophils and 326 exhaled nitric oxide²⁵. In the current study, the serum periostin level, which was associated 327 with the blood eosinophil count, was the sole biomarker that reflected greater decline in FEV₁. 328 Periostin is secreted by airway epithelial cells^{23, 24} and lung fibroblasts²¹ in response to IL-4 329 and IL-13 and is thought to be secreted into the capillary vessels. Downstream of IL-13, 330 which plays a pivotal role in subepithelial airway fibrosis³⁴, airway remodeling³⁵, and steroid 331insensitivity³⁶, periostin mediates collagen synthesis²⁴ and fibrillogenesis^{24, 37} by binding to 332collagen³⁷ and activates TGF- β^{24} . In the asthmatic airway, periostin is deposited in the 333 subepithelial layer, colocalizing with collagens I, III, and V, fibronectin, tenascin-C, and 334periostin itself²¹, which indicates involvement of periostin in airway remodeling in asthma. 335Collectively, periostin may be a key molecule that links eosinophilic inflammation and 336 remodeling via IL-13 in asthmatic airways. Further roles of periostin in allergic inflammation 337and remodeling in the airways remain undetermined because studies using periostin-deficient 338 mice with acute allergen exposure have yielded conflicting findings³⁸⁻⁴⁰; one study showed 339 that periostin facilitates eosinophil infiltration into the lung³⁸, whereas two other studies^{39,40} 340 suggested protective roles of periostin. Meanwhile, a recent study of a chronic mouse model 341of atopic dermatitis demonstrated periostin's role in the chronicity of Th2 inflammation²⁹. 342In the present study, patients on high-dose ICS showed higher serum periostin levels 343 than the other patients. Although a longitudinal study is needed to determine responses of 344serum periostin levels to ICS treatment, we do not think that the high serum periostin levels 345in patients on high-dose ICS were induced by ICS treatment, because periostin expression in 346

the airway epithelium was decreased with ICS treatment²³. Rather, the elevation of serum 347periostin in this population may reflect IL-13-mediated inflammation that is partly refractory 348to ICS, as was reported in a recent study by Jia and colleagues²⁵. They showed that, in 349patients with severe asthma who were treated with high doses ICS (> 1000 µg daily), 350elevation of serum periostin levels was associated with persistent airway tissue eosinophilia, 351concluding that serum periostin is a systemic biomarker of airway eosinophilia refractory to 352high-dose ICS²⁵. Providing further support, among patients with moderate to severe asthma 353who are inadequately controlled despite ICS treatment, patients with high serum periostin 354levels are likely to benefit from anti-IL-13 antibody, lebrikizumab, treatment³³. The novelty 355of the present finding is that high serum periostin is an independent risk factor for greater 356decline in FEV₁, providing the first evidence for the potential association between persistent 357Th2- or IL-13-driven inflammation refractory to ICS treatment and greater decline in FEV₁, a 358functional consequence of airway remodeling. 359

Needless to say, current smokers with asthma have more accelerated FEV_1 decline⁴ than 360 those not smoking, and current smoking impairs the therapeutic response to ICS or oral 361corticosteroids⁴¹. Meanwhile, smoking cessation improves their FEV₁ levels⁴², and ex-362smokers with asthma with 10 pack-years or more show an intermediate response to short-363term oral corticosteroid treatment, between current smokers and never-smokers⁴¹. In the 364present study, rather unexpectedly, ex-smoking with 10 pack-years or less was still an 365independent risk factor for a decline in FEV_1 of 30 mL·yr⁻¹ or greater. It should be recognized 366 that even light ex-smoking increases the risk of airway remodeling in asthmatic patients on 367368 ICS, and its underlying mechanisms should be clarified.

Chronic sinusitis is a well-known comorbidity with severe asthma^{43, 44}. In the present study, rapid decliners were more frequently observed in asthmatic patients with sinusitis than those without sinusitis on univariate analysis, and their periostin levels were higher than in patients without sinusitis. In the present study, polypoid lesions in the sinuses were not evaluated by otolaryngologists at enrollment. However, considering that periostin is upregulated in nasal polyp tissue in patients with chronic rhinosinusitis⁴⁵, asthmatic patients with sinusitis may have had severe upper and lower airway inflammation with persistent increases in periostin expression, which may have resulted in a decline in FEV₁ of 30 mL·yr⁻¹ or greater. Periostin is a potential molecule that unifies sinusitis and severe asthma.

378 Periostin is encoded on the *POSTN* gene, which is located on chromosome 13q13.3. 379 rs3829365, which is located at the 5'UTR region that may contain sequences to regulate translation efficiency or mRNA stability, was associated with serum periostin levels. This 380 finding suggests that, besides IL-13, a master regulator of periostin, genetic background 381382partly determines periostin levels, although a replication study would be necessary to confirm this. The minor A allele of rs9603226, located 66 bp upstream of exon 21 in the C-terminal 383 region, was associated with a decline in FEV₁ of 30 mL·yr⁻¹ or greater. In periostin, FAS I 384domains are thought to be primary binding sites to fibronectin, tenascin-C, and collagen V^{21} , 385whereas the C-terminal region in its intact form may down-regulate the binding activity of 386 periostin to these extracellular matrix proteins²¹. We therefore speculate that the minor A 387allele of rs9603226 might modify the binding activity at the C-terminal region and facilitate 388 airway remodeling, particularly if the airway is in periostin enriched milieu. Further studies 389are needed to clarify if these SNPs are functional variants. 390

The age of patients in this study appears to be older than in other Euro-American studies^{6,7,14,18,20,23,25}. One reason for the age distribution would be the entry criteria of this study. Another reason would be explained by population aging including population with asthma in Japan. According to a patient survey by the Japanese Ministry of Health, Labour and Welfare in 2008, patients aged 70 to 74 years were the most frequent age group of adult patients with asthma⁴⁶, which is still older than the average age of patients in this study. There are several limitations to the present study. First, since this study was

399	the follow-up period. Controllers such as long-acting β_2 agonists were not withdrawn at
400	pulmonary function testing to evaluate function on daily medications, which may have
401	resulted in the small average ΔFEV_1 , -7.8 mL · yr ⁻¹ . Meanwhile, averages of 16.2
402	measurements of FEV_1 and 8.0 years of follow-up were satisfactory for a longitudinal
403	analysis of pulmonary function ⁴⁷ , and ΔFEV_1 was normally distributed. Secondly, serum
404	biomarkers were measured only once at enrollment, but the significant associations between
405	<i>POSTN</i> gene polymorphisms and serum periostin levels or a decline in FEV ₁ of 30 mL·yr ⁻¹
406	or greater may circumvent the inherent insufficiency of single measurement of serum
407	periostin. Thirdly, most of the clinical information, including smoking history and chronic
408	sinusitis, was based on a self-completed questionnaire, which might be biased by recall
409	memory. Despite these limitations, the current findings may provide directions for future
410	research.
411	In conclusion, serum periostin appears to be a useful biomarker that reflects the

development of airflow limitation in patients on prolonged treatment with ICS. *POSTN* gene
polymorphisms may also be helpful for identification of rapid decliners.

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553		
554		

555 Table 1. Contents of the self-completed questionnaire

Asthma-related history

- family history of asthma
- •age of asthma onset
- •history of pediatric asthma
- ·history of admission due to asthma worsening or exacerbation
- •aspirin hypersensitivity
- •asthma deterioration at the working place

Comorbidity or a history of the following diseases			
 allergic dermatitis 	•cardiovascular diseases including ischemic heart disease		
 allergic rhinitis 	 gastrointestinal diseases including GERD 		
 seasonal rhinitis 	•collagen vascular diseases including rheumatoid arthritis		
 allergic conjunctivitis 	• diabetes mellitus		
 chronic sinusitis 	•pulmonary diseases other than asthma		
	•other diseases including malignancy		

Lifestyle and environment

 smoking history 	•a highway near the home
•pet breeding	•age at menopause
• type of occupation	

Adherence to medication, sputum production, and exacerbations

• How often do you forget to take inhaled corticosteroids or other medications?

0: never, 1: seldom, 2: sometimes, 3: often, 4: always

•How often do you produce sputum?

0: never, 1: once in a few days, 2: every morning, 3: every morning and daytime

• How often did you receive systemic steroids due to asthma exacerbations during the recent 6 months?

0: never, 1: once, 2: twice or more

556 GERD: gastro-esophageal reflux disease

558 Table 2. Patients' characteristics

Sex (males/ females), n	53 / 171
Age at enrollment, years	62.3 (13.7)
Age at asthma onset, years	42.0 (19.0)
Body mass index (kg/m ²)	23.1 (3.5)
Smoking history (never), n	181
Atopic predisposition [*] , %	70
Pediatric asthma (none/ recurrent/ persistent), %	81 / 8 / 11
Disease duration, years	20.2 (14.5)
ICS-untreated period, years	9.2 (13.1)
ICS daily maintenance dose [†] , µg	525 (318)
Number of other controller medications, n	1.4 (1.2)
Treatment step $(2/3/4/5)^{\ddagger}$, %	16 / 27 / 49 / 8
Sputum production $(0/1/2/3)^{\$}$, %	54 / 20 / 8 / 18
Asthma Control Test, points	22.6 (3.5)
History of admission due to asthma, n (%)	78 (35)
Allergic rhinitis, n (%)	129 (58)
Chronic sinusitis, n (%)	65 (29)
Blood neutrophils, %	60.1 (10.0)
eosinophils, %	5.2 (4.9)
Serum IgE, IU/mL	180 (0 - 16000)
periostin, ng/mL	92.8 (38.4)
high sensitivity C-reactive protein, mg/L	1341 (3147)
eosinophil cationic protein, µg/L	15.1 (29.3)
FEV_1 at the first measurement, L [¶]	2.11 (0.69)
% predicted FEV ₁ at the first measurement, %	91.9 (19.2)
FEV_1 / FVC at the first measurement, %	73.9 (9.8)
FEV_1 at enrollment, L	2.04 (0.73)
% predicted FEV ₁ at enrollment, %	97.4 (22.2)
FEV ₁ / FVC at enrollment, %	72.2 (10.0)
Reversibility at enrollment, % [#]	3.8 (6.0)
Data at annollment are presented unless otherwise stated. Data are	avaraged as means (SD) avaant for ma

559 Data at enrollment are presented unless otherwise stated. Data are expressed as means (SD) except for median 560 (range) for serum IgE. *Considered atopic when one or more specific IgE antibodies against cat or dog dander, 561 weed, grass, or Japanese cedar pollens, moulds, or house dust mite were positive. [†]Equivalent to fluticasone 562 propionate. [‡]according to the Global Initiative for Asthma 2010 guideline²⁷. [§] 0 = never, the details are shown in 563 Table 1. [¶]The first pulmonary function test was performed at least one year after the commencement of ICS 564 treatment and at 25 years of age or older. [#]n = 206, airway reversibility to 200 µg of inhaled salbutamol.

	Estimates	95% C.I.	p valu
Smoking history, ex vs. never	-8.48	-20.2, 3.27	0.16
Atopic predisposition	-1.10	-6.29, 4.09	0.68
Disease duration, years	-4.79	-18.4, 8.86	0.56
ICS-untreated period, years	0.10	-0.24, 0.45	0.65
ICS daily maintenance dose, µg	-0.01	-0.03, 0.001	0.07
Number of other controller medications, n	-0.36	-4.21, 3.49	0.86
Adherence to medication, incomplete vs. complete*	-4.56	-9.08, -0.04	0.05
Treatment step, 5 vs. 2-4 [†]	-7.77	-15.7, 0.13	0.05
Sputum production, never vs. others [‡]	0.99	-3.53, 5.51	0.67
Asthma Control Test, points	1.53	0.29, 2.77	0.02
History of admission due to asthma	-4.49	-9.45, 0.46	0.08
Aspirin hypersensitivity	-6.52	-20.0, 6.98	0.34
Asthma deterioration at the working place	-12.2	-54.4, 30.0	0.57
Allergic rhinitis	-1.21	-5.88, 3.45	0.61
Allergic dermatitis	4.51	-1.51, 10.5	0.14
Chronic sinusitis	-10.1	-19.8, -0.27	0.04
Ischemic heart disease	3.41	-16.6, 23.4	0.74
Hypertension	-3.79	-9.12, 1.53	0.16
Dyslipidemia	-3.67	-9.42, -2.06	0.21
Diabetes mellitus	-8.03	-15.4, -0.67	0.03
Gastro-esophageal reflux disease	-3.85	-9.89, 2.19	0.21
Malignancy	-3.44	-26.0, 19.1	0.76
Post-menopause	5.05	-14.2, 24.3	0.60
Pet breeding	-0.28	-12.6, 12.0	0.96
Log blood neutrophils, %	-7.40	-69.1, 54.3	0.81
eosinophils, %	-0.67	-1.60, 0.27	0.16
Log serum IgE, IU/mL	-2.85	-9.74, 4.04	0.42
periostin, ng/mL	-29.1	-56.2, -1.97	0.04
high sensitivity C-reactive protein, mg/L	-1.88	-9.85, 6.10	0.64
eosinophil cationic protein, µg/L	-4.47	-15.7, 6.81	0.44
Periostin group, high vs. low [§]	-6.96	-11.4, -2.51	0.002

Estimated effects were adjusted by sex, height, age at enrollment, and FEV₁ at the first measurement. ^{*} "Complete", when patients answered

 $566 \\ 567 \\ 568 \\ 569$ that they never forgot to take ICS or other medications; "incomplete", the remaining cases. [†]according to the Global Initiative for Asthma 2010 guideline²⁷. [‡]The details are shown in Table1. [§] Patients were stratified into two groups according to their serum periostin levels: high \geq 95 ng/mL, low < 95 ng/mL. ICS: inhaled corticosteroids, C.I.: confidence interval

	Univariate analysis			Multivariate analysis		
	Estimates	95% C.I.	p value	Estimates	95% C.I.	p value
Treatment step, 5 vs. $2-4^*$	1.63	0.51, 2.60	0.004	1.24	0.078, 2.30	0.04
History of admission due to asthma	1.09	0.37, 1.90	0.003	0.70	-0.11, 1.50	0.09
ICS daily maintenance dose, µg	0.001	0.00, 0.002	0.01	-		
Chronic sinusitis	0.82	0.11, 1.53	0.03	0.61	-0.15, 1.37	0.12
Smoking history, ex vs. never	0.87	-0.002, 1.74	0.05	0.98	0.030, 1.93	0.04
Log serum periostin, ng/mL	2.96	0.78, 5.13	0.008	-		
Periostin group, high vs. low [†]	1.03	0.33, 1.72	0.004	0.87	0.11, 1.63	0.03

Table 4. Estimated effects of clinical indices and serum periostin on a decline in FEV₁ of 30 mL·yr⁻¹ or greater

572 Estimated effects were adjusted by sex, height, age at enrollment, and FEV_1 at the first measurement.

573 *according to the Global Initiative for Asthma 2010 guideline²⁷.

[†]Patients were stratified into two groups according to their serum periostin levels: high \geq 95 ng/mL, low <

575 95 ng/mL. ICS: inhaled corticosteroids, C.I.: confidence interval

576 ICS daily maintenance dose was excluded from multivariate analysis because of its strong correlation with

treatment step.

					Serum periostin levels		Frequency of rapid decliners	
					p value		p value	
Tag SNP	Genotype	n (%)	Allelic	n (%)	$Dominant^{\dagger}$	Recessive*	$Dominant^{\dagger}$	Recessive*
rs1028728	AA	164 (74)	А	383 (86)				
	AT	55 (25)	Т	63 (14)	0.40	0.46	0.17	0.14
	TT	4 (2)						
rs3829365	GG	113 (51)	G	316 (71)				
	GC	90 (40)	С	130 (29)	0.003	0.70	0.40	0.33
	CC	20 (9)						
rs9603226	GG	107 (48)	G	311 (70)				
	AG	97 (44)	А	135 (30)	0.80	0.33	0.01	0.81
	AA	19 (9)						

Table 5. Frequencies of 3 tag SNPs and analysis results using dominant and recessive models
for serum periostin levels and frequency of rapid decliners*

581 ^{*} defined as patients who showed a decline in FEV₁ of 30 mL·yr⁻¹ or greater

582 [†]Assuming that heterozygotes have the same increased risk as minor homozygous genotypes.

583 ^{*}Assuming that heterozygotes have no increased risk.

585 Figure legends

- 586 Figure 1. Three tag SNPs that determine 4 major haplotypes of the *POSTN* gene and
- 587 haplotype frequencies in the Japanese population are presented.
- ⁵⁸⁸ *at intron 66 bp upstream of exon 21
- 589
- 590 Figure 2. Relationships between serum periostin levels and blood eosinophil counts (left) or
- 591 serum IgE levels (right).
- 592 Presented in logarithmic scales on both the X- and Y-axes.
- 593

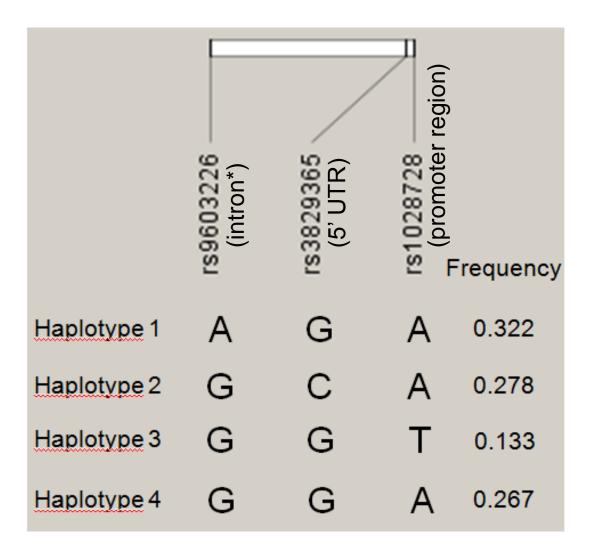


Figure 1.

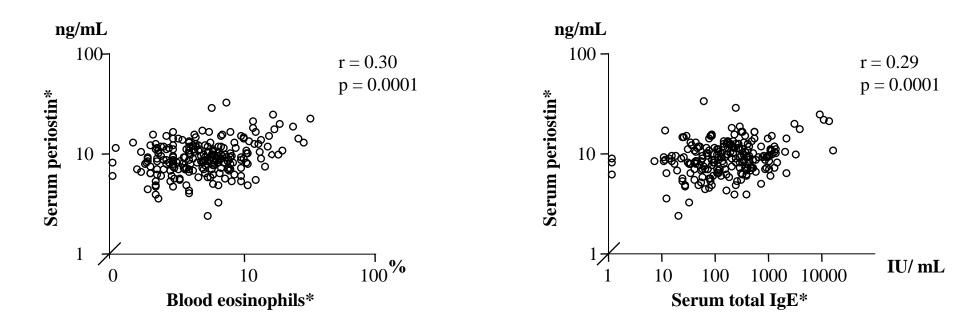


Figure 2.

1 Online Repository

Increased periostin associates with greater airflow limitation in patients receiving inhaled corticosteroids

4

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52 Methods

53 Patients

Patients with asthma were recruited from nine institutions belonging to the Kinki 54Hokuriku Airway disease Conference where asthma specialists manage patients, 5556including six university hospitals, two satellite general hospitals, and one satellite clinic. Asthma was diagnosed according to the American Thoracic Society criteria^{E1} on the 5758basis of a history of recurrent episodes of wheezing and chest tightness with or without cough and documented airway reversibility to a bronchodilator or hyper-responsiveness 5960 to inhaled methacholine. From September 2009 to December 2011, patients were 61 enrolled if they had received ICS treatment for 4 years or more, undergone three or more pulmonary function tests when they were stable, and were free from exacerbations 62 63 for at least one month. The first pulmonary function test was performed at least one year after the commencement of ICS treatment and at 25 years of age or older. Patients who 64 65 had smoked more than 10 pack-years, smoked in the past one year, or had other 66 pulmonary diseases were excluded.

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Self-completed questionnaire and clinical indices

69 The self-completed questionnaire was composed of 4 major items, as presented70 in Table 1.

Adherence to ICS or other medications, frequency of sputum production, and requirement for systemic corticosteroids during the last 6 months were graded as shown in Table 1. The Asthma Control Test (ACT)TM was also scored. Duration of ICS treatment and details on medication at enrollment were recorded from medical charts by patients' physicians. The treatment step at enrollment was determined according to the 76 Global Initiative for Asthma 2010 guideline^{E2}.

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78 Measurement of systemic biomarkers

Blood eosinophil and neutrophil counts, and serum levels of total 79 immunoglobulin E (IgE) (ImmunoCAP[®] total IgE, Phadia K.K., Tokyo, Japan), specific 80 IgE against common inhaled allergens (ImmunoCAP[®] specific IgE), eosinophil cationic 81 protein (ECP) (ImmunoCAP[®] ECP), high sensitivity C-reactive protein (hsCRP) 82 (CardioPhase[®] hsCRP. Siemens Healthcare Diagnostics K.K., Tokyo, Japan), and 83 periostin were determined. 84 Serum periostin levels were measured using an enzyme-linked immunosorbent 85 assay at Shino-test (Kanagawa, Japan), as described previously^{E3}. Briefly, two rat 86 anti-human periostin monoclonal antibodies (SS18A and SS17B) were used. SS18A and 87 SS17B are antibodies against the first and fourth FAS I domains, respectively. Intra- and 88 89 inter-assay coefficients of variation ranged from 1.31% to 2.54% and 1.49% to 2.01%, 90 respectively.

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92 Haplotype analysis, DNA extraction, and genotyping of the POSTN gene

A total of 47 single-nucleotide polymorphisms (SNPs) in the region of the *POSTN* gene and its upstream, total 39 kb, was captured in the HapMap Japanese data set with minor allele frequencies > 0.10. Pairwise tagging was performed at $r^2 > 0.8$ using a tagger in Haploview 4.2 software. Haplotype analysis identified 4 major haplotypes and 2 minor haplotypes. Two minor haplotypes were grouped into the closest major haplotype, and 3 tag SNPs that determined the 4 haplotypes were identified (Figure 1). These 3 tag SNPs were located at promoter region (rs1028728), 5'UTR region (rs3829365), and at intron 66 bp upstream of exon 21 (rs9603226). The
frequencies of the minor alleles in the Japanese population were 0.136 (rs1028728),
0.278 (rs3829365), and 0.330 (rs9603226).

Genomic DNA was isolated from blood cells using a QIAamp DNA Blood Mini
Kit (Qiagen, Tokyo, Japan). SNPs were genotyped using a Taqman genotyping assay
according to the manufacturer's instructions (Applied Biosystems, Tokyo, Japan) and
analyzed using an Applied Biosystems 7300 Real-Time PCR System (Applied
Biosystems).

109 **References**

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124 Figure legends

- 126 Figure E1. The distribution of ΔFEV_1 in the study population
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- 128 Figure E2. ROC curve analysis of serum periostin levels comparing asthmatic patients
- and healthy subjects, in which the cutoffs of 95 ng/mL, 80 ng/mL, 92 ng/mL, and 100
- 130 ng/mL are presented with arrows.

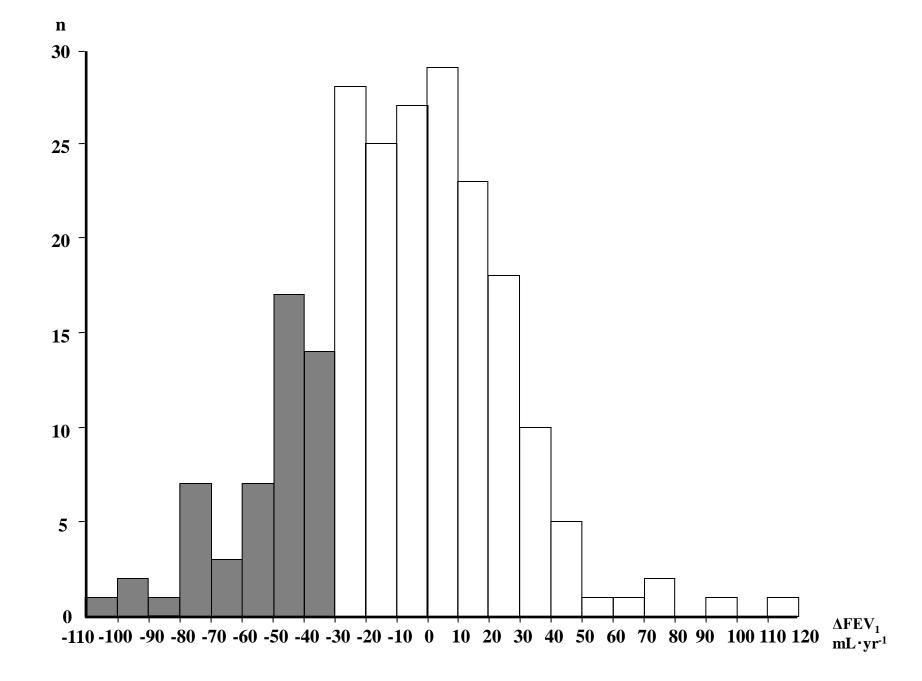


Figure E1.

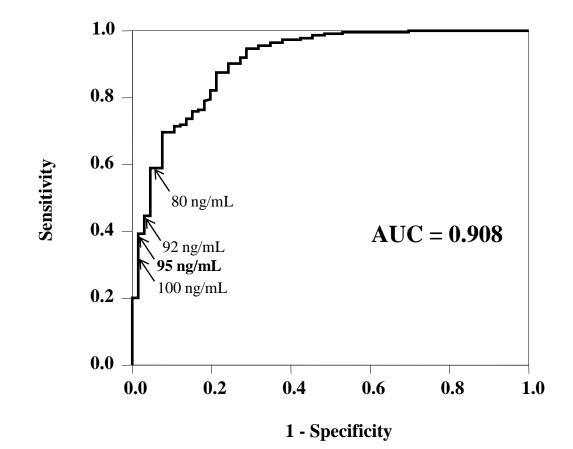


Figure E2.