

1 Higher FVIII:C measured by chromogenic substrate assay than by one-stage assay is  
2 associated with silent hemophilic arthropathy

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4 Mika Ogawa<sup>1\*</sup>, Nobuaki Suzuki<sup>2</sup>, Nobunori Takahashi<sup>3</sup>, Shogo Tamura<sup>4</sup>, Atsuo Suzuki<sup>5</sup>,  
5 Sachiko Suzuki<sup>4†</sup>, Yuua Hattori<sup>4‡</sup>, Misaki Kakihara<sup>4</sup>, Takeshi Kanematsu<sup>6</sup>, Toshihisa  
6 Kojima<sup>3</sup>, Akira Katsumi<sup>7</sup>, Fumihiko Hayakawa<sup>4</sup>, Tetsuhito Kojima<sup>4</sup>, Naoki Ishiguro<sup>3</sup>,  
7 Hitoshi Kiyoi<sup>1</sup>, Tadashi Matsushita<sup>2,6</sup>

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9 <sup>1</sup> Department of Hematology and Oncology, Nagoya University Graduate School of  
10 Medicine, Nagoya, Japan

11 <sup>2</sup> Department of Transfusion Medicine, Nagoya University Hospital, Nagoya, Japan

12 <sup>3</sup> Department of Orthopaedics/Rheumatology Nagoya University Graduate School of  
13 Medicine, Nagoya, Japan

14 <sup>4</sup> Department of Pathophysiological Laboratory Sciences, Nagoya University Graduate  
15 School of Medicine, Nagoya, Japan

16 <sup>5</sup> Department of Medical Technique, Nagoya University Hospital, Nagoya, Japan

17 <sup>6</sup> Department of Clinical Laboratory, Nagoya University Hospital, Nagoya, Japan

18 <sup>7</sup> Department of Hematology, National Center for Geriatrics and Gerontology, Aichi  
19 Prefecture, Japan

20 \*†‡Current affiliation:

21 \*Department of Hematology and Oncology, Japanese Red Cross Nagoya Daini Hospital,

22 Nagoya, Japan

23 †Department of Nursing & Health, Aichi Prefectural University, Aichi Prefecture, Japan

24 ‡Department of Medical Technique, Nagoya University Hospital, Nagoya, Japan

25

26 **Correspondence to:** Dr. Nobuaki Suzuki

27 Department of Transfusion Medicine

28 Nagoya University Hospital

29 65 Tsurumai-cho, Showa-ku, Nagoya, Aichi 466-0065, Japan

30 Tel: +81-52-744-2652; Fax: +81-52-744-2610

31 E-mail: [suzukin@med.nagoya-u.ac.jp](mailto:suzukin@med.nagoya-u.ac.jp)

32 **Keywords:** hemophilia A, hemophilic arthropathy, factor VIII measurement, genetic

33 mutation

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## 35 **1. Introduction**

36 The Joint Outcome Study demonstrated that prophylactic therapy reduced joint bleeding

37 and improved joint status. However, magnetic resonance imaging (MRI) revealed that the

38 joints of patients who had no clinically evident joint bleeding (CEJB) displayed some

39 abnormalities indicative of hemophilic arthropathy [1]. One explanation might be the

40 presence of subclinical joint bleeding that passed unrecognized by patients. However,

41 although many reports have implied their existence [1, 2], such bleeds have remained

42 unproven. If subclinical bleeds are really present, **annualized joint bleeding rate (AJBR)**

43 would be of limited use for monitoring hemophilic arthropathy. Factors more closely  
44 related to development of arthropathy should thus be elucidated.

45 Here, we performed a fact-finding survey of 15 adult patients with hemophilia A who  
46 had never experienced CEJB throughout their lives. The aim was to confirm whether  
47 patients who had never experienced CEJB did not develop hemophilic arthropathy, and  
48 also whether patients can develop hemophilic arthropathy without CEJB, to identify the  
49 characteristics of such patients.

50

## 51 **2. Methods**

### 52 **2.1. Study participants and investigation items**

53 This study included adult hemophilia A patients (age >20 years) who had never  
54 experienced CEJB. To investigate the history of CEJB, patients were specifically  
55 interviewed and their medical records were reviewed as far back as possible. Exclusion  
56 criteria comprised the presence or history of FVIII inhibitor. Investigation items  
57 comprised: age, body mass index (BMI), FVIII:C, FVIII antigen (FVIII:Ag), causative  
58 mutation, and Pettersson score (PS). All patients provided written, informed consent to  
59 participate in this study, which then proceeded according to the Declaration of Helsinki  
60 under the approval of the Nagoya University Hospital Ethics Committee (approval

61 number: 2015-0391).

62

## 63 **2.2. PS**

64 Hemophilic arthropathy was assessed in bilateral knees, ankles and elbows using  
65 Pettersson radiological scores [3]. Having a joint replaced with an artificial prosthesis  
66 was considered a full score (13 points). Only orthopedists estimated joint scores. The PS  
67 employed was the most recent one available. **Any positive score (i.e.,  $\geq 1$ ) for the joint  
68 was judged as representing a positive PS in this study. As a positive PS represents  
69 radiologic changes, such a score indicates the development of arthropathy.**

70

## 71 **2.3. FVIII activity (FVIII:C) and FVIII antigen (FVIII:Ag)**

72 We measured FVIII activity (FVIII:C) using both chromogenic substrate assay (CSA) and  
73 one-stage assay (OSA). CSA measures FVIII activity using Biophen FVIII:C (Hyphen  
74 BioMed, Neuville-sur-Oise, France). The OSA is based on the activated partial  
75 thromboplastin time (APTT) of FVIII immune-depleted plasma using an APTT reagent  
76 provided with an analyzer. The OSA proceeded using Thrombocheck APTT with ellagic  
77 acid activator, Thrombocheck FVIII and a CS-5100 automated coagulation analyzer (all  
78 from Sysmex, Kobe, Japan) [4].

79 Factor VIII antigen (FVIII:Ag) was quantified using the VisuLize™ FVIII Antigen Kit  
80 (Affinity Biologicals, Ancaster, Canada), an enzyme-linked immunosorbent assay, in  
81 accordance with the instructions from the manufacturer.

82

83

### 84 **3. Results**

#### 85 **3.1. Background and clinical stratification of patients**

86 We enrolled 15 male adult patients with hemophilia A (mean age  $\pm$  standard deviation  
87 (SD),  $47.3 \pm 17.1$  years; median age, 37 years). Mean and median BMI were  $23.3 \pm 3.9$   
88  $\text{kg/m}^2$  and  $21 \text{ kg/m}^2$ , respectively (Supplemental table 1). Disease severity was classified  
89 as severe in 1 patient, differed according to determination by CSA or OSA in 3 patients  
90 (moderate by CSA and mild by OSA in 1 patient, mild by CSA and moderate by OSA in  
91 2 patients), moderate in 4 patients, and mild in 7 patients.

92

#### 93 **3.2. Patients with hemophilic arthropathy displayed higher FVIII:C measured by** 94 **CSA than that measured by OSA**

95 Relationships between baseline FVIII:C and PS in the 15 patients are shown in Figure 1A  
96 and 1B. Five of these patients showed positive PS. No patients with baseline FVIII:C

97 >6.2% as measured with CSA showed positive PS, nor did any patients with baseline  
98 FVIII:C >3% as measured with OSA. All patients who showed a positive PS displayed  
99 higher FVIII:C ratio (CSA/OSA) (Fig. 1C). The 15 patients were divided into two groups:  
100 one comprising patients with positive PS; and the other comprising patients with negative  
101 PS. The results were analyzed by non-parametric test (Table 1). Mean and median  
102 FVIII:C ratio (CSA/OSA) were significantly higher for those patients who were PS-  
103 positive than for patients who were PS-negative. In other words, a discrepancy existed in  
104 baseline FVIII:C between CSA and OSA, with higher baseline FVIII:C as measured from  
105 CSA.

106

### 107 **3.3. Causative mutation in PS-positive patients**

108 The causative mutations for hemophilia A were identified in all patients, and the five  
109 patients who were PS-positive displayed missense mutations, as p. Asp182Tyr, p.  
110 Thr314Ile, p. Arg1800His and p. Arg2169His (Supplemental table 2). These mutations  
111 were not located near thrombin cleavage or factor IXa-binding sites that have been  
112 reported to be associated with discrepancies between the two methods with higher  
113 FVIII:C as measured by CSA, not only from the perspective of gene sequences, but also  
114 in terms of spatial location (Supplemental fig. 1).

115

#### 116 **4. Discussion**

117 PS is an additive radiographic scoring system for the severity of hemophilic  
118 arthropathy, and has been adopted by the Orthopedic Advisory Committee of the World  
119 Federation of Hemophilia. However, the score has been reported to vary according to  
120 the evaluator [5]. To reduce variation due to the evaluator, the same expert orthopedist  
121 (X-rays of joints are evaluated by orthopedists in Japan, not by radiologists) performed  
122 all evaluations in this study. As this scoring system is based on radiologic changes, early-  
123 phase hemophilic arthropathy cannot be detected. These issues represent limitations in  
124 the present study. Five of the 15 patients who were free of CEJB showed a positive PS in  
125 the study, providing evidence for the development of hemophilic arthropathy. This result  
126 indicated that lack of CEJB does not represent a necessary condition for a lack of  
127 hemophilic arthropathy. Interestingly, these patients displayed discrepancies in FVIII:C  
128 as measured by CSA and OSA, with higher FVIII:C by CSA than by OSA. The  
129 discrepancy in FVIII:C between CSA and OSA has been discussed for years [6, 7].  
130 Trossaert et al. reported that FVIII:C as measured by CSA better reflected the clinical  
131 bleeding tendency compared with the FVIII:C by OSA [8]. Postgieter et al. also reported  
132 that patients who had higher FVIII:C by CSA than by OSA displayed few or no bleeding

133 symptoms [9]. Considering these reports, these patients might escape CEJB, but not  
134 subclinical bleeding or subsequent hemophilic arthropathy. As the mechanism underlying  
135 such discrepancy, they hypothesized that the genetic mutations causing the discrepancy  
136 clustered around thrombin cleavage or factor IXa binding sites. These mutations would  
137 impair FVIII activation by thrombin or binding of FVIII to factor IX (FIXa). However,  
138 such effects were overcome in CSA by supraphysiological concentrations of thrombin  
139 and FIX, as well as the longer reaction time compared to OSA [9]. Considering these  
140 findings, FVIII that is higher by CSA than by OSA can prevent CEJB, but not subclinical  
141 joint bleeding, because the hemostatic effects are slow. However, this theory does not  
142 explain all cases. We actually identified four kinds of missense mutations in these patients.  
143 These mutations were not located near thrombin cleavage or factor IXa-binding sites, not  
144 only from the viewpoint of gene sequence, but also in terms of spatial location. As for p.  
145 Arg1800His, Yada et al. reported that the mild clinical phenotypes of patients with p.  
146 Arg1800His were associated with enhanced binding affinity of FVIII for FX [10].  
147 However, the mechanisms underlying these discrepancies are far from being completely  
148 elucidated, and further investigations are required.

149 When we estimated hemophilic arthropathy, variations in activity should have been  
150 considered in greater detail. Unfortunately, physical loading by lifestyle (such as



151 participation in sports) was not investigated in the study. Such differences might influence  
152 the development and progression of hemophilic arthropathy. Two of the PS-positive  
153 patients showed the p. Arg2169His mutation. These patients were almost the same age,  
154 but PS status differed. One possibility is that differences in the physical load according to  
155 lifestyle among patients with the same causative mutation might cause variations in PS.

156

## 157 **5. Conclusions**

158 Our study indicated that even hemophilia A patients who had never experienced CEJB  
159 risked developing hemophilic arthropathy. A distinct trend was seen for these patients to  
160 show higher FVIII:C by CSA than by OSA. Although patients who show higher FVIII:C  
161 by CSA than by OSA are generally known to display mild bleeding tendency, subclinical  
162 bleeding may still occur and lead to the development of silent hemophilic arthropathy.

163

## 164 **Conflicts of interest**

165 The authors have no interests that might be perceived as posing a conflict of interest or  
166 bias.

167

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177

#### 178 **Author contribution**

179 MO, AS and TK performed the research. NS analyzed the data and wrote the paper. NT  
180 estimated the Pettersson score. MO, ST, SS, YH, MK, FH and TK performed the genetic  
181 analyses. TK, NI, AK, HK and TM designed the research study and reviewed the paper.

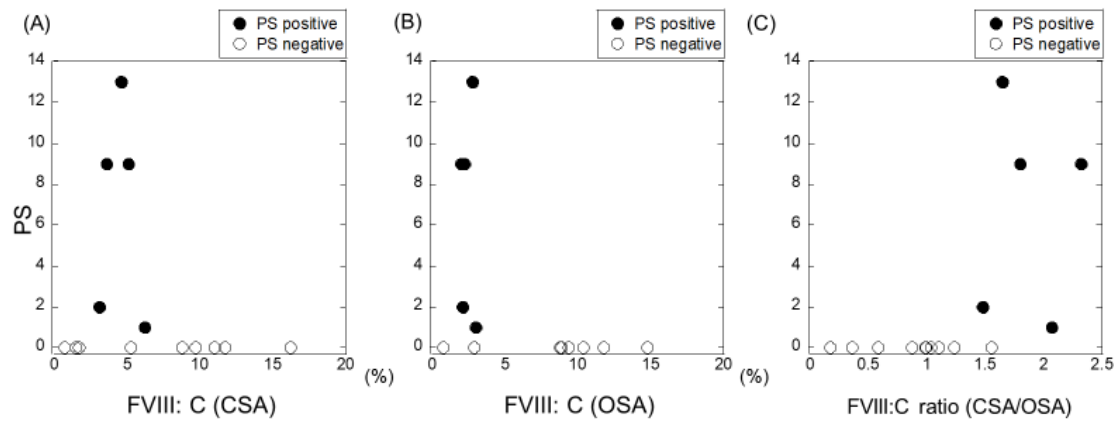
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Fig. 1.



220

221 **Fig. 1. Five of 15 patients were PS-positive**

222 PS of all enrolled patients by baseline FVIII:C by CSA (A) or OSA (B). Five patients

223 were PS-positive. They all displayed a discrepancy in FVIII:C between CSA and OSA,

224 with higher FVIII:C by CSA, Their FVIII:C ratio (CSA/OSA) was thus also higher than

225 that of the others (C).

226 PS: Pettersson score, CSA: chromogenic substrate assay, OSA: one-stage assay

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**Table 1. Comparison of the two groups**

	PS negative			PS positive			P-value
	n	mean±SD	median [IQR]	n	mean±SD	median [IQR]	
Age (years)	10	42.2±16.1	34 [31.8, 56.8]	5	57.4±15.7	67 [40.5, 69.5]	0.342
FVIII: C(CSA) (%)	10	7.6±5.1	9.3 [1.7, 11.2]	5	4.5±1.2	4.6 [3.4, 5.7]	>0.999
FVIII: C(OSA) (%)	10	8.6±4.0	8.9 [7.3, 10.8]	5	2.4±0.4	2.2 [2.1, 2.9]	0.118
FVIII: C ratio(CSA/OSA)	10	0.9±0.4	1.0 [0.5, 1.1]	5	1.9±0.3	1.8 [1.6, 2.2]	<b>0.020</b>
FVIII:Ag (%)	10	19.3±20.5	10.8 [6.9, 25.2]	5	5.2±2.9	4.9 [2.4, 8.2]	0.120
PS	10	0.0	0.0 [0.0, 0.0]	5	6.8±5.1	9.0 [1.5, 11.0]	—

Mann-Whitney U test (Bonferroni correction)

PS: Pettersson score

**Supplemental table 1. The feature of the patients included in the study**

n=15	mean±SD	median [IQR]
Age (years)	47.3±17.1	37 [33, 66]
BMI (kg/m <sup>2</sup> )	23.3±3.9	21 [20, 25]
FVIII: C(CSA) (%)	6.6±4.4	5.3 [3.4, 9.7]
FVIII: C(OSA) (%)	6.5±4.4	8.8 [2.5, 9.15]
FVIII: C ratio(CSA/OSA)	1.2±0.6	1.1 [0.9, 1.6]
FVIII:Ag (%)	14.6±17.9	8.3 [6.3, 12.1]

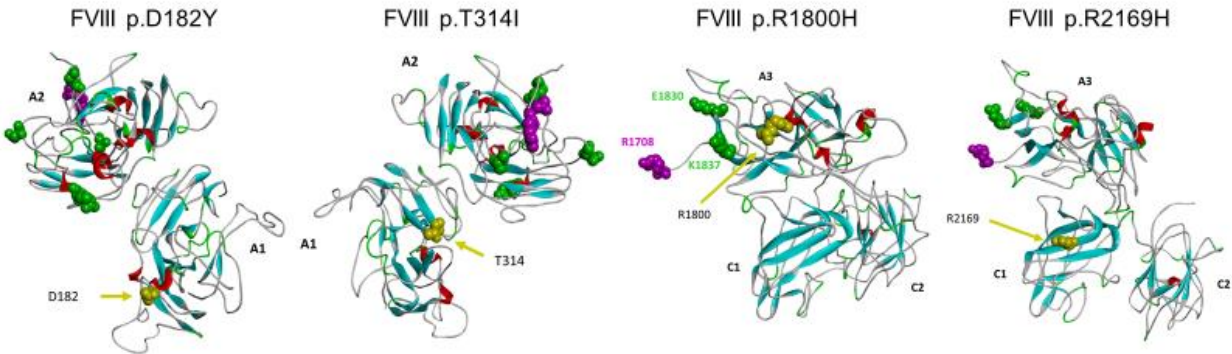
**Supplemental Table 2. The feature of the patients who were positive PS**

Age	BMI	FVIII: C (CSA)	FVIII: C (OSA)	FVIII: C ratio (CSA/OSA)	FVIII:Ag	PS	Location	DNA level	Protein level
69	20.6	4.6	2.8	1.64	8.3	13	Exon4	c.544G>T	p.Asp182Tyr
44	25	3.6	2	1.8	2.9	9	Exon16	c.5399G>A	p.Arg1800His
70	25.1	5.1	2.2	2.32	8.0	9	Exon23	c.6506G>A	p.Arg2169His
67	32.7	3.1	2.1	1.48	1.8	2	Exon23	c.6506G>A	p.Arg2169His
37	21.6	6.2	3	2.07	4.9	1	Exon7	c.941C>T	p.Thr314Ile

BMI: Body Mass Index, CSA: chromogenic substrate assay, OSA: one-stage assay, PS:Pettersson score

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Supplemental fig. 1.



244

245 **Supplemental fig. 1. Crystal structure analysis of coagulation factor VIII (human)**

246 **PDBj: 2R7E Simulation software: Discovery studio**

247 Thrombin cleavage sites are indicated as purple circles. FIXa-binding sites are indicated

248 as green circles. These mutations are not located near thrombin cleavage or factor IXa-

249 binding sites, not only from the viewpoint of gene sequences, but also in terms of spatial  
250 location.